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THE UNIVERSITY OF ALBERTA

ECOLOGY AND TAXONOMY OF HYBRIDIZING CYPRINID FISHES FROM  
UPPER AND LOWER KANANASKIS RESERVOIRS, ALBERTA

by



GEORGE A. BUTCHER

A THESIS

SUBMITTED TO THE FACULTY OF GRADUATE STUDIES AND RESEARCH  
IN PARTIAL FULFILMENT OF THE REQUIREMENTS FOR THE DEGREE  
OF MASTER OF SCIENCE

DEPARTMENT OF ZOOLOGY

EDMONTON, ALBERTA

SPRING, 1979

THE UNIVERSITY OF ALBERTA  
FACULTY OF GRADUATE STUDIES AND RESEARCH

The undersigned certify that they have read, and recommend to the Faculty of Graduate Studies and Research, for acceptance, a thesis entitled ECOLOGY AND TAXONOMY OF HYBRIDIZING CYPRINID FISHES FROM UPPER AND LOWER KANANASKIS RESERVOIRS, ALBERTA submitted by GEORGE A. BUTCHER in partial fulfilment of the requirements for the degree of MASTER OF SCIENCE.

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## ABSTRACT

Principal components and canonical variates analyses of hybridizing Couesius plumbeus (lake chub) and Rhinichthys cataractae (longnose dace) from Upper and Lower Kananaskis Reservoirs, Alberta, using 21 morphological characters, revealed two parental groups bridged continuously by an intermediate hybrid group. This morphological overlap between the hybrids and the parental species indicates that it is not always possible to positively identify 'pure' parental individuals from the Kananaskis. Neither backcross nor hybrid F2 individuals could be identified from the analyses, although the Kananaskis R. cataractae showed effects suggestive of introgression, relative to known R. cataractae from widespread Alberta localities. R. cataractae has also declined in numbers relative to C. plumbeus since last studied in 1961. C. plumbeus does not show similar effects suggestive of introgression.

C. plumbeus, R. cataractae, and their F1 hybrids were electrophoretically examined for five different protein systems. Only the serum prealbumen bands proved to be a reliable biochemical marker for the distinction of the two species. F1 hybrids can be identified by the presence of both parental prealbumen bands.

The effectiveness of the species' reproductive isolating mechanisms was studied in Lower Kananaskis Reservoir. Segregation in spawning time and habitat was not apparant. There was complete overlap in the spring spawning

period of the two species, and habitat isolation was absent in all areas studied except one inlet stream. Artificial crosses of the species, examination of hybrid sex products, and life history data showed no effective postmating isolation. The existence of distinct species' specific mating behaviors was not confirmed. However, ethological isolation is inferred to be important in minimizing this hybridization. Hybrid fertility was not experimentally confirmed.

All of the recognized environmental factors which facilitate the dissolution of species' isolation appear to be important in causing this hybridization. This includes environmental disturbance, paucity of spawning areas, species introductions, and rarity of one parental species. It does not appear possible to single out any of these factors as having the major role in facilitating hybridization.

## PREFACE

"That old and true method of Natural History - observation - must ever have a large share in the study of living things. Together they are omnipotent; disjoined they become impotent fetishes. Biology of today, as we are beginning to realize, has not too much laboratory but too little of living Nature."

C.O. Whitman (1902)

Biological Farm.

Biol. Bull. Mar. Biol. Lab.,

Woods Hole 3: 214-224.

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Environment Canada (Inland Water Directorate), Calgary, performed the water quality analyses.

Mr. J. Abelseth, Calgary Power Ltd. provided data on reservoir levels and assisted by winching the truck from



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## I. GENERAL INTRODUCTION

Interspecific and intergeneric hybridization are frequent phenomena among northern temperate, freshwater fish species, especially the cyprinids (Hubbs, 1955).

Hybridization - defined here as the successful reproduction between two species (sensu Mayr, 1969) following secondary contact - poses interesting problems and makes contributions to several levels of study, ranging from systematics and evolution to ecology and habitat management.

At the taxonomic level, putative hybrid individuals pose a methodological problem of identification. Traditionally, hybrids have been identified by their intermediacy in those characters which discriminate the parental species (Hubbs, 1955). However, the wild-caught hybrid individuals are not always strictly intermediate between that of the parental species. Certain morphological characters of hybrids may be identical to, or approximate those of one parental type, or they may be beyond the range of either parental type (Hubbs and Strawn, 1957). The difficulty of discriminating hybrids in a wild-caught collection from possible uncommon parental variants is additionally complicated by the possible presence of backcrosses and further hybrid generations. These difficulties have been largely ignored in most previous taxonomic investigations. The inadequacy of traditional hybrid index statistical techniques (Smith, 1973) may have contributed to this situation. Multivariate statistical

techniques have been advanced to overcome these problems of hybrid identification, by maximizing the separation of parental taxa, and by revealing the possible presence of backcrosses (Colgan et al., 1976; Smith, 1973). Such claims have not yet been subjected to intense scrutiny.

Apart from a purely morphological identification of hybrid individuals, the taxonomist is presented with the possibility of discriminating hybrids using single isoenzyme characters. This offers a second set of data with which to test the initial identification.

At the evolutionary level of study, two hybridizing populations may be expected to undergo observable and statistically significant evolutionary changes if hybrid fertility occurs (i.e., fusion or character displacement). The investigator is presented with the problem of determining the magnitude and direction of such changes if they exist. Such an approach can contribute to our understanding of evolution, but only with the availability of adequate past specimens from the area of hybridization.

At the ecological level, it is necessary to determine if and how environmental factors were responsible for the dissolution of the species' reproductive isolating mechanisms. Hybridization may indicate a wide overlap in those components of the species' niches concerning habitat preferences and spawning habits. Also, the long-term viability of hybrid individuals has implications for the strength of competitive interactions between the species in

an area of hybridization. Accumulating evidence suggests that hybrids succeed in environments where competition from parental phenotypes is weak (Moore, 1977).

Disturbance of the aquatic environment, either naturally or man-induced, is a primary cause of fish hybridization (Hubbs, 1955). The escalating manipulation of our lakes and rivers for reservoirs and other development purposes poses a threat to the integrity of the resident fish species. At the practical level then, hybridization studies improve our understanding of how environmental changes directly affect species, and may contribute to the mitigation of effects of future development.

Several problems at these levels were undertaken in this study of hybridization between Couesius plumbeus (Agassiz) and Rhinichthys cataractae (Valenciennes) in Upper and Lower Kananaskis Reservoirs, Alberta. At the taxonomic level, this study presents the first multivariate statistical analyses of external morphological data; it examines hybrid variability and investigates the possibility of hybrid fertility. In so doing, the effectiveness of multivariate techniques to yield such information can be practically evaluated. To corroborate the hybrid identification by morphology, an electrophoretic analysis of the population was initiated. The possibility of phenotypic changes in the Kananaskis population, evolving over 17 years of known hybridization, is also examined. The ecology of spawning populations, the effectiveness of the isolating

mechanisms, and the results of artificial cross experiments are described in order to ascertain the causes of hybridization. As a consequence, this research provides the first thorough reproductive and life history data for these species from Alberta.

## II. STUDY SITE

The Upper and Lower Kananaskis Reservoirs are located approximately 50 km south of the confluence of the Kananaskis and Bow Rivers, and within 5 km of the headwaters of the Kananaskis River at the continental divide (Fig. 1). Both reservoirs cover original mountain lake basins. The Upper Lake was the first to be modified with the construction of two hydroelectric dams (Interlakes) in 1936 and 1942. The Lower Lake was impounded in 1955, by the Pocaterra Dam. A detailed description of the history of hydroelectric development and the storage characteristics of these reservoirs is provided by Nelson (1962).

Water is stored during the spring and summer run-off and released through the fall and winter to generate electricity. The Upper Reservoir is the last to be filled and the first to be drawn down. Both reservoirs follow the same annual cycle of level fluctuations: extreme high in October and extreme low in April. The amount of water drawn down in late winter is regulated according to the amount of snowfall accumulation. Thus, the amplitude of reservoir level fluctuations may be quite variable from year to year and may differ between the two reservoirs.

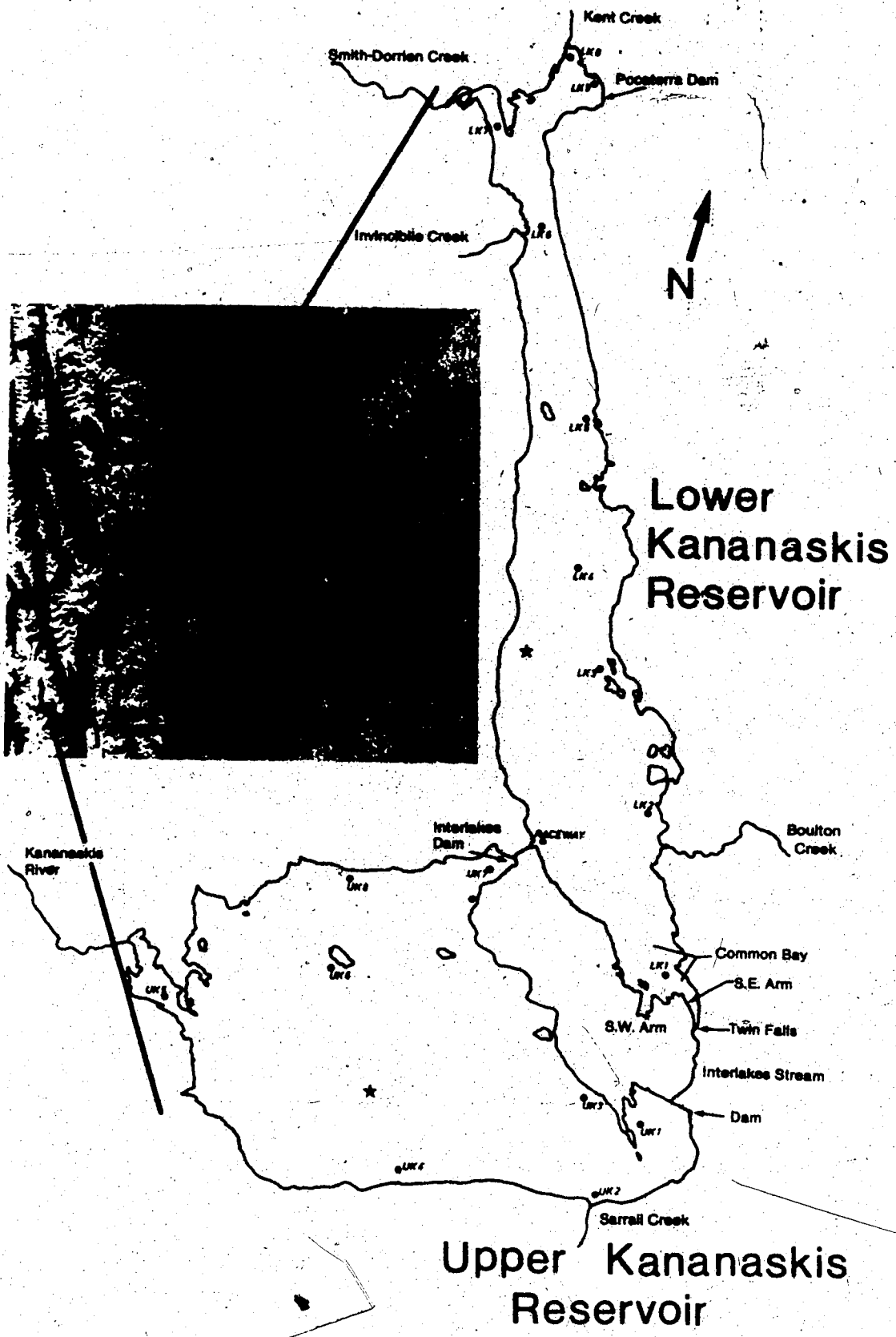
The surface area of the Lower Reservoir varies from 283 hectares at low supply level to 648 hectares at full supply level. The surface area of the Upper Reservoir varies from 633 to 855 hectares (data from Nelson, 1962). The extensive area of flooded shoreline is littered with the stump and

2

Figure 1. Map of Upper and Lower Kananaskis reservoirs (1 cm = 0.55 km).

The stars in each reservoir mark positions of water quality sampling stations. Fish sampling sites are marked with dots and site numbers (corresponding to the numbers in Appendix Tables 1. and 2.). The inset photo is copied from a Landsat I false color image shot on 5 October 1972, at an altitude of 920 km and processed by the Canada Centre for Remote Sensing, Ottawa. The image format centre is Lat.N. 51°15', Long.W. 155°00' and corresponds to the National Topographic Survey (Canada) Map 820.





root remnants of trees removed prior to dam construction. The Government of Alberta began partial removal of stumps and roots in June, 1978. The substrate of this 'fluctuation' zone consists of unconsolidated gravel and boulders interspersed with large areas of thick eroded silt (Plate 8). The only benthic fauna to utilize this 'fluctuation' zone are the chironomids (Fillion, 1963).

In the spring, as the reservoirs fill, the spawning fish are presented with flooded shoreline areas and with upstream areas of original inlet creeks. By the time spawning commences in late May (suckers) or early June (cyprinids), the presumed original spawning beds have been flooded.

All inlet streams of the Upper Kananaskis Reservoir flow unrestricted in their natural beds. The flow of Boulton Creek and Invincible Creek into the Lower Reservoir has remained unchanged since pre-impoundment days. However, due to dam construction, Smith-Dorrien Creek and the diverted Kent Creek, which formerly drained into the Kananaskis River, now drain into the Lower Reservoir. All four streams flow unregulated with spring and summer run-off.

The original outlet flow (at the extreme south end of the Lower Reservoir), from the Upper Lake into the Lower Lake, has been reduced to seepage through the earthfill dam (here designated as Interlakes Stream). Similarly, the man-made outlet of the Upper Reservoir into the Lower Reservoir at Interlakes Dam is controlled so that there is

no flow during the spring and summer (i.e., from end of drawdown in April, to its recommencement in October).

### III. PAST STUDIES

Previous pre- and post-impoundment studies on the limnology, productivity, and fishes of the Upper and Lower Kananaskis Reservoirs were reviewed by Nelson (1962) in order to establish the chronology of biological changes. Nelson deduced that both C. plumbeus and P. cataractae were introduced to the Upper and Lower Kananaskis Reservoirs as bait fishes, probably from Bow River populations (P. cataractae probably in the 1930's; C. plumbeus probably in the late 1950's). The occurrence of hybridization between these two species in Upper Kananaskis Reservoir was noted by Nelson (1962). The external morphology of hybrids from Upper Kananaskis Reservoir was described by Nelson (1966). The hybrids from both reservoirs were later examined using osteological characters and a hybrid index technique (Nelson, 1973). One juvenile specimen appeared to be intermediate between the majority of hybrids and one parental phenotype.

Hybridization between C. plumbeus and P. cataractae has been noted in several localities outside Alberta (Simon, 1946, in Wyoming; Hubbs and Lagler, 1949, in Lake Superior; Taylor, 1954, in Michigan), but without the substantiating evidence provided by Nelson (1966, 1973).

Identification of C. plumbeus x P. cataractae hybrids by numerical taxonomic or electrophoretic techniques has not been reported. Additionally, the reproductive ecology of the Kananaskis populations has not been studied.

## IV. FISH SAMPLING

### METHODS

Fish were collected between May 10 and August 29, 1977, from Upper and Lower Kananaskis Reservoirs and between May 1 and July 27, 1978, from Lower Kananaskis Reservoir.

C. plumbeus, and R. cataractae and their hybrids were sampled from the reservoirs proper, using monofilament gillnets (18 mm, 26 mm, and 38 mm stretch mesh), beach seines (8 mm mesh), dipnets, and baited minnow traps. This variety of sampling methods assured the capture of all sizes of specimens. Sampling sites are shown in Figure 1 and site descriptions provided in Appendix Table 1 and Table 2. The relative abundance of the hybridizing species was established for all localities. Data on depth, substrate type, and distance from shore were recorded for all captures. The study of the breeding biology of the cyprinids in 1978, involved intensive sampling of only the Lower Reservoir. The Upper Reservoir presented boat access problems in the early spring.

Inlets of Lower Kananaskis Reservoir were electroshocked (Smith-Root Electroshocker Model VII) beginning in late April 1978, when the majority of the reservoir was ice-covered (95%). A beach seine was stretched across each inlet stream mouth and the stream was electroshocked and the bottom disturbed downstream into the seine (refer to Plate 7). Smith-Dorrien Creek, Boulton Creek, Kent Creek, and the original outlet stream of Upper

Kananaskis Lake (here designated as Interlakes Stream) were sampled in this manner. As spring run-off increased, it became impossible to sample inlet streams (except Interlakes Stream), and gillnets were placed across the stream mouths and minnow traps placed along the edges.

Standard length and sex were recorded for all fish sampled. A representative sample was returned to the laboratory for weighing, gonad and scale removal, and preservation in 10% formalin. Samples of the two species and their hybrids were deposited in the University of Alberta Museum of Zoology (UAMZ).

#### GENERAL CATCH RESULTS

Cyprinid catch data from 1977 and 1978 for Upper and Lower Kananaskis Reservoirs are summarized in Appendix Tables 1 and 2. Capture method, locality, date, depth, and numbers of specimens are provided.

The ratios of parental species to hybrid individuals captured in each year were as follows: 1977, 133 C. plumbeus to 7 R. cataractae to 1 putative C. plumbeus x R. cataractae; 1978, 174 C. plumbeus to 12 R. cataractae to 1 putative C. plumbeus x R. cataractae. A total of 56 putative hybrids were captured in 1977 and a total of 9 putative hybrids were captured in 1978. A disproportionate cyprinid species' ratio in favour of C. plumbeus was also recorded by Nelson (1962). He found C. plumbeus to be 4.0 to 5.6 times as numerous as R. cataractae in gillnet catches for Lower

and Upper Kananaskis Reservoirs, respectively. Results of the present study (C. plumbeus 14 - 19 times as numerous as R. cataractae) may indicate an increase in the disproportion of species' numbers over the intervening 17 years. It is not known if C. plumbeus is numerically increasing, or if R. cataractae is numerically declining.

Suspected cyprinid hybrids were captured throughout both reservoirs and were not concentrated in a limited number of locations. Relatively large hybrid capture numbers (Appendix Table 1 and Table 2) at certain sites are due to more intensive sampling at these sites. Hybrids were captured alone and with individuals of each species. Capture of the hybrids was both infrequent and unpredictable.

The cyprinids sporadically swarmed in large schools with juvenile suckers over recently flooded inshore areas. The schools were observed only in midsummer on sunny days when the water temperature exceeded 15°C; schooling did not occur on cloudy, windy, or cool days. The majority of the large seined collections came from the infrequent occasions on which such large schools were observed. Hybrids were often collected from such schools. Except for these occasions, the cyprinids generally remained offshore. The mixed schools were involved neither with reproduction (since they occurred after spawning) nor with feeding (since the flooded areas are assumed not to have a developed bottom fauna). The reasons for this mixed schooling remains unstudied.

## V. LIMNOLOGY

### INTRODUCTION

Changes in environmental factors can trigger fish hybridization (Hubbs, 1955). The modification of a lake to a reservoir is accompanied by an obvious annual change in surface water elevation and basin morphometry. This change may directly affect fish spawning habitat so that miscegenation results.

Less visible are the possible indirect effects of changes in water quality and the thermal regime. Small changes in these factors may upset the competitive balance existing between the species and result in a disproportion of one species (a possible precursor to hybridization). An altered temperature regime may affect the timing of species' spawning activities thereby destroying a temporal barrier to reproductive isolation. Changes in these environmental factors may contribute to what Anderson (1949) termed the "hybridization of the environment" (Anderson, 1949).

Apart from accurate water level statistics, there are only few and rudimentary limnological data available for Upper and Lower Kananaskis Lakes prior to impoundment (Rawson, 1937, 1948; Miller, 1954; Thomas, 1955).

The purpose of this section is to: 1) document the degree of water level fluctuations; 2) document present water chemistry and temperature conditions; and 3) to speculate as to whether the latter have undergone significant changes since impoundment.



## METHODS

Physico-chemical parameters were measured mid-monthly during June, July, and August, 1977, from three depths (surface, mid-depth, and 1 m off bottom) at one sampling station on each of the Upper and Lower Kananaskis Reservoirs (Fig. 1). The water samples were shipped within 24 hours to Environment Canada, Inland Waters Branch, Calgary, Alberta, for analysis of alkalinity, color, pH, residue (non-filterable), turbidity, total Kjeldahl nitrogen, total phosphorus, and total carbon. Water quality samples were not taken during the summer of 1978. Water temperature profiles were recorded when water samples were collected, and also sporadically through the summer. The dates on which water temperature profiles were taken from Lower Kananaskis Reservoir in 1977, are as follows: May 9, 17, 31; June 20, 29; July 15; Aug. 8, 16; Sept. 3, 17. Water temperature was measured at 1 m intervals using a Y.S.I. telethermometer. Reservoir elevations were provided by Calgary Power Ltd.

## RESULTS AND DISCUSSION

Results of the water quality analysis for Upper and Lower Kananaskis Reservoirs are presented in Appendix Tables 3 and 4. Most of these water quality parameters are reported for the first time from these reservoirs.

Both reservoirs have similar basic pH values. These do not appear to have changed since last studied by Fillion (1963) and Nelson (1962). According to Nelson, no pH change

accompanied construction of Lower Kananaskis Reservoir.

Specific conductance values are high for each reservoir as a result of spring run-off. The values for the Lower Kananaskis Reservoir are consistently higher than those of the Upper Reservoir, a trend shown by Pillion. Inorganic carbon values are also higher for the Lower Kananaskis Reservoir.

The remaining parameters - color, residue, turbidity, nitrogen, phosphorus, and organic carbon - show no difference between the reservoirs. An examination of all the above parameters does not reveal any change attributable to reservoir construction. Both reservoirs are characterized by the clarity and nutrient-poor state expected for that of the original lakes.

Spring and summer isotherms from Lower Kananaskis Reservoir, 1977, are shown in Figure 2. These were constructed from the water temperature profile data. Additional temperature data from 1978, are provided in Appendix Table 5. Some thermal stratification occurs during the summer. There have been no thermal changes since the studies by Nelson and Pillion. This probably represents the original thermal situation in the Lower Lake for the spring and summer months. Since Calgary Power Ltd. does not lower the level of Upper Kananaskis Reservoir in the summer, there is no unnatural input of cool hypolimnetic waters into the Lower Reservoir in the summer. There is no reason to believe that the water temperature cue for spawning has been


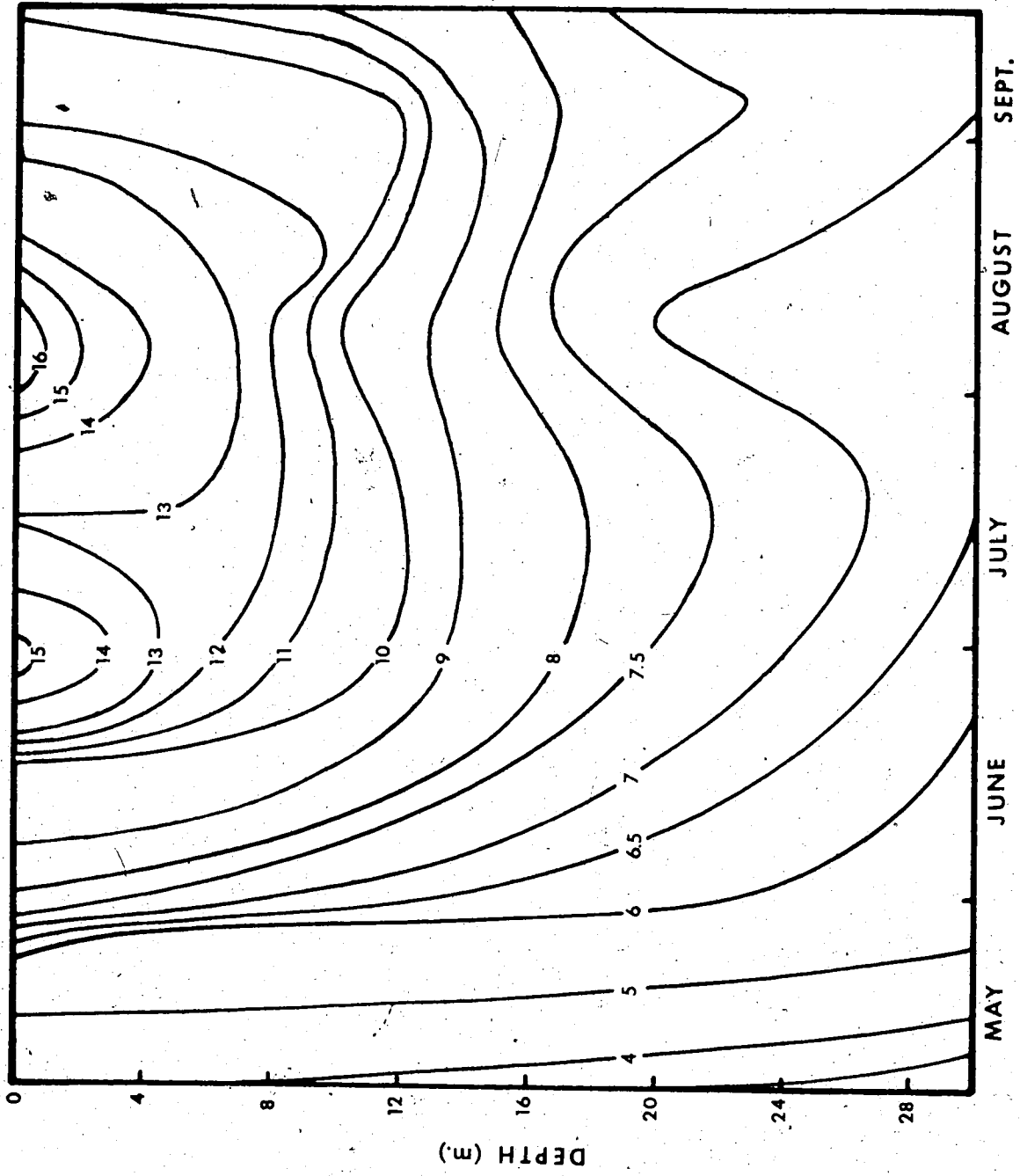


Figure 2. Spring and summer, 1977 isotherms (in °C) from the water quality sampling site, Lower Kananaskis Reservoir.

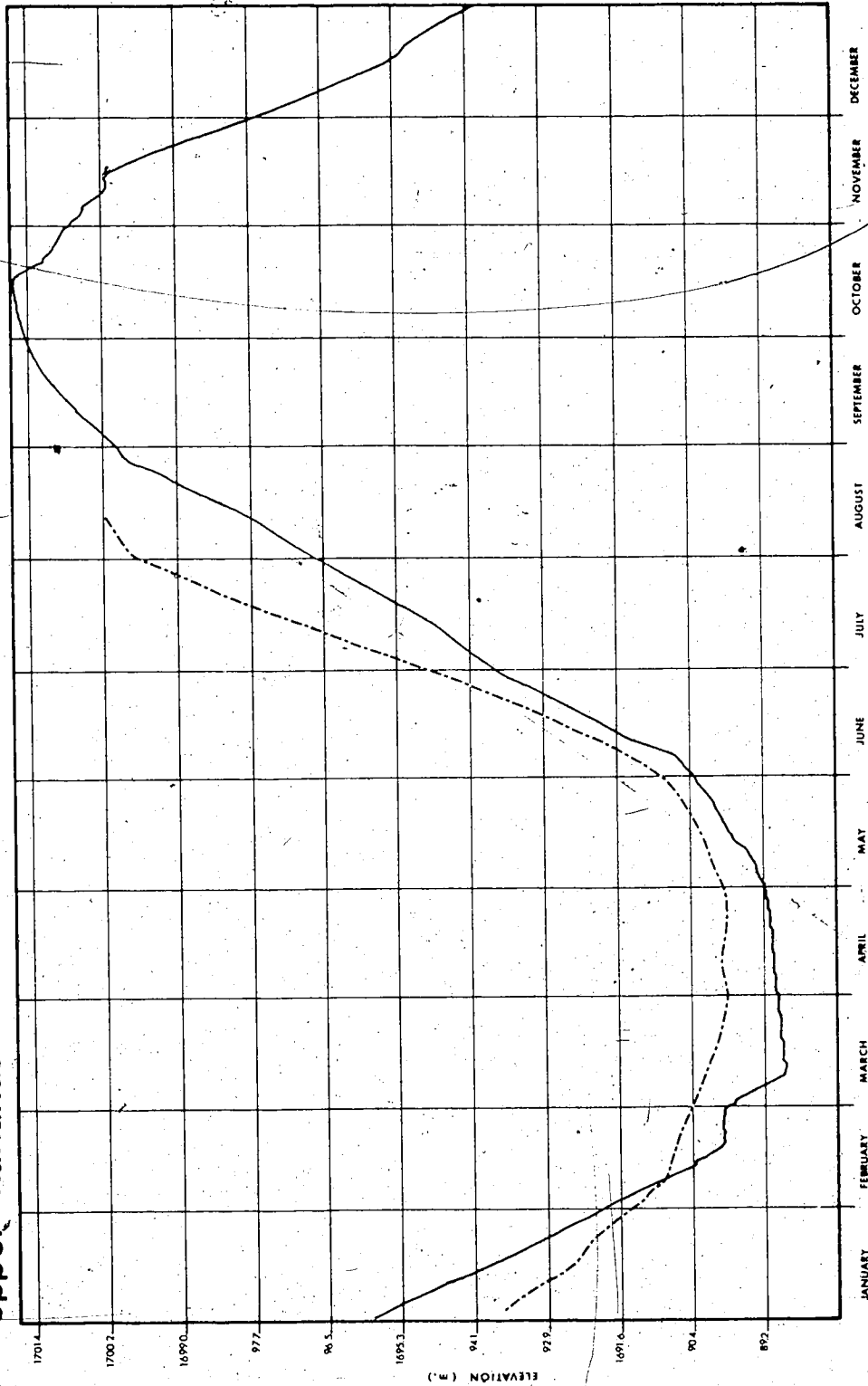


altered. However, the effects of hypolimnetic input from Upper Kananaskis Reservoir during fall drawdown on fish biology remains unstudied.

Annual water level fluctuations for 1977 and 1978, are shown graphically in Figures 3 and 4 for Upper and Lower Kananaskis Reservoirs, respectively. There is much unpredictability in water level from year to year; the amount of drawdown is adjusted according to the amount of snowfall accruing through the winter. Of more importance, there is significant annual variation in water levels at spawning time: in Upper Kananaskis Reservoir a 0.6 m difference and in Lower Kananaskis Reservoir a 2.0 m difference. These differences are enough to radically alter shoreline characteristics from year to year. Permanent spawning areas cannot be established under this regime.

Figure 3. Annual water level fluctuations in the Upper Kananaskis Reservoir during 1977 and 1978. Elevation is given as above sea level. Data provided by Calgary Power Ltd.

# Upper Kananaskis Reservoir

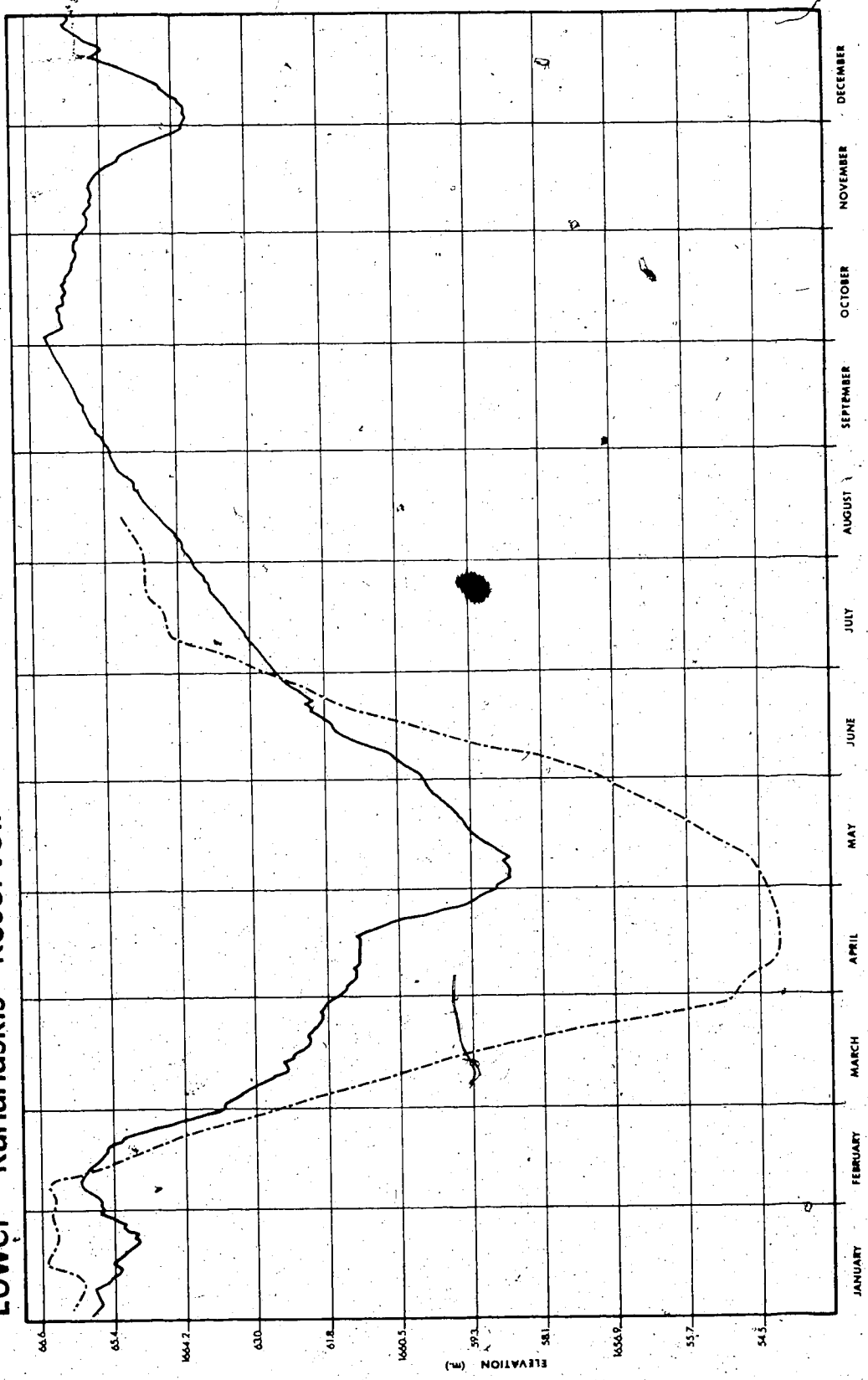


1977  
1978

Figure 4. Annual water level fluctuations in the Lower Kananaskis Reservoir during 1977 and 1978. Elevation is given as above sea level. Data provided by Calgary Power Ltd.



# Lower Kananaskis Reservoir



1977 ———  
1978 - - - -



## VI. IDENTIFICATION OF HYBRIDS USING MULTIVARIATE ANALYSIS

### INTRODUCTION

Although the Kananaskis cyprinid hybrids were identified and morphologically examined by Nelson (1966, 1973), including analysis by hybrid index, there have been no numerical taxonomic studies attempted.

The advantages of multivariate statistical techniques relative to traditional methods have been reviewed by Smith (1973). Of singular importance is elimination of the necessity to prejudge hybrid or parental origin of the specimens. This is of particular value when hybrids are variable and hybrid fertility is possible. In such cases, it may be impossible to discriminate hybrids from parental individuals.

This section of the study describes principal components and canonical variates analyses of numerous morphological characters of both hybrids and parental individuals. The methods do not require assumptions as to the identity of any Kananaskis specimens. Hybrid variability, possible hybrid fertility, and the effects of hybridization over time on the populations are described.

### METHODS

#### SPECIMENS EXAMINED

The external morphology of representative Kananaskis specimens was compared to: 1) that of museum specimens from Alberta localities where hybridization is presumed not to

occur; and to 2) that of museum specimens collected from these reservoirs in previous years.

All Kananaskis specimens examined in the morphological studies were deposited in the University of Alberta Museum of Zoology (UAMZ numbers 4145-4201).

There was a collecting bias in the field directed equally toward the less common specimens, i.e., typical *R. cataractae* and all putative hybrids. Typical *C. plumbeus*, being highly abundant, were randomly collected. A random sample of 137 specimens was chosen from the field collections of each reservoir, irrespective of sex and presumed identity, and were used in the analyses (designated here as the Kananaskis specimens). The sex ratio for the entire Kananaskis sample was 62 females to 75 males.

Fifty *C. plumbeus* and fifty *R. cataractae* were selected from widespread Alberta localities in UAMZ collections encompassing several drainage systems (designated here as the Alberta specimens). These control specimens were selected such that the size (standard length) range was similar to that of the Kananaskis specimens. The catalogue numbers, locations, and sex ratios of these collections are given in Table 1. Unfortunately, due to the scarcity of large male *R. cataractae* in the UAMZ collections, the sex ratio of Alberta *R. cataractae* was predominantly female. The sex ratio for the entire Alberta sample was 73 females to 27 males.

Past collections of *C. plumbeus* from Upper and Lower

TABLE 1. Locations and University of Alberta Museum of Zoology catalogue numbers for Alberta control specimens of *Couesius plumbeus* and *Rhinichthys cataractae*.

Species	Locality	Lat. N/Long. W	UAMZ#	Number of Specimens Examined
<i>C. plumbeus</i>	Brazeau Reservoir	52°57' 115°35'	566	8
	Red Deer River	52°19' 113°06'	3326	8
	Abraham Reservoir	52°16' 116°23'	3350	8
	Little Smokey River	54°44' 110°10'	1654	8
	Pothole Creek	49°25' 112°54'	3843	8
	Winefred River	56°02' 110°36'	3451	2
	Bare Creek	49°14' 110°00'	3231	3
	McIvor River	58°18' 118°03'	3418	3
	N. Saskatchewan River	52°25' 116°04'	3325	2
Total sex ratio of 27 ♀:23 ♂				
<i>R. cataractae</i>	N. Saskatchewan River	52°16' 116°23'	3381	7
	N. Saskatchewan River	53°30' 113°34'	2044	2
	N. Saskatchewan River	52°16' 116°23'	3322	1
	Graburn Creek	49°38' 110°01'	1620	7
	Cutbank River	54°43' 118°32'	3903	2
	Horse River	56°43' 111°23'	3435	2
	Abraham Reservoir	52°16' 116°23'	3369	5
	Gregoire River	56°29' 110°48'	3414	6
	Pothole Creek	49°25' 112°54'	4000	2
	Pothole Creek	49°25' 112°54'	3843	5
	Bare Creek	49°14' 110°00'	3231	5
	Willson Creek	51°49' 115°13'	3873	4
	Wapiti River	55°08' 118°18'	269	1
	Pinto Creek	54°58' 119°28'	272	1
Total sex ratio of 46 ♀:4 ♂				

Kananaskis Reservoirs were subjected to morphological study to determine any effects of hybridization over time. The 1961 collection of 11 C. plumbeus from Upper Kananaskis Reservoir (UAMZ 863) were analyzed. Twenty-five C. plumbeus were examined from the 1969 collections (UAMZ 2706, 2707, 2518) and 25 from the 1973 collections (UAMZ 3345, 3346) including specimens from both Upper and Lower Kananaskis Reservoirs. Only specimens within the size range of the Kananaskis specimens were selected.

Past Alberta collections of R. cataraetae in the UAMZ holdings were few and contained only small specimens beyond the size range comparable to the Kananaskis specimens. These were deemed inadequate for comparison to recent Kananaskis collections. Thus, it was only possible to study the effects of hybridization over time on C. plumbeus.

#### CHARACTERS USED

The following 21 morphological and meristic characters were recorded for all 298 specimens. These were chosen because all were good descriptors of the fishes' shape and because some are known (Nelson, 1966) to be efficient discriminators of the parental species and of the putative hybrids. Counts and measurements follow the descriptions and diagrams in Hubbs and Lagler (1964) except as otherwise defined. Measurements were made using needle point dial calipers, read to the nearest 0.1 mm. Microscope magnification was used when required. Abbreviations for each

character follow in parentheses.

1. standard length (SL)
2. head length (HL)
3. orbit length (OL)
4. snout length, (SN)
5. interorbital width (IN) - (the least fleshy width of the interorbital)
6. head depth at nostril (HDN) - the distance from the nostril downward to the ventral contour of the head.
7. head width at eye (HWE) - the greatest dimension from one lateral side of the head to the other lateral side on a line below the orbits.
8. mouth length (ML) - (length of upper jaw)
9. mandible width (MAN) - (width of gape)
10. snout overhang (SOV) - the distance from the tip of the snout along the midline to the most anterior portion of the lower jaw.
11. frenum width (FRE) - the distance between the grooves separating the upper jaw from the snout.
12. isthmus width (IW) - the distance between the gill membranes at point of attachment to isthmus.
13. body width at dorsal fin origin (WDO) - the maximum distance from one lateral side of the body to the other lateral side on a line below the dorsal fin origin.
14. snout to dorsal fin origin (SDO) - (pre-dorsal length)
15. caudal peduncle length (CP)
16. snout to anus (STA) - distance from the tip of the snout to the anal opening.
17. pectoral fin origin to anus (PTA)
18. depth of caudal fin fork (CF) - the distance from the apex of the caudal fork outward to a straight line connecting the tips of each outstretched caudal fin lobe.

19. number of lateral line scales (LS)
20. number of scales from dorsal fin origin along diagonal to lateral line (ALS) - (number of scales above lateral line).
21. number of scales around caudal peduncle (CPS)

This is the order in which the characters were used in all following morphological analyses.

Ratios were specifically avoided in order to eliminate problems with allometry and compounded variance (Varr, 1955; Atchley et al., 1976). Multivariate analyses were run with untransformed data and square-root transformation, since size differences could lead to vectors not being coplanar (Pimental, 1975). Frenum width for known C. plumbeus was always 0.00, and since several other characters also had values < 1.0, a logarithmic transformation of the data set was inappropriate. The effectiveness of using transformed data was assessed.

These 21 characters were chosen from an original 31 characters because of their significant contribution toward group discrimination. The 10 characters discarded prior to in-depth analysis included: mouth to eye length (distance from corner of the mouth along midline to a vertical line marking the anterior margin of the orbit), cheek depth, dorsal fin origin to pelvic fin origin, dorsal fin height, anal fin height, pelvic fin axillary process (absence, presence), number of gillrakers, number of pectoral rays, number of anal rays, and number of dorsal rays.

Some of these characters, e.g., dorsal fin origin to

pelvic fin origin, number of gillrakers, and number of fin rays were discarded because they were ineffective at group discrimination. The rest were removed because they were confounding sources of variation. The characters, mouth to eye length, and cheek depth could not be accurately measured, pelvic fin axillary process could not be accurately scored, and the fin height characters are sexually dimorphic (Wells, 1978). Since each sex was not separately analyzed, characters of known sexual dimorphism were eliminated.

#### STATISTICAL TECHNIQUES

##### Principal Components Analysis

Principal components analysis (PCA) (Morrison, 1967; Cooley and Lohnes, 1971; Pimental, 1976) was used to display patterns of morphological similarity of individuals from mixed species' populations with hybrids, relative to populations without hybrids. This method finds the orthogonal axes through the n-dimensional character space in directions of greatest variance. The new axes provide new directions from which to view the group relationships, if any, within the data. The plot of the specimens on the first three components will usually display a great amount of the total variance within the data set.

PCA scores were calculated from the character correlation matrix using the Clustan 1C computer program (Wishart, 1975) for the following two separate data sets: 1)



all 137 Kananaskis specimens; and 2) 100 control Alberta specimens. The projection scores of the specimens in the two data sets were plotted against the first and second axes and also against the second and third axes. Untransformed and square-root transformed data were analyzed. The assumption of a single multivariate normal population for a PCA has been ignored for these reasons: 1) no statistical inferences were drawn from these analyses; and 2) departures from normality are not thought to invalidate the results (Dudzinski et al., 1975).

#### Canonical Variates Analysis

The same two data sets (untransformed and square-root transformed) were subjected to canonical variates analysis (CVA) in order to quantify the relationship between individuals (Pimental, 1976). This method requires the a priori identification of two or more known groups within the data set. Axes are calculated which maximally separate these groups while minimizing the within-groups variance. Specimens of unknown origin can then be assigned a position on this axis (calculated using the discriminant function) relative to their similarity to either group.

The BMD 07M computer program (Dixon, 1973) was used to assess the relationship between the Kananaskis specimens (the a priori unknown group) and the control Alberta specimens (the a priori known group). The canonical scores were plotted against the first canonical axis and graphed as

frequency histograms. An examination of further canonical axes did not discriminate more groups than was revealed by the first canonical axis.

In addition to correct a priori identification, CVA assumes a multivariate normal distribution with equal variance-covariance matrices. A Bartlett's test of homogeneity of dispersions (DERS program; Bay, 1969) on the two known groups showed a significant difference between the matrices ( $F = 2.6$ ,  $F_{.05}(210, 29352) = 1.00$ ). However, it is generally held that CVA has sufficient robustness, so that departures from the assumptions will not be serious (Pimental, 1976).

## RESULTS

### PRINCIPAL COMPONENTS ANALYSIS

The principal component scores for the Alberta specimens, calculated using untransformed data, were plotted against the first three axes (Fig. 5). The results confirm the initial identification of 'pure' specimens representing the two species, with no morphological intermediates.

Component I represents overall size differences among the specimens, ranging from small on the left to large on the right. Component II represents between-species discrimination and Component III represents the largest component of within-species variation (perhaps due to sexual dimorphism or population differences). The variation expressed by the three components represents 91.3% of the




Figure 5. PCA Using Untransformed Data; Frenum Included

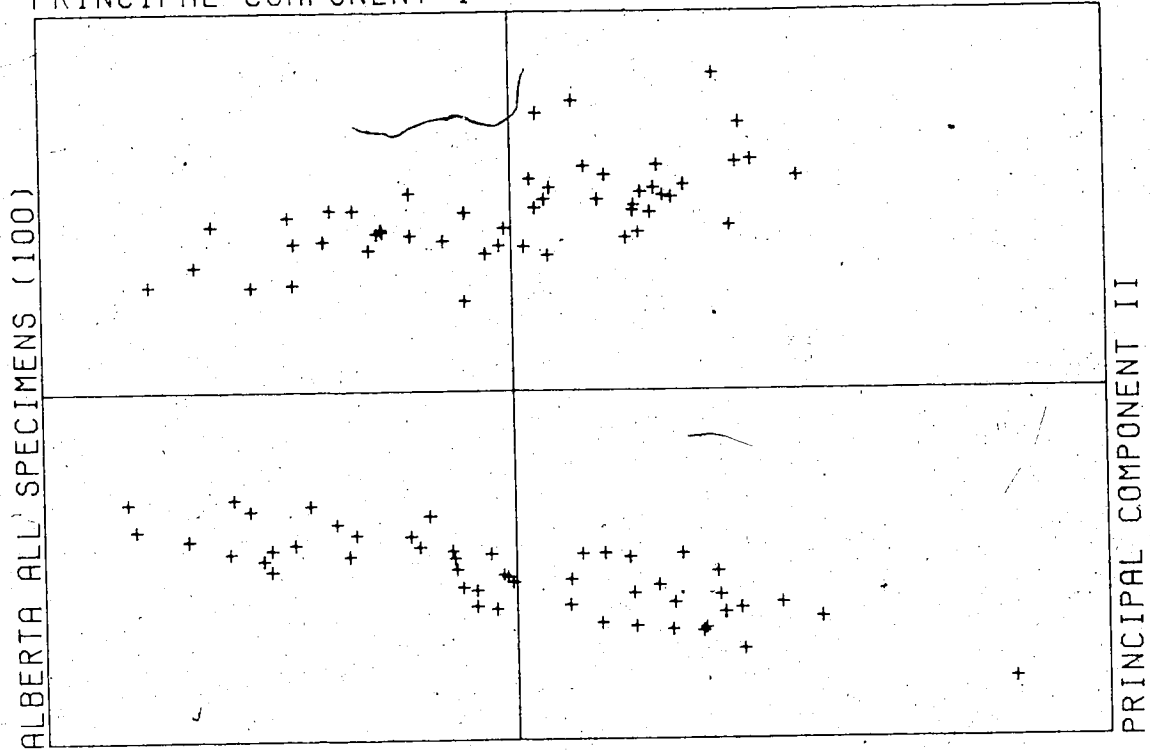
- A. Plot of first and second principal component scores for Alberta, 'control' specimens. *Couesius plumbeus*, top cluster; *Rhinichthys cataractae*, bottom cluster.

Component I = 59.4% of total variation in 21 character correlation matrix; Component II = 28.3% of variation.

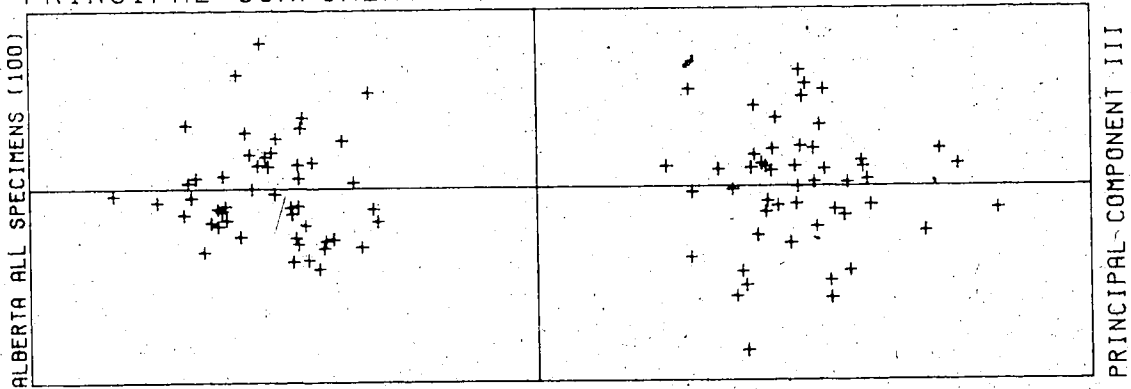
- B. Plot of second and third principal component scores for Alberta 'control' specimens. *R. cataractae* on left; *C. plumbeus* on right. Component III = 3.5% of variation.

**A.**

PRINCIPAL COMPONENT I

**B.**

PRINCIPAL COMPONENT II



variance in the total data matrix. Investigation of additional components would have been redundant. A summary of the computer results on which these plots are based, is given in Appendix Table 6.

The principal component scores for all Kananaskis specimens, calculated using untransformed data, were plotted against the first three axes (Fig. 6). The projection of scores on components I and II show the elongate clusters, as expected from the control group, but with a dissimilar bridging of the two species' clusters. The specimens of this bridge represent the putative hybrid specimens. The absence of larger hybrid specimens from the Kananaskis sample accounts for the clear discrimination between the two species' clusters to the right of the origin. Large hybrids were available in the collections, and if included in the analysis would presumably occupy this space.

The variation expressed by the three components represents 91.5% of the variance in the total data matrix. The difference in % variation represented by component I between the Alberta group (59.4%) and the Kananaskis group (66.4%) is due to the difference in range of standard lengths between the Alberta group (4.1-11.0 cm) and the Kananaskis group (3.4-12.3 cm).

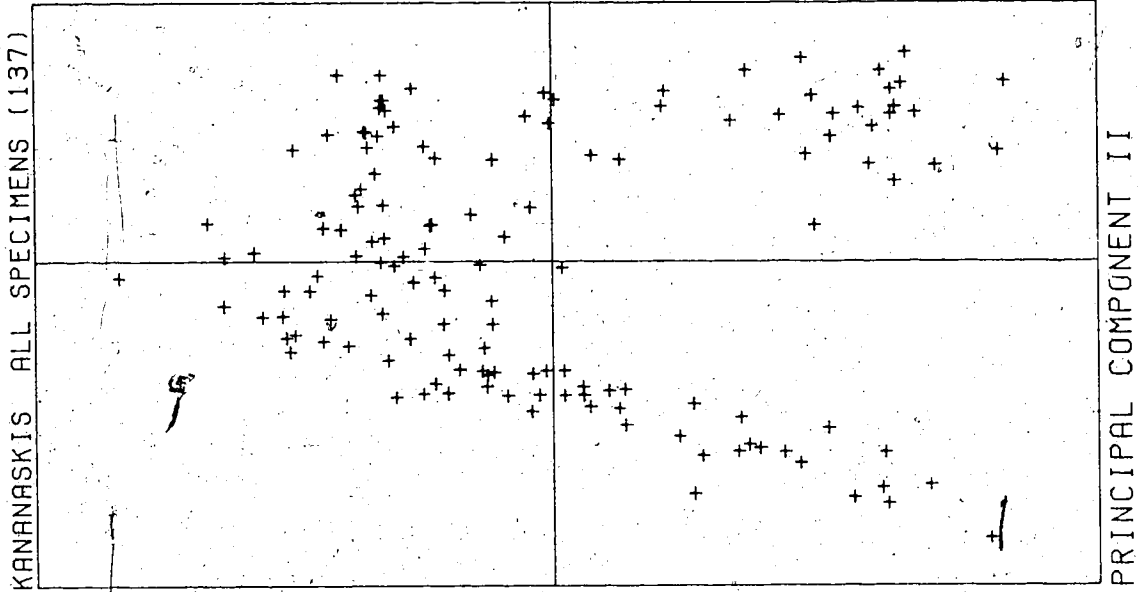
The intervening hybrid complex, shown on the Kananaskis projection of the second and third axes, obscures discrimination between the two species. This analysis provides evidence that the putative hybrids are not all

Figure 6. PCA Using Untransformed Data; Frenum Included

- A. Plot of first and second principal component scores for all Kananaskis specimens. *Couesius plumbeus* -like specimens towards bottom; *Rhinichthys cataractae* -like specimens towards top, with a hybrid complex bridging the 2 species clusters. Component I = 66.4% of variation.
- B. Plot of second and third principal component scores for all Kananaskis specimens. *C. plumbeus* -like towards left; *R. cataractae* -like towards right. Component III = 4.9% of variation.

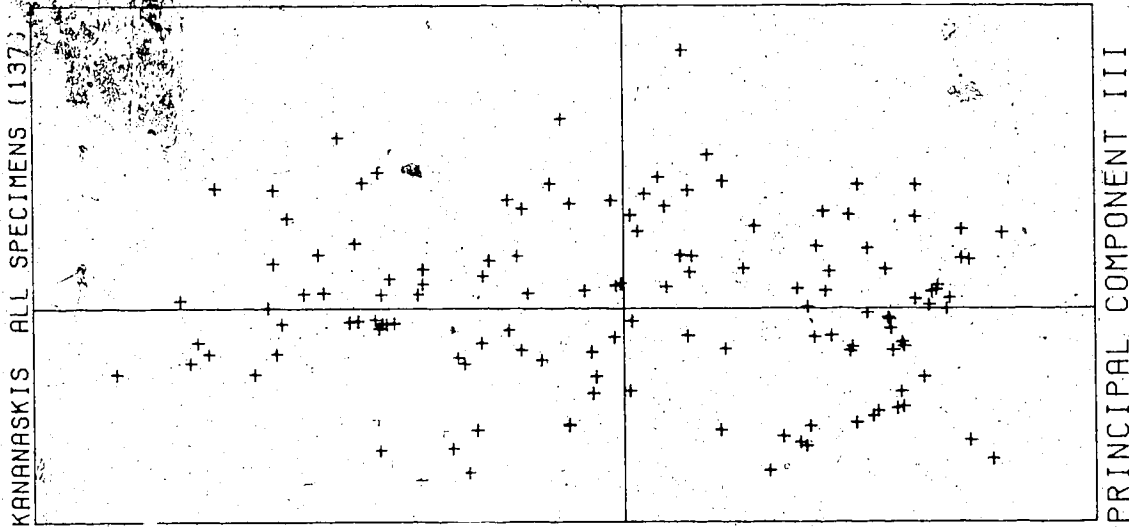
A.

PRINCIPAL COMPONENT I



B.

PRINCIPAL COMPONENT II



strictly intermediate. They cover the entire phenotypic range between the parentals suggesting mixed genetic origins, i.e., backcrossing of F1 hybrids to each parental species. A summary of the computer results on which these plots are based is given in Appendix Table 7.

Similar analyses were again performed separately on Alberta specimens, and Kananaskis specimens using square-root transformed data. The resulting plots were similar to those generated using untransformed data. Transformation produced a slightly better discrimination between the smallest members of each species, but produced no new insights. The Kananaskis parental species remain continuously bridged to the same degree by the hybrid complex. Computer results for these analyses are summarized in Appendix Tables 8 and 9.

#### CANONICAL VARIATES ANALYSIS

The relationship between the Kananaskis specimens including hybrids, and the Alberta 'control' species can be expressed in a quantitative manner by a frequency distribution of their canonical scores. The results of a CVA using untransformed data with the full character set are given in Figure 7.

The two distributions for each Alberta species are separated by a large gap of approximately 14 standard deviations between their means. The Kananaskis specimens form a continuous distribution with no gap between the



Figure 7. CVA Using Untransformed Data; Frenum Included

Frequency histograms of canonical scores on first canonical axis of specimens from Alberta 'control' group (upper histogram) and Kananaskis group (inverted histogram). Alberta species formed 2 *a priori* known groups; all Kananaskis specimens ran as *a priori* unknown groups. Units on x-axis are in standard deviations; y-axis is number of specimens. Initial field identifications of Kananaskis specimens are shaded as follows:



*Couesius plumbeus*

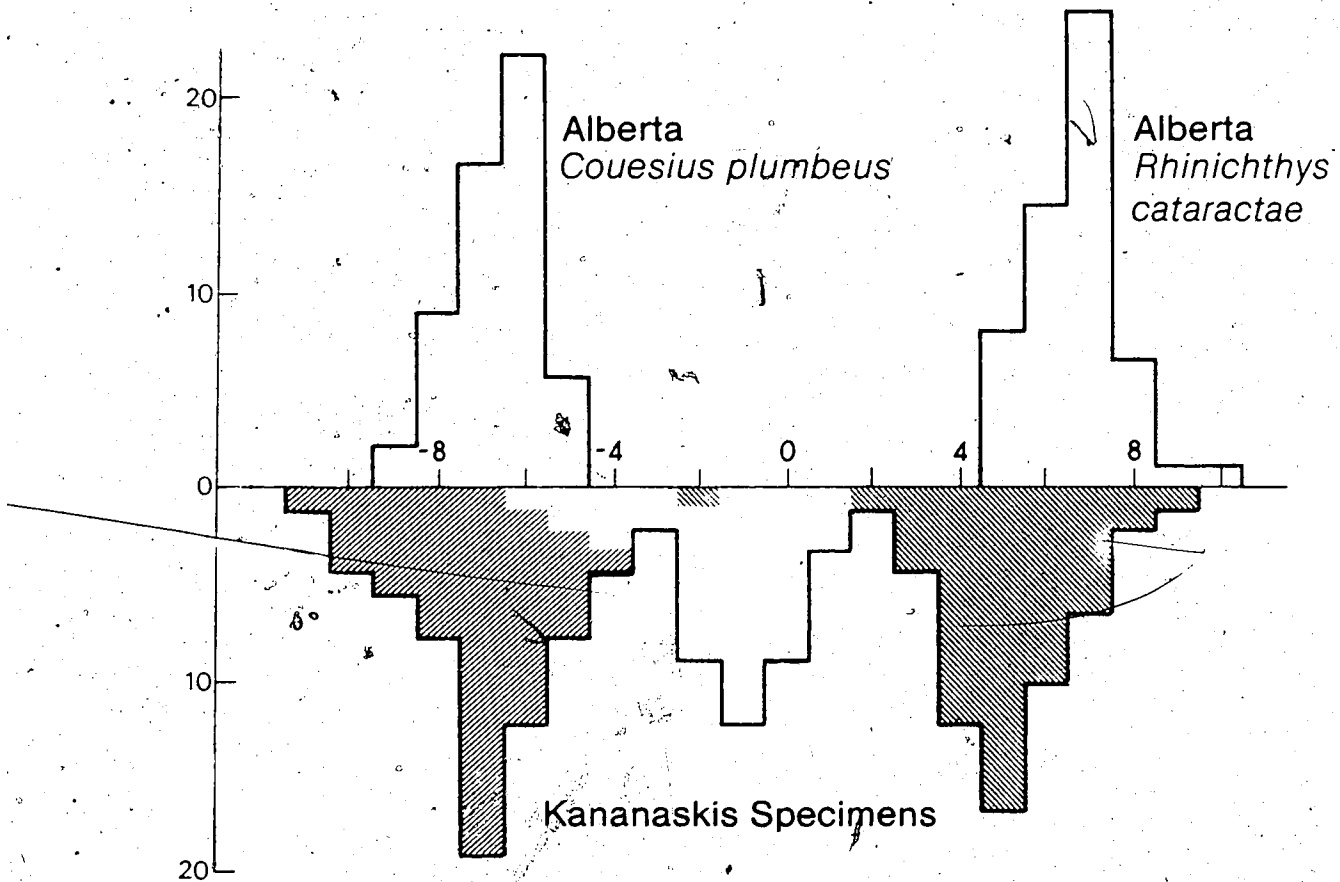


*C. plumbeus* X *R. cataractae*.



*Rhinichthys cataractae*

### Untransformed Data, Frenum Character Included



typical parental species.

The apparent trimodality of the Kananaskis distribution was confirmed using Cassie's (1963) graphic method of polymodal frequency separation ( $\chi^2$  test for goodness of fit,  $\chi^2 = 25.0$ ; d.f. = 15,  $p > 0.05$ ). The two outer modes of Kananaskis specimens are composed of the corresponding typical parental species. The intermediate mode is composed of the hybrid complex. Although the majority of sampled Kananaskis hybrids fall into the intermediate group, as expected for F1 hybrids, the specimens intermediate between the parental modes and the hybrid mode are of contentious origin. It appears from this analysis that the variance of the typical Kananaskis C. plumbeus and R. cataractae modes (taken as the far left and far right modes, respectively) are greater than the variances of their respective Alberta control distributions.

The shading patterns in the Kananaskis distribution represent the initial field identifications of specimens, prior to the analysis (Fig. 7). All specimens initially identified as "good" C. plumbeus fell within the outer C. plumbeus mode. All specimens initially identified as "good" R. cataractae, except one, fell within the outer R. cataractae mode. A re-examination of this one misidentified R. cataractae specimen confirmed its hybrid appearance. Most specimens, initially identified as putative hybrids, were classified in an intermediate position by the CVA. However, several putative hybrids fell within the outer C. plumbeus

mode. A re-examination of these putative hybrids (with a canonical score  $> -3.0$ ) revealed the absence of any frenum in all specimens, suggesting an initial misidentification.

The 12 characters contributing most to species' separation in this CVA are given in Table 2 (along with their coefficients and constants for the discriminant functions and their F ratios). The remaining 9 characters with F values  $> 1.0$  contributed little to the discrimination and could have been eliminated from the analysis. The frenum was the most discriminating character with a high F value (F = 1048.10). The computer results for this analysis are summarized in Appendix Table 10.

It should be recalled (see Methods) that abundant C. plumbeus were randomly collected, whereas all R. cataractae and all putative hybrids captured were kept in the field collections (specimens were identified in the field as putative hybrids by their possession of a partial frenum). The objection might be raised that the formation of the intermediate peak on the histogram is an artifact of selective collecting of hybrids. It is true that the relative number of typical C. plumbeus in the population is far greater than indicated by the histogram. However, it is unlikely that the distinctiveness of the intermediate hybrid group was exaggerated since only a random sample of the field collections was analyzed. Moreover, the distinctiveness of this group relative to C. plumbeus was likely minimized since possible backcross individuals (in

TABLE 2. The 12 most discriminating characters in a CVA using untransformed data, including frenum. Characters are in order of decreasing importance with F ratios (1, 98 df) for inclusion and coefficients and constant for canonical variables (1st canonical axis).

Rank	Character	F	Coefficient
1	FRE	1048.10	-6.65
2	OL	60.69	30.74
3	IW	41.81	-12.41
4	HWE	14.27	7.88
5	CF	13.24	6.08
6	STA	12.09	-2.29
7	HL	3.18	-3.10
8	CP	2.00	-1.76
9	CPS	1.53	-0.07
10	ML	1.41	9.14
11	SN	1.06	5.88
12	ALS	1.05	0.09
	Constant		-1.97

the direction of C. plumbeus) were collected more rigorously than typical C. plumbeus.

Although the frenum character is an efficient discriminator, its removal from the analysis does not seriously affect the results. The CVA histogram using untransformed data without the frenum character shown in Figure 8, is essentially identical to the histogram in Figure 7. Again, there appears to be greater variance in the modes of the typical Kananaskis parental species compared to the control distributions. The computer results for this analysis are summarized in Appendix Table 11. The 12 most discriminating characters in this analysis are given in Table 3. A comparison of Tables 2 and 3 illustrates that CVA is heavily weighted to a dependence on only a few characters.

The results of a CVA using square-root transformed data, with frenum character in the data set, are given in Figure 9. Transformation of the character data results in increased discrimination between the two species. The means of the two Alberta species' distributions are separated by approximately 23 units of standard deviation. Although an intermediate hybrid mode remains, its mean is 2/ standard deviations to the left of strict intermediacy (toward R. cataractae). This is due to a heavy weighting on the transformed frenum character ( $F = 3893.55$ ). (Table 4). The troughs between the hybrid and parental modes are spread thinner along the first canonical axis than seen previously.

Figure 8. CVA Using Untransformed Data; Without Frenum

Frequency histograms of canonical scores on first canonical axis of specimens from Alberta 'control' group (upper histogram) and Kananaskis group (inverted histogram). Alberta species formed 2 *a priori* known groups; all Kananaskis specimens ran as *a priori* unknown groups. Units on x-axis are in standard deviations; y-axis is number of specimens.

### Untransformed Data, Without Frenum Character

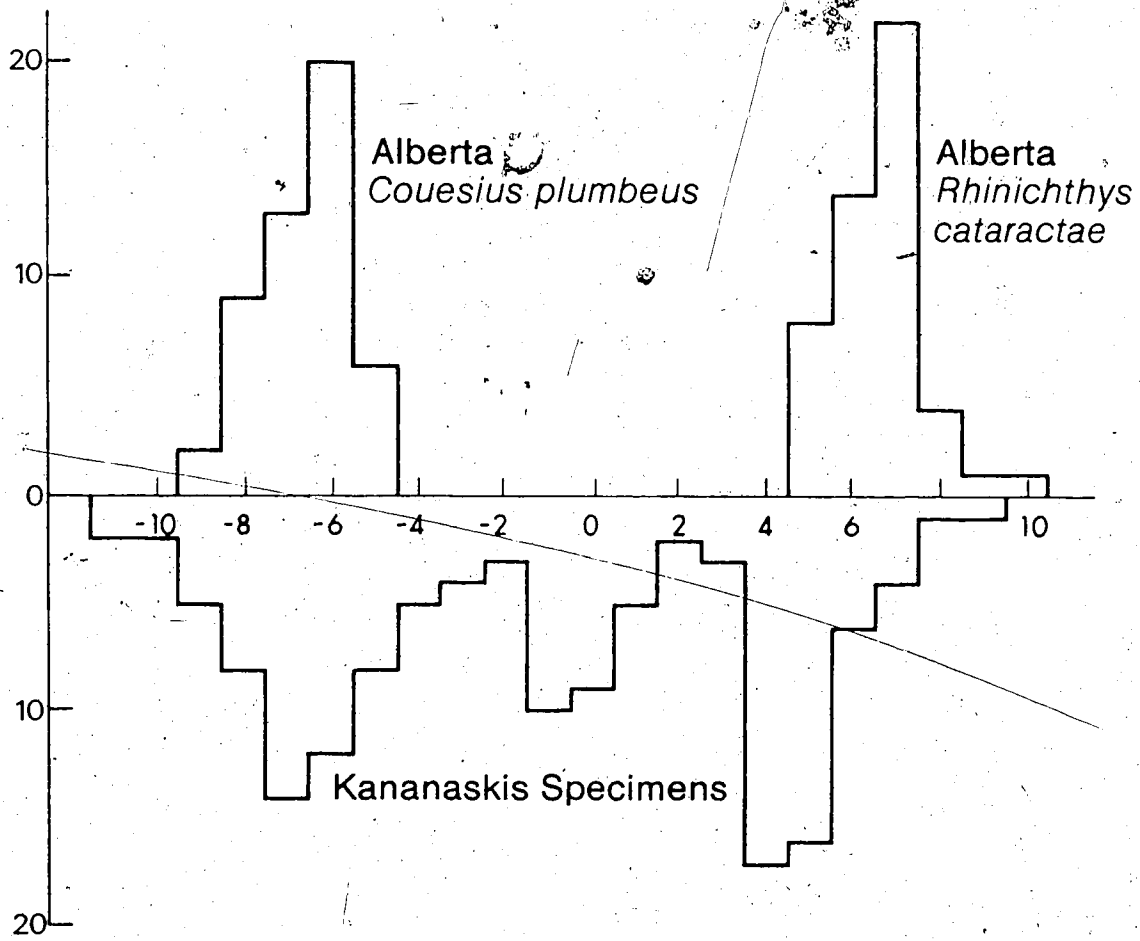




TABLE 3. The 12 most discriminating characters in a CVA using untransformed data, without frenum. Characters are in order of decreasing importance with F ratios (1, 98 df) for inclusion and coefficients and constant for canonical variables (1st canonical axis).

Rank	Character	F	Coefficient
1	IW	682.35	-13.53
2	OL	207.69	32.87
3	CF	13.65	6.93
4	PTA	12.92	0.04
5	HL	8.41	-3.66
6	HWE	8.00	8.88
7	ML	4.95	11.59
8	CP	3.30	-2.07
9	CPS	1.58	-0.08
10	STA	1.52	-2.04
11	MAN	0.84	-4.48
12	HDN	0.51	-2.58
	Constant		-1.06

Figure 9. CVA Using Square-root Transformed Data; Frenum Included

Frequency histograms of canonical scores on first canonical axis of specimens from Alberta 'control' group (upper histogram) and Kananaskis group (inverted histogram). Alberta species formed 2 *a priori* known groups; all Kananaskis specimens ran as *a priori* unknown groups. Units on x-axis are in standard deviations; y-axis is number of specimens.

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## Square-root Transformed Data, Frenum Character Included

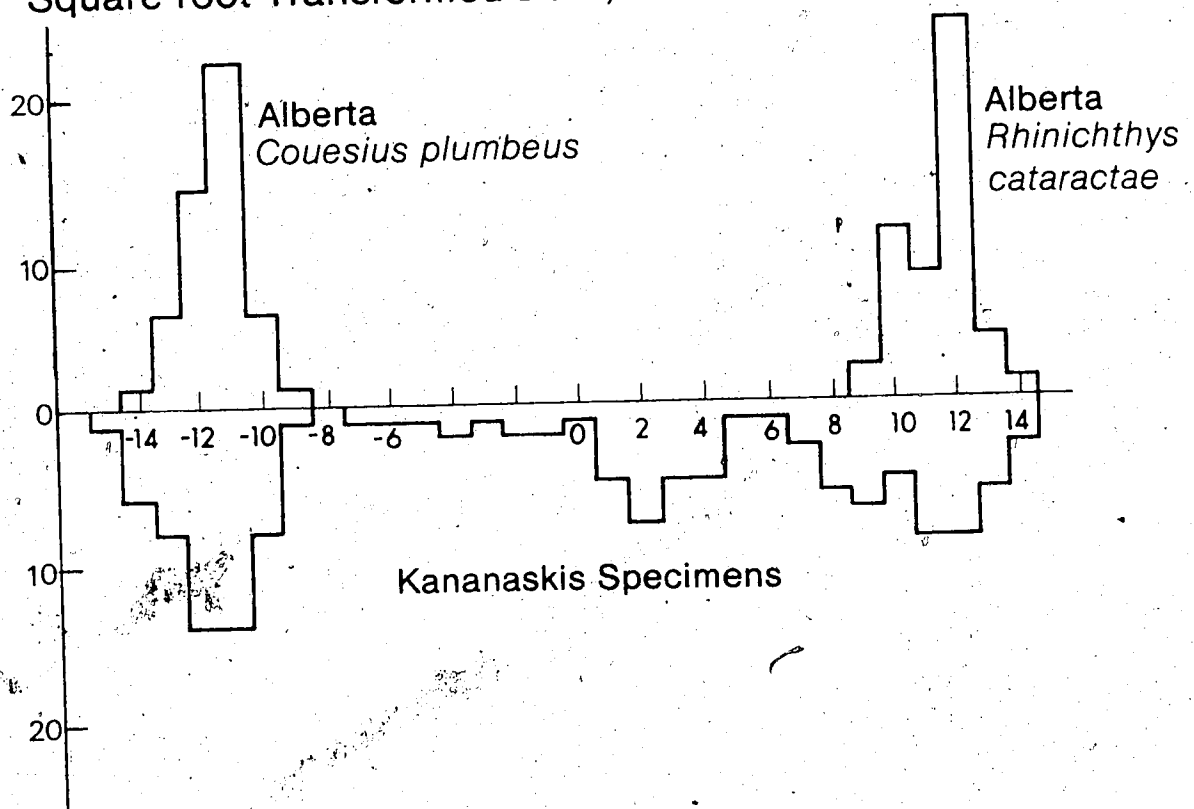


TABLE 4. The 12 most discriminating characters in a CVA using square-root transformed data, including fremum. Characters are in order of decreasing importance with F ratios (1, 98 df) for inclusion and coefficients and constant for canonical variables (1st canonical axis).

Rank	Character	F	Coefficient
1	FRE	3893.55	24.36
2	OL	49.30	-26.72
3	IW	20.49	10.31
4	CF	17.34	-9.33
5	HWE	13.12	-13.82
6	STA	7.94	14.96
7	SOV	3.98	5.82
8	CP	3.10	3.46
9	SN	2.72	-12.63
10	ALS	1.63	-0.13
11	PTA	1.39	-9.24
12	SDO	1.15	7.68
			Constant 6.39

This indicates that there may be specimens which cannot be accounted for as either good parentals or F<sub>1</sub> hybrids.

Unlike the results with untransformed data, the CVA using transformed data does not show as great a difference between the variances of the typical Kananaskis species' modes and the variances of the Alberta control species' distributions. Computer results for the CVA using transformed data are summarized in Appendix Table 12.

Have the two Kananaskis species been modified by this hybridization relative to the 'control' specimens? If it is assumed that the Kananaskis parental modes (Fig. 7) end at the troughs (tails of the mode) next to the hybrid mode, it becomes apparent that: 1) the Kananaskis parental species have greater variances than the 'control' species; and that 2) the means of the Kananaskis species' modes are shifted toward each other, relative to the means of the 'control' species' distributions. This leads to speculation that the cause of this could be backcrossing between hybrids and parentals, i.e., introgression (sensu Mayr, 1969).

#### CVA- VARIANCE TESTS

The differences in variance were investigated in order to determine if this could be a reliable indicator of introgression. It is known that the CVA procedure minimizes the within-group variance of the a priori known group, at the expense of the variance of the unknown group (Pimental, 1976). To determine how this affected the variances, the

known and unknown groups were switched: the assumed Kananaskis parental species (without presumed hybrids) were run as the a priori known group. The results of Variance Test I (Fig. 10) show a far narrower variance for the Kananaskis parentals, relative to the Alberta parentals, than previously seen (Table 5). Thus, the variance of the a priori unknown group cannot be directly compared to the variance of the a priori known group. The difference in variance (Fig. 7) is an artifact of CVA procedures.

What then is an accurate estimate of the Kananaskis species' variance which can be compared to the Alberta species' variance arrived at in the original CVA (Fig. 7)? Are the Kananaskis species' variances significantly different from the Alberta species' variances? The variance for the Kananaskis parentals from Test I are not truly representative because: 1) the Kananaskis group is the a priori unknown group; and because 2) there is little difference between the variances of the Alberta and Kananaskis groups in Test I, even though the Kananaskis variances are being minimized - indicating that the Kananaskis variances are wider than the Test I estimates.

Variance Test II (Fig. 11) was performed in order to improve the variance estimate for the Kananaskis groups. All specimens of the same species were pooled into two groups irrespective of membership in the Alberta or Kananaskis groups. This was done to relate each specimen to all other specimens of the same species irrespective of group

Figure 10. Variance Test I. Untransformed Data; Frenum Included.

Frequency histograms of canonical scores on first canonical axis of presumed pure specimens of Kananaskis species without hybrids (upper histogram) and Alberta 'control' group (inverted histogram). Kananaskis specimens formed 2 *a priori* known groups; Alberta species ran as *a priori* unknowns. Units on x-axis are in standard deviations; y-axis is number of specimens.

### Variance Test I

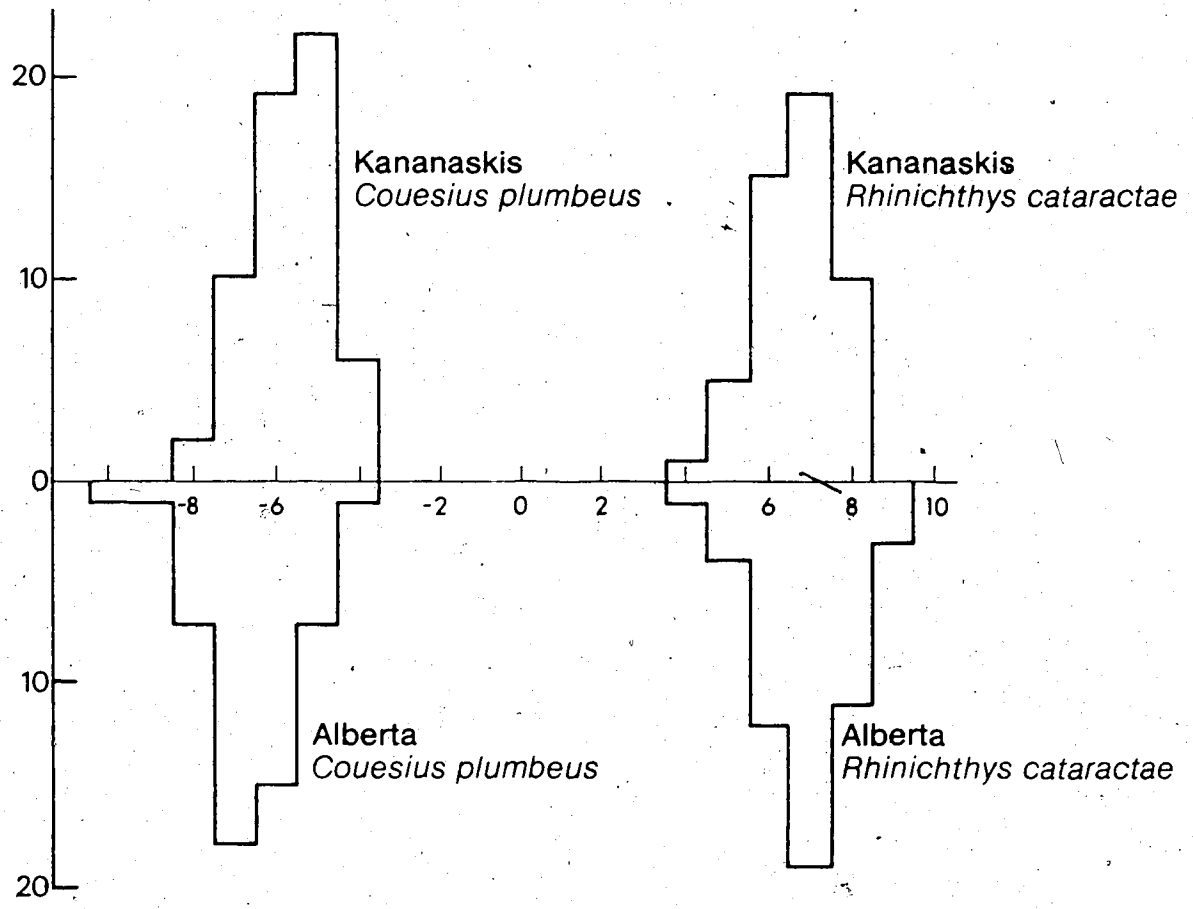




TABLE 5. A comparison of variance estimates for the 2 species from Kananaskis and Alberta groups.

	Kananaskis		Alberta	
	<i>C. plumbeus</i>	<i>R. cataractae</i>	<i>C. plumbeus</i>	<i>R. cataractae</i>
Original CVA (Fig. 7)	3.21	1.95	0.90	1.07
Variance Test I	0.96	1.03	1.27	1.09
Variance Test II	0.92	1.39	0.71	0.89

Tests for significance of differences between Alberta and Kananaskis species' variances,

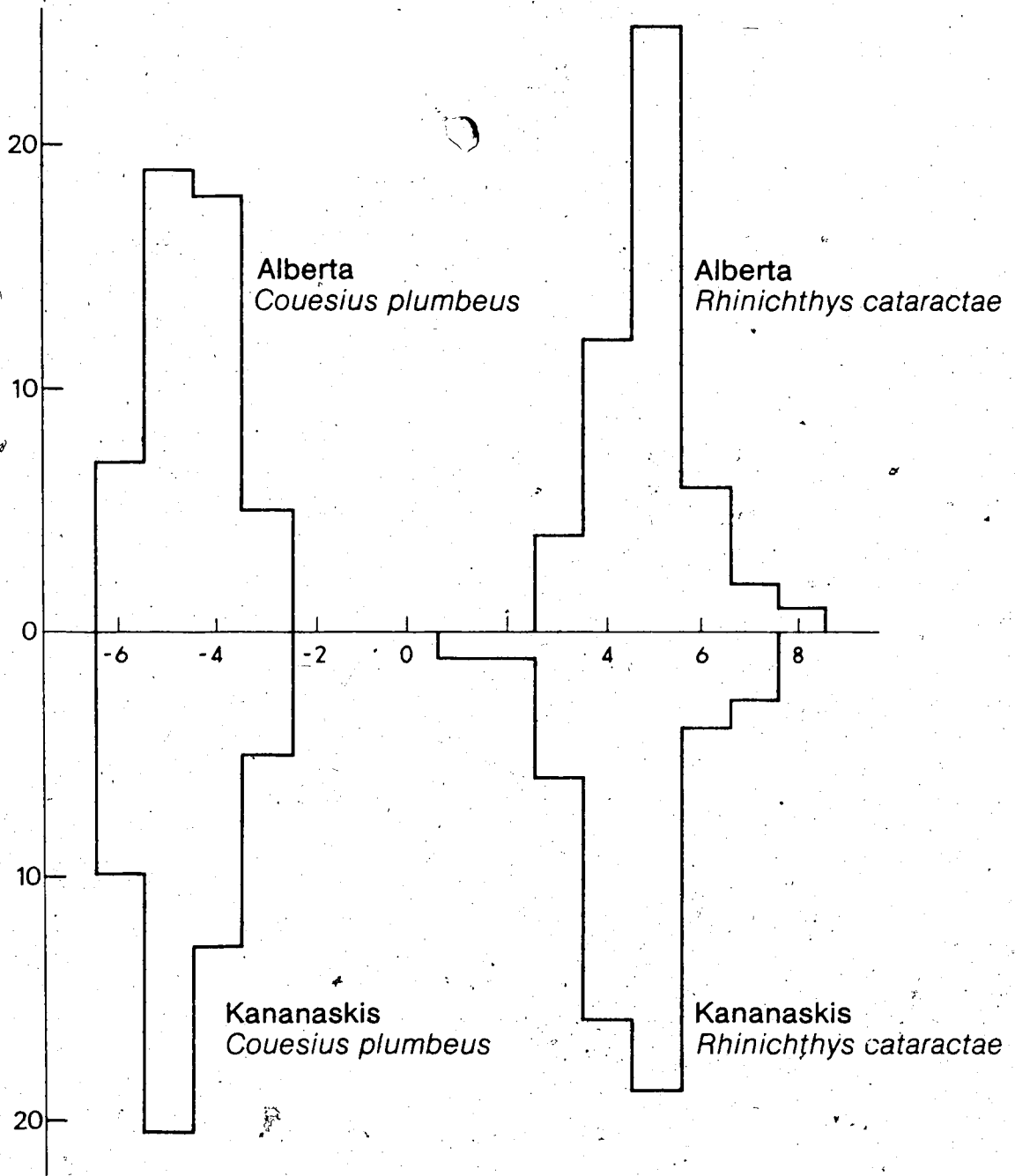
from Test II: *C. plumbeus*  $F = \frac{0.92}{0.71} = 1.30$ ;  $F_{.05 (49, 49)} = 1.67$ ,  $P > 0.05$ .

*R. cataractae*  $F = \frac{1.39}{0.89} = 1.56$ ;  $F_{.05 (49, 49)} = 1.67$ ,  $P > 0.05$ .

Figure 11. Variance Test II. Untransformed Data; Frenum Included.

Frequency histograms of canonical scores on first canonical axis of presumed pure specimens of Kananaskis species without hybrids (upper histogram) and Alberta 'control' group (inverted histogram). Kananaskis and Alberta groups of same species were combined to form 2 *a priori* known groups; there was no unknown group. Alberta and Kananaskis specimens are separated as upper and lower histograms.

# Variance Test II



membership. The two groups were run in a CVA as a priori knowns. No a priori unknowns were included. Once computed, the canonical scores of the Kananaskis group were separated from the Alberta group and plotted (Fig. 11). The variances of the Kananaskis species' distributions were calculated separately from the Alberta species' distributions. The results of Variance Test II show that the variances of the Kananaskis parental groups are not significantly different ( $P > 0.05$ ) from the variances of their respective controls (see Table 5). There does not appear to be a significantly greater variability in the Kananaskis parentals than in the Alberta parentals, which could be ascribed to introgression. It is unsafe to draw conclusions concerning introgression from direct comparisons of the variances of species' canonical score distributions. The CVA must be manipulated in order to obtain comparable variance estimates.

The computer results for both Variance Tests are summarized in Appendix Tables 13 and 14.

#### CVA - COMPARISON OF GROUP MEANS

To avoid the confusion and difficulties associated with performing variance tests, it may be simpler to compare group means in order to determine the presence of introgressive effects. It appears that there has been a significant shift of the Kananaskis R. cataractae group mean away from the control mean (Figs. 7, 8, 9) ( $\chi^2 = 31.8, P < 0.05$ ). The mean of the Kananaskis C. plumbeus group is

similar to the control group mean (Figs. 7, 8, 9) ( $\chi^2 = 13.1, P > 0.05$ ). The shift of Kananaskis R. cataractae toward the hybrid group may indicate: 1) the presence of backcrosses to R. cataractae; or 2) phenotypic changes in R. cataractae due to hybrid fertility, i.e., introgression. Conversely, this would indicate an absence of phenotypic change in Kananaskis C. plumbeus.

#### CVA - PAST COLLECTIONS

The group means of the canonical score distributions for typical C. plumbeus, collected in 1961, 1969, 1973, and 1977, were compared (Fig. 12) to determine if this Kananaskis species had been modified by introgressive hybridization over time. The temporal distributions of canonical scores indicates a significant shift in means in 1961, 1969, C. plumbeus toward R. cataractae and a return shift in 1973, 1977, toward the control group mean. There is a significant difference between the control distribution and the 1961 Kananaskis distribution ( $\chi^2 = 31.0, P < 0.05$ ), no significant difference between the 1961 and 1969 Kananaskis distributions ( $\chi^2 = 3.9, P > 0.05$ ), no significant difference between the 1969 and 1973 Kananaskis distributions ( $\chi^2 = 2.9, P > 0.05$ ), and a significant difference between the 1973 and 1977 Kananaskis distributions ( $\chi^2 = 58.8, P < 0.05$ ). One specimen in the 1969 UAMZ collection is revealed to be of hybrid origin. The computer results for this analysis are given in Appendix

Figure 12. CVA of Past Collections of Kananaskis *Couesius plumbeus*.

Using Square-root Transformed Data; Frenum Included.

Frequency histograms of canonical scores on first canonical axis of specimens from Alberta 'control' group 1977 (upper histogram) and Kananaskis groups, 1961, 1969, 1973, 1977 (inverted histogram). Alberta group formed *a priori* known group; all Kananaskis specimens ran as *a priori* unknowns.

Units on x-axis are in standard deviations; y-axis is number of specimens. Sample size and group centroid are shown for each collection.

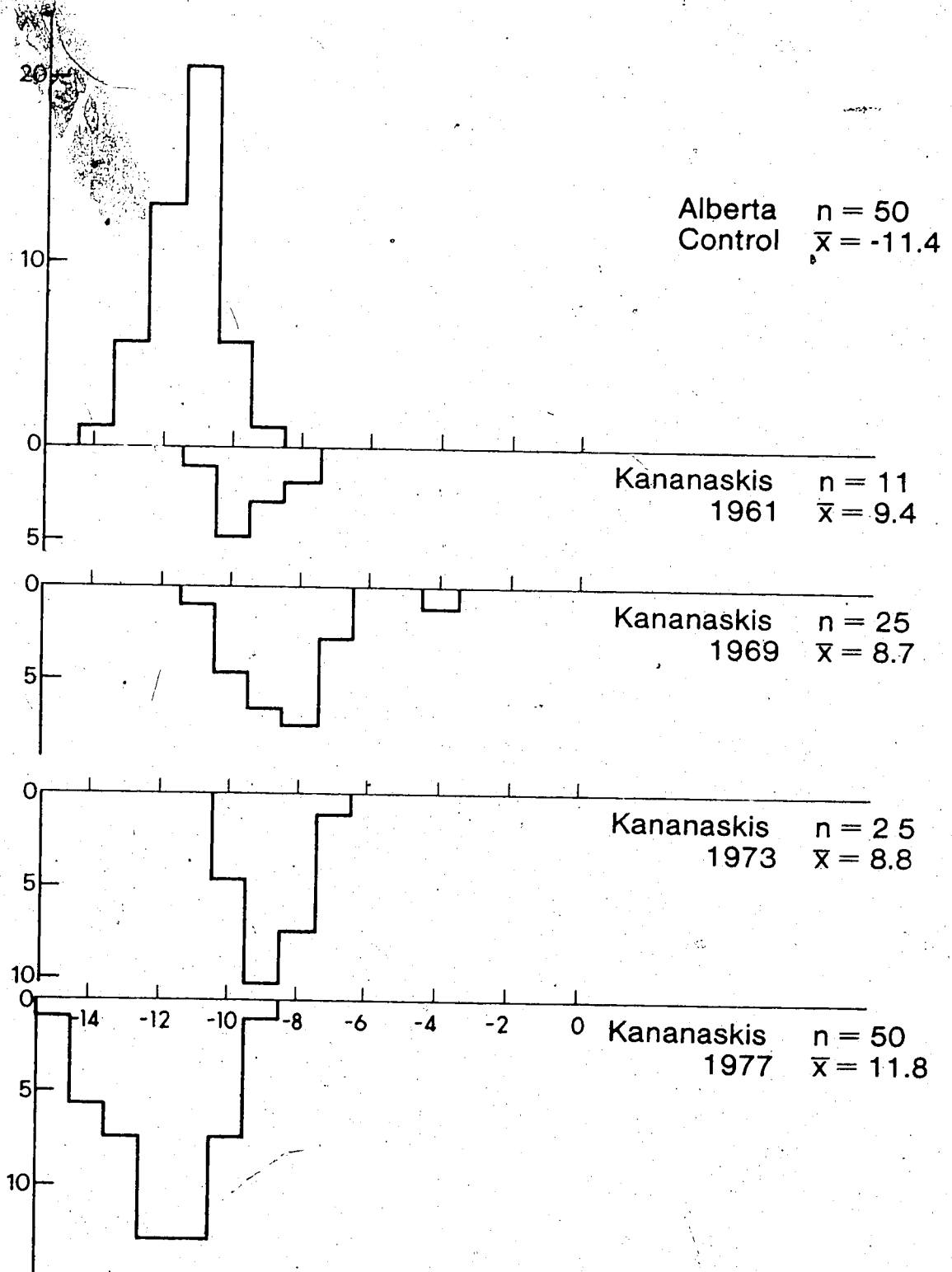


Table 15.

## PCA - INTRA-GROUP VARIATION

It was necessary to determine if either sex or unknown inter-population differences were confounding species' discrimination. To test this, two PCAs were performed: one on Alberta 'control' C. plumbeus; another on Alberta 'control' R. cataractae. The results shown in Figures 13 and 14 demonstrate that there are no subgroups within each species' cluster. An examination of the sex and population membership of each point in the clusters did not reveal any pattern attributable to these variables. PCA was not performed separately on typical individuals of the two Kananaskis species. The reason for this was that CVA results indicate that it is impossible to positively discriminate some individuals of the Kananaskis parental species from variant hybrid individuals on a morphological basis.

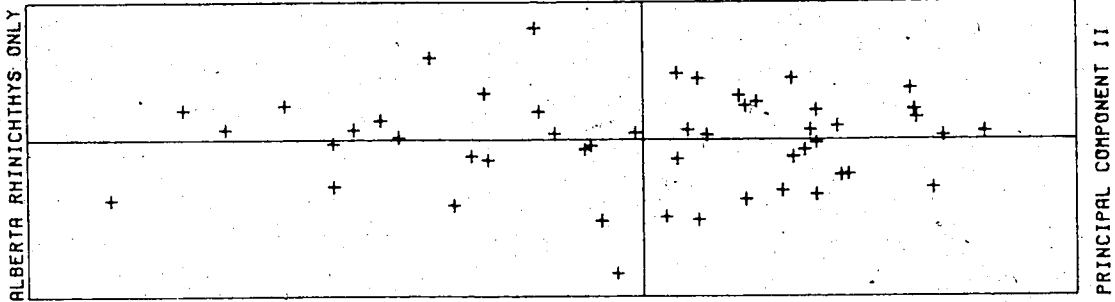
It is unlikely that the inclusion of individuals from Upper and Lower Kananaskis Reservoirs in the Kananaskis groups would increase the within-groups variability and reduce species' discrimination. The previous inclusion of diverse populations in the Alberta groups did not result in such subgroups reducing species' separation. In addition, the two Kananaskis populations analyzed (Upper and Lower Kananaskis Reservoirs) have a probably similar and recent Bow River origin. Computer results for these analyses are summarized in Appendix Tables 16 and 17.



Figure 13. PCA of Alberta *Rhinichthys cataratae* alone. Using Square-root Transformed Data; Frenum Included.

- A. Plot of first and second principal component scores.  
Component I = 77.5% of variation; Component II = 6.2% of variation.
- B. Plot of second and third principal component scores.  
Component III = 4.9% of variation.

PRINCIPAL COMPONENT I



PRINCIPAL COMPONENT II

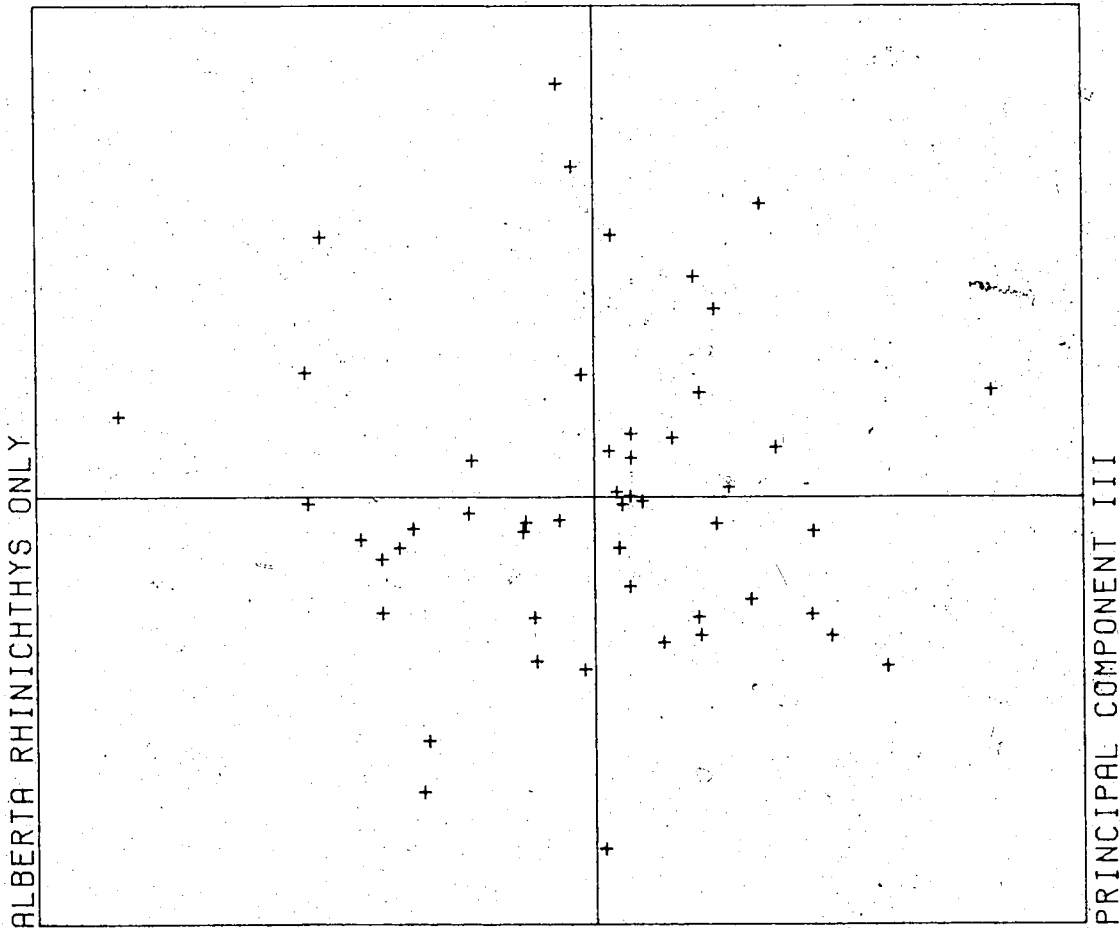


Figure 14. PCA of *Alphacoccus Couesius plumbeus* alone. Using Square-root Transform. Frenum Included.

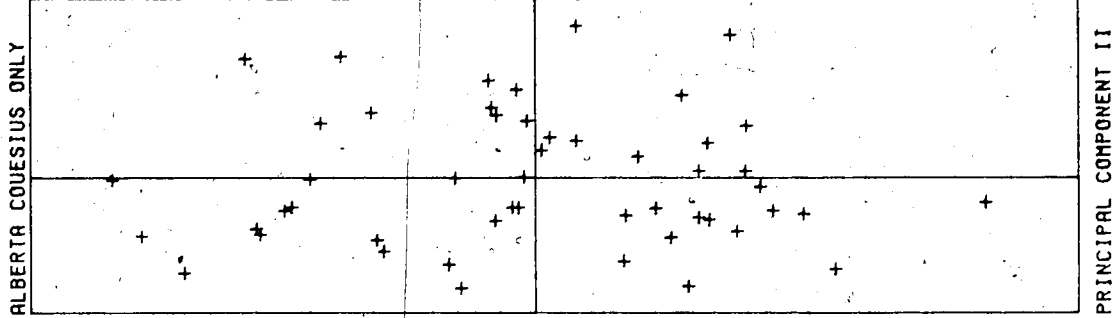
A. Plot of first and second principal component scores.

Component I = 72.0% of variation; Component II = 6.6% of variation.

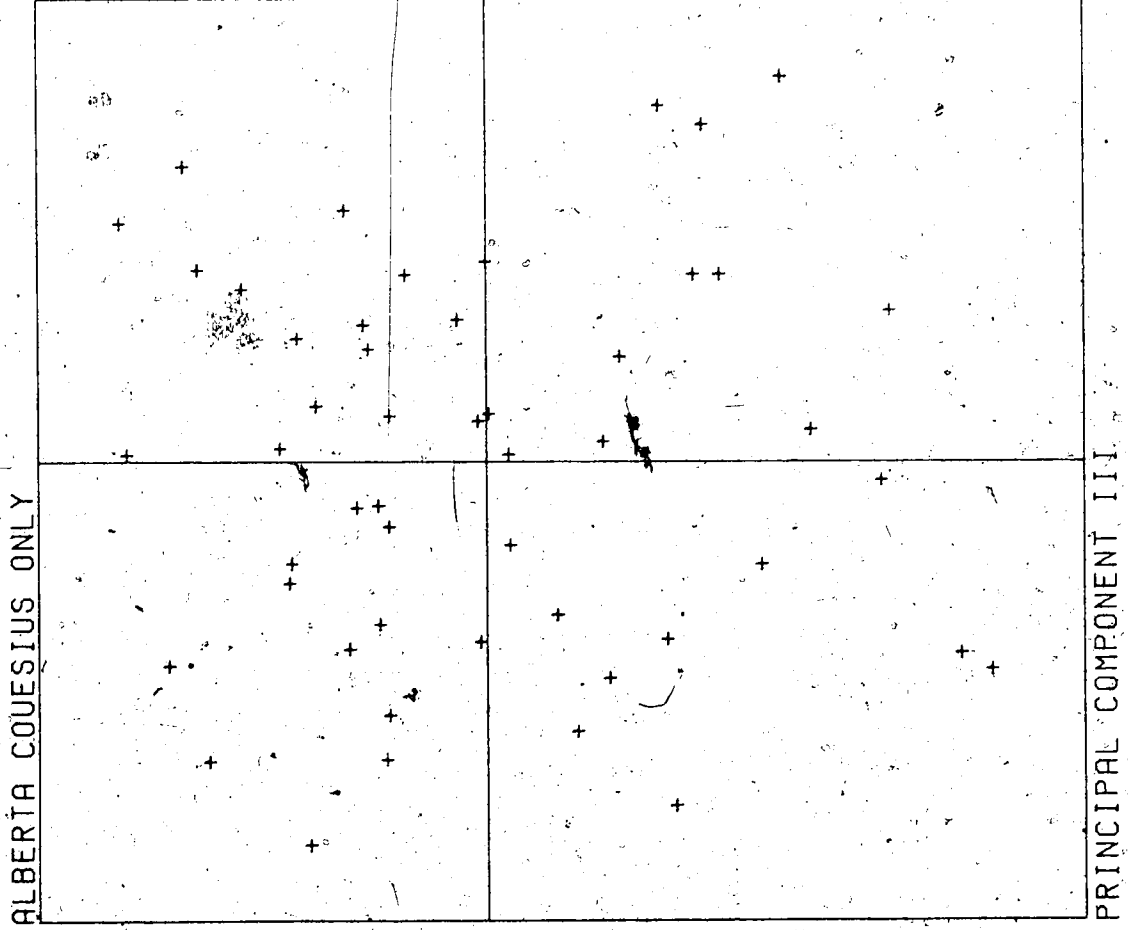
B. Plot of second and third principal component scores.

Component III = 5.2% of variation.

PRINCIPAL COMPONENT I



PRINCIPAL COMPONENT II



A listing of actual character values for all specimens used in all above multivariate statistical analyses is given in Appendix Tables 18-23.

#### DISCUSSION

Central to most modern studies of fish hybridization has been the desire to reach a conclusion on the fertility of hybrids from an analysis of morphological data. This stems from the assertion by Hubbs (1955) that not only are F1 hybrids intermediate between their parental species, except for some features reflecting hybrid vigor, but they are also of the same order of variability as are the parental species. Subsequently, when wild-caught hybrids showed increased degrees of morphological variability relative to the parental species, several types of hybrids were reported to be fertile without experimental evidence (Hubbs and Strawn, 1957). More recently, Schueler and Rising (1976) have stated that when individuals representing F2 or backcross generations are present, the hybrid population is found to be more variable than the parental populations.

The assumption of low F1 hybrid morphological variability has been proven false when the morphology of lab-reared F1 hybrids is compared to the morphology of lab-reared controls of the parental species. Hubbs and Strawn (1957) found the artificial F1 hybrids between *Etheostoma spectabile* and *Percina caprodes*, from the San Gabriel River, were more variable in all characters analyzed

than their parental controls. Greenfield and Greenfield (1972) found that artificial F1 hybrids between Gila orcutti and Hesperoleucus symmetricus were not strictly intermediate between the parental species; rather the distribution of hybrid index values for known hybrids filled the intervening space between the parental species' distributions. Neff (personal communication, 1977) analyzed numerous morphological characters from 2 separate groups of artificial F1 hybrids (Lepomis cyanella x L. macrochirus and Notropus whipplei x N. spilopterus). Using CVA and PCA, she found greater variation in the known F1 hybrids than in the parental groups. This would preclude the confident identification of backcrosses (if they were present) from hybrid individuals. Without knowing the genetic history of the hybrids in the above cases, one would have identified non-intermediates as backcrosses or F2 individuals under the erroneous assumption of low F1 hybrid variability.

It is likewise not possible to decide on C. plumbeus x R. cataractae hybrid fertility or to positively identify backcross individuals from the multivariate analyses of Kananaskis cyprinids performed in this study. Non-intermediate, putative hybrid individuals in the trough region of the Kananaskis CVA distribution could represent F2 hybrids, variant F1 hybrids, or backcross individuals.

Smith (1973) using PCA found lone hybrids occupying the area between strict intermediacy and one parental type. These were identified as backcross individuals. Without

knowing the variance of the F1 hybrids, such a conclusion is uncertain.

It may be possible to determine F1 hybrid variance from a morphological study without resorting to the study of artificial F1 hybrids. This may be possible given a large phenotypic distance between parental distributions, and a large hybrid sample forming one intermediate distribution and two smaller non-intermediate distributions toward the parental types. In this idealized case, the variability of the intermediate F1 hybrid group would be established and a backcross interpretation would be warranted for non-intermediate individuals. The results of my study do not reveal such backcross distribution peaks. Moreover, the F1 ~~cannot~~ cannot be known with such a continuous phenotypic bridging. In a study where the hybrids bridged the gap to one species and not to the other, a backcrossing interpretation would be warranted, without the need to show F1 variance.

This continuous distribution of hybrids bridging C. plumbeus and P. cataractae indicates that it is not always possible to positively identify 'pure' individuals of the Kananaskis parental species. This strengthens the need to objectively classify all individuals in an area of hybridization relative to known control specimens from areas where hybridization is not known to occur.

It is also possible to use the control specimens as an objective standard against which to compare the morphology

of specimens collected over a long time period in a hybridizing area. If the area is not a stable hybrid zone (sensu Moore, 1977) the following alternatives of evolutionary change may be observed over time: 1) a trend toward fusion of the species by introgression - recognized by a progressive increase in hybrid frequency, skewing of parental distributions toward each other, or an increasing variance of parental distributions relative to the control distribution; 2) a strengthening of species' isolating mechanisms and character displacement - recognized by a progressive decrease in hybrid frequency, skewing of parental distributions away from each other, and no increase in parental variance; 3) extirpation of one species by competitive exclusion and hybridization; or 4) continued low level hybridization as long as the environment remains modified.

Jones (1973) observed the effects of thirty years of hybridization on the toad species, Bufo americanus, and Bufo woodhousii, in this manner. Hybridization with introgression can be best recognized temporally relative to known control groups. This would appear to be a rewarding approach for studies of fish hybridization, provided adequate past collections are available. In my study there were only sufficient C. plumbeus in past collections for morphological comparisons.

Three approaches were used to determine if the Kananaskis populations had changed due to hybridization.



First, the greater variances of the Kananaskis groups compared to the variances of the control groups suggested the effects of introgression. However, using the Variance Tests (I and II), this difference was found to be a property of CVA which obscured the equality of variances between the Kananaskis and Alberta groups. There has been neither an increase nor a decrease in population variance, evidence that neither introgression nor character displacement has occurred. Comparisons of variances (of CVA distributions) is not a preferred indicator of morphological change due to introgression.

The second approach was to compare the position of the Kananaskis group mean (1977) to that of the Alberta group mean, separately for each species. The analysis revealed no difference in the positions of the means for C. plumbeus, but a definite shift in the Kananaskis R. cataractae group mean away from the control mean toward the hybrid mode. This evidence suggests that Kananaskis R. cataractae has been morphologically changed due to hybridization.

The third approach was to determine if the modes of the two Kananaskis parental species were moving toward each other relative to the controls (introgression) or receding from each other (character displacement). The mode of canonical scores for Kananaskis C. plumbeus (presumably pure parental types) does reveal a shift in 1961, and 1969 means toward R. cataractae away from the control C. plumbeus distribution. Subsequently in 1973, 1977, a shift back

toward the control group mean occurred. The shift in means may indicate the initial effects of introgression followed by character displacement. However, this seems implausible for such a short time span. Without investigating the fertility potential of the hybrids by experimental means, this remains speculation. It is also reasonable to attribute the morphological shifts to selection by some unknown environmental changes through the intervening 17 year period.

In summary, there is no clear evidence of introgressive effects of hybridization upon Kananaskis C. plumbeus. However, Kananaskis R. cataractae does show a morphological change suggestive of introgression, or the presence of backcrosses. It is also possible for this difference in means to be a result of population variation due to the Founder Effect (sensu Mayr, 1969). Further evidence, especially from breeding tests, is required to test this hypothesis. Apart from this slight change, there does not appear to be a significant trend toward breakdown of the species' integrity.

Both R. cataractae and individuals of hybrid origin have remained at low frequencies during the past 17 years. Although the rate of hybridization has remained at a low level, there is some evidence of a trend toward extirpation of R. cataractae from the reservoirs (alternatives 3 and 4 from above).

This study has assessed the relative importance of the

morphological characters of C. plumbeus and R. cataractae. Nelson's (1966) finding that the best hybrid discriminators are from the head , i.e., frenum width, snout length, isthmus width, and orbit length has been confirmed. Future workers may find the canonical coefficients and constants given in Tables 2-4 to be valuable in classifying putative hybrid specimens.

## VII. ELECTROPHORETIC IDENTIFICATION OF HYBRIDS

### INTRODUCTION

Electrophoretic results can provide a second set of data to test the hypothesis of hybrid origin of certain specimens based on morphological criteria. The advantage of electrophoresis over a purely morphological analysis is that the genetic contributions from each parent may be detected in a single individual.

Nyman (1970) has listed all possible types of protein patterns found in fish hybrids, reflecting simple codominant expression of allelic genes. Hybrid protein patterns can be: 1) identical to one or the other parental species; 2) a complete summation of the two species; 3) a partial summation; or 4) the hybrid pattern and the two parental species' patterns can all be identical. For a protein to be a useful discriminator of hybrids, it must be present in a high frequency as a variant form in each parental species. The hybrid then, inherits a summation of the parental banding patterns (Reinitz, 1977). The confirmation of "hybridity" in this study was an empirical search for such hybrid specific biochemical markers.

### METHODS

Adult C. plumbeus, and R. cataractae and their putative, wild hybrids were collected from Lower Kananaskis Reservoir during the summer of 1978 for electrophoretic examination. Specimens from outside the system were not

examined. Adult fish of comparable age were chosen in order to exclude electrophoretic variation due to developmental differences.

To ensure that individuals representing the parental species were 'pure', the same 21 morphological characters used for PCA and CVA were measured, and the canonical scores calculated. Individuals with canonical scores within one standard deviation unit of the control parental means (from the multivariate morphological analysis Fig. 7) were used to represent 'pure' parental types.

Only hybrids with intermediate canonical scores (in the presumed F1 portion of the Kananaskis distribution on the first canonical axis, Fig. 7) were electrophoresed. Putative hybrids, with canonical scores intermediate between the parental species and the presumed F1 mode (possible backcrosses) were unfortunately not available in either a fresh or frozen state necessary for electrophoresis. The progeny of the experimental crosses had not reached sufficient size to permit electrophoretic examination.

Protein extracts were obtained from blood sera and skeletal muscle of the same specimens. Blood sera were obtained from live fish in the laboratory by severing the caudal peduncle, withdrawing the blood into heparinized capillary tubes, and spinning down the corpuscles in a Hematocrit centrifuge for 4 min.. The corpuscles were discarded and the pure sera frozen at  $-5.0^{\circ}\text{C}$ . The fish were then frozen.

Skeletal muscle samples from all frozen fish were electrophoresed within six weeks of collection. Fish skeletal muscle, frozen more than a few days, has been found to be readily extractable, and muscle proteins quite stable under frozen storage (Utter *et al.*, 1974).

One cc of the frozen dorsal musculature from each specimen was ground in a glass tissue grinder with an equivalent volume of distilled water. The creamy muscle homogenate was then centrifuged for 15 min. at X20,000 g in a Dupont Ultracentrifuge. The resulting clear supernatant was stored frozen at -5.0°C. Frozen samples of this supernatant gave identical electrophoretic patterns to freshly prepared ones and they were used as convenience dictated.

The number of typical parental specimens and hybrid specimens used for the electrophoresis of the two tissue types is given in Table 6.

Vertical polyacrylamide disc gel electrophoresis was performed at room temperature with a Buchler power supply (set on constant current) using the techniques described by Davis (1964) and Smith (1969). Seven percent polyacrylamide gels were made without sample or spacer gels according to a recipe modified from Clarke (1964) (refer to recipe in Appendix). A cold, continuous tris-glycine buffer (pH 8.3) was used for the electrode buffer in order to minimize column heating.

It was found experimentally that 8  $\mu$ l was the optimal

TABLE 6. List of proteins examined in specific tissues. Numbers of fish specimens analysed for each protein is given. R - R. cataractae; C - C. plumbeus; H - C. plumbeus x R. cataractae hybrid. LDH: lactate dehydrogenase; GPD: glycerophosphate dehydrogenase; MDH: malate dehydrogenase.

Proteins	Serum	Muscle	Replicates		
LDH		X	15C	9R	3H
$\alpha$ GPD		X	8C	8R	2H
MDH		X	3C	2R	2H
Serum protein	X		14C	12R	5H
Muscle protein		X	14C	12R	5H

amount of extract of both tissue types for electrophoresis. An equal volume of a 40% sucrose solution was mixed with the sample extract. The entire solution was then layered between the top of the gel surface and the less dense buffer above. All samples were applied within 5 min..

One ml of 0.001 percent Bromophenyl Blue in water was added to the upper reservoir as a tracking dye. Simultaneous electrophoresis of 9 gels was begun with a current of 1.5 mA/tube for the first 10 min. to prevent undue diffusion of the sample. This was increased to 4.0 mA/tube for the remainder of the run once the tracking dye was 2 mm into the gel. Electrophoresis was completed in approximately 75 min. when the tracking dye had reached the end of the gel. Only the anodic portion of the samples was studied.

Gels were stained with amido black 10B (1 gm in 100 ml of 7% acetic acid) for one hour to discern the general

protein patterns of both the sera and muscle extracts from the two species and their hybrid. Destaining was accomplished in the same apparatus using a 7% acetic acid wash solution with a current of 10 mA/tube. Destaining was complete in 60-90 min..

Specific enzymes, lactate dehydrogenase (LDH),  $\alpha$ -glycerophosphate dehydrogenase ( $\alpha$ -GPD) and malate dehydrogenase (MDH) were visualized using the staining procedures of Shaw and Prasad (1970). However, the gels used to separate these enzymes were not run with the recommended electrode buffers. Rather, the gel pH was adjusted closer to the stain pH by immersing the gels in the appropriate stain buffer for 15 min. prior to staining.

Gels were stored in test tubes containing 7% acetic acid and were photographed (Kodak Plus-X film) directly in the destaining tubes with no liquid interface using diffused back lighting.

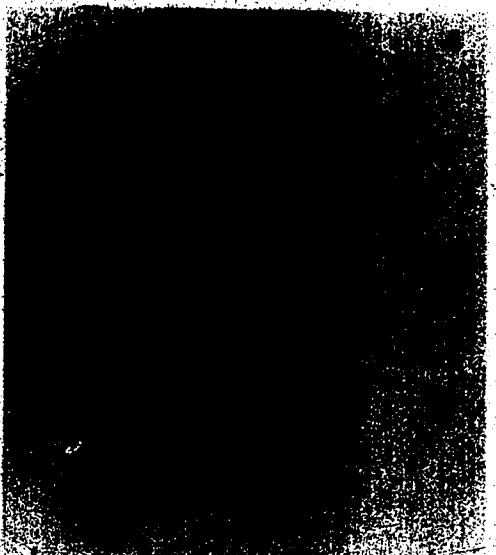
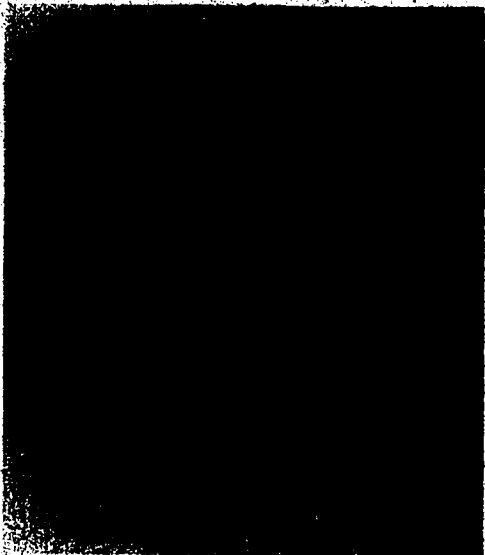
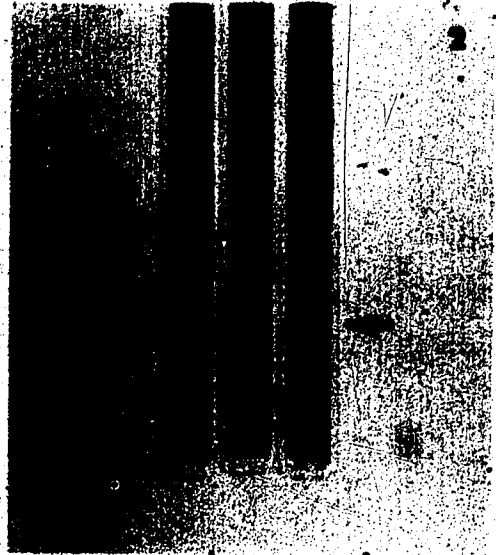
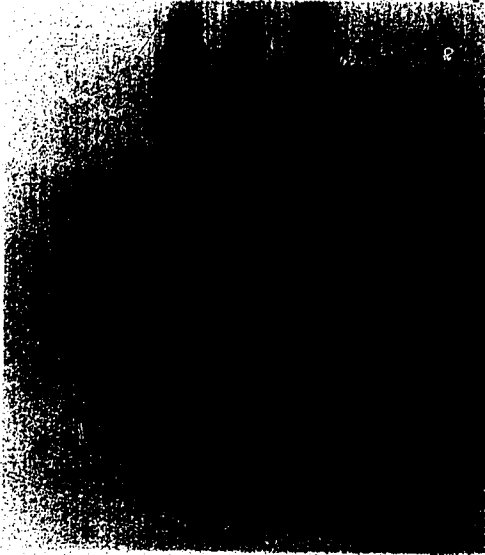
## RESULTS

The polymorphic banding pattern of muscle LDH was identical in all individuals of each species and hybrids that were tested (Fig. 15, #1). The appearance of three muscle LDH isozyme bands in this study is contrary to the results of Clayton and Gee (1969) who found five muscle LDH isozyme bands in both Rhinichthys atratulus and P. cataractae collected in Manitoba. The species were distinguished by the occurrence of a variant muscle LDH in



Figure 15. Acrylamide gel electrophoretic patterns of muscle and serum tissues from *Rhinichthys cataractae* (R.c.), *Couesius plumbeus* (C.p.) and their F. hybrid (CxR) sampled from Lower Kananaskis Reservoir, 1978.

- 1 - muscle myogen extract stained for lactate dehydrogenase (LDH)
  - 2 - serum proteins stained with amido black, a general protein stain. Arrows indicate the 2 variant forms of prealbumen.
  - 3 - muscle myogen extract stained for  $\alpha$  glycerophosphate dehydrogenase (GPD)
  - 4 - muscle myogen extract stained with amido black, a general protein stain.
- Anode is towards the bottom.



R. cataractae. Although our techniques differed, it is likely that my results are misleading. Given a longer electrophoretic running time, the large, most-cathodal band may have resolved into three bands of differing mobility, thereby producing a five band pattern. This was not tested in this study.

Muscle GPD (Fig. 15, #3) was polymorphic in each parental species and the F1 hybrids. Each possessed identical variant forms. Muscle MDH was expressed as a single nonvariant band with equivalent electrophoretic mobilities in each parental species and their F1 hybrids.

Muscle myogen phenotypes have been touted as an efficient method to identify hybrid individuals between other species (Tsuyuki and Roberts, 1965; Aspinwall and Tsuyuki, 1968). This was not the case in this study. All individuals of each species and all hybrids had identical three band patterns (Fig. 15, #4) for muscle extract visualized with amido black. There was some variation in the faint minor bands between the species but these were not reproducible.

The banding patterns of serum proteins for the two species differ in the possession of dissimilar, fast moving prealbumen bands (Fig. 15, #2). All specimens of R. cataractae tested, were characterized by the front band of slower anodal mobility, whereas C. plumbeus was characterized by a faster variant. All 'pure' F1 hybrids tested, were characterized by the possession of both bands,

i.e., a summation of the parental bands. These bands clearly defined a difference between the species. Although there are probably similar differences in the slowest bands, these were not investigated. The differences in banding patterns were independent of sex and were repeatable with several subsamplings of the same tissue of one individual. Electrophoretic phenotypes were not observed in the major bands of the two species' serum protein patterns.

Serum proteins appear to be influenced by environmental factors e.g., diet and physiology. It is unlikely that this could be a confounding variable in this study since the fish were sampled from the same reservoir, at the same time of the year, and presumably matured under the same conditions.

#### DISCUSSION

Electrophoretic serum protein analysis indicates that putative C. plumbeus x R. cataractae hybrids possess two prealbumen bands. These correspond in electrophoretic mobility to the single prealbumen bands of each of the parental species. This substantiates the morphological and reproductive evidence of hybridization between R. cataractae and C. plumbeus.

Different hybrid specific protein patterns have been similarly described in only two other North American cyprinid hybrids: redbreasted shiner (Richardsonius balteatus) x peamouth chub (Mylocheilus caurinus) (Aspinwall and Tsuyuki, 1968), and longnose dace (Rhinichthys cataractae) x

blacknose dace (R. atratulus) (Clayton and Gee, 1969).

Any R. cataractae, C. plumbeus or F1 hybrid specimens from Lower Kananaskis Reservoir could be classified as such solely by their serum phenotype. It would seem reasonable to assume that this is true for other populations of these species. Thus it should be possible to test the purity of sympatric C. plumbeus and R. cataractae populations and to confirm F1 "hybridity" by the use of this single biochemical character.

Nyman (1970) compared the efficiencies of electrophoretic and morphological techniques in detecting natural fish hybrids. He found that electrophoresis was more reliable in identifying F1 hybrids, but that the two methods were equally inefficient in identifying F2 hybrids. This is a result of the almost complete dominance of one parent's genes over the other, in the F2. However, Nyman did find 4 out of 32 protein systems in F2 hybrids (Salmo salar x S. trutta) possessing distinct banding patterns. A thorough electrophoretic survey should be undertaken by future workers in order to test the hypothesis of introgression between C. plumbeus and R. cataractae in Lower Kananaskis Reservoir.

## VIII. REPRODUCTIVE ISOLATING MECHANISMS

### GENERAL INTRODUCTION

Hybridization results from the dissolution of the reproductive isolating mechanisms which formerly preserved the integrity of the species' gene pools. An understanding of which isolating mechanisms are inoperative can suggest the environmental factors which facilitated the hybridization.

Reproductive isolation between sympatric C. plumbeus and R. cataractae populations has not been previously studied. In fact, published material on the breeding biology of these species is scant. The reproductive habits of R. cataractae have been comprehensively described only from Manitoba (Bartnik, 1970). The most comprehensive report for C. plumbeus is from Saskatchewan (Brown, 1969). However, it should be noted that according to Wells (1978), the Kananaskis C. plumbeus are of the 'dissimilis' morphological form while Brown's population consisted of the 'plumbeus' form. Whether there are significant biological differences between these forms is not presently known.

This section of the study investigates the effectiveness of the species' pre mating and post mating isolating mechanisms through ecological field studies and artificial crossing experiments. Ecological and life history data for the Kananaskis cyprinids are presented for the first time.

## STUDY OF PREMATING ISOLATION

### INTRODUCTION

Premating isolating mechanisms prevent the wastage of gametes and are thus more efficient than postmating isolating mechanisms. Premating isolation involves the reduction of contact between the species (i.e., spatially and temporally) and the reduction of interspecific mating by species' specific behaviors. The latter, ethological isolation, has been found to be generally more important than the other isolating mechanisms (Littlejohn, 1969).

The purpose of this section is to determine the following biological characteristics: 1) extent of differences in spawning habitat and spawning time between the two species; and 2) degree of interspecific mate preference.

### METHODS

#### Temporal Isolation

The time of spawning for each of the parental species was followed using three different methods. The first method from Nelson (1968) entailed making a qualitative judgement on the degree of female gonad development. The following qualitative criteria were used to describe female gonad maturity:

ripe - eggs near maximal size; cream colored; extruded only under firm finger pressure.

fully ripe - eggs at maximal size; golden colored;

extruded spontaneously due to handling by experimenter.

spent - few or no golden eggs extruded from body cavity; abdomen visibly sunken.

The peak time in which fully ripe females were present in the Lower Kananaskis Reservoir was compared between the two species.

The second method was quantitative. The spawning period for each species was followed by determining the percentage contribution of female gonads to body weight. Total body weight and gonad weight (both sides) were measured fresh to the nearest 0.01 gm. These gonadosomatic indices ( $\frac{\% \text{ gonad weight}}{\text{total body weight}}$ ) of each species were pooled for sampling dates and plotted over the spawning period.

The third method was also quantitative. This involved measuring the mean diameter of 10 ova per sample female using needle point dial calipers under magnification (read to the nearest 0.01 mm). The grand mean of ova diameters from all sample females was graphed over time.

Fish sampled from three different locations on Lower Kananaskis Reservoir were used for all three methods and were included in the graphic analysis, irrespective of origin. The three locations were: Interlakes Raceway, mouth of Boulton Creek, and Common Bay. The purpose of pooling locations was to generate a large sample size, and to give a general picture of the spawning period for the entire Lower Reservoir, without specific site variability. Time



limitations prevented an in-depth study of spawning periods on Upper Kananaskis Reservoir. The occurrence of diel (temporal) isolation was not investigated.

#### Habitat Isolation

The concurrent abundance of the two species and their putative hybrids at the sampling sites was monitored using all capture techniques throughout the spawning period. Catch data were reviewed to see if there was habitat segregation by depth, current or substrate.

#### Behavioral Isolation

Several ripe males and females of each parental species, captured between May 26 and June 5, 1978, were returned alive to the laboratory and were kept in separate tanks. On June 6, 1978, the following reciprocal, interspecific matches were established for observations of behavioral interactions and mating:

Tank A C. plumbeus - 4 females

R. cataractae - 2 males

Tank B C. plumbeus - 3 males

R. cataractae - 2 females

Two ripe male hybrids captured on June 7, 8, 1978, were put into the same aquaria with three ripe female C. plumbeus on June 10, 1978:

Tank C R. cataractae x C. plumbeus hybrids 2  
males

C. plumbeus - 3 females

All aquaria were kept in a controlled environment chamber which replicated the reservoir's light and temperature regime. The three 20 l tanks were started at 10°C on June 6, 1978, and increased to 14°C by June 15, 1978, (similar to the temperature rise in the reservoir). The photoperiod was a constant 15.5 hours of daylight (0600-2130 hr). Throughout the holding period, fish were fed TetraMin flakes and frozen brine shrimp.

Descriptions of breeding colors and tubercle patterns were made for each sex of the two species immediately after capture, and upon return to the laboratory. Color photographs were taken.

## RESULTS

### Temporal Isolation

Spawning at different times of the year constitutes an important deterrent to interspecific breeding in sympatric fish species. At present, there exist only a few records of spawning times for R. cataractae and C. plumbeus from widespread localities across Canada (Scott and Crossman, 1973). What are the spawning times of these species in Lower Kananaskis Reservoir? To what extent is temporal isolation operative?

Temperature appears to be the single, most important factor inducing spawning in these two species, as is true for most temperate, freshwater fishes. Ahsan (1966) showed

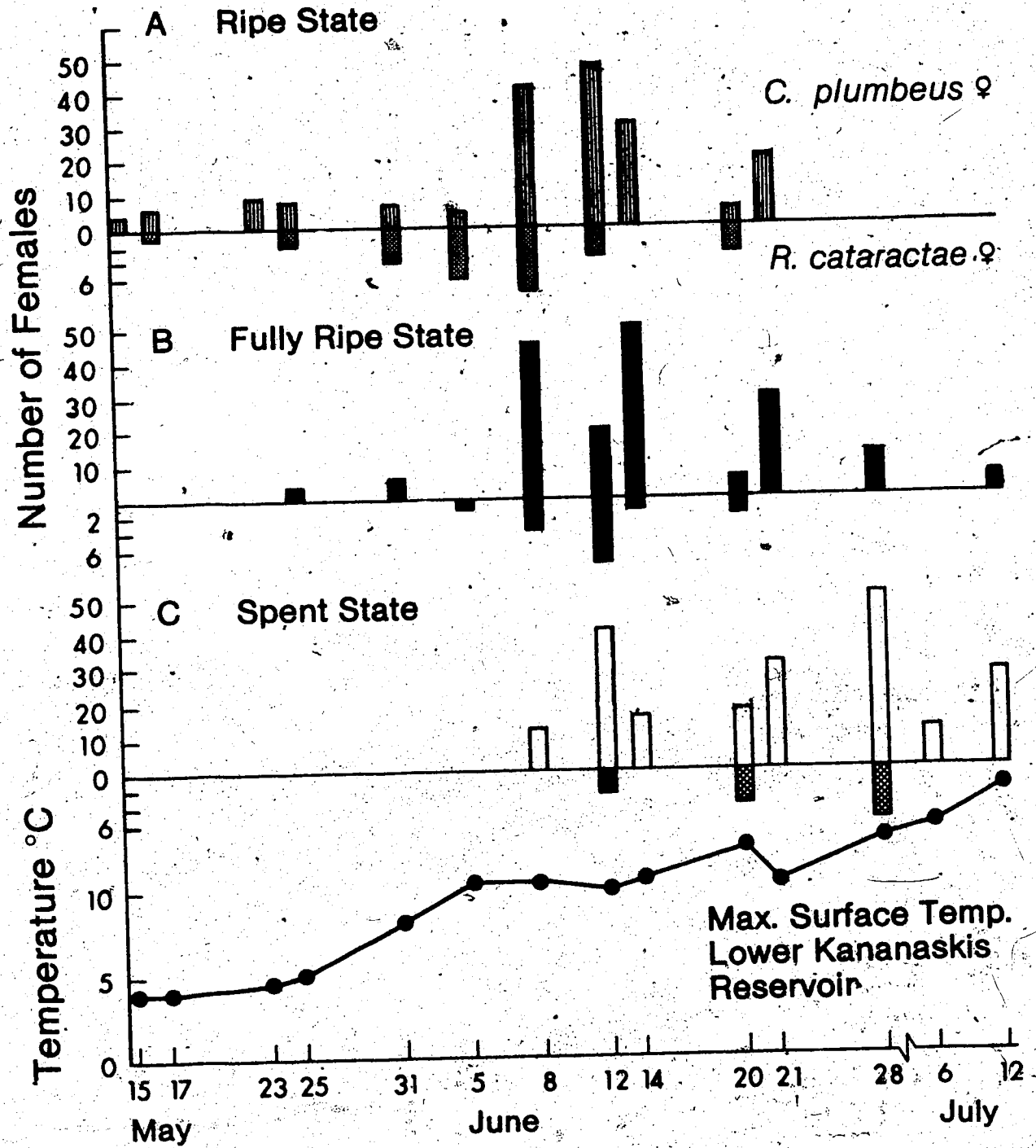
that temperature was the dominant regulator of the spermatogenic activity of male C. plumbeus. Females, judged to be in a fully ripe state, were first collected in late May and early June after the rapid increase in surface water temperature from 5°C, May 25 to 10°C, June 4, 1978. Data from at least two years would be required to show a strict correlation of temperature with initiation of spawning.

Neither R. cataractae nor C. plumbeus were observed to undergo spawning migrations up any of the flowing inlets to the ~~Lower~~ Kananaskis Reservoir. Thus, it was not possible to use timing of spawning migrations as a measure of temporal isolation.

A meaningful measure of temporal isolation is overlap in the time of fully ripe females in the spawning area (Nelson, 1968). The results of the first of the three methods, involving a qualitative judgement as to the state of female gonad maturity, are presented in Figure 16 and in Appendix Table 24.

Fully ripe females of the two species were present for the first three weeks in June, 1978, during the height of spawning. C. plumbeus females outnumbered R. cataractae females in all three different gonad states. Fully ripe male C. plumbeus similarly outnumbered fully ripe male R. cataractae. The first male C. plumbeus with freely flowing milt was captured May 17; the first such male R. cataractae was captured May 18, 1978. The last fully ripe male C.

Figure 16. Change in female gonad maturity for *Couesius plumbeus* and *Rhinichthys cataractae*, sampled from Lower Kananaskis Reservoir, during the 1978 spawning period. Maximum surface temperatures (mercury thermometer) are also presented.



plumbeus and R. cataractae were captured July 11 and June 29, 1978, respectively.

The results of the second method, involving changes in the actual gonad weight of the females of each species are shown in Figure 17 and Appendix Table 25. There is complete overlap in the period of maximum gonad weight (or spawning period) between the curves of the two species. Similarly, the results of the third method (Fig. 18 and Appendix Table 26) show synchrony between the species for a period of maximum ova size (spawning period).

Spawning activities were not actually observed due to the depth at which it occurred. However, gillnet capture records indicate that numbers of C. plumbeus increased in the early morning and late afternoon, over the spawning sites. Comparable data are not available for R. cataractae.

It is not known if the two species spawn at different times of the day. These results indicate that, apart from diel isolation, there is little temporal isolation between the species in Lower Kananaskis Reservoir.

#### Habitat Isolation

There is little known of the type of spawning habitat selected by the two species. R. cataractae probably spawn in riffles over a gravel bottom (Scott and Crossman, 1973), and C. plumbeus are known to spawn either along lakeshores or in streams (Geen, 1955) amid large rocks (Brown et al., 1970). The intention of this portion of the study was to determine

Figure 17. Change in gonadosomatic values (gonad weight/body weight%) of female *Couesius plumbeus* (upper graph) and female *Rhinichthys cataractae* (lower graph) sampled from Lower Kananaskis Reservoir during the 1978 spawning period. Points on hatched line represent sample means of gonadosomatic values, bars represent sample standard deviations and lines represent range of sample values. Sample range and standard deviation are not given when  $n < 5$ .

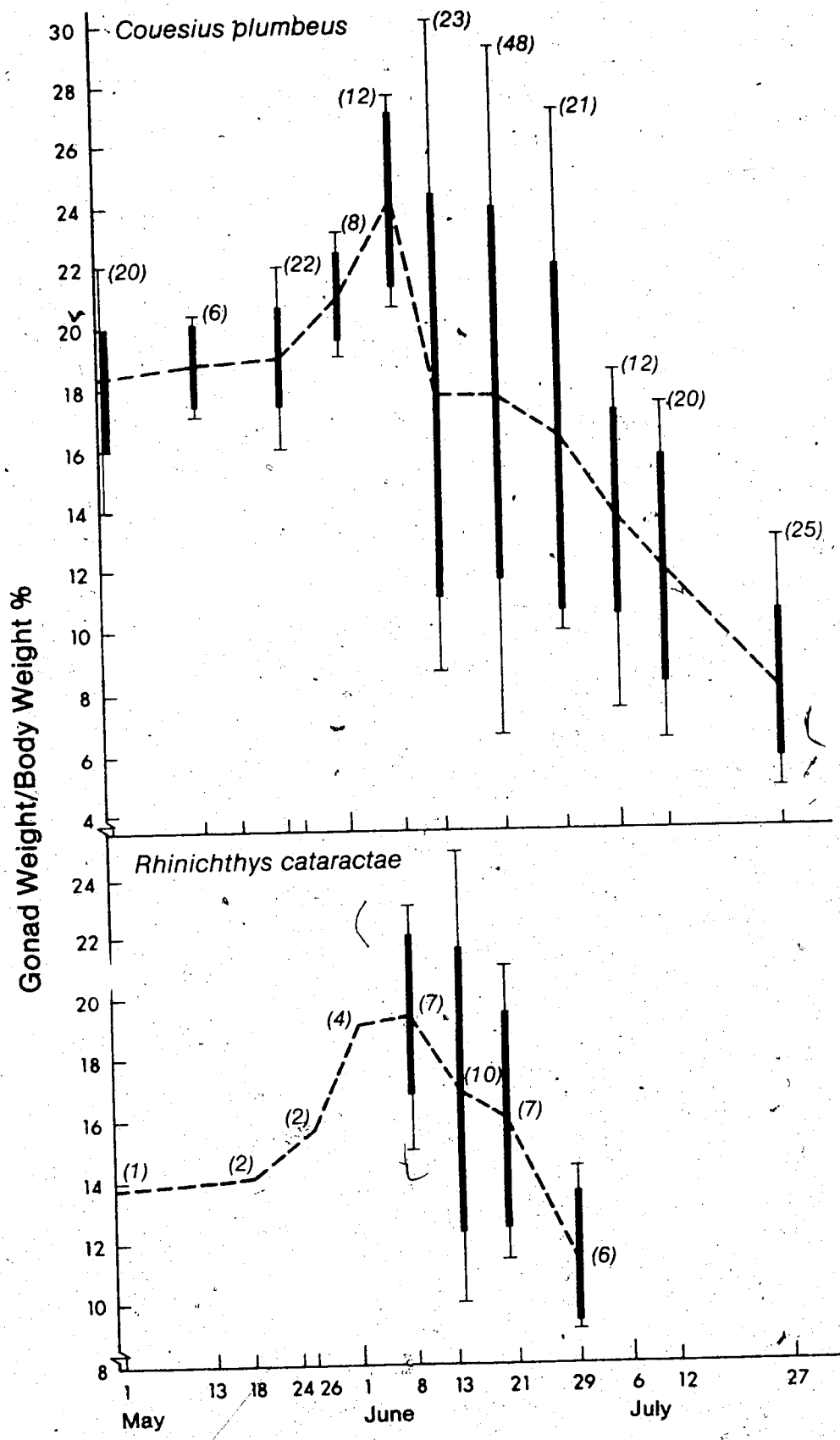
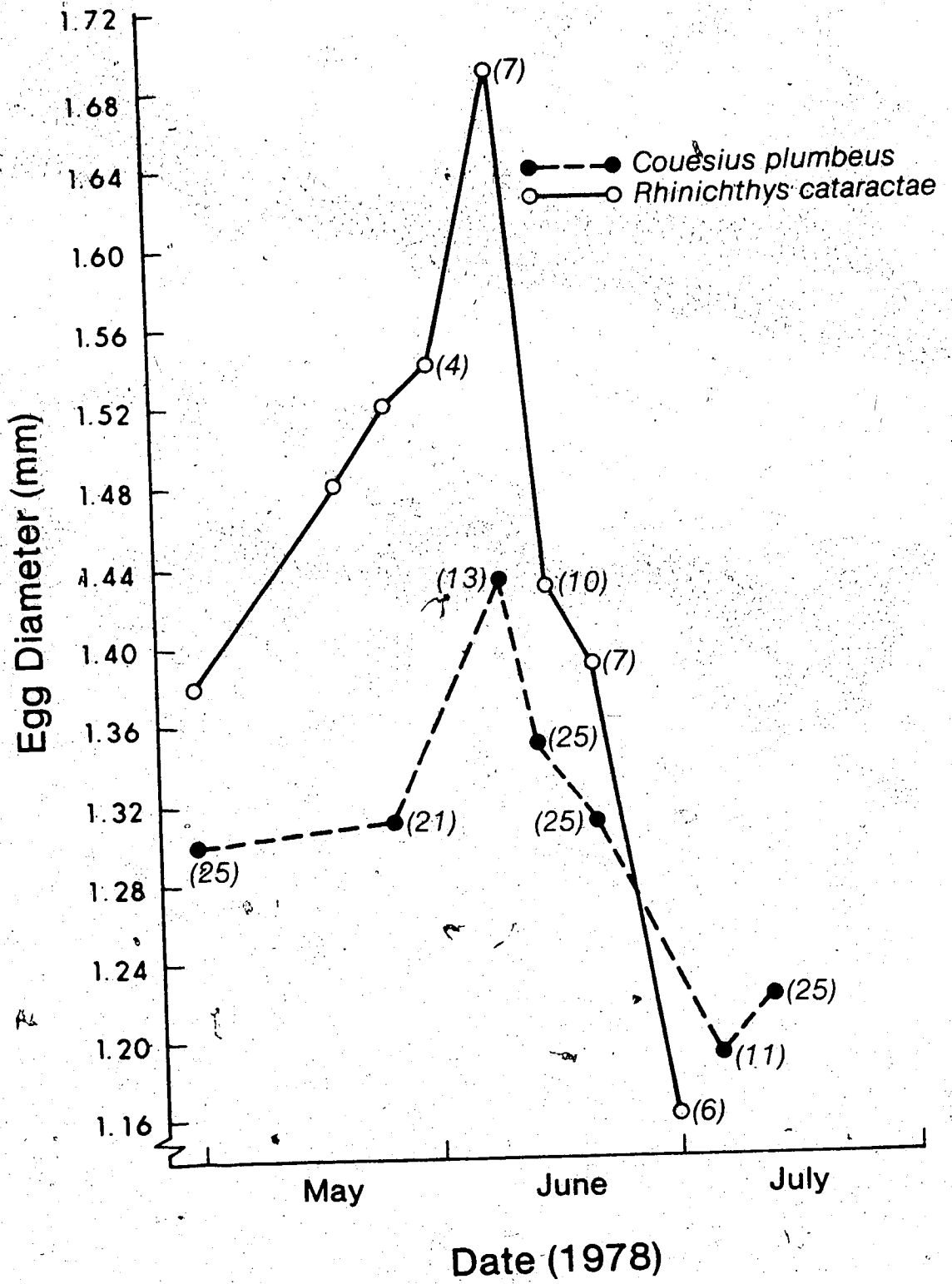




Figure 18. Trends in mean ova diameter during spawning period for *Couesius plumbeus* and *Rhinichthys cataractae*, Lower Kananaskis Reservoir, 1978. Sample size in parentheses (not presented if n = 1).





the degree of habitat segregation between these species in Lower Kananaskis Reservoir.

The inlet streams of Lower Kananaskis Reservoir were carefully monitored from when they were still ice-covered until the end of spawning in order to observe species' spawning movements and areas occupied. Due to spring-time access problems, the streams of Upper Kananaskis Reservoir were not similarly studied. There was no observed spawning migration of either species into Smith-Dorrien Creek during May, June, or early July. Only a few C. plumbeus and one B. cataractae were gillnetted at the mouth of the creek during this period (Fig. 19; Appendix Table 27). Neither species were captured upstream of the mouth. Electroshocking the creek (Plate 7) yielded migrating Salmo gairdneri, and baited minnow traps captured small Salvelinus malma (< 10 cm standard length). The current velocity of Smith-Dorrien Creek increased rapidly from 0.8 m/s on April 30 to a torrential 2.5 m/s on June 8 (Appendix Table 28) due to the spring melt. It became impossible to sample this creek for cyprinids except in the lower reaches at the mouth. High current velocities probably make this creek unsuitable for the cyprinids during their spawning season. There may be some shoreline spawning by the two species in the flooded inlet at the mouth of this creek, although my limited collections do not indicate this.

There were no observed spawning migrations of either species within Kent Creek, or in the vicinity of its mouth.

Figure 19. Catch data for adult *Couesius plumbeus*, *Rhinichthys cataractae*, and their putative hybrids from 7 selected sampling sites, Lower Kananaskis Reservoir. The time interval shown includes the period of maximal spawning activity for these species in 1978. Actual numbers of individuals caught by gillnets, minnow traps, and seines are shown by size of histograms.



The creek is steep and swift during the spawning season (> 2.0 m/s; Appendix Table 28), with little suitable spawning habitat. The mouth area of this creek was not extensively sampled (Appendix Table 1), and it may be used by one or both species for spawning.

Cyprinid spawning migrations were not observed during June in Invincible Creek due to its swift current (Appendix Table 28), and silty nature (Secchi disc = 5 cm, June 8, 1978). Fully ripe individuals of each species were captured in the silty outfall of the creek beyond its mouth during the peak spawning period (Fig. 19). Although C. plumbeus outnumbered R. cataractae, the two species appeared to be spawning in a common area without habitat segregation. The capture of one putative hybrid at this site on June 8 might suggest that miscegenation has occurred here.

No spawning individuals of either cyprinid species were found migrating up Boulton Creek in May or June, 1978. Intensive collecting effort up the Creek from its mouth to the edge of the forest did not yield a single cyprinid. Individuals of each species in spawning condition were captured beyond the mouth of Boulton Creek and nearby vicinity (Fig. 19 and Appendix Table 1). An overlap of the species' spawning habitats near Boulton Creek mouth may be indicated by the capture of one hybrid specimen on May 26, 1978.

The outfall of Boulton Creek is often extremely silty due to intermittent rainstorm run-offs and increased melt

flow (see Secchi disc results, Appendix Table 18). This factor is also the case in Invincible Creek and may abet the interspecies hybridization occurring in both locations.

The substrate over which the spawning C. plumbeus individuals were captured was silty mud of a uniform fine texture. This was not original lake bottom but part of the reservoir's littoral flood plain. The flooded meandering bed of Boulton Creek presented the only original pre-reservoir substrate. Capture rate did not appear to rise with increasing proximity to the original creek bed. E. cataractae were captured here in disproportionately greater numbers than in other nearby shoreline areas of equal small size.

Mature cyprinids were common within the confines of Power Plant Raceway (Interlakes Dam). There was no flow of water through the Power Plant during May, June, or July; however, there was a small flow of runoff via a drainage conduit. The raceway forms a deep trench into the original lake basin and the resultant thermal stability may be attractive to the spawners. Fully ripe males and females of each species were captured within the raceway on June 8, 1978, without any observed habitat segregation (Fig. 19; Appendix Table 1). The substrate was a uniform mixture of large boulders (mean diameter 0.3 m) over gravel.

The south end of Lower Kananaskis Reservoir was the most extensively collected area on the entire reservoir. A gently flowing stream (Interlakes Stream), the original

outlet for the Upper Kananaskis Lake (Plate 8) with riffle habitat eminently appropriate for R. cataractae, emptied into the reservoir in this area. Its flow was controlled so that it did not increase with spring run-off beyond the suitability for spawning habitat. Under present reservoir conditions, this stream flows across an extensive area of exposed mudflats, which are progressively flooded as the reservoir level rises in the spring (Fig. 20).

The mouth of Interlakes Stream is constantly receding relative to the advance of Common Bay. The following dynamic habitat types were sampled at this end of the reservoir prior to and during the cyprinid spawning season.

- 1) Interlakes Stream - was sampled as one habitat in its entirety from source to mouth, wherever the latter occurred.
- 2) Interlakes Stream Mouth - the immediate vicinity of the entrance of Interlakes Stream into Common Bay was sampled as one habitat wherever it occurred.
- 3) Common Bay - an area at least 100 m beyond the mouth of Interlakes Stream was sampled as one habitat.

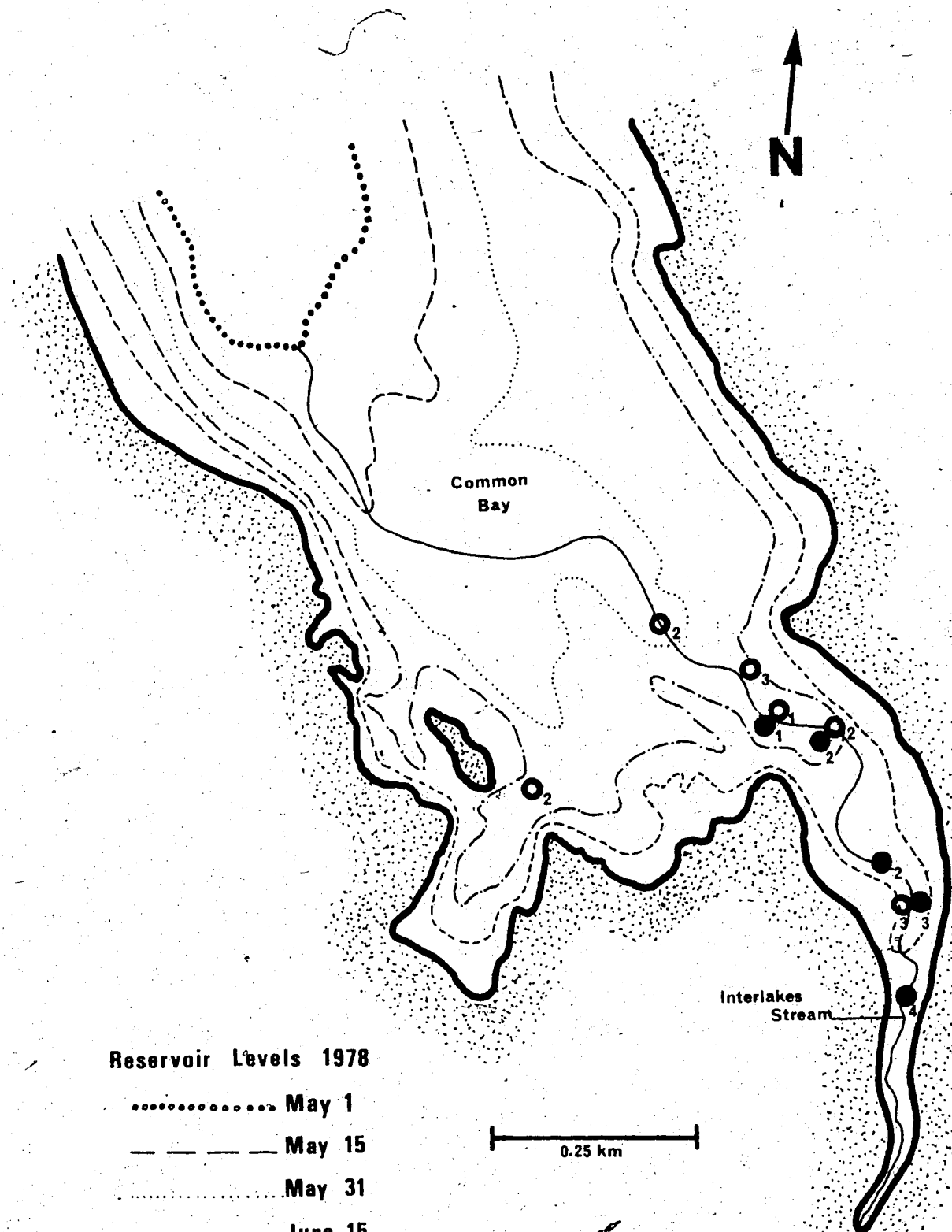
The sampling sites for each habitat type were continuously moving upstream (southwards) as the reservoir filled, but the position of each habitat type remained fixed relative to the others. The types of fish occupying each moving habitat could then be compared.

C. plumbeus was abundant in the deeper waters of Common Bay, but only one hybrid and one R. cataractae were captured. The C. plumbeus captured here appeared to be spawning in the absence of R. cataractae. The substrate



Figure 20. Extreme south end of Lower Kananaskis Reservoir showing the approximate progression of shoreline changes during the summer, 1978, filling of the reservoir and the consequent flooding of Interlakes Stream (1 cm = 0.07 km). Common Bay is designated as the open water beyond the mouth of Interlakes Stream. The capture location of fully ripe and/or spent individuals of each species are presented for selected dates as follows:

- *C. plumbeus*
  - 1 - June 8
  - 2 - June 12, 13
  - 3 - June 20, 21
- *R. cataractae*
  - 1 - June 8
  - 2 - June 12, 13
  - 3 - June 19, 20
  - 4 - June 29



**Reservoir Levels 1978**

- ..... May 1
- May 15
- ..... May 31
- June 15
- June 30

0.25 km

ranged from gravel in the flooded Interlakes Stream bed to silty mud on the flooded mudflats in a depth of 2.5 to 3.6 m (Fig. 19).

In contrast to Common Bay, Interlakes Stream was predominantly occupied by R. cataractae during the May, June, 1978 period. C. plumbeus were occasionally captured here (Fig. 19). A low population of adult R. cataractae of both sexes remained in the diminishing length of the stream throughout the spawning period and likely spawned here. The substrate of the stream was highly variable, ranging from large patches of silt and detritus in areas of slow current to gravel and boulders in the riffle areas. Most R. cataractae were sighted and captured among rocks (10 cm mean diameter) in riffles under a depth of less than 15 cm. Presumably, this is where R. cataractae were spawning. The 52 C. plumbeus captured on May 18, 1978, were mainly small immature specimens (mean standard length 34.3 mm) as were the two C. plumbeus captured May 22, 1978, (17 mm and 25 mm SL). The C. plumbeus captured in the stream on June 5, 1978, were larger specimens (mean SL 63.5 mm) including one fully ripe male and several ripe females (Fig. 19). The R. cataractae specimens captured on the same date were all larger females (mean SL 58.3 mm), some of which were ripe. Spawning among these fish was not observed, nor were any females sufficiently ripe on June 5, 1978, for spawning. Intensive sampling of the stream did not yield further C. plumbeus specimens until June 21, 1978, when two fully ripe

males were collected (70.0 and 72.0 mm SL) at the peak of the spawning period. In late June the reservoir level had flooded approximately half of the Interlakes Stream length within the S.E. Arm. These two C. plumbeus males were found 15 m upstream of the mouth and likely represent exploratory movements, but not a typical pattern. Their presence does suggest the possibility of interspecific spawning with the resident R. cataractae females. Spawning was not observed at any time in Interlakes Stream.

The Interlakes Stream mouth habitat was intermediate between the other two habitats in terms of the simultaneous presence of C. plumbeus and R. cataractae males and females in fully ripe and spent spawning conditions (Fig. 19; Appendix Table 1). There was no noticeable habitat separation between the two species. They were captured in the same gillnets over a substrate ranging from gravel, in the original stream bed, to silt on the flooded mudflats. The depth of the water ranged from 1.7 to 2.6 m. There was a slight, but continual current of  $< 1.0$  m/s at the mouth of the creek. Thus, it was impossible to determine if the species segregated themselves according to current regimes. One C. plumbeus x R. cataractae hybrid specimen was captured on June 8, 1978, along with individuals of the two species.

C. plumbeus numerically outnumbered R. cataractae in the Interlakes Stream mouth habitat. However, both are represented in an intermediate frequency relative to the C. plumbeus -dominated deeper waters of Common Bay and the R.

cataractae -dominated riffle habitat of Interlakes Stream. Partial habitat isolation is achieved in the latter two habitats, but not in the intermediate habitat where the rising reservoir level brings the two species into contact.

It is reasonable to hypothesize that C. plumbeus spawners, in maintaining their spawning proximity to the stream mouth, advance into the newly flooded lower reaches of the stream. Here, R. cataractae maintain their spawning positions over the gravel and cobble rock of the flooded stream bed. Where habitat segregation occurred in the south end of the Lower Reservoir, C. plumbeus presumably spawned offshore in a depth of 2.0 - 3.0 m over a barren silt substrate while R. cataractae are inferred to spawn in very shallow riffles of slight current amid cobble-sized rock.

#### Behavioral Isolation

Behavioral blocks to hybridization form a final line of pre-mating defense in the event of habitat and temporal overlap between reproducing species. However, Hubbs (1961) noticed that while a behavioral barrier to miscegenation occurs in an area where species have been sympatric for a long time, these behaviors may break down in areas where they have recently come together.

Little information concerning reproductive behaviors of the two species is available. C. plumbeus neither builds a nest nor guards the eggs, whereas R. cataractae may both establish a territory and guard the nest (McPhail and

Lindsey, 1970). Nests of R. cataractae were not found in the intensively sampled Interlakes Stream where R. cataractae was inferred to spawn in the absence of C. plumbeus. Nest guarding behaviors were not observed in members of R. cataractae, although some site specificity was indicated. When disturbed from one area, they would invariably be observed returning to the same site. Turning over the rocks and sieving the substrate in such locations did not yield eggs or nests. Nest-guarding adults were not observed when the flooded portion of Interlakes Stream in the S.E. Arm was observed underwater using S.C.U.B.A. on June 28 and July 12, 1978.

Laboratory, aquaria-held fish showed no meaningful behavioral differences when held in the reciprocal, interspecific matchings as opposed to conspecific situations. Neither mating behaviors nor aggressive interactions were observed at any time. Individuals of each sex of each species, while held in tanks, displayed similar fright behaviors and escape responses.

The male hybrids and female C. plumbeus in Tank C were not observed to elicit mating behaviors and tank breeding did not occur. When autopsied on June 28, the females had regressed gonads suggesting that tank conditions were inappropriate for breeding. Although the male hybrids were of equivalent size to the females, they reacted more vigorously to tank disturbance, swam faster, and dominated the feeding by rapidly snatching up the food.

The existence of distinct species' specific breeding behaviors in C. plumbeus and R. cataractae cannot be confirmed in this study. Their isolative value remains unknown. However, it is possible to evaluate species' differences in breeding coloration and nuptial tubercles (Plates 1-6). These differences may provide criteria for favoring conspecific mates. The results of this comparison (Table 7, Plates 1-6), reveal many species' differences, which may act to increase the species' reproductive isolation. Similarly, there are many morphological differences between the species, especially in head characters (previously discussed in text) of potentially high isolative value, assuming such characters are used for mate recognition. Confounding these differences is that in each species smaller males mate with larger females. Thus, mate selection on this criterion would not disfavor interspecific matings.

Without successful, controlled mating experiments, these hypothesized isolating factors remain untested speculations.

## DISCUSSION

Temporal isolation may be inoperative in preventing hybridization in Lower Kananaskis Reservoir (although diel isolation was not investigated). There is total overlap in the period of fully ripe females between the two species. The peak of the spawning period for each species occurred in

TABLE 7. Comparison of Breeding Coloration and Nuptial Tubercles. Refer to Plates 1-6.

<i>Couesius plumbeus</i>	<i>Rhinichthys cataractae</i>
<p>MALE</p> <ul style="list-style-type: none"> <li>- vivid red patch at pectoral fin base, spreading posteriorly. No coloration at bases of pelvic or anal fins.</li> <li>- red patch on top of opercle.</li> <li>- may have red pelvic axillary processes, if these are present.</li> <li>- small red patch at corners of mouth.</li> <li>- no coloration near isthmus.</li> <li>- all fin rays, membranes without color.</li> <li>- fine white tubercles on dorsal surface of pectoral rays.</li> <li>- always brighter red coloration than females.</li> </ul> <p>FEMALE</p> <ul style="list-style-type: none"> <li>- red patch at base of pectoral fin, spreading posteriorly.</li> <li>- red patches at top of opercle and at corners of mouth (on cheek) may be present.</li> <li>- fin rays, membranes without coloration.</li> <li>- color patches are yellow to light red, never vivid red.</li> <li>- fine tubercles may be present on dorsal surface of head and posteriorly to dorsal origin.</li> </ul>	<p>MALE</p> <ul style="list-style-type: none"> <li>- vivid red-orange patches at bases of pectoral, pelvic and anal fins.</li> <li>- no red patch on top of opercle.</li> <li>- red pelvic axillary processes.</li> <li>- red-orange patch at corners of mouth (upper mandible may be entirely colored while lower mandible is less extensively colored).</li> <li>- red-orange color scattered beneath gills at isthmus.</li> <li>- all fin rays, membranes may have a red-orange tinge.</li> <li>- fine white tubercles on dorsal surface of head (may extend posteriorly to origin of dorsal fin) and on dorsal surface of pectoral rays. (Tubercles usually restricted to head only).</li> <li>- usually brighter red coloration than females.</li> </ul> <p>FEMALE</p> <ul style="list-style-type: none"> <li>- may have a small red-orange patch at base of pectoral fin.</li> <li>- no red patch on top of opercle, or on cheek at corner of mouth.</li> <li>- all fin rays, membranes usually with a red-orange tinge.</li> <li>- color patches are orange-red.</li> <li>- tubercles not present.</li> </ul>



mid-June at approximately 10°C and ended prior to July in 1978: spawning did not continue into late August in either 1977 or 1978 as has been reported for Kananaskis cyprinids by McPhail and Lindsey (1970) and Scott and Crossman (1973) (probably based on Nelson's 1965 observation of ripe females being found on 8 August in 1961).

Previous reports have shown different temperature responses for the two species. R. cataractae in Manitoba reached a spawning peak on May 16 at 15°C in Mink River (Bartnik, 1970) and during mid-July at 15°C in Lake Winnipeg (Gee and Machniak, 1972). Kuehn (1949) reported that R. cataractae in Minnesota spawned from late June through August. Brazo et al. (1978), found the peak spawning period for Lake Michigan R. cataractae to be late June and early July at 14-19°C. C. plumbeus have spawned in the Montreal River, Sask. in early May at 4-8°C, Lac la Ronge, Sask. in mid-June at 10°C (Brown et al., 1970), and Lac Saugay, Que. in early June at 19°C (Scott and Crossman, 1973). These may be population differences.

There appears to be no effective habitat isolation between the species. They swarm in mixed schools over the mud flats at Pocaterra Dam in the summer and were captured together at spawning time along the shoreline and at the mouths of all inlet streams. However, habitat isolation was achieved by spawning R. cataractae migrating alone in modest numbers into Interlakes Stream. Presumably no other inlet stream offered the appropriate current, temperature, and

substrate regimes to allow species' segregation. There are no other reports in the literature outside Kananaskis of such close habitat sharing between C. plumbeus and R. cataractae.

Satisfactory evidence regarding the strength of the behavioral barrier between the two species is lacking in this study. However, the failure of the other possible isolating mechanisms, and the low frequency of hybrids, suggests that with C. plumbeus and R. cataractae, as with most species (Mayr, 1959), ethological isolation is of primary importance. Differences in morphology, color, tuberculation, and the suggested territoriality of R. cataractae (McPhail and Lindsay, 1970) are possibly involved. Ethological barriers are also inferred to be of major importance in maintaining species' identity in hybridizing populations of Catostomus commersonii and C. macrocheilus (Nelson, 1968).

## STUDY OF POSTMATING ISOLATION

### INTRODUCTION

Postmating isolating mechanisms involve the reduction of hybrid survival and the reduction of gene flow through the hybrids. Effective postmating isolation is not a general characteristic of sympatric species (Littlejohn, 1969) as it is not open to the operation of natural selection.

The purpose of this section is to determine the importance of postmating isolating mechanisms in maintaining

species' distinctness. The mortality and viability of artificially reared hybrid embryos is compared relative to that of parental embryos. Certain life history characteristics of wild-caught hybrids are compared to those of the parental populations in order to assess their relative survival success under natural conditions.

## METHODS

### Hybrid Embryo Mortality

Embryo mortality experiments were conducted in a controlled environment chamber commencing June 8 and terminating July 1, 1978, with the final hatching of all embryos. Fully ripe adults, caught in Lower Kananaskis Reservoir using gillnets, dipnets, and seines were returned to the laboratory where they were stripped of eggs or milt. Some adults were held in aquaria for as long as two days prior to stripping, while waiting to collect the parental type with which they were crossed. Fertilization procedures were adapted from Davis (1953) and Strawn and Hubbs (1956).

No attempt was made to extract the same number of eggs for all crosses. Each cross received the maximum number of eggs which could be expressed from the female. The fertilized eggs of each cross were placed in separate glass finger bowls and each was covered with a fine Nitex mesh (36 meshes/cm<sup>2</sup>). As many as three finger bowls of the same cross type were submerged into the same 20 l aquarium. This procedure allowed simultaneous experiments to be run without

the danger of accidental interchange of embryos.

Once hatching began, each cross was separated into its own aquarium. All aquaria were held at a constant temperature of 15°C and were constantly aerated. There was no continuous water flow into the aquaria, rather the water was changed approximately every five days. Chemical fungal retardants were not used.

There was a total of 21 crosses (9 homospecific crosses, 11 heterospecific crosses, and 1 cross of two F1 hybrid individuals). Each cross involved different parents. Hybrid and parental crosses were treated identically. Prior to hatching, eggs of all crosses were examined once every 24 to 36 hours. Dead embryos, identified by their opaque, white color, were counted daily and discarded using a pipette. When hatching began, the crosses were examined twice daily and the number of hatched fry recorded. A mean hatching date was determined for each cross. Hatching time data were analyzed using a single classification analysis of variance (Sokal and Rohlf, 1969) and a Duncan's multiple range test (Zalik, 1977).

The diameters of 15 embryos (five days post-fertilization) from four crosses (the two parental cross types and the two reciprocal hybrid cross types) were measured under a microscope using needle-point dial calipers (read to the nearest 0.1 mm). Egg diameter data were analyzed using a single classification analysis of variance (Sokal and Rohlf, 1969) and a Duncan's multiple range test

(Zalik, 1977).

The % pre-hatching mortality results for all crosses were arcsin transformed (Zalik, 1977,) and analyzed using a single classification analysis of variance (with unequal sample sizes) (Sokal and Rohlf, 1969).

#### Hybrid Inviability

##### Immediate Post-hatching Survival

The products of all four artificial cross types were reared beyond the hatched fry stage in several 8 l aquaria. Space limitations necessitated the pooling of replicates from the same cross type. All aquaria were held under the conditions of a constant 15°C temperature, continual aeration and water changes as needed. The developing fry were fed Liquifry, TetraMin baby fish food, and live zooplankton from Lower Kananaskis Reservoir.

The number of live fry 4 days post-hatching was recorded for each cross type replicate and the data were analyzed using a single classification analysis of variance (Sokal and Rohlf, 1969). Thereafter, the number of live fry was recorded once per month until 3 November, 1978. These long-term survival data were not statistically analyzed but simply compared to determine if any differences existed in survivorship between the cross types in the first months of life.

### Long-term Survival and Life History Characteristics

The success and long-term viability of the wild-caught putative hybrids was compared to that of the two parental species by their age, growth, and length-weight data.

Age was determined by scales, read under magnification independently by two people. Where disagreement occurred, the readers conferred until a consensus was reached. Scales were removed from preserved fish in the small area above the lateral line and below the dorsal midline in front of the dorsal fin. Thirty-six putative hybrids with intermediate canonical scores (in the presumed F<sub>1</sub> portion of the Kananaskis CVA distribution) were aged. Aging by the scale method was corroborated using the graphic method of polymodal separation of length frequencies (Cassie, 1963).

Growth rates of these representatives of the hybrid group and the two typical parental species were calculated by plotting the mean size at each age, as derived from the scale analysis, against age-class. Growth rates were then estimated from regressions of logarithms of length on age-class. An analysis of covariance was performed on the same data using the SPSS computer program (Nie et al., 1975). This provided an objective method of determining significant differences between regression lines.

Length-weight relationships for the same subsamples of the three groups were determined according to the methods of Ricker (1971). The relationship was described by a regression of logarithm of length on logarithm of weight. An

analysis of covariance was performed on these data using the SPSS program.

Female fecundity was determined for 25 C. plumbeus, 20 R. cataractae, and 7 hybrids. All ova from one female were spread evenly in a dish divided into quadrats; only one quadrat was actually counted. This value was multiplied by four to give the total number of ova. The fecundity relationship was described by a regression of logarithm of ova number on logarithm of standard length for C. plumbeus and R. cataractae. The hybrid sample size was too small to generate a meaningful regression.

## RESULTS

### Hybrid Embryo Mortality

Successful artificially induced hybridization may confirm the origin of putative hybrids. It can also provide insight into the relative degree of selection against hybrid cross products. The hypothesis of equal hatching success between hybrid crosses and parental crosses was tested (Table 8).

Each of the heterospecific crosses suffered 100% pre-hatching mortality in one replicate, as did the only hybrid backcross replicate. All replicates of the homospecific crosses had some hatching. However, there is no significant difference in pre-hatching mortality between the four different cross types, excluding the hybrid backcross ( $P > 0.05$ , Table 9). Given only one hybrid backcross

TABLE 8. Hatching results for experimental crosses, 1978, at 15°C. C - *Cowpea plumbeus* R - *Rhinichthys cataractas*  
 H - *C. plumbeus* x *H. cataractas* F<sub>1</sub> Hybrid

	<u>Heterospecific Crosses</u>											
	<u>♂ C x ♀ R</u>				<u>♀ C' x ♂ R</u>							
	19 June	19 June	19 June	19 June	13 June	13 June	16 June	8 June	20 June	20 June	20 June	20 June
Date Started	19 June	19 June	19 June	19 June	13 June	13 June	16 June	8 June	20 June	20 June	20 June	20 June
No. Started	181	90	90	205	500	438	373	200	123	400	220	
No. Hatched	128	0	73	156	227	326	0	7	49	177	48	
$\bar{x}$ Pre-hatch Mortality	29.3	100.0	18.8	23.9	54.6	25.6	100.0	96.5	60.2	55.8	78.2	
$\bar{x}$ Hatch Time (days)	10.0	-	10.0	11.0	9.5	10.0	-	10.0	10.0	9.5	10.0	
$\bar{x}$ 4 Day Post-hatch Mortality	98.4	-	2.7	7.0	1.3	0.6	-	0	8.2	42.3	100.0	
No. Alive 4 Day Post-hatching	2	0	71	145	224	324	0	3	45	102	0	

	<u>Homospecific Crosses</u>										Hybrid
	<u>C x C</u>					<u>R x R</u>					<u>Backcross</u>
	13 June	13 June	8 June	20 June	20 June	19 June	19 June	20 June	20 June	8 June	♂ H x ♀ C
Date Started	13 June	13 June	8 June	20 June	20 June	19 June	19 June	20 June	20 June	8 June	
No. Started	280	380	240	240	136	43	90	246	115	62	
No. Hatched	59	98	38	219	94	12	17	84	33	0	
$\bar{x}$ Pre-hatch Mortality	78.9	74.2	84.2	8.8	30.9	72.1	81.1	65.9	71.3	100	
$\bar{x}$ Hatch Time (days)	9.0	9.0	9.5	9.0	8.5	11.0	12.0	12.0	11.0		
$\bar{x}$ 4 Day Post-hatch Mortality	1.4	3.0	0	49.3	3.2	0	5.9	28.6	6.1		
No. Alive 4 Day Post-hatching	57	95	38	111	91	12	16	60	31		



TABLE 9. % Pre-hatching Mortality. Single classification analysis of variance with unequal sample sizes. The single replicate of the hybrid backcross suffered 100% pre-hatching mortality and was not included in this analysis.

Arcsin Transformed Data				ANOVA TABLE				
CC	$\delta CqR$	$qCqR$	RR					
62.7	32.8	47.6	58.1	Source of Variation	SS	df	MS	F
59.5	90.0	30.4	64.2	Among Crosses	743.6	3	247.9	0.59
66.6	25.7	90.0	54.3	Within Crosses	7185.6	17	422.7	
17.3	29.3	79.2	57.6	Total	7929.2	20		
33.8		50.9		$F_{.05(3,17)} = 3.2$				
		48.3						
		62.2		$P > 0.5$				

% 4 Day Post-hatching Mortality. Single classification analysis of variance with unequal sample sizes.

Arcsin Transformed Data				ANOVA TABLE				
CC	$\delta CqR$	$qCqR$	RR					
10.6	82.7	6.6	0.0	Source of Variation	SS	df	MS	F
10.0	0.0	4.4	14.1	Among Crosses	449.4	3	149.8	0.20
0.0	9.5	0.0	32.3	Within Crosses	12479.0	17	743.1	
44.6	15.3	0.0	14.3	Total	12928.4	20		
10.3		16.6		$F_{.05(3,17)} = 3.2$				
		40.6						
		90.0		$P > 0.5$				

replicate, it is not possible to make any conclusions on the sterility of hybrids. I suspect the eggs used in this cross were not fully ripe.

Hybrid crosses hatch as successfully as do control crosses. On the basis of this experiment, hybrid embryo mortality cannot be operative as a postmating isolating mechanism.

There is a significant difference in hatching times for cross types ( $F = 28.0$ ,  $F.05(3,14) = 3.34$ ). The hatching times for pure R. cataractae embryos and pure C. plumbeus embryos are significantly different from all other cross types. The hatching time for embryos of the two reciprocal hybrid cross types are not significantly different (latter results from Duncan's test). Mean hatching time for pure C. plumbeus crosses is 9.0 days: relatively short compared to pure R. cataractae at 12 days. The hatching times of the heterospecific crosses are intermediate: male C. plumbeus x female R. cataractae cross, 10.5 days; female C. plumbeus x male R. cataractae, 9.8 days. This may show a trend for the hybrids to develop at rates similar to that of their maternal controls, as reported by Hubbs and Strawn (1957).

Hybrid fertilized egg sizes are also intermediate between the parental values (Table 10). There is a significant difference in embryo sizes for cross types ( $F=26.9$ ,  $F.05(3,56) = 2.76$ ). Pure C. plumbeus embryos are significantly smaller than embryos from the other three cross types, which are not significantly different from each

other in size (latter results from Duncan's test). There may be a causal relationship here, i.e., the larger R. cataractae egg requiring a longer development time than the smaller egg of C. plumbeus.

#### Hybrid Inviability

It is probable that hybrids are occasionally produced in nature by an accidental mechanism such as drifting sperm. In a natural environment, as opposed to a modified environment, these individuals may be at a selective disadvantage and may die prior to reproducing. The hypothesis that artificially produced hybrids of Kananaskis origin were not selected against was tested.

#### Immediate Post-hatching Survival

Percent 4 day post-hatching mortality results (Table 9) show no significant difference ( $P > 0.05$ ) between the replicates of the four cross types. The highest percent mortality is in two heterospecific cross replicates: more than twice the highest percent mortality recorded for the homospecific crosses. There appeared to be no difference between the cross types in the number of highly deformed individuals that contributed to these mortality figures.

The results of the four month rearing experiment (Table 11) show unexpected differences in viability between the cross types. All R. cataractae control cross individuals died between two and three months after hatching. No other cross type suffered total mortality during the rearing

Table 10. Diameters of Eggs (5 days post-fertilization) for each artificial cross type.

Cross Type	No. of Eggs	mean diameter(mm)	Std. Deviation
C C	15	2.03	0.08
C♀ R♂	15	2.28	0.12
R♀ C♂	15	2.23	0.11
R R	15	2.35	0.08

TABLE 11. Survival of offspring from experimental crosses over a 4 month period (post-hatching). Some replicates of the same cross type were lumped together for economy during rearing as indicated. Survival values for each time period are given as absolute number of live individuals. The mean hatching date for all crosses is taken as June 27, 1978.

R:- *R. cataractae*; C:- *C. plumbeus*.

Cross Type	Replicates		Survival (days post-hatching)			
	(identified by no. hatched)	No. lumped 4th day Post-hatching	35	65	91	124
♀C ♂C	59, 98, 38 219, 94	190 202	131 (55.1) 137 (41.4)	102 (46.3) 127 (39.6)	64 (35) 104 (35.2)	62 (34.3) 82 (30.1)
♀R ♂R	17, 12 84 33	28 60 31	24 (65.5) 0 0	0 0 0	0 0 0	0 0 0
♀R ♂C	73 156 128	71 145 2	62 (67.2) 0 0	12 (23.9) 0 0	9 (20.6) 0 0	8 (19.3) 0 0
♀C ♂R	227 326 7	224 324 3	183 (63.9) 260 (63.3) 0	142 (52.3) 83 (30.3) 0	99 (41.3) 74 (28.5) 0	56 (29.8) 67 (26.9) 0
	49, 177 48	147 0	137 (51.1) 0	78 (36.0) 0	33 (22.5) 0	30 (21.4) 0

period. The hybrid cross of R. cataractae maternal descent also suffered 100% mortality in two of its three replicates in the first month of rearing.

In contrast, the hybrid cross of C. plumbeus maternal descent had higher viability success with 100% mortality in only two of its five replicates. There were no complete die-offs in either replicate of the C. plumbeus control cross.

The results suggest disproportionate selection against R. cataractae control specimens and hybrids of R. cataractae maternal descent under artificial conditions.

#### Long-term Survival and Life History Characteristics

##### a) Age

The presence of five age-classes in the C. plumbeus sample is indicated by scale annuli. The lengths of the fish at each annulus formation are presented in Table 12. Separation of polymodal length frequencies by probability coordinates verified the presence of five age-classes (Fig. 21). A chi-square test showed no significant differences ( $P > 0.05$ ) between the observed and expected normal distributions of length frequencies in each of the age-classes. There is close correspondence between the results of the two methods. The size ranges at each age, determined by the scale method, fell within the size ranges at each age determined by the polymodal probability method. Bruce and Parsons (1976) found no spawning C. plumbeus from Labrador older than 4+ years. However, Brown (1969) found

TABLE 12. Length distribution of the age-groups of *Couesius plumbeus* from May 1, 1978 sample, Lower Kananaskis Reservoir (n = 137). (Numbers in parentheses represent the % of each age-group in each length range.)

Standard Length Interval (cm)	Age-group and Sex												
	0		1		2		3		4				
	M	F	M	F	M	F	M	F	M	F	M	F	
1.0-1.9	1(50)												
2.0-2.9	1(50)												
3.0-3.9			7(30)	8(36)									
4.0-4.9			11(48)	8(36)		1(4)							
5.0-5.9			5(22)	5(23)		4(11)	7(25)						
6.0-6.9			1(5)			14(38)	9(32)						
7.0-7.9						16(43)	10(36)						
8.0-8.9						3(8)	1(4)		4(50)	5(42)			
9.0-9.9									4(50)	7(58)			1(25)
10.0-10.9													1(100) 2(50)
11.0-11.9													1(25)
Subtotals	2	0	23	22	37	28	8	12	1	4			

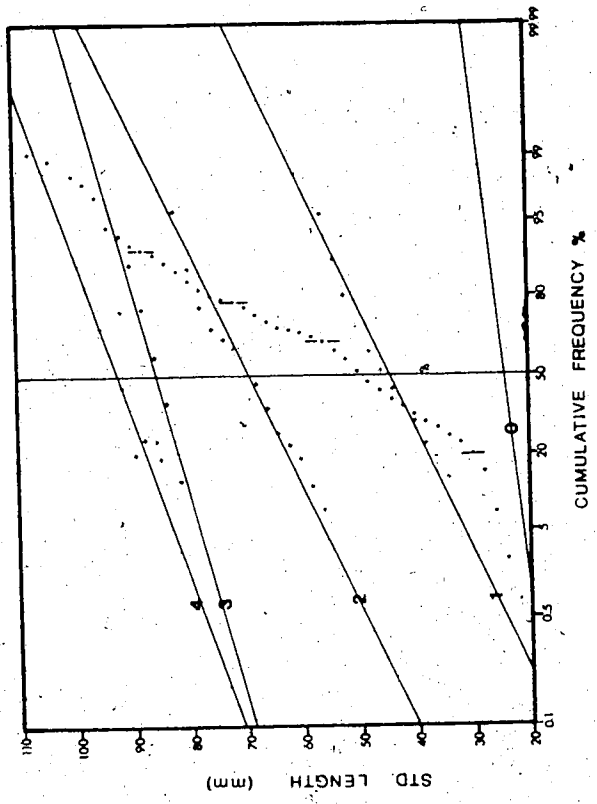
Figure 21. Graphic separation of polymodal length frequencies by probability coordinates for:

- A. 343 *Couesius plumbeus* from Lower Kananaskis Reservoir.  
 $\chi^2$  test for departure from normality ( $\chi^2 = 19.90$ ;  
 $\chi^2_{.05(43df)} = 55.7$ ,  $P > 0.05$ ).
- B. 241 *Rhinichthys cataractae* from Lower Kananaskis Reservoir.  $\chi^2$  test for departure from normality ( $\chi^2 = 3.32$ ;  $\chi^2_{.05(40df)} = 55.8$ ,  $P > 0.05$ ).

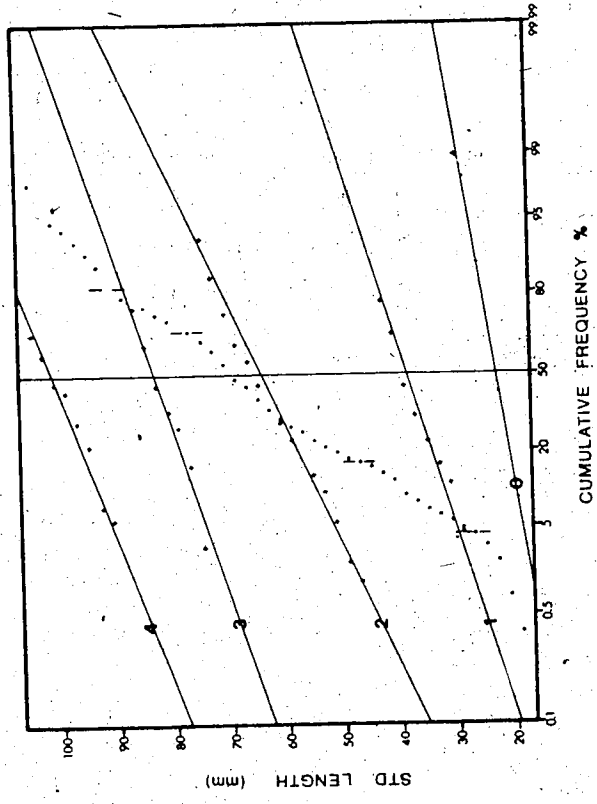
Numerals are age-class designations. Solid dots are total cumulative frequencies and starred circles are cumulative frequencies within size groups.



B RHINICHTHYS CATARACTAE



A COUESIUS PLUMBEUS



spawning C. plumbeus 5 years and older in Saskatchewan.

Six age-classes are indicated for R. cataractae from scale annuli. The lengths of the fish at each annulus formation are represented in Table 13. Separation of polymodal length frequencies by probability coordinates revealed the presence of only 5 age-classes (Fig. 21). However, this is likely due to the low frequency of older 5+ R. cataractae. A chi<sup>2</sup>-square test showed no significant differences ( $P > 0.05$ ) between the observed and expected normal distribution of length frequencies in each of the age-classes. Six age-classes have also been reported from Lake Michigan populations of R. cataractae (Brazo et al., 1978).

Four age-classes are indicated for the hybrid sample by counts of the scale annuli (Table 14). Although age-class 0 hybrid individuals were available they are not included in the analysis. No hybrid individuals were older than 4 years and there were no male hybrids older than 3 years. The hybrid sample is small ( $n = 36$ ) and contains mostly small hybrids in the younger age-classes. The low frequency of hybrid individuals precludes age verification by polymodal probability method. Age analysis has not been previously reported for C. plumbeus x R. cataractae hybrids.

#### b) Growth Rates

Growth rates of the two species and the hybrids are described by the following equations:

$$\underline{C. plumbeus} \log L = 0.513 + 0.147T$$

TABLE 13. Length distribution of the age-groups of *Rhinichthys cataractae* from 1977, 1978 collections, Lower Kananaskis Reservoir (n = 131). Numbers in parentheses represent the % of each age-group in each length range.

Standard Length Interval (cm)	Age-group and Sex											
	0		1		2		3		4		5	
	M	F	M	F	M	F	M	F	M	F	M	F
1.0-1.9												
2.0-2.9	2(100)	2(100)	6(40)									
3.0-3.9			6(40)	9(64)								
4.0-4.9			2(13)	5(36)	7(33)	6(25)						
5.0-5.9			1(7)		12(57)	11(46)						
6.0-6.9					2(10)	4(17)	9(82)	4(14)				
7.0-7.9					3(12)		2(18)	11(39)				
8.0-8.9							10(36)		2(17)			
9.0-9.9							3(11)	1(100)	9(75)			
10.0-10.9									1(8)			1(100)
Subtotals	2	2	15	14	21	24	11	28	1	12	0	1

TABLE 14. Length distribution of the age-groups of *Couesius plumbeus* x *Rhinichthys cataractae* hybrids, from 1977, 1978 collections, Lower Kanawasis Reservoir (n = 36). Numbers in parentheses represent the % of each age-group in each length range.

Standard Length Interval (cm)	Age-group and Sex									
	0		1		2		4			
	M	F	M	F	M	F	M	F		
1.0-1.9										
2.0-2.9										
3.0-3.9										
4.0-4.9										
5.0-5.9										
6.0-6.9										
7.0-7.9										
8.0-8.9										
9.0-9.9										
10.0-10.9										
11.0-11.9										
-Subtotals	0	0	5	3	12	8	5	1	0	2

$$\underline{C. plumbeus} \times \underline{R. cataractae} \log L = 0.494 + 0.124T$$

$$\underline{R. cataractae} \log L = 0.425 + 0.145T$$

where L = standard length, T = age-class and the regression coefficients equal the instantaneous growth rates. Data from each sex were pooled to calculate the regression for each species.

C. plumbeus appears to have the fastest growth rate. The growth rates for C. plumbeus were determined from Saskatchewan (Brown et al., 1970) using the Peterson method and from Labrador (Brazo et al., 1978) using back calculations. Thus, their results are not directly comparable to those in this study.

Comparison of the regression lines by inspection reveals that R. cataractae has the slowest growth rate for fish younger than 3 years. The hybrids have a generally intermediate growth rate between the parental species, but after 3 years age it declines below the growth rate of R. cataractae (Fig. 22). If the growth rates are extended back beyond the Y-axis to include all growth in the first year, it appears that the hybrid growth rate initially exceeds that of the two parentals. The hybrid offspring of the experimental crosses also showed a faster growth rate than that of the parental offspring. This apparent decline in the hybrid growth rate may suggest the accumulative effects of some selective disadvantage.

However, an analysis of covariance (Nie et al. ., 1975) was performed to test for significant differences between

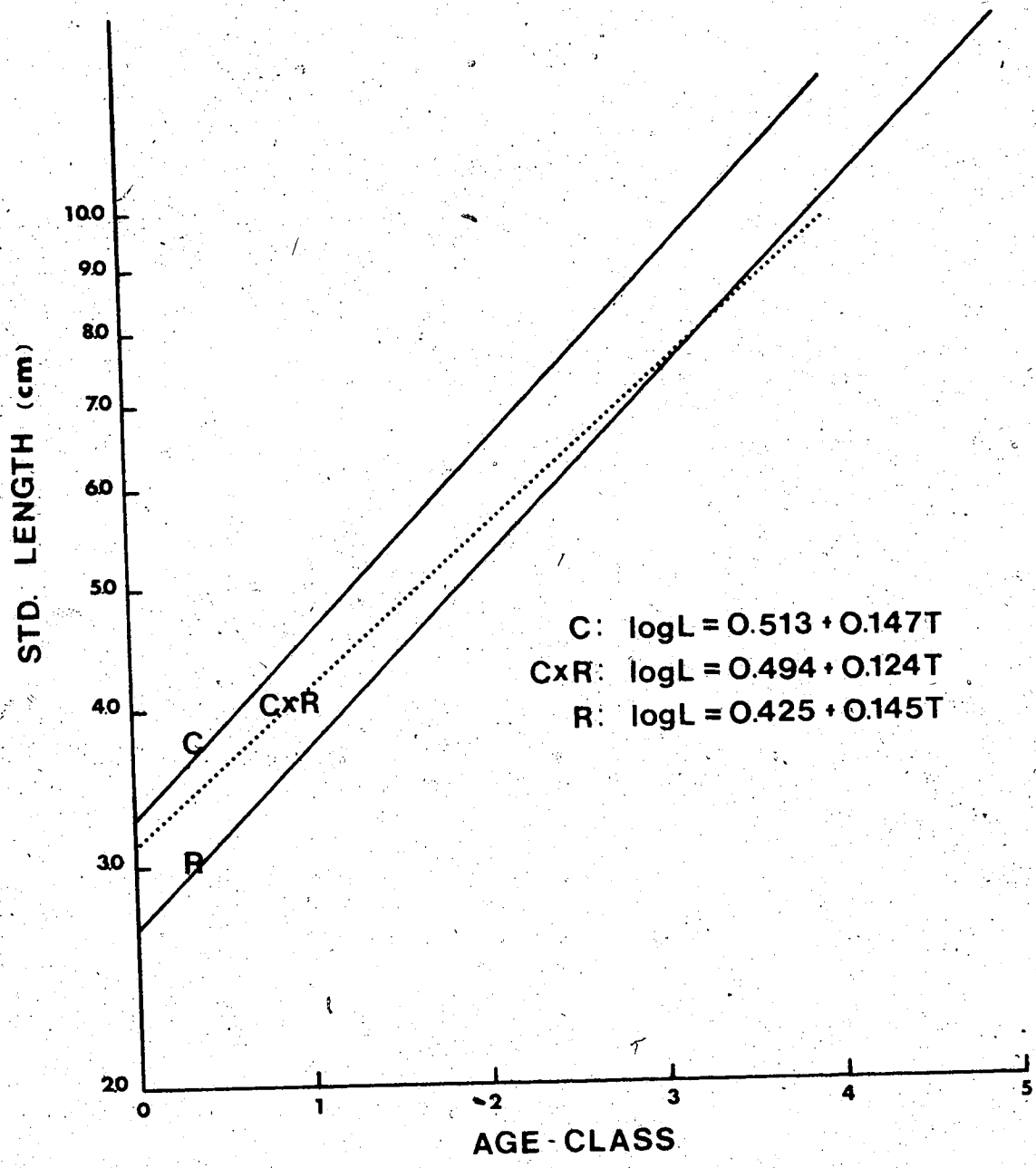
Figure 22. Growth rate regressions for:

C - 140 *Couesius plumbeus* ( $r = 0.90$ )

R - 130 *Rhinichthys cataractae* ( $r = 0.91$ )

C x R - 36 *C. plumbeus* x *R. cataractae* hybrids ( $r = 0.93$ )

from Lower Kananaskis Reservoir.



the regressions (Appendix Table 29). Statistically, there is no significant difference ( $P > 0.05$ ) between the slopes of the three regression lines: there is significant difference ( $P < 0.05$ ) between the intercepts of R. cataractae and C. plumbeus and between R. cataractae and the hybrid but no significant difference ( $P > 0.05$ ) between the intercepts of C. plumbeus and the hybrid.

c) Length-Weight Relations

C. plumbeus, R. cataractae and the hybrid populations each approximate isometric growth, as described by the equations:

$$\text{C. plumbeus } \log W = -1.781 + 3.064 \log L$$

$$\text{C. plumbeus} \times \text{R. cataractae } \log W = -1.725 + 2.813 \log L$$

$$\text{R. cataractae } \log W = -1.899 + 3.120 \log L$$

where  $W$  = weight and  $L$  = standard length. Data from both sexes were pooled. Comparison of the three length-weight regressions (Fig. 23) by inspection reveals no apparent differences between the three populations. Analysis of covariance between the regressions shows no significant differences ( $P > 0.05$ ) between the slopes but significant differences ( $P < 0.05$ ) between the intercepts. Length-weight regressions calculated separately for the sexes of each population are given in Appendix Table 30.

C. plumbeus from Saskatchewan (Brown, 1969) are heavier for any given length ( $\log = -3.00 \times 10^{-6} + 3.20 \log L$ ) than the Kananaskis C. plumbeus. Additionally, C. plumbeus from Labrador (Bruce and Parsons, 1976) weigh less for any given



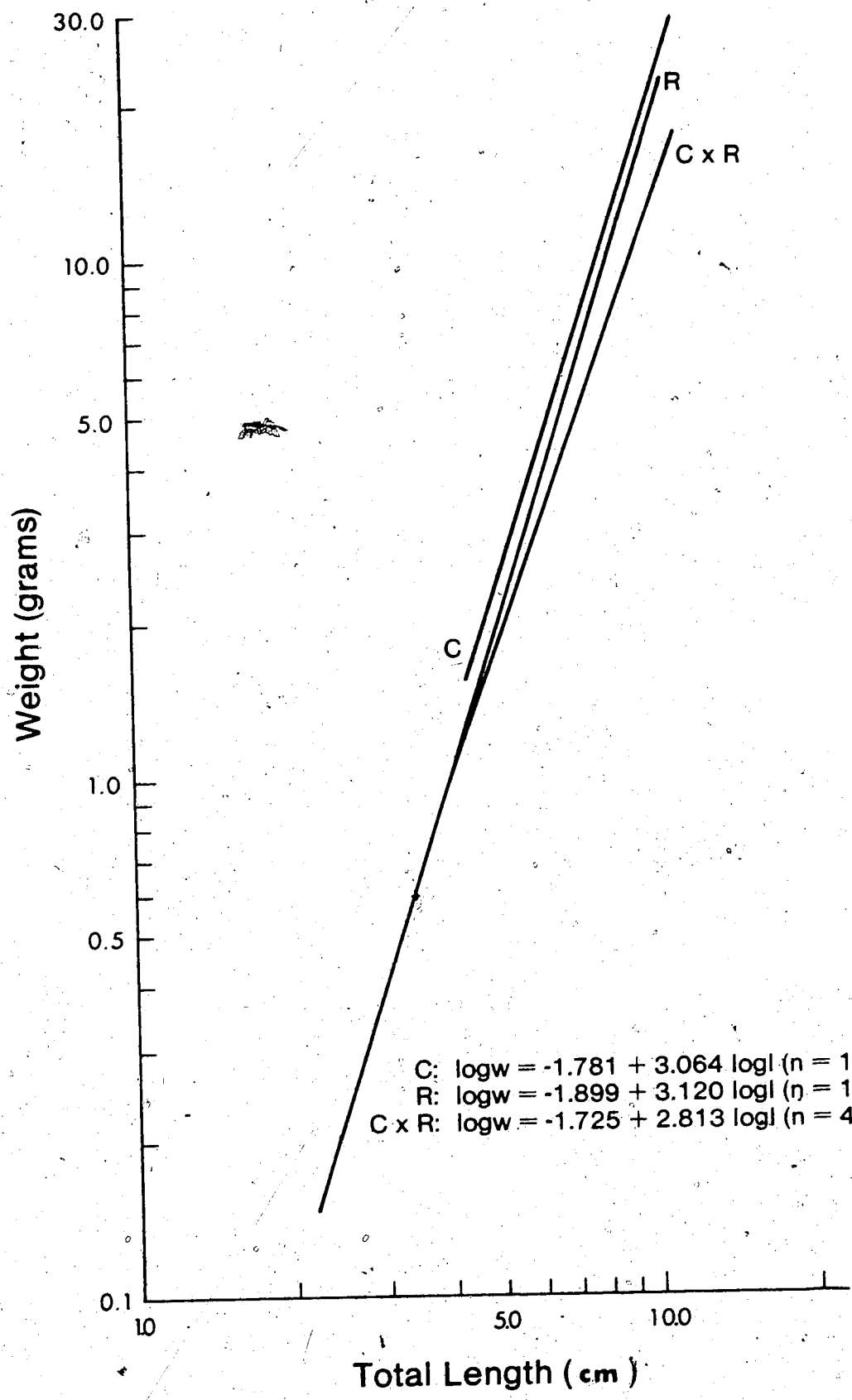
Figure 23. Length-weight regressions for:

C - 170 *Couesius plumbeus* ( $r = 0.96$ )

R - 157 *Rhinichthys cataractae* ( $r = 0.99$ )

C x R - 33 *C. plumbeus* x *R. cataractae* hybrids ( $r = 0.98$ )

from Lower Kananaskis Reservoir.



length ( $\log W = -1.88 + 2.88 \log L$ ) than the Kananaskis C. plumbeus. Length-weight regressions have not been previously reported for R. cataractae or C. plumbeus x R. cataractae hybrids.

#### d) Fecundity

The relationship of fecundity to standard length for C. plumbeus and R. cataractae can be described by the following equations:

$$\text{C. plumbeus } \log F = 0.673 + 2.534 \log L$$

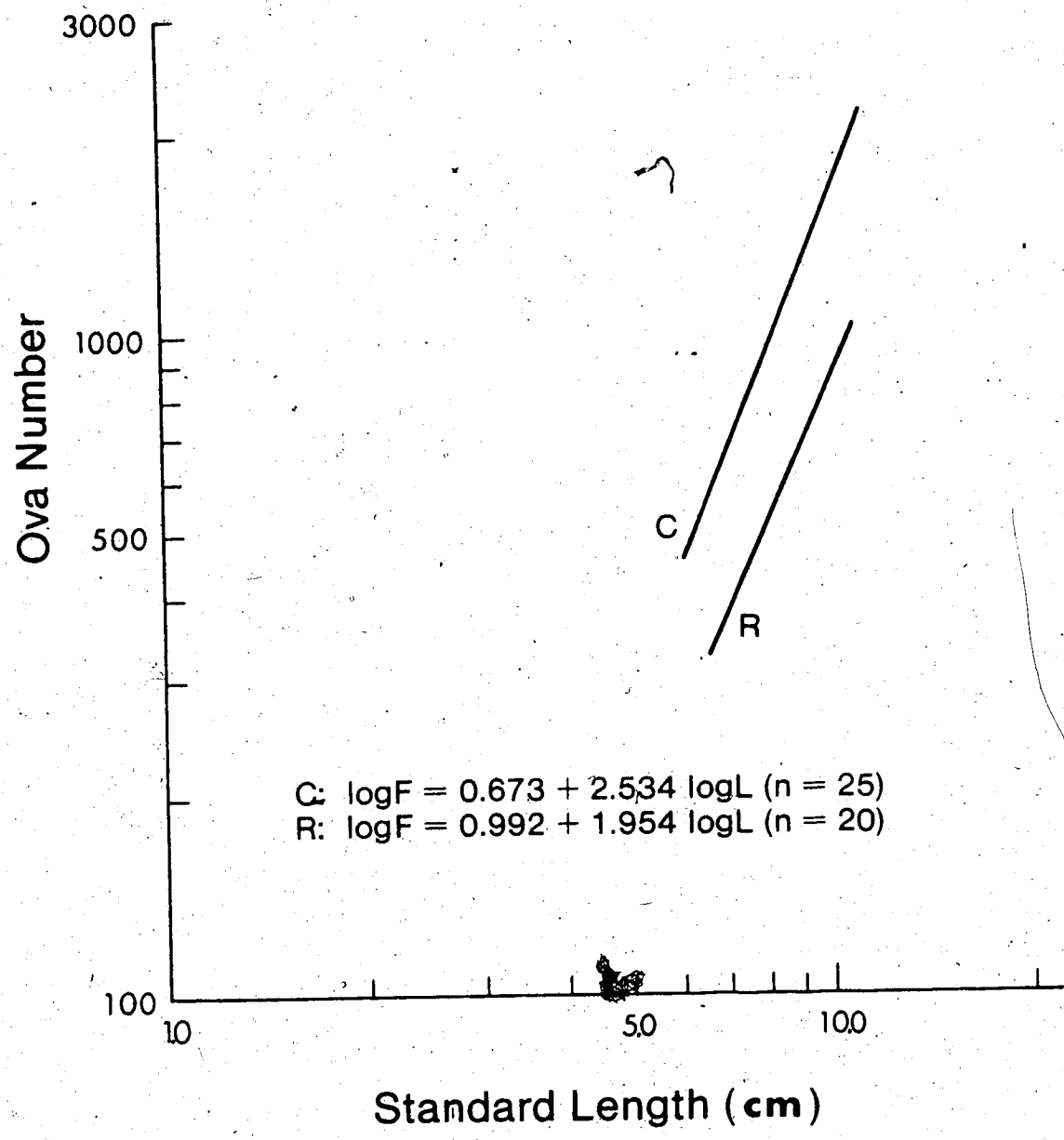
$$\text{R. cataractae } \log F = 0.922 + 1.954 \log L$$

Analysis of covariance shows significant differences ( $P < 0.05$ ) between these regressions in both slopes and intercepts (Appendix Table 29). The fecundity regressions of the parental species are presented as a contribution to our knowledge of their life histories (Fig. 24).

Of the seven adult, female hybrids examined, four had regressed gonads with very small, indistinguishable eggs. These were captured July 27, 1977, and were probably in a post-spawning state. Another post-spawning hybrid female (SL = 62 mm) had 92 ova. Two hybrid females were captured during the spawning season: one was captured June 16, 1977, (SL = 96 mm) and was fully spent with only 38 ova remaining; the other female was captured May 26, 1978, (SL = 114 mm) and was fully ripe with 1038 ova.

Brown et al. (1970) gave a range of 450-2450 eggs for 20 C. plumbeus with a total length range of 9.0-13.2 cm. The fecundity regression for this Saskatchewan population was

Figure 24. Fecundity regressions for 25 *Couesius plumbeus* (C)  
( $r = 0.80$ ) and 20 *Rhinichthys cataractae* (R) ( $r = 0.70$ )  
from Lower Kananaskis Reservoir.



$\log F = -3.243 + 3.096 \log L$ . Additionally, Bruce and Parsons (1976) gave a range of 573-1158 eggs for 10 C. plumbeus with a fork length range of 9.2-11.2 cm. The fecundity regression for this Labrador population was  $\log F = -0.441 + 3.356 \log L$ . The number of eggs for C. plumbeus in this study ranged from 380-2316 eggs for 25 fish with a standard length range of 6.1-11.2 cm.

Brazo et al. (1978) gave a range of 870-9953 eggs for 33 R. cataractae from Lake Michigan with a total length range of 7.4-11.7 cm. The number of eggs for R. cataractae in this study ranged from 364-1121 eggs for 20 fish with a standard length range of 6.6-10.0 cm.

#### DISCUSSION

Embryo mortality tests revealed no difference in hatching success between homospecific and heterospecific crosses. The artificial backcross was a failure, but conclusions as to hybrid fertility from one replicate would not be sound. Also, there was no difference in long-term survival between artificially produced hybrids and individuals of parental species origin. In fact, hybrid offspring survived in aquaria better than did pure R. cataractae individuals.

Wild-caught hybrids have similar growth rates and attain a similar age structure compared to the parental species. Also, there is no statistical difference between the length-weight relationships of the hybrids and the

parental species. The life history data suggest that the hybrids may not be facing excessive competition from the parental species. Thus, neither genetic mortality nor hybrid inviability are operative barriers preventing hybridization between C. plumbeus and R. cataractae.

Hybrid gonads were not examined histologically, but the hybrid gonads did not appear to be deformed or superficially different from those of the parental species. Fully ripe male and female hybrids were captured and exuded sex products similar to individuals of the parental species. Hybrid sperm was observed microscopically and found to be motile. Hybrid eggs appeared to be of similar shape and condition as parental eggs. Additionally, it is not known experimentally if hybrids are sterile, but it can be inferred from the results of the morphological analysis that at least the barrier is low. Introgression does not appear to put the species' status in peril. The sex ratio of the hybrids was biased toward males.

f

## IX. GENERAL DISCUSSION

The occurrence of this hybridization between C. plumbeus and R. cataractae does not alter the systematic position of these two species within cyprinid genera that are generally held to be distantly related. Fish hybridization in areas of environmental disturbance no more imperils the generic or specific status of the animals involved than do the multitude of possible artificial crosses (Hubbs, 1961). The fact that these two types of cyprinid remain distinct in most other localities of their range overlap confirms the recognition of the two types as distinct species (sensu Mayr, 1969).

Certain recognized environmental factors, which facilitate the dissolution of species' reproductive isolating mechanisms (Hubbs, 1955; Mayr, 1969), all appear to be important in causing hybridization between C. plumbeus and R. cataractae in Lower Kananaskis Reservoir.

1) Environmental Disturbance; Paucity of Spawning Areas

Disturbance of the environment is often alluded to as the most frequent cause of hybridization of both plants and animals (Mayr, 1969; Anderson, 1949, 1953; Hubbs et al., 1943; Sibley, 1961; Mechan, 1960). However, its method of operation has not been demonstrated in any rigorous way.

The Lower Kananaskis Reservoir is a disturbed environment in terms of the annual level fluctuations associated with filling and lowering. The change from a lake to a reservoir may indirectly contribute to hybridization.



Spawning species may be brought together because flooding causes the loss and reduction of originally distinct spawning habitats. Also, with variability in drawdown levels from year to year, there is unpredictability in normal shoreline features which could be used for new spawning sites. Without predictable flooding during the spawning season, it is likely that species' habitat segregation breaks down.

C. plumbeus is an adaptable species which can lake spawn and undergo spawning migrations up rivers and streams (Brown et al., 1970). R. cataractae probably prefers spawning in riffles (Scott and Crossman, 1973). With the possible exception of R. cataractae moving into Interlakes Stream, neither species underwent spawning migrations up inlet streams. Although each species has a preference for inlet stream spawning, it appears that the streams are unavailable (except Interlakes Stream) during the spawning season due to fast currents and low temperatures. (Why the unflooded portions of Interlakes Stream were not utilized more for spawning by the two species, remains unknown.) It is likely that the two species have been restricted to spawning along the shoreline and at stream mouths since their initial introduction to the lakes. Reservoir construction could neither have caused nor altered this spawning relationship with inlet streams. Lower Kananaskis Lake was equally depauperate in appropriate spawning areas, as the Reservoir is today.

However, while spawning habitat segregation was possible under original, predictable lake conditions, fluctuating reservoir levels may have disrupted this tenuous segregation. Evidence provided in this study indicates that rising reservoir levels at Interlakes Stream mouth brought together fully ripe individuals of each species.

Hybridization may result because of the action of an hypothesized difference between the two species in their reaction to advancing water levels. C. plumbeus, in maintaining their depth distribution and their proximity to the stream mouth, advance into the newly flooded lower reaches of the stream. Here, P. cataractae maintain their positions relative to the stream perhaps due to spawning territoriality, as cited by McPhail and Lindsey (1970). This series of events may be inferred to occur at the other stream mouths where habitat segregation is not so apparent. The generality of this model requires verification from other reservoirs where these species are sympatric.

Without pre-impoundment information on where the species were spawning or whether hybridization occurred, it is difficult to evaluate the contribution of reservoir construction to fish hybridization. Environmental impact statements must provide these data to rigorously assess both the beneficial and adverse effects of future reservoir construction.

## 2) Species Introduction; Rarity of One Parental Species

Nelson (1966) advanced the idea that the two species were introduced as live bait into the original Kananaskis Lakes; R. cataractae prior to 1936 and C. plumbeus prior to 1961 (in the late 1950's). The introduction of one or both species to a new environment appears to be one of the many precursors to hybridization (Hubbs et al., 1943; Hubbs, 1961; Greenfield and Greenfield, 1972). The effects of introduction will depend upon the abundance of each species, their genetic similarity, and the strength of the species' isolating mechanisms.

In Lower Kananaskis Reservoir, the introduction of C. plumbeus has successfully resulted in development of a large population size. Introduced R. cataractae have not adapted as well and have attained a small population size relative to that of C. plumbeus. R. cataractae are possibly beyond the margin of their normal habitat range in Lower Kananaskis Reservoir, which is without abundant riffle habitat. In Lower Kananaskis Reservoir, this low frequency of R. cataractae relative to C. plumbeus is a contributing factor to hybridization. In the absence of appropriate mating stimuli, individuals of the less common species may have no alternative but to respond to inappropriate stimuli from individuals of the abundant species (Mayr, 1969). Hybridization has long been attributed to a disproportion in the abundance of species (Hubbs et al., 1943; Mecham, 1960; Stebbins, 1959). Whether R. cataractae will mate with C. plumbeus in the absence of conspecific mates is unknown.

Such mating did not occur during the mating tests conducted in this study.

### 3) Overview of Environmental Factors

Given the absence of appropriate historical information on the ecology and abundance of these two species, it is not possible to single out one of the above factors as having the major role in facilitating hybridization.

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Plate 1. Male *Couesius plumbeus* captured at height of spawning period, mid-June, 1978, from Lower Kananaskis Reservoir.

Plate 2. Female *Couesius plumbeus* captured at height of spawning period, mid-June, 1978, from Lower Kananaskis Reservoir.



Plate 3. Male *Rhinichthys cataractae* captured at height of spawning period, mid-June, 1978, from Lower Kananaskis Reservoir (Interlakes Stream).




Plate 4. Female *Rhinichthys cataractae* captured at height of spawning period, mid-June, 1978, from Lower Kananaskis Reservoir (Interlakes Stream).



Plate 5. Adult male hybrid, *Couesius plumbeus* x *Rhinichthys cataractae*,  
captured at height of spawning period, mid-June, 1978, from  
Lower Kananaskis Reservoir.

Plate 6. Top: *Couesius plumbeus* (male)

Middle: *Couesius plumbeus* x *Rhinichthys cataractae* hybrid  
(male)

Bottom: *Rhinichthys cataractae* (male)

All specimens captured in 1977, Lower Kananaskis Reservoir.

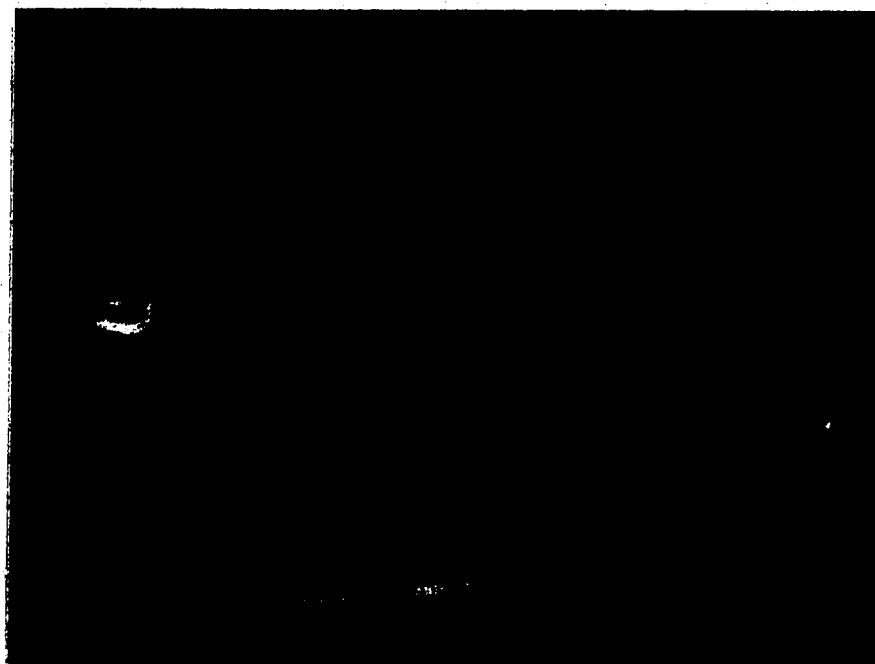
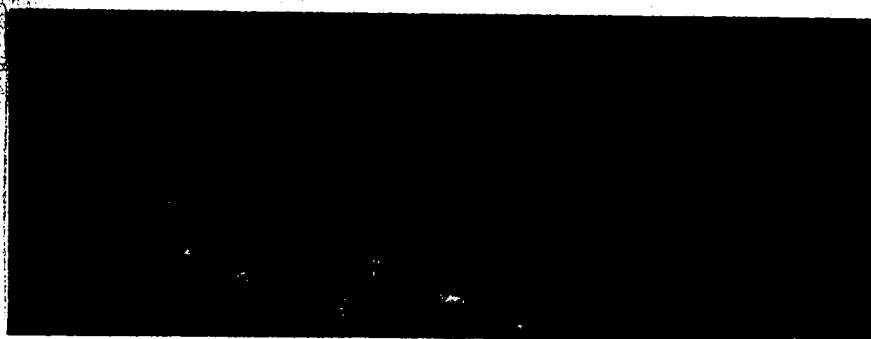
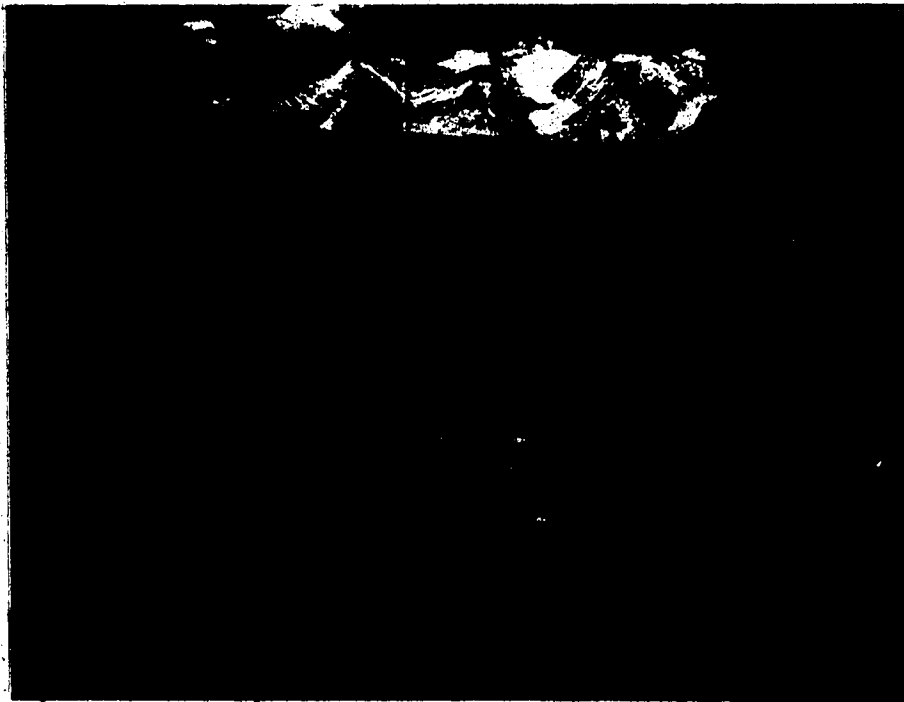




Plate 7. Electroshocking the lower reaches of Smith-Dorrien Creek near its mouth on Lower Kananaskis Reservoir, early May 1978 (looking east).

Plate 8. Interlakes Stream at south end of Lower Kananaskis Reservoir (looking south), in early June prior to cyprinid spawning peak. This illustrates Interlakes stream habitat before it is flooded with the rising reservoir level.



APPENDIX TABLE 1. Summary of cyprinid catch data from Upper and Lower Kananaskis Reservoirs, 1977. Size of gillnet mesh given in mm followed by number of hours set (in parentheses).

Capture Site Description	Date	Method of Capture	Depth	Numbers of Specimens Captured		
				<i>C. plumbeus</i>	<i>R. cataractae</i>	Putative Hybrids
UPPER KANANASKIS RESERVOIR						
UK1 - East Bay near main dam	May 25	Gillnet 26 mm(24)	4.5 m	9		
UK1 - East Bay near main dam	May 27	Gillnet 26 mm(48)	2.0 m	2		
UK2 - mouth of Sarrail Cr.	June 14	Gillnet 26 mm(24)	3.5-5.5 m	27	4	
UK2 - mouth of Sarrail Cr.	June 16	Gillnet 26 mm(48)	6.0-7.0 m	7		1
UK2 - mouth of Sarrail Cr.	June 21	Gillnet 18 mm(24)		10		
UK2 - mouth of Sarrail Cr.	June 29	Gillnet 26 mm(43)	4.5-6.5 m	29	7	
UK3 - E. shore; 300 m N of rocky pt. in East Bay	May 27	Gillnet 26 mm(24)		1		
UK5 - mouth of Kananaskis R.	June 21	Gillnet 26 mm(23)	3.0-4.0 m	6		1
UK5 - mouth of Kananaskis R.	July 21	Seine		10		
UK7 - inlet to Interlakes penstock	July 21	Seine		182	1	
UK7 - inlet to Interlakes penstock	Aug. 9	Seine		378	2	7
UK8 - N. shore; midway from Interlakes to Kananaskis R.	July 21	Seine		1		
LOWER KANANASKIS RESERVOIR						
LK1 - extreme S. end of reservoir	May 10	Gillnet 26 mm(24)	3.0 m	9	1	
LK1 - extreme S. end of reservoir	May 11	Gillnet 26 mm		4		
LK1 - as above & off E. shore	May 27	Gillnet 18 mm(31)	1.5-2.5 m	8		
LK1 - extreme S. end of reservoir	June 2	Gillnet 18 mm		12		
LK1 - as above & mouth to S.E. Arm	June 14	Gillnet 26, 18 mm(27)		74	4	1
LK1 - as above & mouth to S.E. Arm	June 15	Gillnet 18 mm(26)	2.0-4.0 m	11		
LK1 - as above & mouth to S.E. Arm	June 15	Gillnet 26 mm(26)	2.0-4.0 m	22		
LK1 - as above & mouth to S.E. Arm	June 23	Gillnet 26 mm(19)	4.0-6.0 m	23		
LK1 - as above & mouth to S.W. Arm	June 23	Gillnet 26, 18 mm(18)		8		
LK1 - as above & mouth to S.W. Arm	June 23	Seine		75		
LK1 - as above & mouth to S.W. Arm	June 24	Gillnet 26, 18 mm(21)	5.5-6.0 m	41		
LK1 - as above & mouth to S.E. Arm	July 6	Gillnet 26, 18 mm(20)	5.8-6.5 m	54		
LK1 - as above & mouth to S.W. Arm	July 8	Seine		522		2
LK1 - as above & mouth to S.W. Arm	July 15	Seine		28		
LK1 - as above & near campsites W. shore	July 21	Seine		998	17	13
LK1 - as above & near campsites W. shore	July 27	Gillnet 26 mm(21)	4.5 m	28		
LK1 - as above & mouth to S.E. Arm	Aug. 9	Gillnet 26, 18 mm(22)	8.0 m	50		

APPENDIX TABLE 1. Continued

Capture Site Description	Date	Method of Capture	Depth	Numbers of Specimens Captured		
				<i>C. plumbeus</i>	<i>B. cataractae</i>	Putative Hybrids
LOWER KANANASKIS RESERVOIR						
LK2 - E. shore; N. of Boulton Cr.	June 3	Gillnet 26, 18 mm(24)	2.0 m	13		1
LK2 - E. shore; N. of Boulton Cr.	June 14	Seine		14		
LK2 - E. shore; N. of Boulton Cr.	June 15	Gillnet 18 mm(26)		40		
LK2 - E. shore; N. of Boulton Cr.	June 16	Gillnet 18 mm(25)		7		
LK2 - E. shore; N. of Boulton Cr.	June 22	Gillnet 26 mm(24)	2.0-2.6 m	7		
LK2 - E. shore; N. of Boulton Cr.	June 24	Gillnet 26 mm(22)	4.0-5.0 m	11		
LK2 - E. shore; N. of Boulton Cr.	July 27	Seine		110		
LK2 - E. shore; N. of Boulton Cr.	July 27	Seine		3		2
LK4 - E. shore; near cottages	July 27	Seine		530	149	2
LK4 - E. shore; near cottages	July 27	Seine		3		
LK5 - mouth of Muskeg Cr.	June 1	Gillnet 26, 18 mm(29)	2.5-12 m	24	1	
LK6 - mouth of Invincible Cr.	June 1	Gillnet 26 mm(28)	2.0 m	18		
LK6 - mouth of Invincible Cr.		Gillnet 26, 18 mm		11		
LK7 - mouth of Smith-Dorrien Cr.		Gillnet 26, 18 mm(25)	2.0 m	5		
LK7 - mouth of Smith-Dorrien Cr.		Gillnet 18 mm(24)	3.0 m		1	
LK7 - mouth of Smith-Dorrien Cr.		Gillnet 26 mm		2		
LK7 - mouth of Smith-Dorrien Cr.		Seine		62		
LK8 - mouth of Kent Cr.	July 11	Seine		199		2
LK8 - mouth of Kent Cr.	July 11	Seine		62		
LK8 - mouth of Kent Cr.	July 20	Seine		732	94	12
LK9 - flats N. of Pocaterra Dam	July 11	Seine		1227	95	12
LK9 - flats N. of Pocaterra Dam	July 27	Seine		1271		
LK9 - flats N. of Pocaterra Dam	July 27	Seine		29		
LK9 - flats N. of Pocaterra Dam	Aug. 8	Seine		344		
Interlakes P.P. Raceway	June 16			97	5	

APPENDIX TABLE 2. Copepod captures from Kennecott Reservoir, Utah (LKI Stream - Interlakes Stream)

Date	Location	Capture Method	Depth	Substrate	May			June			Total					
					Sex	Total	Size	Sex	Total	Size	Sex	Total	Size			
May 1	Common Bay (E. shore)	seine	0.1 m	fine silt	♀	250	250	♀	3	3	♀	3	3	♀	3	3
May 7	Smith-Dorrien Cr. Mouth	minnow trap	1.0	coarse silt	♂	1	1	♂	1	1	♂	1	1	♂	1	1
May 9	Common Bay (E. shore)	minnow trap	0.5 m	silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
May 11	LKI Stream	seine	0.4 m	coarse silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
May 15	LKI Stream	seine	0.2 m	fine silt	♀	3	3	♀	3	3	♀	3	3	♀	3	3
May 16	Smith-Dorrien Cr. Mouth	alinet 18, 26 mm (28.5)	1 m	silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
May 17	Smith-Dorrien Cr. Mouth	alinet 26 mm (24)	1.25 m	silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
May 18	Smith-Dorrien Cr. Mouth	alinet 18, 26 mm (22)	1.25 m	silt	♀	6	6	♀	6	6	♀	6	6	♀	6	6
May 18	LKI Stream	seine	0.5 m	silt	♀	3	3	♀	3	3	♀	3	3	♀	3	3
May 22	LKI Stream	seine	0.1 m	silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
May 22	LKI Stream Mouth	seine	0.1 m	silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
May 23	Common Bay	alinet 18, 26 mm (21.5)	1.0 m	silt	♀	46	46	♀	46	46	♀	46	46	♀	46	46
May 24	Common Bay	alinet 18, 26 mm (24)	1 m	silt	♀	6	6	♀	6	6	♀	6	6	♀	6	6
May 24	Smith-Dorrien Cr. Mouth	alinet 18, 26 mm (24)	1 m	silt & gravel	♀	4	4	♀	4	4	♀	4	4	♀	4	4
May 25	Interlakes Runway	alinet 18 mm (23)	1.0 m	boulder	♀	1	1	♀	1	1	♀	1	1	♀	1	1
May 25	Smith-Dorrien Cr. Mouth	alinet 18, 26 mm (22)	1 m	silt	♀	9	9	♀	9	9	♀	9	9	♀	9	9
May 26	Smith-Dorrien Cr. Mouth	alinet 18, 26 mm (23)	1.1 m	gravel	♀	1	1	♀	1	1	♀	1	1	♀	1	1
May 26	Boulton Cr. Mouth	alinet 18, 26 mm (20)	1-3.0 m	silt, stones	♀	9	9	♀	9	9	♀	9	9	♀	9	9
May 27	Smith-Dorrien Cr. Mouth	alinet 18 mm (22)	1-3.0 m	silt & gravel	♀	3	3	♀	3	3	♀	3	3	♀	3	3
May 27	Boulton Cr. Mouth	alinet 18, 26 mm (24)	1-3.0 m	silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
June 1	Interlakes Runway	alinet 26 mm (28)	1.1-3.2 m	boulder	♀	13	13	♀	13	13	♀	13	13	♀	13	13
June 1	LKI Stream Mouth	alinet 18 mm (22)	1.2 m	silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
June 1	Kent Cr. Mouth	alinet 18, 26 mm (21)	1-3.0 m	gravel	♀	2	2	♀	2	2	♀	2	2	♀	2	2
June 5	LKI Stream	seine, dipnet	0.1 m	boulder & silt	♀	9	9	♀	10	10	♀	10	10	♀	10	10
June 5	LKI Common Bay	alinet 18, 26 mm (18)	17.5 m	silt	♀	2	2	♀	2	2	♀	2	2	♀	2	2
June 5	LKI Common Bay	alinet 18 mm (46)	17 m	silt	♀	2	2	♀	2	2	♀	2	2	♀	2	2
June 7	Boulton Cr. Mouth	alinet 26 mm (48)	1.5 m	silt	♀	10	10	♀	10	10	♀	10	10	♀	10	10
June 7	LKI Stream	seine, dipnet	0.05-0.5 m	boulder & silt	♀	1	1	♀	1	1	♀	1	1	♀	1	1
June 7	LKI Stream Mouth	alinet 26 mm (22)	1.8 m	silt	♀	11	11	♀	11	11	♀	11	11	♀	11	11

2 = sex unknown





APPENDIX TABLE 3. Water quality data, Upper Kanana Reservoir (1977). S - surface; M - mid-depth; B - 1 m off bottom;

N.A. - not available.

PARAMETER	June 20			July 15			August 15		
	S	M	B	S	M	B	S	M	B
Total Alkalinity (mg/l CaCO <sub>3</sub> )	83.9	84.4	84.6	83.0	85.0	86.0	79.0	84.0	85.0
Color	5	5	5	5	5	5	5	5	5
pH	8.3	8.2	8.0	8.3	8.1	8.1	8.5	8.1	8.0
Residue, Non-filterable (mg)	15	15	15	15	15	15	15	15	15
Specific Conductance (us/cm)	173	177	174	N.A.	N.A.	N.A.	155	166	167
Turbidity (JTU)	1.4	1.2	2.8	1.5	2.5	3.5	1.5	3.1	2.0
Dissolved Nitrogen (NO <sub>3</sub> NO <sub>2</sub> mg/l N)	0.02	0.05	0.05	0.03	0.05	0.05	0.01	0.06	0.01
Total Kjeldahl Nitrogen (mg/l)	0.28	0.21	0.23	0.10	0.10	0.10	0.10	0.10	0.10
Total Phosphorus (mg/l)	0.007	0.005	0.009	0.014	0.005	0.010	0.003	0.005	0.009
Total Organic Carbon (mg/l)	4	5	4	1	3	1	2	1	1
Total Inorganic Carbon (mg/l)	17	17	18	19	19	20	17	19	19



APPENDIX TABLE 4. Water quality data, Lower Kananaskis Reservoir (1977). S - surface; M - mid-depth; B - 1 m off bottom;

N.A. - not available.

PARAMETER	June 20			July 15			August 15		
	S	M	B	S	M	B	S	M	B
Total Alkalinity (mg/l CaCO <sub>3</sub> )	102.0	104.0	101.0	103.0	103.0	100.0	104.0	107.0	103.0
Color	< 5	< 5	< 5	< 5	< 5	< 5	< 5	< 5	< 5
pH	8.4	8.3	8.3	8.6	8.2	7.9	8.5	8.3	7.9
Residue Non-filterable (mg)	15	15	15	15	15	15	15	15	15
Specific Conductance (us/cm)	217	232	218	N.A.	N.A.	N.A.	216	230	218
Turbidity (JTU)	1.6	2.2	2.8	1.5	1.5	2.5	1.5	2.5	2.0
Dissolved Nitrogen (NO <sub>3</sub> , NO <sub>2</sub> mg/l N)	< 0.01	0.03	0.02	0.01	0.01	0.02	< 0.01	0.03	0.02
Total Kjeldahl Nitrogen (mg/l)	0.27	0.21	0.27	< 0.10	0.10	< 0.10	< 0.10	< 0.10	< 0.10
Total Phosphorus (mg/l)	0.006	0.007	0.013	0.010	0.006	0.005	0.005	0.008	0.013
Total Organic Carbon (mg/l)	4	3	6	1	2	2	3	3	3
Total Inorganic Carbon (mg/l)	21	22	21	24	24	23	23	24	23

APPENDIX TABLE 5. Temperature profiles of Lower Kananaskis Reservoir, 1978. Taken using a YSI telethermometer at Water Quality Sampling Site.

Depth (m)	Temperature °C		
	May 18	June 8	June 29
0	3.0	10.5	13.0
1	4.0		13.0
2	4.0		12.6
3	4.0	8.8	11.6
4	4.0		11.2
5	4.0		10.5
6	4.0	7.1	9.9
7	3.8		9.0
8	3.5		8.4
9	3.4		7.7
10	3.4		7.5
11	3.4		7.3
12	3.3	6.0	7.0
13	3.2		-6.9
14	3.1		6.9
15	3.0	5.1	6.4
16	2.9		6.2
17	2.9		6.1
18	3.0	5.0	5.9
19	3.0		5.9
20	3.0		5.9
21		5.0	5.9
24		4.5	5.8
25			5.5

APPENDIX TABLE 6.

ALBERTA ALL SPECIMENS (100)

PRINCIPAL COMPONENTS ANALYSIS - CLUSTAN 1C

NUMBER OF CASES = 100  
 NUMBER OF BINARY VARIABLES = 0  
 NUMBER OF NUMERIC VARIABLES = 21

(UNTRANSFORMED DATA)

NUMERIC MEANS AND STANDARD DEVIATIONS

VARIABLE MINIMUM VALUE

MAXIMUM VALUE

1	7.2409	1.4340
2	1.7565	0.3337
3	0.3299	0.0864
4	0.5926	0.1791
5	0.5907	0.1215
6	0.5604	0.1241
7	0.4187	0.1781
8	0.4321	0.0953
9	0.5906	0.1371
10	0.1184	0.0949
11	0.1890	0.1986
12	0.4265	0.2826
13	1.0150	0.2361
14	3.7543	0.7397
15	1.7702	0.3736
16	4.5683	0.9460
17	2.9569	0.6750
18	0.5628	0.1435
19	65.0900	4.1734
20	12.1700	1.3078
21	25.9200	3.0309

1	4.3000	11.0000
2	1.0300	2.7100
3	0.1800	0.5700
4	0.2700	1.0000
5	0.3400	0.9600
6	0.2900	0.8700
7	0.4500	1.3200
8	0.2500	0.7200
9	0.3000	0.9100
10	0.0000	0.3000
11	0.0000	0.5700
12	0.0700	0.3800
13	0.5000	1.9900
14	2.2200	5.4400
15	1.9400	2.5500
16	2.6000	7.4300
17	1.5800	5.1900
18	0.2400	0.8900
19	58.0000	74.0000
20	9.0000	15.0000
21	21.0000	33.0000

EIGENVALUES

12.48	5.95	0.74	0.55	0.34	0.25	0.19	0.31	0.07	0.06	0.05	0.05	0.04	0.03	0.03	0.02
0.02	0.01	0.01	0.01	0.00											

PERCENTAGE VARIANCE

59.41	28.31	3.53	2.60	1.61	1.20	0.92	0.51	0.36	0.30	0.25	0.23	0.19	0.13	0.12	0.10
0.09	0.06	0.04	0.03	0.02											

CUMULATIVE VARIANCE

59.41	87.72	91.25	93.85	95.46	96.66	97.57	98.08	98.44	98.73	98.98	99.21	99.41	99.54	99.66	99.76
99.85	99.91	99.96	99.98	100.00											

EIGENVECTORS - BY ROWS

VECTOR 1

0.279 0.277 0.148 0.204 0.245 0.274 0.272 0.249 0.258 0.117 0.051 0.085 0.242 0.280 0.272 0.276 0.270 0.241  
 0.037 0.010 0.028

VECTOR 2

-0.046 0.022 -0.339 0.270 -0.188 0.017 -0.030 -0.164 0.147 0.361 0.396 0.362 -0.020 -0.019 0.022 -0.365 -0.085 -0.057  
 0.273 -0.277 0.354

VECTOR 3

-0.001 -0.051 0.012 -0.116 0.006 0.078 -0.048 -0.001 -0.051 -0.119 0.012 -0.037 0.231 0.009 -0.016 -0.304 0.074 -0.216  
 0.711 0.566 0.160

FIRST 3 FACTOR SCORES

S	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	S	22	23	24	25	26	27	28	29	30	31	32	33
S 1	4.145	-2.872	-0.505	S 34	0.051	-2.940	-0.962	S 67	26.154	1.644	0.284																							
S 2	-0.071	-2.394	0.522	S 35	2.279	-2.749	-0.247	S 68	-7.859	1.398	-1.080																							
S 3	-0.879	-2.168	-0.270	S 36	2.513	-3.350	-0.184	S 69	-2.476	2.113	0.266																							
S 4	-1.593	-1.603	0.121	S 37	1.361	-2.103	-0.247	S 70	-4.204	1.934	0.306																							
S 5	1.135	-2.766	0.225	S 38	-0.632	-2.778	-0.323	S 71	-4.235	1.407	-0.067																							
S 6	2.405	-3.050	-0.113	S 39	-3.181	-2.123	-0.747	S 72	-5.049	1.381	1.545																							
S 7	-3.034	-1.847	-0.801	S 40	-4.721	-2.035	-0.542	S 73	0.792	1.769	-0.028																							
S 8	-1.152	-2.060	1.132	S 41	3.767	-3.075	0.715	S 74	-2.561	2.066	0.327																							
S 9	-0.307	-2.815	-0.285	S 42	1.154	-2.434	2.304	S 75	-0.835	1.182	0.334																							
S 10	-1.069	-2.299	0.813	S 43	4.009	-2.336	0.593	S 76	-1.889	2.031	0.371																							
S 11	-4.725	-2.310	-0.058	S 44	3.710	-3.116	-0.378	S 77	-2.716	1.650	-1.311																							
S 12	-0.950	-2.533	0.566	S 45	3.327	-2.106	0.395	S 78	4.766	3.001	-0.268																							
S 13	-7.352	-1.778	-0.772	S 46	4.466	-2.814	-0.558	S 79	3.034	2.544	0.979																							
S 14	-0.455	-2.369	0.367	S 47	2.853	-2.513	-0.026	S 80	3.206	2.517	-0.607																							
S 15	-6.332	-1.909	-1.239	S 48	2.370	-2.467	-0.723	S 81	2.853	2.637	-1.452																							
S 16	-0.623	-2.574	0.903	S 49	0.025	-2.463	0.368	S 82	2.303	1.991	0.743																							
S 17	4.054	-2.646	1.810	S 50	2.295	-2.183	-1.115	S 83	2.449	2.348	0.012																							
S 18	1.813	-2.181	-0.843	S 51	-0.052	-2.724	0.605	S 84	2.610	2.588	0.283																							
S 19	-1.110	-3.095	1.025	S 52	-0.815	-2.333	-0.250	S 85	4.337	2.153	1.089																							
S 20	-3.162	-2.740	-0.7464	S 53	-1.254	1.964	0.514	S 86	4.474	2.969	0.123																							
S 21	-2.055	-1.969	-0.416	S 54	-3.489	-2.371	0.644	S 87	2.787	2.325	0.331																							
S 22	-4.133	-1.525	-0.889	S 55	-4.314	1.285	-0.866	S 88	1.770	2.487	0.610																							
S 23	-4.535	-2.079	-0.972	S 56	4.542	3.484	-0.675	S 89	2.476	2.416	1.622																							
S 24	-1.513	-1.421	-0.297	S 57	1.517	2.916	0.422	S 90	0.734	2.497	0.086																							
S 25	-1.734	-3.308	0.193	S 58	2.932	2.931	0.324	S 91	0.558	2.388	1.421																							
S 26	-0.406	-2.094	0.185	S 59	0.604	3.615	0.608	S 92	3.457	2.680	-0.341																							
S 27	-1.109	-2.155	-0.369	S 60	4.044	4.124	-0.324	S 93	2.559	2.066	-0.379																							
S 28	-1.962	-1.865	-0.912	S 61	1.307	3.773	-0.869	S 94	0.465	2.760	-0.426																							
S 29	-1.789	-2.008	-1.089	S 62	-2.451	-2.082	-0.207	S 95	0.843	2.645	-1.725																							
S 30	-3.942	-1.400	1.521	S 63	-1.900	2.560	1.534	S 96	-0.152	1.897	-2.542																							
S 31	-5.451	-1.379	-0.491	S 64	-3.024	2.363	1.836	S 97	-0.426	1.797	-1.695																							
S 32	-3.425	-1.702	0.777	S 65	-3.624	1.961	1.282	S 98	30.336	1.885	-1.528																							
S 33	5.841	-3.740	-0.084	S 66	-5.815	2.173	-0.279	S 99	2.914	2.807	-1.298																							
								S 100	15.572	2.788	0.083																							

APPENDIX TABLE 7.

HAWAIIANIS & SPECIMENS (137)  
 NUMBER OF CASES = 137  
 NUMBER OF BINARY VARIABLES = 21  
 NUMBER OF BURNIC VARIABLES = 0

PRINCIPAL COMPONENTS ANALYSIS - CLUSTAN 1C  
 (UNTRANSFORMED DATA)

NUMERIC MEANS AND STANDARD DEVIATIONS					VARIABLE MINIMUM VALUE					MAXIMUM VALUE				
1	7.4072	1.9825	1	3.4000	12.3000	2	6.9000	2.6900	3	0.1800	0.6000	4	0.2200	1.1100
2	1.7632	0.4117	3	0.2000	0.6000	5	0.3100	1.0500	6	1.2800	1.1000	7	0.4000	1.5800
3	0.3419	0.0951	4	0.2200	1.1100	8	0.2300	0.9600	9	0.2600	1.0500	10	0.0000	0.3700
4	0.5959	0.2058	5	0.3100	1.0500	11	0.0000	0.6200	12	0.0800	0.9200	13	0.4000	1.7800
5	0.5976	0.1584	6	1.2800	1.1000	14	1.8000	6.2600	15	0.7500	2.8800	16	2.1900	8.1500
6	0.6171	0.1974	7	0.4000	1.5800	17	1.3500	5.7200	18	0.2100	1.0000	19	59.0000	73.0000
7	0.8950	0.2665	8	0.2300	0.9600	20	16.0000	16.0000	21	21.0000	32.0000			
8	0.4772	0.1638	9	0.2600	1.0500									
9	0.4323	0.1058	10	0.0000	0.3700									
10	0.1104	0.0950	11	0.0000	0.6200									
11	0.1746	0.1833	12	0.0800	0.9200									
12	0.1923	0.2401	13	0.4000	1.7800									
13	1.0393	0.3338	14	1.8000	6.2600									
14	3.8056	1.0151	15	0.7500	2.8800									
15	1.7936	0.4759	16	2.1900	8.1500									
16	4.7460	1.3169	17	1.3500	5.7200									
17	3.0867	0.9467	18	0.2100	1.0000									
18	0.5325	0.1482	19	59.0000	73.0000									
19	65.4234	3.5144	20	16.0000	16.0000									
20	12.9927	1.2237	21	21.0000	32.0000									
21	25.9854	2.7813												

EIGENVALUES															
13.24	4.26	1.02	0.56	0.33	0.20	0.14	0.10	0.08	0.06	0.06	0.05	0.05	0.04	0.03	0.02
0.02	0.01	0.01	0.00	0.00											

PERCENTAGE VARIANCE															
66.36	20.31	4.87	2.68	1.55	0.96	0.66	0.50	0.39	0.31	0.30	0.25	0.24	0.17	0.13	0.10
0.08	0.06	0.04	0.02	0.02											

CUMULATIVE VARIANCE															
66.36	86.67	91.53	94.21	95.76	96.72	97.38	97.88	98.27	98.58	98.88	99.13	99.37	99.54	99.68	99.78
99.86	99.93	99.96	99.98	100.00											

EIGENVECTORS - BY ROWS

VECTOR 1

0.260 0.264 0.144 0.244 0.246 0.259 0.263 0.250 0.262 0.173 0.115 0.146 0.259 0.262 0.255 0.262 0.255 0.216  
 0.007-0.068 0.065

VECTOR 2

-0.104-0.036-0.387 0.179-0.166 0.059-0.013 0.001 0.061 0.337 0.422 0.379-0.037-0.074-0.086-0.084-0.121-0.195  
 0.189-0.279 0.385

VECTOR 3

0.052 0.015 0.006-0.018 0.050-0.027-0.019 0.015-0.006-0.065-0.024-0.068-0.003 0.037-0.031 0.027 0.032-0.022  
 0.826 0.501 0.204

FIRST	3	FACTOR SCORES	FIRST	3	FACTOR SCORES	FIRST	3	FACTOR SCORES			
S 1	6.445	2.416	-0.487	S 47	-2.826	2.453	-1.171	S 93	-2.898	-0.808	0.176
S 2	6.093	1.320	-1.915	S 48	-2.903	2.014	0.126	S 94	-2.145	-1.145	0.178
S 3	4.649	2.691	-0.796	S 49	-2.690	2.191	-0.037	S 95	-2.665	-2.158	-0.118
S 4	3.005	2.916	0.141	S 50	-3.834	2.060	-0.441	S 96	-1.738	-1.476	-1.653
S 5	5.708	2.193	0.732	S 51	4.134	-3.046	0.551	S 97	-1.068	-1.989	-0.165
S 6	5.022	2.399	-0.223	S 52	6.635	-3.558	1.445	S 98	-4.417	-1.148	0.585
S 7	5.440	3.084	0.602	S 53	7.741	-4.427	-0.768	S 99	-5.659	-0.693	-0.603
S 8	6.024	2.796	0.237	S 54	5.952	-3.670	0.105	S 100	-1.121	-1.371	-0.646
S 9	3.863	4.374	-0.094	S 55	3.584	-2.924	1.085	S 101	0.280	-1.726	-1.479
S 10	4.075	4.134	0.973	S 56	4.481	-3.217	-0.766	S 102	-0.565	-0.987	1.310
S 11	5.469	2.488	0.973	S 57	2.545	-2.266	1.510	S 103	-2.023	0.617	0.446
S 12	6.794	1.571	0.251	S 58	2.700	-3.098	0.022	S 104	-2.354	-0.310	0.225
S 13	6.207	3.383	0.918	S 59	1.357	-2.667	0.198	S 105	-0.038	-1.731	0.484
S 14	6.046	2.506	-1.152	S 60	0.629	-2.129	1.634	S 106	-0.946	-0.613	1.455
S 15	4.968	4.032	1.130	S 61	2.555	-3.715	-0.392	S 107	-0.251	-0.471	1.333
S 16	0.777	1.726	-0.122	S 62	-0.710	-2.134	-1.102	S 108	-1.807	-0.417	1.264
S 17	-2.160	1.871	-0.306	S 63	-1.061	-1.808	1.186	S 109	-2.527	0.099	1.124
S 18	-0.956	1.651	-1.400	S 64	-1.970	-1.941	-0.881	S 110	-2.859	0.398	1.233
S 19	-3.015	1.440	-1.508	S 65	-0.945	-1.789	-2.028	S 111	-3.363	1.097	0.493
S 20	1.270	1.665	0.034	S 66	-1.149	-1.715	0.101	S 112	-3.927	0.359	3.088
S 21	3.486	3.094	-1.553	S 67	-1.549	-1.712	-1.034	S 113	-3.070	0.364	1.575
S 22	4.471	3.274	-1.776	S 68	-0.572	-1.204	0.399	S 114	-5.123	0.161	0.935
S 23	2.058	2.711	0.215	S 69	-2.801	-1.504	-1.443	S 115	-4.311	0.911	1.530
S 24	6.024	2.386	-0.117	S 70	3.330	-3.025	-0.328	S 116	-4.637	-0.855	1.202
S 25	5.687	1.538	-1.476	S 71	3.371	-2.485	4.051	S 117	-4.036	-0.200	-0.395
S 26	2.009	2.513	-0.436	S 72	5.255	-3.621	-0.529	S 118	-3.262	1.194	0.395
S 27	0.037	2.240	-1.264	S 73	5.904	-3.049	1.428	S 119	-4.815	-0.445	-1.165
S 28	4.533	1.744	0.755	S 74	1.331	-2.031	0.363	S 120	-1.824	-0.578	-0.241
S 29	-2.918	2.497	-0.388	S 75	1.070	-2.057	-0.172	S 121	-1.165	-0.030	0.286
S 30	-2.861	2.617	1.103	S 76	-2.411	-1.218	-0.430	S 122	-3.352	0.111	-0.134
S 31	-2.907	2.023	0.957	S 77	2.295	-2.788	-0.187	S 123	-2.694	-0.045	-0.325
S 32	-2.566	2.039	-0.495	S 78	0.612	-1.993	-0.614	S 124	-3.945	-1.458	-3.433
S 33	-3.146	1.652	0.460	S 79	0.292	-1.126	-0.224	S 125	-3.102	-0.515	2.274
S 34	-3.200	2.110	1.489	S 80	-0.267	-1.780	0.184	S 126	-5.647	0.091	-0.184
S 35	-2.881	0.932	-0.485	S 81	-1.751	-2.093	-0.169	S 127	-7.469	-0.249	-0.505
S 36	-4.449	1.019	0.127	S 82	1.244	-2.335	0.791	S 128	-3.301	0.779	1.848
S 37	-3.650	3.021	0.613	S 83	4.917	-2.657	0.056	S 129	-2.152	0.224	1.375
S 38	3.210	2.284	-1.206	S 84	0.737	-2.309	-0.139	S 130	-2.921	0.015	-0.338
S 39	0.208	2.869	0.011	S 85	-0.219	-2.384	-0.149	S 131	-4.163	-0.447	-3.379
S 40	-0.368	1.355	0.482	S 86	-0.104	-2.121	-1.670	S 132	-3.412	-0.539	0.687
S 41	-0.294	0.889	-1.428	S 87	-3.802	-0.902	0.641	S 133	-5.946	0.030	0.639
S 42	0.134	2.622	1.400	S 88	-4.980	-0.866	-0.480	S 134	-1.970	-0.233	-0.934
S 43	-2.367	4.810	0.289	S 89	5.157	-1.779	-0.637	S 135	42.056	0.661	-0.310
S 44	-3.170	2.093	-1.339	S 90	3.707	-2.977	-0.163	S 136	4.688	0.600	1.421
S 45	-1.955	1.685	-1.383	S 91	-4.531	-1.440	-4.579	S 137	-0.744	0.410	0.270
S 46	-0.043	2.733	0.057	S 92	-3.466	-1.331	-1.937				



### APPENDIX TABLE 9.

KANAWASKIS ALL SPECIMENS (137)

NUMBER OF CASES = 137  
 NUMBER OF BINARY VARIABLES = 21  
 NUMBER OF NUMERIC VARIABLES = 21

PRINCIPAL COMPONENTS ANALYSIS - CLUSTAN 1C

(SQUARE-ROOT TRANSFORMED DATA)

NUMERIC MEANS AND STANDARD DEVIATIONS			VARIABLE MINIMUM VALUE			MAXIMUM VALUE		
1	2.6979	0.3595	1	1.8439	3.5071			
2	1.3196	0.1515	2	4.9487	1.6401			
3	0.5792	0.0804	3	0.4243	0.7746			
4	0.7611	0.1296	4	0.4690	1.0536			
5	0.7664	0.1011	5	0.5568	1.3247			
6	0.7760	0.1222	6	0.5292	1.0438			
7	0.9356	0.1391	7	0.6325	1.2570			
8	0.6815	0.1137	8	0.4796	0.9798			
9	0.7855	0.1241	9	0.5099	1.0247			
10	0.4932	0.1567	10	0.0000	0.8083			
11	0.3134	0.2774	11	0.0000	0.7874			
12	0.5980	0.1641	12	0.2626	0.4592			
13	1.0066	0.1623	13	0.6633	1.3342			
14	1.3339	0.2569	14	1.3416	2.5020			
15	1.2775	0.1773	15	0.6660	1.6971			
16	2.1575	0.3027	16	1.4799	2.8548			
17	1.7369	0.2654	17	1.1619	2.3917			
18	0.7229	0.1000	18	0.4589	1.0000			
19	65.4234	3.5143	19	50.0000	70.0000			
20	12.9927	1.2217	20	10.0000	16.0000			
21	25.9854	2.7813	21	20.0000	32.0000			

EIGENVALUES

13.83	4.24	1.01	0.57	0.30	0.24	0.20	0.13	0.10	0.07	0.06	0.06	0.05	0.04	0.03	0.02
0.02	0.01	0.01	0.00	0.00											

PERCENTAGE VARIANCE

65.87	20.19	4.79	2.69	1.44	1.16	0.94	0.62	0.44	0.33	0.30	0.27	0.24	0.18	0.14	0.11
0.10	0.06	0.04	0.02	0.02											

CUMULATIVE VARIANCE

65.87	86.06	90.85	93.55	94.99	96.15	97.08	97.71	98.18	98.52	98.81	99.09	99.33	99.50	99.65	99.76
95.86	99.92	99.96	99.98	99.99	100.00										

EIGENVECTORS - BY ROWS

VECTOR 1

0.263 0.265 0.154 0.244 0.250 0.259 0.264 0.262 0.262 0.166 0.008 0.133 0.201 0.265 0.259 0.265 0.258 0.221  
 0.005-0.064 0.059

VECTOR 2

-0.067-0.020-0.379 0.183-0.150 0.063-0.010-0.010 0.072 0.321 0.448 0.389-0.027-0.059-0.072-0.069-0.103-0.175  
 0.203-0.274 0.397

VECTOR 3

0.052 0.010 0.031-0.023 0.058-0.025-0.014 0.007-0.010-0.043-0.611-0.073-0.014 0.034 0.023 0.025 0.031-0.027  
 0.612 0.542 0.166

FIRST 3 FACTOR SCORES

S 1	6.033	2.286	-0.518	S 47	-2.880	2.657	-1.309	S 93	-2.783	-1.071	0.187
S 2	5.738	1.257	-1.878	S 48	-1.451	2.784	-0.030	S 94	-1.920	-2.269	0.306
S 3	4.453	2.442	-0.763	S 49	-2.669	2.390	-0.154	S 95	-2.473	-2.387	0.025
S 4	7.271	2.607	0.151	S 50	-4.006	2.292	-0.545	S 96	-1.502	-1.744	-1.637
S 5	5.368	1.980	0.806	S 51	4.257	-2.787	0.554	S 97	-0.751	-2.130	-0.100
S 6	4.849	2.372	-0.284	S 52	6.450	-3.070	1.418	S 98	-4.561	-1.510	0.680
S 7	5.522	2.938	0.535	S 53	7.295	-3.797	-0.775	S 99	-6.292	-1.320	-0.539
S 8	5.669	2.855	0.172	S 54	5.826	-3.408	0.106	S100	-0.806	-1.480	-0.652
S 9	7.286	1.881	1.125	S 55	3.709	-2.642	1.103	S101	0.458	-1.785	-1.394
S10	4.019	2.274	-0.173	S 56	4.508	-2.950	-0.784	S102	-0.759	-1.094	1.314
S11	5.200	2.351	-1.018	S 57	2.775	-2.171	1.466	S103	-1.836	0.889	0.429
S12	6.384	1.583	0.234	S 58	2.945	-2.927	0.061	S104	-2.181	-1.189	0.199
S13	5.642	3.034	0.694	S 59	1.691	-2.582	0.255	S105	0.338	-1.501	0.588
S14	5.688	2.257	-1.114	S 60	0.984	-2.194	1.673	S106	-0.732	-0.431	1.166
S15	4.803	2.016	1.120	S 61	2.809	-3.583	-0.302	S107	0.604	0.215	1.104
S16	0.990	1.623	-0.346	S 62	-0.407	-2.329	-1.264	S108	-1.523	-0.491	1.277
S17	-2.140	1.972	-0.283	S 63	-0.760	-1.994	-1.144	S109	-2.410	0.146	1.100
S18	-0.790	1.733	-1.647	S 64	-1.760	-2.248	-0.619	S110	-2.739	0.566	1.223
S19	-3.033	1.645	-1.532	S 65	-0.829	-1.929	-2.081	S111	-3.389	1.240	0.418
S20	1.813	1.643	0.101	S 66	-0.827	-1.851	0.356	S112	-3.596	0.734	3.097
S21	3.407	2.929	-1.581	S 67	-1.283	-1.917	-0.997	S113	-2.973	0.549	1.590
S22	4.272	3.026	-1.794	S 68	-4.827	-1.706	0.493	S114	-5.461	0.222	0.963
S23	2.094	2.678	0.209	S 69	-2.773	-2.044	-1.374	S115	-3.292	1.103	1.475
S24	5.656	2.202	-0.092	S 70	3.557	-2.811	-0.494	S116	-4.849	-1.107	1.275
S25	5.371	1.453	-1.492	S 71	3.583	-2.308	2.078	S117	-4.337	-0.558	-0.731
S26	-2.124	2.505	-0.505	S 72	5.794	-3.161	-0.540	S118	-3.224	1.414	0.919
S27	0.205	2.333	-1.361	S 73	5.804	-2.607	1.493	S119	-4.948	-0.728	-1.137
S28	4.434	1.716	0.780	S 74	1.674	-2.039	0.355	S120	-1.751	-1.117	-0.180
S29	-2.971	2.654	-0.512	S 75	1.410	-2.069	-0.154	S121	-0.861	0.197	0.268
S30	-2.975	2.860	0.967	S 76	-2.217	-1.416	-0.373	S122	-3.322	0.257	-0.114
S31	-3.026	3.191	0.859	S 77	2.559	-2.716	0.228	S123	-2.531	0.108	-0.337
S32	-3.014	2.192	-0.560	S 78	0.962	-1.997	-0.572	S124	-4.099	-1.670	-1.382
S33	-3.210	2.025	0.413	S 79	0.666	-2.146	-0.162	S125	-3.139	-0.674	2.348
S34	-3.283	2.339	1.431	S 80	0.049	-1.973	0.217	S126	-6.298	-0.436	-0.987
S35	-2.870	1.096	-0.420	S 81	-1.481	-2.287	-0.041	S127	-6.766	-0.516	-0.434
S36	-4.748	0.035	0.176	S 82	-1.593	-2.319	0.853	S128	-1.108	0.980	1.758
S37	-3.840	0.218	0.463	S 83	4.940	-2.234	0.509	S129	-1.971	-0.402	1.367
S38	3.259	2.267	-1.281	S 84	1.099	-2.307	-0.075	S130	-2.955	-0.130	0.315
S39	5.595	2.799	-0.063	S 85	0.051	-2.535	-0.060	S131	-4.286	-0.475	-1.374
S40	-0.206	2.493	0.402	S 86	0.201	-2.250	-1.613	S132	-3.852	0.615	0.591
S41	-0.128	0.733	-1.415	S 87	-3.887	-1.329	0.661	S133	-6.580	0.755	0.634
S42	0.263	2.620	1.488	S 88	-5.401	-1.888	-0.448	S134	-1.735	-0.052	-0.944
S43	-2.372	2.666	0.161	S 89	5.301	-3.165	-0.028	S135	-1.868	0.801	-0.349
S44	-3.274	1.256	-1.412	S 90	3.904	-2.733	-0.082	S136	-4.644	0.967	1.344
S45	-1.834	1.777	-1.416	S 91	-4.607	-2.052	-0.464	S137	-0.433	0.651	0.219
S46	0.004	2.716	0.014	S 92	-3.497	-1.730	-1.900				

APPENDIX TABLE 10.

MOORE - STERILIZATION DISCUSSION ANALYSIS  
PLANT SCIENCES COMPUTING FACILITY, UCL

NUMBER OF VARIABLES 41  
NUMBER OF GROUPS 5  
NUMBER OF CASES IN EACH GROUP 50 517  
PULSE PROBABILITIES 0.1011 0.1011 0.1011 0.0  
VARIABLES

Table with columns: VARIABLE, GROUP, F, C, E. Lists 41 variables and their group assignments.

(UNTRANSFORMED DATA)  
ALBERTA GROUPS P, A, B, C, D, E  
MANITOWA GROUP P, A, B, C, D, E

STEREOTYPIC STATISTICS

Table with columns: VARIABLE, GROUP, F, C, E. Lists 41 variables and their group assignments.

SUBROUTINE TABLE

STEP VARIABLE DELETED TESTS OF HOMOGENEITY NUMBER OF VARIABLES INCLUDED

Table with columns: STEP, VARIABLE, DELETED, TESTS OF HOMOGENEITY, NUMBER OF VARIABLES INCLUDED. Lists steps 1 through 21.

EIGENVALUES

Table with columns: EIGENVALUES. Lists eigenvalues for steps 1 through 21.

CORRELATION COEFFICIENTS

Table with columns: CORRELATION COEFFICIENTS. Lists correlation coefficients for steps 1 through 21.

COEFFICIENTS FOR LOGNORMAL TRANSFORM

Table with columns: COEFFICIENTS FOR LOGNORMAL TRANSFORM. Lists coefficients for steps 1 through 21.

1. FIRST LOGNORMAL TRANSFORM

Table with columns: GROUP, CASE, F, REAR COEFFICIENTS, W, X, Y, Z, S, T. Shows data for the first lognormal transform.

2. SECOND LOGNORMAL TRANSFORM

Table with columns: GROUP, CASE, F, REAR COEFFICIENTS, W, X, Y, Z, S, T. Shows data for the second lognormal transform.

3. FIRST LOGNORMAL TRANSFORM

Table with columns: GROUP, CASE, F, REAR COEFFICIENTS, W, X, Y, Z, S, T. Shows data for the third lognormal transform.

APPENDIX TABLE 11.

PROBABILITY ESTIMATION ANALYSIS  
HEALTH SCIENCES COMPUTING FACILITY, UCRL

NUMBER OF VARIABLES 20  
NUMBER OF CASES IN EACH GROUP 50 50 1117  
GROUP PROBABILITIES 0.1114 0.1113 0.1117

COMPONENTS: TRACK, MIND, CT, TALKING, CHARACTER, AGENCY, TONE, A, THERM, PERSN, DIST, PARADOX, ORIENT, A, A, THERM, TALKING, ORIENT

FIRST CARNIVAL VARIABLE  
SECOND CARNIVAL VARIABLE

Table with columns: GROUP, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION. Rows 1-10.

FIRST CARNIVAL VARIABLE  
SECOND CARNIVAL VARIABLE

Table with columns: GROUP, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION. Rows 1-10.

FIRST CARNIVAL VARIABLE  
SECOND CARNIVAL VARIABLE

Table with columns: GROUP, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION. Rows 1-10.

FIRST CARNIVAL VARIABLE  
SECOND CARNIVAL VARIABLE

Table with columns: GROUP, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION. Rows 1-10.

FIRST CARNIVAL VARIABLE  
SECOND CARNIVAL VARIABLE

Table with columns: GROUP, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION, CASE, F, REAR CORRELATION. Rows 1-10.

FIRST CARNIVAL VARIABLE  
SECOND CARNIVAL VARIABLE

SUPPLEMENT TABLE

STEP VARIABLE NUMBER OF VARIABLES INCLUDED

Table with columns: STEP, VARIABLE, NUMBER OF VARIABLES INCLUDED. Rows 1-20.

REGRESSIONS

Table with columns: REGRESSION, REGRESSION, REGRESSION, REGRESSION, REGRESSION, REGRESSION. Rows 1-20.

CORRELATIONS

Table with columns: CORRELATION, CORRELATION, CORRELATION, CORRELATION, CORRELATION, CORRELATION. Rows 1-20.

Coefficients for Canonical Variables

Table with columns: COEFFICIENT, COEFFICIENT, COEFFICIENT, COEFFICIENT, COEFFICIENT, COEFFICIENT. Rows 1-20.

CONSTANT

Table with columns: CONSTANT, CONSTANT, CONSTANT, CONSTANT, CONSTANT, CONSTANT. Rows 1-20.



APPENDIX TABLE 12.

GROUP 1 - FIRST CHEMICAL TABLE  
GROUP 2 - SECOND CHEMICAL TABLE

Table with columns: CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T. Rows 1-10.

GROUP 3 - FIRST CHEMICAL TABLE  
GROUP 4 - SECOND CHEMICAL TABLE

Table with columns: CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T. Rows 1-10.

GROUP 5 - FIRST CHEMICAL TABLE  
GROUP 6 - SECOND CHEMICAL TABLE

Table with columns: CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T. Rows 1-10.

GROUP 7 - FIRST CHEMICAL TABLE  
GROUP 8 - SECOND CHEMICAL TABLE

Table with columns: CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T. Rows 1-10.

GROUP 9 - FIRST CHEMICAL TABLE  
GROUP 10 - SECOND CHEMICAL TABLE

Table with columns: CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T. Rows 1-10.

GROUP 11 - FIRST CHEMICAL TABLE  
GROUP 12 - SECOND CHEMICAL TABLE

Table with columns: CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T, CASE, I, T. Rows 1-10.





APPENDIX TABLE 15.

1. QUANTUM TRANSFORMER DATA  
A. DATA SOURCE: 1. 2. 3. 4. 5. 6. 7. 8. 9. 10. 11. 12. 13. 14. 15. 16. 17. 18. 19. 20. 21. 22. 23. 24. 25. 26. 27. 28. 29. 30. 31. 32. 33. 34. 35. 36. 37. 38. 39. 40. 41. 42. 43. 44. 45. 46. 47. 48. 49. 50. 51. 52. 53. 54. 55. 56. 57. 58. 59. 60. 61. 62. 63. 64. 65. 66. 67. 68. 69. 70. 71. 72. 73. 74. 75. 76. 77. 78. 79. 80. 81. 82. 83. 84. 85. 86. 87. 88. 89. 90. 91. 92. 93. 94. 95. 96. 97. 98. 99. 100. 101. 102. 103. 104. 105. 106. 107. 108. 109. 110. 111. 112. 113. 114. 115. 116. 117. 118. 119. 120. 121. 122. 123. 124. 125. 126. 127. 128. 129. 130. 131. 132. 133. 134. 135. 136. 137. 138. 139. 140. 141. 142. 143. 144. 145. 146. 147. 148. 149. 150. 151. 152. 153. 154. 155. 156. 157. 158. 159. 160. 161. 162. 163. 164. 165. 166. 167. 168. 169. 170. 171. 172. 173. 174. 175. 176. 177. 178. 179. 180. 181. 182. 183. 184. 185. 186. 187. 188. 189. 190. 191. 192. 193. 194. 195. 196. 197. 198. 199. 200. 201. 202. 203. 204. 205. 206. 207. 208. 209. 210. 211. 212. 213. 214. 215. 216. 217. 218. 219. 220. 221. 222. 223. 224. 225. 226. 227. 228. 229. 230. 231. 232. 233. 234. 235. 236. 237. 238. 239. 240. 241. 242. 243. 244. 245. 246. 247. 248. 249. 250. 251. 252. 253. 254. 255. 256. 257. 258. 259. 260. 261. 262. 263. 264. 265. 266. 267. 268. 269. 270. 271. 272. 273. 274. 275. 276. 277. 278. 279. 280. 281. 282. 283. 284. 285. 286. 287. 288. 289. 290. 291. 292. 293. 294. 295. 296. 297. 298. 299. 300. 301. 302. 303. 304. 305. 306. 307. 308. 309. 310. 311. 312. 313. 314. 315. 316. 317. 318. 319. 320. 321. 322. 323. 324. 325. 326. 327. 328. 329. 330. 331. 332. 333. 334. 335. 336. 337. 338. 339. 340. 341. 342. 343. 344. 345. 346. 347. 348. 349. 350. 351. 352. 353. 354. 355. 356. 357. 358. 359. 360. 361. 362. 363. 364. 365. 366. 367. 368. 369. 370. 371. 372. 373. 374. 375. 376. 377. 378. 379. 380. 381. 382. 383. 384. 385. 386. 387. 388. 389. 390. 391. 392. 393. 394. 395. 396. 397. 398. 399. 400. 401. 402. 403. 404. 405. 406. 407. 408. 409. 410. 411. 412. 413. 414. 415. 416. 417. 418. 419. 420. 421. 422. 423. 424. 425. 426. 427. 428. 429. 430. 431. 432. 433. 434. 435. 436. 437. 438. 439. 440. 441. 442. 443. 444. 445. 446. 447. 448. 449. 450. 451. 452. 453. 454. 455. 456. 457. 458. 459. 460. 461. 462. 463. 464. 465. 466. 467. 468. 469. 470. 471. 472. 473. 474. 475. 476. 477. 478. 479. 480. 481. 482. 483. 484. 485. 486. 487. 488. 489. 490. 491. 492. 493. 494. 495. 496. 497. 498. 499. 500. 501. 502. 503. 504. 505. 506. 507. 508. 509. 510. 511. 512. 513. 514. 515. 516. 517. 518. 519. 520. 521. 522. 523. 524. 525. 526. 527. 528. 529. 530. 531. 532. 533. 534. 535. 536. 537. 538. 539. 540. 541. 542. 543. 544. 545. 546. 547. 548. 549. 550. 551. 552. 553. 554. 555. 556. 557. 558. 559. 560. 561. 562. 563. 564. 565. 566. 567. 568. 569. 570. 571. 572. 573. 574. 575. 576. 577. 578. 579. 580. 581. 582. 583. 584. 585. 586. 587. 588. 589. 590. 591. 592. 593. 594. 595. 596. 597. 598. 599. 600. 601. 602. 603. 604. 605. 606. 607. 608. 609. 610. 611. 612. 613. 614. 615. 616. 617. 618. 619. 620. 621. 622. 623. 624. 625. 626. 627. 628. 629. 630. 631. 632. 633. 634. 635. 636. 637. 638. 639. 640. 641. 642. 643. 644. 645. 646. 647. 648. 649. 650. 651. 652. 653. 654. 655. 656. 657. 658. 659. 660. 661. 662. 663. 664. 665. 666. 667. 668. 669. 670. 671. 672. 673. 674. 675. 676. 677. 678. 679. 680. 681. 682. 683. 684. 685. 686. 687. 688. 689. 690. 691. 692. 693. 694. 695. 696. 697. 698. 699. 700. 701. 702. 703. 704. 705. 706. 707. 708. 709. 710. 711. 712. 713. 714. 715. 716. 717. 718. 719. 720. 721. 722. 723. 724. 725. 726. 727. 728. 729. 730. 731. 732. 733. 734. 735. 736. 737. 738. 739. 740. 741. 742. 743. 744. 745. 746. 747. 748. 749. 750. 751. 752. 753. 754. 755. 756. 757. 758. 759. 760. 761. 762. 763. 764. 765. 766. 767. 768. 769. 770. 771. 772. 773. 774. 775. 776. 777. 778. 779. 780. 781. 782. 783. 784. 785. 786. 787. 788. 789. 790. 791. 792. 793. 794. 795. 796. 797. 798. 799. 800. 801. 802. 803. 804. 805. 806. 807. 808. 809. 810. 811. 812. 813. 814. 815. 816. 817. 818. 819. 820. 821. 822. 823. 824. 825. 826. 827. 828. 829. 830. 831. 832. 833. 834. 835. 836. 837. 838. 839. 840. 841. 842. 843. 844. 845. 846. 847. 848. 849. 850. 851. 852. 853. 854. 855. 856. 857. 858. 859. 860. 861. 862. 863. 864. 865. 866. 867. 868. 869. 870. 871. 872. 873. 874. 875. 876. 877. 878. 879. 880. 881. 882. 883. 884. 885. 886. 887. 888. 889. 890. 891. 892. 893. 894. 895. 896. 897. 898. 899. 900. 901. 902. 903. 904. 905. 906. 907. 908. 909. 910. 911. 912. 913. 914. 915. 916. 917. 918. 919. 920. 921. 922. 923. 924. 925. 926. 927. 928. 929. 930. 931. 932. 933. 934. 935. 936. 937. 938. 939. 940. 941. 942. 943. 944. 945. 946. 947. 948. 949. 950. 951. 952. 953. 954. 955. 956. 957. 958. 959. 960. 961. 962. 963. 964. 965. 966. 967. 968. 969. 970. 971. 972. 973. 974. 975. 976. 977. 978. 979. 980. 981. 982. 983. 984. 985. 986. 987. 988. 989. 990. 991. 992. 993. 994. 995. 996. 997. 998. 999. 1000.

STATISTICAL DATA  
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Table with 10 columns: GROUP, CASE, T, CASE, T, CASE, T, CASE, T, CASE. Rows 1-100.

Table with 10 columns: GROUP, CASE, T, CASE, T, CASE, T, CASE, T, CASE. Rows 1-100.

Table with 10 columns: GROUP, CASE, T, CASE, T, CASE, T, CASE, T, CASE. Rows 1-100.

Table with 10 columns: GROUP, CASE, T, CASE, T, CASE, T, CASE, T, CASE. Rows 1-100.

Table with 10 columns: GROUP, CASE, T, CASE, T, CASE, T, CASE, T, CASE. Rows 1-100.

Table with 10 columns: GROUP, CASE, T, CASE, T, CASE, T, CASE, T, CASE. Rows 1-100.

Table with 10 columns: GROUP, CASE, T, CASE, T, CASE, T, CASE, T, CASE. Rows 1-100.

APPENDIX TABLE 16.

ALBERTA PHOSPHATES ONLY  
 NUMBER OF CASES = 50  
 NUMBER OF BINARY VARIABLES = 0  
 NUMBER OF NUMERIC VARIABLES = 21

PRINCIPAL COMPONENTS ANALYSIS - CLUSTAN 1C  
 (SQUARE-ROOT TRANSFORMED DATA)

NUMERIC MEANS AND STANDARD DEVIATIONS

Variable	Mean	Standard Deviation
1	2.6650	0.2303
2	1.3307	0.1113
3	0.5091	0.0350
4	0.8367	0.0965
5	0.7320	0.0598
6	0.7509	0.0743
7	0.8537	0.1032
8	0.6270	0.0600
9	0.7963	0.0827
10	0.4419	0.0706
11	0.6110	0.0693
12	0.8268	0.0749
13	1.0013	0.0969
14	1.9286	0.1677
15	1.3361	0.1297
16	2.1034	0.1903
17	1.6755	0.1658
18	0.7290	0.1046
19	67.7000	3.6100
20	11.5400	1.1466
21	26.4600	2.0001

VARIABLE MINIMUM VALUE

Variable	Minimum Value	Maximum Value
1	2.1213	3.0822
2	1.0583	1.5166
3	0.4243	0.5745
4	0.6245	1.0000
5	0.5831	0.8307
6	0.5916	0.9055
7	0.6708	1.0630
8	0.5000	0.7211
9	0.6000	0.9381
10	0.2646	0.5477
11	0.4359	0.7550
12	0.6557	0.9895
13	0.7810	1.1747
14	1.5904	2.2204
15	1.0000	1.5362
16	1.6793	2.4352
17	1.2728	1.9698
18	0.4899	0.9055
19	66.0000	74.0000
20	9.6000	14.0000
21	25.0000	33.0000

EIGENVALUES

16.28	1.31	1.02	0.79	0.52	0.26	0.26	0.12	0.09	0.08	0.07	0.06	0.06	0.04	0.04	0.02
0.62	0.01	0.01	0.00	0.00											

PERCENTAGE VARIANCE

77.51	6.22	4.86	3.74	2.47	1.22	0.96	0.56	0.42	0.40	0.35	0.28	0.27	0.21	0.19	0.11
0.09	0.07	0.05	0.02	0.02											

CUMULATIVE VARIANCE

77.51	83.72	88.59	92.33	94.80	96.02	96.98	97.54	97.97	98.36	98.71	98.99	99.26	99.47	99.65	99.76
99.84	99.91	99.96	99.98	100.00											

EIGENVECTORS - BY FOLK

VECTOR 1

0.245	0.242	0.226	0.240	0.240	0.238	0.238	0.240	0.241	0.234	0.212	0.232	0.226	0.243	0.236	0.243	0.238	0.208
0.033	0.015	0.076															

VECTOR 2

-0.033	-0.015	0.000	-0.087	0.025	0.051	-0.049	0.017	-0.009	-0.094	0.166	-0.041	0.089	0.024	-0.069	-0.051	0.038	-0.210
0.569	0.501	0.558															

VECTOR 3

-0.049	0.056	-0.054	0.063	0.003	-0.068	0.031	-0.033	0.030	0.080	0.165	-0.009	-0.113	-0.076	-0.058	-0.019	-0.082	0.162
-0.648	0.673	0.106															

FIRST 3 FACTOR SCORES

Case	Factor 1	Factor 2	Factor 3
S 1	-0.072	0.193	-0.322
S 2	-1.010	-0.211	-0.140
S 3	-1.569	0.180	0.043
S 4	-4.475	0.091	-2.293
S 5	-5.679	-1.103	-1.586
S 6	5.667	0.128	0.309
S 7	1.982	0.845	1.244
S 8	3.697	0.353	-0.017
S 9	1.096	1.504	-0.753
S 10	5.129	0.751	-0.893
S 11	1.861	1.093	-0.657
S 12	-2.818	-0.455	-0.215
S 13	-1.934	2.738	0.711
S 14	3.893	2.020	-1.093
S 15	-2.815	0.499	-0.941
S 16	-2.701	0.279	0.265
S 17	-8.445	0.748	0.692
S 18	-9.846	-1.421	-0.396
S 19	-2.887	1.159	1.936
S 20	-5.318	0.277	0.422
S 21	-5.692	-0.057	0.812
S 22	-6.601	0.856	-0.162
S 23	1.266	0.145	1.729
S 24	-3.121	-0.372	-1.063
S 25	-0.888	-0.118	2.176
S 26	-1.670	-0.707	1.455
S 27	-3.453	-1.561	-0.268
S 28	5.461	-1.136	-1.921
S 29	3.297	0.736	-0.775
S 30	3.766	-0.627	-0.097
S 31	3.300	-1.300	-0.322
S 32	2.673	-1.208	-0.197
S 33	2.879	-0.384	-0.777
S 34	3.193	0.271	0.016
S 35	5.057	1.261	0.334
S 36	5.169	0.558	0.397
S 37	3.302	-0.042	-1.119
S 38	2.194	0.943	0.077
S 39	2.847	1.516	-0.211
S 40	0.911	0.268	-0.575
S 41	0.712	1.639	-0.895
S 42	3.904	-0.804	0.248
S 43	3.089	-0.215	2.719
S 44	0.714	-0.439	-0.157
S 45	1.114	-1.922	-0.036
S 46	-0.412	-3.208	0.547
S 47	-0.704	-3.941	0.824
S 48	0.610	-1.837	1.716
S 49	2.001	-1.419	-0.747
S 50	6.448	0.215	-0.039

APPENDIX TABLE 17.

ALBERTA COVIDIUS ONLY

NUMBER OF CASES = 50  
 NUMBER OF BINARY VARIABLES = 0  
 NUMBER OF NUMERIC VARIABLES = 21

PRINCIPAL COMPONENTS ANALYSIS - CLUSTAN 1C  
 (SQUARE-ROOT TRANSFORMED DATA)

NUMERIC MEANS AND STANDARD DEVIATIONS			VARIABLE MINIMUM VALUE		MAXIMUM VALUE	
1	2.6897	0.3089	1	2.0736	3.3166	
2	1.3078	0.1425	2	1.0189	1.6462	
3	0.6300	0.0503	3	0.5292	0.7550	
4	0.6853	0.0809	4	0.5196	0.8485	
5	0.7971	0.0834	5	0.6083	0.9798	
6	0.7369	0.0937	6	0.5385	0.9327	
7	0.9050	0.0970	7	0.7071	1.1489	
8	0.6798	0.0746	8	0.5196	0.8485	
9	0.7301	0.0870	9	0.5477	0.9539	
10	0.1688	0.0909	10	0.0000	0.3606	
11	0.0	1.0000	11	0.0	0.0	
12	0.3966	0.0803	12	0.2686	0.6164	
13	1.0002	0.1349	13	0.7071	1.4107	
14	1.9271	0.2183	14	1.4500	2.4166	
15	1.3043	0.1582	15	0.9695	1.5969	
16	2.1478	0.2547	16	1.6125	2.7258	
17	1.7367	0.2257	17	1.2570	2.2782	
18	0.7588	0.0885	18	0.5831	0.9434	
19	62.4800	2.8734	19	58.0000	70.0000	
20	13.2060	0.8572	20	11.0000	15.0000	
21	23.4400	1.4168	21	21.0000	26.0000	

EIGENVALUES

15.13	1.38	1.08	0.73	0.55	0.44	0.21	0.12	0.10	0.06	0.05	0.04	0.03	0.02	0.02	0.01
0.01	0.00	0.00	0.00	0.00											

PERCENTAGE VARIANCE

72.04	6.57	5.16	3.49	2.60	2.11	1.01	0.56	0.47	0.29	0.23	0.19	0.15	0.12	0.10	0.06
0.04	0.02	0.01	0.01	0.00											

CUMULATIVE VARIANCE

72.04	78.61	83.77	87.27	89.86	91.98	92.99	93.55	94.01	94.31	94.54	94.73	94.87	94.99	95.09	95.15
95.20	95.22	95.23	95.24	95.24											

EIGENVECTORS - BY POWS

VECTOR 1

0.255	0.254	0.248	0.251	0.253	0.251	0.252	0.246	0.252	0.177	0.0	0.190	0.220	0.255	0.249	0.254	0.250	0.241
0.043	0.026	-0.006															

VECTOR 2

0.006	-0.025	-0.005	-0.055	-0.019	0.032	-0.017	-0.014	-0.042	-0.246	0.0	-0.073	0.117	0.010	0.017	0.014	0.029	0.014
0.650	0.648	0.127															

VECTOR 3

-0.061	0.026	0.071	0.037	0.033	-0.036	0.037	-0.044	0.017	0.272	0.0	0.052	0.000	-0.065	-0.088	-0.054	-0.058	-0.007
-0.230	-0.196	0.693															

FIRST 3 FACTOR SCORES

S 1	4.036	-0.913	-2.034	S 26	-0.140	0.037	0.264
S 2	-0.080	1.023	-1.824	S 27	-0.704	-0.729	1.351
S 3	-4.715	-0.496	-1.343	S 28	-1.636	-1.499	1.032
S 4	-1.497	0.023	1.075	S 29	-1.389	-1.911	1.282
S 5	0.896	0.676	-1.146	S 30	-3.756	2.158	0.814
S 6	2.854	-0.510	-1.578	S 31	-5.419	-0.876	0.304
S 7	-2.925	-1.268	0.929	S 32	-3.168	1.168	1.807
S 8	-0.837	1.735	0.180	S 33	8.932	-0.408	1.004
S 9	-0.251	-0.496	-0.338	S 34	5.962	-1.573	1.585
S 10	-0.784	1.257	1.006	S 35	5.337	-0.607	0.609
S 11	-4.362	-0.009	-0.951	S 36	4.491	-0.129	0.767
S 12	-0.684	1.120	1.006	S 37	1.868	-0.633	0.739
S 13	-7.669	-1.007	-0.537	S 38	-0.365	-0.494	0.251
S 14	0.377	0.732	0.569	S 39	-3.058	-1.069	0.079
S 15	-6.821	-1.653	-1.080	S 40	-4.867	-0.555	-0.226
S 16	-0.287	1.578	2.063	S 41	4.195	0.143	0.047
S 17	3.901	2.537	-1.012	S 42	0.896	2.698	-1.099
S 18	1.827	-1.441	-1.588	S 43	4.222	-0.941	1.908
S 19	2.963	-1.474	-0.540	S 44	3.500	-0.708	-0.991
S 20	3.296	-0.666	-0.237	S 45	3.463	0.644	0.114
S 21	-4.149	0.980	-0.938	S 46	4.737	-0.549	-0.859
S 22	-5.336	-0.978	0.668	S 47	3.293	0.150	-0.437
S 23	-5.634	2.112	-0.089	S 48	2.751	-1.021	-0.639
S 24	-8.253	-0.020	0.224	S 49	0.211	0.506	-1.429
S 25	2.161	0.402	-0.808	S 50	3.087	-1.477	0.046



APPENDIX TABLE 19. Alberta control *Rhinichthys cataractae*; Raw character values, n = 50.

ALBERTA CONTROL	RHINICHTHYS	ALABAMA	ALABAMA	N 50	0.75	0.40	0.62	0.71
401A	0.37	1.50	1.73	0.49	0.65	1.25	0.94	0.75
402A	0.41	7.00	1.01	2.78	1.92	4.45	2.84	3.52
403A	0.37	6.70	0.92	0.74	0.68	0.55	0.57	0.75
404A	0.30	0.57	1.68	3.46	1.89	4.15	2.55	0.45
405A	0.28	6.00	1.02	3.81	1.69	4.29	2.82	0.43
409A	0.57	0.57	1.86	0.23	0.52	0.47	0.45	0.61
410A	0.51	5.50	0.83	3.09	1.43	3.61	2.24	0.35
411A	0.50	0.52	1.35	0.21	0.49	0.44	0.45	0.62
412A	0.45	8.50	0.77	2.85	1.48	3.36	2.20	0.30
415A	0.52	0.88	2.30	0.32	1.20	0.69	0.71	0.94
416A	0.47	7.40	1.27	4.48	2.05	5.24	3.44	0.78
417A	0.36	0.68	2.05	0.74	0.83	0.58	0.56	0.81
419A	0.41	8.10	1.00	4.01	1.76	4.78	3.09	0.69
426A	0.27	0.81	2.04	0.29	0.87	0.63	0.63	0.87
427A	0.30	7.10	1.16	4.28	1.90	5.02	3.15	0.76
428A	0.25	0.77	1.89	0.28	0.77	0.54	0.61	0.79
429A	0.20	8.60	1.00	3.74	1.84	4.34	2.73	0.65
431A	0.19	0.86	2.23	0.29	0.94	0.65	0.66	0.90
433A	0.28	7.40	1.00	3.90	1.45	4.58	2.81	0.69
434A	0.27	6.20	1.54	0.26	0.56	0.47	0.49	0.80
435A	0.40	0.59	0.92	3.19	1.54	3.89	2.49	0.65
436A	0.42	6.30	1.53	0.26	0.59	0.47	0.52	0.74
437A	0.39	6.00	0.89	3.26	1.53	4.02	2.56	0.41
438A	0.25	0.67	1.84	0.22	0.51	0.45	0.47	0.63
439A	0.20	0.60	0.99	3.07	1.49	3.57	2.42	0.40
440A	0.30	5.80	1.39	0.20	0.40	0.41	0.45	0.55
441A	0.25	0.59	0.95	3.14	1.35	3.64	2.42	0.39
442A	0.20	5.00	1.29	0.20	0.46	0.39	0.36	0.52
443A	0.19	0.50	0.61	2.56	1.29	2.85	1.62	0.36
444A	0.39	4.80	1.21	0.18	0.41	0.37	0.36	0.50
445A	0.28	0.46	0.72	2.53	1.11	2.94	1.88	0.31
446A	0.27	4.50	1.12	0.19	0.39	0.34	0.35	0.45
447A	0.25	0.43	0.62	2.56	1.00	2.82	1.66	0.24
448A	0.20	0.43	1.48	0.22	0.63	0.50	0.50	0.42
449A	0.19	0.55	0.84	3.30	1.38	3.69	2.32	0.42
450A	0.38	5.60	1.41	0.22	0.52	0.46	0.41	0.36
451A	0.45	0.53	0.78	2.89	1.47	3.36	2.15	0.42
452A	0.42	5.60	1.40	0.21	0.48	0.43	0.42	0.36
453A	0.39	0.45	0.81	2.91	1.34	3.52	2.27	0.48
454A	0.27	5.00	1.40	0.23	0.47	0.44	0.42	0.32
455A	0.25	0.49	0.72	2.78	1.26	3.09	1.88	0.48
456A	0.20	7.10	1.75	0.29	0.72	0.61	0.61	0.60
457A	0.19	0.79	1.19	3.66	1.76	4.65	2.92	0.60
458A	0.17	6.00	1.50	0.26	0.55	0.42	0.50	0.59
459A	0.16	0.64	0.99	3.16	1.52	3.74	2.45	0.46
460A	0.15	6.80	1.75	0.26	0.64	0.51	0.55	0.73
461A	0.14	0.63	1.03	3.53	1.72	4.30	2.81	0.49
462A	0.13	6.40	1.69	0.26	0.63	0.48	0.50	0.67
463A	0.12	0.70	0.83	3.31	1.62	4.11	2.57	0.51
464A	0.11	6.00	1.51	0.25	0.56	0.46	0.42	0.63
465A	0.10	0.55	0.82	3.09	1.52	3.80	2.44	0.52
466A	0.09	8.40	2.19	0.32	0.97	0.66	0.78	1.04
467A	0.08	0.88	1.38	4.80	2.23	5.44	3.56	0.63
468A	0.07	8.50	1.90	0.27	0.80	0.54	0.69	0.91
469A	0.06	0.77	1.27	4.38	2.31	5.28	3.56	0.69
470A	0.05	8.30	2.03	0.29	0.80	0.60	0.67	1.06
471A	0.04	0.79	1.22	4.44	2.11	5.20	3.46	0.66
472A	0.03	8.50	1.98	0.30	0.76	0.58	0.61	0.99
473A	0.02	0.80	1.17	4.35	2.36	5.12	3.32	0.70
474A	0.01	0.20	1.86	0.30	0.78	0.60	0.68	0.90
475A	0.00	0.75	1.09	4.29	2.21	5.05	3.35	0.57
476A	0.00	0.76	1.93	0.29	0.79	0.59	0.65	0.92
477A	0.00	0.80	1.28	4.37	2.07	5.35	3.44	0.54
478A	0.00	0.83	1.18	4.17	2.04	4.91	3.20	0.68
479A	0.00	0.85	0.82	3.09	1.52	3.80	2.44	0.52
480A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
481A	0.00	0.90	2.19	0.32	0.97	0.66	0.78	1.04
482A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
483A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
484A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
485A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
486A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
487A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
488A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
489A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
490A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
491A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
492A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
493A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
494A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
495A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
496A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
497A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
498A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
499A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63
500A	0.00	0.88	1.38	4.80	2.23	5.44	3.56	0.63



APPENDIX TABLE 2J. Kananaskis typical *Couesius*

*plumbeus*; Raw character

values, n = 50.

KANANASKIS	APPARANT	COUESIUS PLUMBEUS									
601	0.0	16.20	2.31	0.47	0.67	0.87	0.74	1.17	0.64	0.57	0.00
602	0.0	12.16	2.57	0.58	0.75	0.92	0.89	1.32	0.75	0.88	0.11
603	0.0	0.22	1.78	6.10	2.72	7.63	5.11	0.85	68.00	18.00	25.00
604	0.0	12.30	2.19	0.60	0.79	1.05	1.02	1.58	0.70	1.00	0.07
605	0.0	6.26	1.75	6.28	2.75	8.15	5.72	0.84	61.00	13.00	22.00
606	0.0	11.50	2.48	0.55	0.76	0.91	0.87	1.26	0.71	0.99	0.04
607	0.0	10.00	2.21	0.50	0.61	0.83	0.77	1.10	0.58	0.78	0.05
608	0.0	0.32	1.46	5.16	2.33	6.38	4.40	0.71	65.00	15.00	26.00
609	0.0	10.60	2.35	0.51	0.72	0.85	0.74	1.20	0.69	0.63	0.04
610	0.0	0.24	1.38	5.25	2.40	6.78	4.87	0.78	61.00	13.00	24.00
611	0.0	9.80	1.97	0.50	0.60	0.79	0.71	1.07	0.55	0.70	0.05
612	0.0	0.12	1.21	4.73	2.28	5.86	3.90	0.77	70.00	13.00	27.00
613	0.0	9.50	2.06	0.52	0.60	0.78	0.75	1.13	0.54	0.76	0.03
614	0.0	6.28	1.27	4.85	2.11	6.08	4.20	0.71	63.00	14.00	24.00
615	0.0	8.70	2.02	0.48	0.58	0.73	0.62	1.07	0.53	0.70	0.05
616	0.0	0.19	1.20	4.32	1.97	5.42	3.52	0.62	64.00	14.00	24.00
617	0.0	4.20	1.85	0.66	0.57	0.65	0.60	1.04	0.49	0.69	0.03
618	0.0	0.22	1.10	4.12	1.92	5.10	3.27	0.65	64.00	15.00	26.00
619	0.0	9.80	2.15	0.52	0.61	0.78	0.67	1.11	0.56	0.71	0.04
620	0.0	0.20	1.26	4.93	2.20	6.11	4.14	0.71	62.00	14.00	21.00
621	0.0	7.40	1.68	0.44	0.48	0.64	0.47	0.92	0.49	0.58	0.03
622	0.0	0.10	0.99	1.70	1.70	3.76	3.76	0.58	62.00	12.00	22.00
623	0.0	7.00	1.72	0.43	0.49	0.62	0.55	0.88	0.45	0.52	0.04
624	0.0	0.17	0.81	3.59	1.84	4.42	2.78	0.50	60.00	13.00	25.00
625	0.0	6.60	1.55	0.40	0.43	0.58	0.47	0.75	0.44	0.48	0.02
626	0.0	0.15	0.92	3.27	1.54	4.26	2.86	0.58	62.00	13.00	23.00
627	0.0	6.90	1.59	0.41	0.44	0.60	0.56	0.78	0.48	0.50	0.05
628	0.0	6.20	1.03	3.46	1.84	4.47	3.03	0.63	61.00	11.00	22.00
629	0.0	7.10	1.66	0.40	0.49	0.64	0.57	0.79	0.43	0.52	0.05
630	0.0	6.22	0.98	3.52	1.78	4.47	2.87	0.54	65.00	14.00	24.00
631	0.0	7.00	1.60	0.36	0.46	0.63	0.44	0.75	0.42	0.49	0.05
632	0.0	0.17	0.57	3.45	1.80	4.32	2.80	0.57	61.00	13.00	24.00
633	0.0	5.30	1.32	0.32	0.32	0.46	0.38	0.60	0.33	0.35	0.01
634	0.0	0.11	0.10	2.70	1.35	3.33	2.70	0.37	66.00	14.00	21.00
635	0.0	6.00	1.47	0.38	0.43	0.50	0.49	0.64	0.40	0.45	0.0
636	0.0	0.25	0.86	3.09	1.44	3.75	2.35	0.51	60.00	13.00	23.00
637	0.0	10.00	2.19	0.50	0.67	0.80	0.71	1.18	0.59	0.74	0.10
638	0.0	0.24	1.32	4.56	2.49	6.19	4.22	0.74	63.00	13.00	22.00
639	0.0	5.80	2.23	0.48	0.67	0.80	0.72	1.00	0.68	0.76	0.11
640	0.0	0.17	1.23	4.10	2.53	6.83	4.61	0.61	65.00	15.00	26.00
641	0.0	11.00	2.44	0.53	0.76	0.97	0.76	1.31	0.71	0.91	0.10
642	0.0	0.22	1.57	5.38	2.52	7.03	4.95	1.00	62.00	13.00	24.00
643	0.0	11.00	2.53	0.52	0.84	0.82	0.92	1.26	0.72	0.99	0.07
644	0.0	0.27	1.48	5.60	2.55	7.76	4.77	0.99	66.00	15.00	27.00
645	0.0	8.50	1.93	0.44	0.58	0.72	0.63	1.00	0.64	0.64	0.04
646	0.0	0.14	1.35	4.44	2.18	5.38	3.63	0.60	64.00	13.00	25.00
647	0.0	4.20	1.53	0.46	0.59	0.68	0.56	1.01	0.53	0.60	0.08
648	0.0	0.17	1.24	4.40	2.04	5.32	3.56	0.62	64.00	13.00	25.00
649	0.0	6.80	1.54	0.37	0.43	0.50	0.42	0.73	0.40	0.47	0.05
650	0.0	0.21	0.75	3.41	1.53	4.23	2.78	0.50	64.00	13.00	24.00
651	0.0	6.20	2.18	0.46	0.56	0.65	0.60	0.85	0.72	0.80	0.11
652	0.0	0.15	1.32	4.01	2.07	4.05	4.39	0.64	66.00	13.00	25.00
653	0.0	7.70	1.80	0.44	0.55	0.74	0.69	1.00	0.46	0.64	0.37
654	0.0	6.25	1.30	3.59	2.12	4.82	3.43	0.51	61.00	13.00	23.00
655	0.0	7.70	1.76	0.40	0.55	0.67	0.59	0.93	0.46	0.65	0.07
656	0.0	6.25	1.26	3.92	1.94	4.46	3.38	0.61	63.00	14.00	23.00
657	0.0	0.11	0.73	1.77	1.1	3.1	2.48	0.85	0.46	0.54	0.07
658	0.0	7.10	1.14	4.11	1.37	4.63	3.34	0.70	60.00	13.00	24.00
659	0.0	6.50	1.14	4.41	1.42	4.63	3.41	0.76	0.42	0.52	0.04
660	0.0	6.18	1.38	4.11	2.00	5.34	3.56	0.66	67.00	14.00	23.00
661	0.0	10.80	2.25	0.47	0.70	0.87	0.75	1.14	0.62	0.79	0.09
662	0.0	6.38	1.72	5.49	2.86	6.39	4.30	0.94	67.00	13.00	25.00
663	0.0	7.70	1.80	0.45	0.56	0.71	0.66	1.00	0.47	0.68	0.07
664	0.0	6.21	1.15	3.94	2.07	4.87	3.39	0.65	63.00	14.00	24.00
665	0.0	7.10	1.80	0.43	0.51	0.66	0.50	0.90	0.50	0.60	0.05
666	0.0	6.14	1.37	3.93	1.73	4.97	3.43	0.51	63.00	14.00	23.00
667	0.0	7.60	1.77	0.42	0.54	0.60	0.55	0.84	0.45	0.62	0.09
668	0.0	6.16	1.09	3.86	1.93	4.80	3.49	0.62	61.00	12.00	21.00
669	0.0	5.90	1.26	0.44	0.38	0.43	0.43	0.65	0.37	0.41	0.01
670	0.0	0.12	0.72	2.88	1.24	3.43	2.71	0.55	69.00	13.00	23.00
671	0.0	4.70	1.15	0.30	0.27	0.41	0.39	0.60	0.33	0.38	0.0
672	0.0	6.15	0.88	2.50	1.22	3.05	1.93	0.37	62.00	14.00	25.00
673	0.0	11.50	2.51	0.54	0.69	0.90	0.87	1.32	0.60	0.89	0.06
674	0.0	10.26	1.50	0.80	2.07	7.52	5.36	0.80	62.00	13.00	22.00
675	0.0	10.00	2.22	0.48	0.70	0.79	0.67	1.15	0.55	0.87	0.10
676	0.0	6.37	1.39	5.24	2.22	6.51	4.39	0.67	61.00	15.00	23.00
677	0.0	5.40	1.35	0.36	0.33	0.40	0.38	0.64	0.36	0.32	0.0
678	0.0	0.15	0.59	2.62	1.38	3.35	2.14	0.40	62.00	14.00	23.00
679	0.0	5.80	1.42	0.35	0.39	0.45	0.43	0.67	0.19	0.40	0.01
680	0.0	0.19	0.68	2.89	1.42	3.64	2.27	0.48	60.00	12.00	22.00
681	0.0	6.00	1.45	0.34	0.40	0.51	0.50	0.68	0.38	0.46	0.02
682	0.0	0.23	0.73	3.13	1.43	3.85	2.63	0.47	64.00	12.00	22.00
683	0.0	6.50	1.51	0.38	0.48	0.54	0.48	0.77	0.43	0.48	0.05
684	0.0	6.74	0.85	3.74	1.62	4.12	2.44	0.54	64.00	15.00	21.00
685	0.0	6.30	1.58	0.39	0.45	0.50	0.48	0.68	0.42	0.45	0.03
686	0.0	0.21	0.85	3.21	1.43	4.08	2.60	0.48	60.00	16.00	24.00
687	0.0	6.50	1.55	0.41	0.45	0.56	0.54	0.79	0.40	0.53	0.04
688	0.0	0.13	0.90	3.38	1.50	4.60	2.85	0.55	60.00	12.00	25.00
689	0.0	7.10	1.70	0.42	0.47	0.58	0.54	0.82	0.47	0.54	0.05
690	0.0	0.19	0.81	3.61	1.93	4.35	2.82	0.57	63.00	14.00	26.00
691	0.0	5.40	1.22	0.37	0.34	0.44	0.42	0.62	0.35	0.41	0.02
692	0.0	0.15	0.67	2.73	1.28	3.26	2.06	0.38	65.00	15.00	24.00
693	0.0	4.40	1.03	0.31	0.25	0.39	0.35	0.54	0.29	0.35	0.0
694	0.0	0.16	0.62	2.24	1.05	2.01	1.80	0.34	64.00	13.00	23.00
695	0.0	6.80	1.73	0.42	0.46	0.58	0.56	0.83	0.48	0.52	0.05
696	0.0	6.28	0.99	3.43	1.81	4.20	2.87	0.57	65.00	12.00	23.00



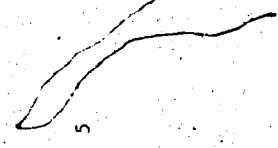
APPENDIX TABLE 22. Kananaskis suspected hybrids *Couesius plumbeus* x *Rhinichthys cataractae*; Raw character values, n = 37.

KANANASKIS	SUSPECTED	HYBRIDS	COUESIUS PLUMBEUS X		RHINICHTHYS CATARACTAE (1977)						
501		7.60	1.72	0.43	0.46	0.61	0.71	1.00	0.52	0.69	0.0
	0.21	0.40	1.08	4.00	1.89	4.69	2.85	0.63	60.00	13.00	22.00
502		7.10	1.69	0.32	0.55	0.62	0.60	0.79	0.46	0.52	0.0
	0.03	0.35	0.83	3.68	1.81	4.61	3.00	0.60	69.00	14.00	25.00
503		6.30	1.63	0.29	0.54	0.51	0.50	0.72	0.36	0.51	0.09
	0.22	0.41	0.80	3.26	1.44	4.16	2.63	0.48	68.00	13.00	25.00
504		6.20	1.58	0.31	0.51	0.48	0.52	0.69	0.41	0.50	0.03
	0.08	0.32	0.82	3.19	1.55	3.86	2.52	0.49	67.00	13.00	24.00
505		8.10	1.70	0.43	0.51	0.64	0.57	0.85	0.47	0.54	0.10
	0.04	0.31	1.05	4.04	2.17	4.95	3.28	0.51	66.00	14.00	23.00
506		7.10	1.52	0.39	0.45	0.59	0.54	0.77	0.42	0.55	0.03
	0.21	0.38	1.10	3.63	1.79	4.56	3.19	0.50	70.00	14.00	25.00
507		7.60	1.82	0.33	0.64	0.66	0.68	0.89	0.47	0.65	0.11
	0.17	0.35	1.02	3.88	1.87	4.87	3.23	0.49	69.00	14.00	26.00
508		6.60	1.66	0.29	0.53	0.58	0.58	0.78	0.40	0.55	0.08
	0.0	0.31	0.81	3.43	1.38	4.05	2.67	0.52	67.00	15.00	27.00
509		6.20	1.52	0.33	0.43	0.53	0.52	0.73	0.38	0.48	0.02
	0.18	0.22	0.86	3.15	1.50	3.92	2.48	0.39	70.00	13.00	26.00
510		6.10	1.48	0.25	0.50	0.52	0.46	0.67	0.39	0.47	0.07
	0.16	0.24	0.81	2.97	1.48	3.74	2.37	0.40	69.00	14.00	26.00
511		5.80	1.29	0.28	0.45	0.45	0.51	0.61	0.16	0.46	0.05
	0.20	0.29	0.94	2.73	1.23	3.44	2.28	0.38	67.00	13.00	26.00
512		5.70	1.30	0.24	0.40	0.48	0.42	0.63	0.33	0.45	0.05
	0.17	0.27	0.67	2.78	1.19	3.34	2.17	0.42	73.00	16.00	28.00
513		5.80	1.44	0.26	0.48	0.49	0.45	0.67	0.35	0.48	0.07
	0.16	0.36	0.81	2.98	1.39	3.58	2.35	0.43	69.00	15.00	26.00
514		4.80	1.17	0.25	0.40	0.40	0.37	0.54	0.26	0.36	0.03
	0.15	0.22	0.57	2.38	1.19	3.00	1.93	0.35	69.00	14.00	23.00
515		5.60	1.35	0.25	0.44	0.45	0.47	0.65	0.32	0.47	0.06
	0.18	0.33	0.72	2.95	1.36	3.51	2.31	0.47	70.00	14.00	28.00
516		4.90	1.22	0.34	0.33	0.44	0.42	0.72	0.25	0.40	0.03
	0.0	0.22	0.74	2.49	1.07	3.00	2.00	0.39	68.00	15.00	24.00
517		5.60	1.26	0.34	0.35	0.43	0.46	0.58	0.29	0.40	0.0
	0.24	0.14	0.70	2.83	1.30	3.50	2.27	0.36	62.00	13.00	26.00
518		5.60	1.39	0.24	0.52	0.40	0.49	0.69	0.34	0.45	0.10
	0.12	0.32	0.68	2.86	1.31	3.61	2.24	0.45	70.00	13.00	27.00
519		5.20	1.23	0.26	0.34	0.42	0.40	0.58	0.26	0.38	0.0
	0.06	0.20	0.69	2.59	1.27	3.20	2.20	0.40	60.00	13.00	25.00
520		6.70	1.60	0.41	0.45	0.54	0.48	0.80	0.38	0.50	0.0
	0.20	0.27	0.87	3.53	1.52	4.15	2.73	0.52	62.00	14.00	27.00
521		6.70	1.69	0.33	0.56	0.58	0.56	0.82	0.43	0.58	0.07
	0.15	0.39	0.84	3.57	1.45	4.45	2.78	0.52	67.00	13.00	25.00
522		5.80	1.41	0.23	0.43	0.48	0.45	0.69	0.32	0.44	0.06
	0.18	0.26	0.76	2.93	1.44	3.58	2.23	0.45	61.00	15.00	28.00
523		6.10	1.43	0.25	0.43	0.54	0.47	0.71	0.38	0.53	0.04
	0.13	0.32	0.82	3.00	1.50	3.88	2.53	0.45	62.00	14.00	27.00
524		5.60	1.35	0.33	0.36	0.48	0.44	0.68	0.29	0.42	0.0
	0.02	0.15	0.71	2.85	1.40	3.56	2.26	0.36	60.00	13.00	23.00
525		6.00	1.42	0.36	0.40	0.51	0.49	0.70	0.32	0.48	0.0
	0.10	0.21	0.77	3.07	1.49	3.87	2.46	0.38	71.00	15.00	27.00
526		4.40	1.20	0.23	0.26	0.38	0.33	0.50	0.29	0.35	0.0
	0.0	0.23	0.57	2.27	1.01	2.81	1.73	0.37	64.00	12.00	24.00
527		3.40	0.90	0.24	0.22	0.31	0.28	0.40	0.23	0.26	0.01
	0.02	0.10	0.44	1.80	0.75	2.19	1.35	0.21	63.00	14.00	23.00
528		7.20	1.76	0.31	0.55	0.47	0.58	0.79	0.41	0.60	0.06
	0.16	0.20	0.80	3.57	1.85	4.49	2.86	0.36	72.00	13.00	29.00
529		6.70	1.65	0.31	0.50	0.57	0.51	0.68	0.43	0.49	0.05
	0.18	0.30	0.68	3.35	1.59	4.20	2.48	0.39	69.00	14.00	27.00
530		6.00	1.49	0.32	0.44	0.48	0.46	0.70	0.39	0.52	0.0
	0.17	0.32	0.71	3.10	1.46	3.71	2.28	0.41	67.00	13.00	25.00
531		5.50	1.33	0.26	0.39	0.42	0.49	0.62	0.32	0.39	0.02
	0.06	0.23	0.59	2.86	1.42	3.21	1.94	0.40	63.00	12.00	21.00
532		5.60	1.34	0.27	0.41	0.46	0.45	0.57	0.34	0.41	0.03
	0.16	0.24	0.82	2.93	1.29	3.49	2.25	0.42	68.00	13.00	27.00
533		4.30	1.11	0.25	0.33	0.38	0.32	0.46	0.23	0.32	0.05
	0.09	0.26	0.48	2.21	1.08	2.65	1.55	0.28	67.00	14.00	26.00
534		6.50	1.62	0.27	0.54	0.52	0.49	0.71	0.39	0.52	0.12
	0.14	0.34	0.76	3.42	1.51	4.25	2.73	0.50	60.00	14.00	25.00
535		6.00	1.58	0.25	0.52	0.43	0.50	0.62	0.55	0.64	0.12
	0.09	0.44	0.76	3.04	1.48	3.90	2.48	0.52	65.00	13.00	25.00
536		10.00	2.41	0.42	0.82	0.86	0.91	1.20	0.71	0.95	0.15
	0.35	0.44	1.42	5.21	2.31	6.47	4.27	0.39	72.00	12.00	27.00
537		6.90	1.56	0.31	0.53	0.54	0.64	0.86	0.41	0.63	0.13
	0.12	0.38	0.95	3.44	1.73	4.39	2.76	0.60	69.00	12.00	25.00



APPENDIX TABLE 24. Summary of changes in female gonad maturity for both *Couesius plumbeus* and *Rhinichthys cataractae* and surface water temperature during spawning period, 1978, Lower Kananaskis Reservoir. These data were extracted from total catch data in Table 2 in Appendix.

Date	<i>C. plumbeus</i>			<i>R. cataractae</i>			Temperature °C
	Ripe	Fully Ripe	Spent	Ripe	Fully Ripe	Spent	
May 15, 16	3						4.0
17, 18	6			1			4.0
23, 24	9						4.5
25, 26	8	3		2			5.0
27	1						
May 31, June 1	7	6		4			8.2
June 5	5			6	1		10.5
6							10.5
7, 8	44	44	13	9	4		10.0
12, 13	48	20	41	4	8	3	10.5
14	32	50	16		1		12.5
19, 20	15	7	17	3	2	4	10.5
21, 22	20	31	31				
28, 29		13	52			5	13.0
July 6			12				13.7
12		7	26				16.0



APPENDIX TABLE 25. A. Data summary for female *R. cataractae*  
Gonadosomatic Index values (GSI), Lower Kananaskis  
Reservoir, 1978.

Date	n	$\bar{X}$	$s_D$	Range
May 1	1	13.8	-	-
May 18	2	14.1	-	12.6 - 15.6
May 26	2	15.6	-	14.9 - 16.2
June 1	4	19.1	0.3	18.8 - 19.5
June 8	7	19.4	2.6	15.0 - 23.0
June 14	10	17.0	4.6	10.0 - 24.8
June 20	7	16.0	3.5	11.4 - 21.0
June 29	6	11.4	2.1	9.1 - 14.4

APPENDIX TABLE 25. B. Data summary for female *C. plumbeus*  
Gonadosomatic Index values (GSI), Lower Kananaskis  
Reservoir, 1978.

Date	n	$\bar{X}$	$s_D$	Range
May 1	20	18.4	2.2	14.0 - 22.0
May 13	6	18.8	1.4	17.1 - 20.4
May 24, 25	22	19.0	1.6	16.6 - 22.0
June 1	8	21.0	1.4	19.0 - 23.2
June 8	12	24.2	2.9	20.6 - 27.6
June 13	23	17.7	6.6	8.7 - 30.1
June 21	48	17.7	6.1	6.7 - 29.2
June 29	21	16.3	5.6	9.9 - 27.1
July 6	12	13.7	3.3	7.4 - 18.4
July 12	20	11.9	3.6	6.4 - 17.3
July 27	25	8.1	2.3	4.8 - 12.9

APPENDIX TABLE 26. Summary of ova diameter statistics for both  
*Couesius plumbeus* and *Rhinichthys cataractae* from  
 Lower Kananaskis Reservoir, 1978.

Date	Single Values (mm)	n	$\bar{X}$ (mm)	$s_D$	Range
<i>Couesius plumbeus</i>					
May 1		25	1.30	0.11	1.01 - 1.48
May 24		21	1.31	0.07	1.17 - 1.39
June 8		13	1.43	0.07	1.31 - 1.53
June 13		25	1.35	0.23	0.95 - 1.88
June 21		25	1.31	0.23	0.88 - 1.70
July 6		11	1.19	0.15	0.94 - 1.36
July 12		25	1.22	0.16	0.88 - 1.46
<i>Rhinichthys cataractae</i>					
May 1	1.38	1			
May 18	1.48	1			
May 26	1.52	1			
June 1		4	1.54	0.05	1.49 - 1.60
June 8		7	1.69	0.05	1.62 - 1.78
June 14		10	1.43	0.27	0.91 - 1.68
June 20		7	1.39	0.26	0.88 - 1.69
June 29		6	1.16	0.22	0.96 - 1.48

APPENDIX TABLE 27. Summary of catch data for *Coxesius flumbeus* (C), *Rhinichthys cataractae* (R) and their putative hybrids (H) from 7 different selected habitats on Lower Kananaskis Reservoir. All samples of these species taken during the time period of maximal spawning activity in 1978 are listed below.

Date	Sampling (Interlakes Stream)	Lk1 Stream (Interlakes Stream)	Habitat Locations				Smith-Dorrrien
			Lk1 Common (Interlakes) Bay	Interlakes Raceway	Boulton Cr. Mouth	Invincible Cr. Mouth	
May 18	C-52, R-2						C-8, R-1
22	C-2	C-58					C-4
24		C-6					C-9
25			C-3				
26					C-12, R-5, H-1		
June 1		C-3, R-1	C-20, R-4				
5	C-18, R-12	C-2					
7	R-7, H-1	C-15, R-4		C-11			
8	R-4	C-101, R-7, H-2	C-25, R-3			C-81, H-1	C-5
12	R-8	C-14					
13	R-1	C-69, R-13	C-23		C-55, R-1		C-68, R-3
14	R-1		C-31, R-1	C-71			C-18, R-1
19	R-3						
20	R-9			C-4, R-1	C-32		C-8
21	C-5, R-3		C-38	C-71			C-2
29	R-14		C-64, H-1		C-15		
July 6							



APPENDIX TABLE 28. Lower Kananaskis Reservoir inlet stream surface temperatures ( $^{\circ}\text{C}$ , Hg thermometer), approximate current speeds (calculated from timed passage of a floating object over 10 m) and Secchi disc readings. CPL:- data from Calgary

Power L. flow station.

Date	Smith-Dorrien Creek	Kent Creek	Invincible Creek	Boulton Creek	LXI Stream
April 28	5.5	5.0		3.0	
29	4.5				
30	6.0	0.9 m/s		3.0	
May 1					7.0
7	5.5				0.5 m/s
10					4.4
13					4.0
15	3.3	1.0 m/s			0.7 m/s
16	4.0				7.0
22					8.0
25	4.9 - mouth	7.2.7			
	3.3 - upstream				
	1.7 m/s				
26	3.8		1.0 m/s	5.0	Secchi = 0.2 m
30		2.5			
31					4.0
June 1					
5	6.5			4.5	
7				3.5	
8	4.3	2.5 m/s	4.0		17.5
		3.8	2.0 m/s		16.5
			Secchi = 1.4 m/s		
			Secchi = 0.05 m		
9		64 cfs (CPL)			
12			12.5	8.2	Secchi = 0.4 m
13					10.5
20	5.2				0.2 m/s
21	4.8	7.2	2.2 m/s	12.0	

APPENDIX TABLE 29. Analysis of Covariance of Viability Regressions.

## A. Growth Rate Regressions (Fig. 23 in text)

		F	F <sub>.05(df 1, α)</sub>	P
<i>R. cataractae</i> + <i>C. plumbeus</i>	slope	0.017	3.84	P > 0.05
	intercepts	21.20	3.84	P < 0.05
<i>R. cataractae</i> + Hybrid	slope	2.27	3.84	P > 0.05
	intercepts	4.68	3.84	P < 0.05
<i>C. plumbeus</i> + Hybrid	slope	2.83	3.84	P > 0.05
	intercepts	0.407	3.84	P > 0.05

## B. Length-Weight Regressions (Fig. 24 in text)

		F	F <sub>.05(df 1, α)</sub>	P
<i>R. cataractae</i> + <i>C. plumbeus</i>	slope	0.86	3.84	P > 0.05
	intercepts	48.66	3.84	P < 0.05
<i>R. cataractae</i> + Hybrid	slope	-0.19	3.84	P > 0.05
	intercepts	13.69	3.84	P < 0.05
<i>C. plumbeus</i> + Hybrid	slope	0.91	3.84	P > 0.05
	intercepts	14.62	3.84	P < 0.05

## C. Fecundity Regressions (Fig. 25 in text)

		F	F <sub>.05(df 1, 41)</sub>	P
<i>R. cataractae</i> + <i>C. plumbeus</i>	slope	12.959	4.08	P < 0.05
	intercept	6.878	4.08	P < 0.05

APPENDIX TABLE 30. Length-weight regression summary.

Species & Sex	Sample Size	Correlation Coefficient	Regression Equation
C x R combined ♀ & ♂	42	0.980	$\log w = -1.725 + 2.813 \log l$
C x R ♂ only	24	0.975	$\log w = -1.759 + 2.858 \log l$
C x R ♀ only	18	0.986	$\log w = -1.698 + 2.774 \log l$
C combined	162	0.962	$\log w = -1.781 + 3.064 \log l$
C ♂ only	65	0.957	$\log w = -1.710 + 2.967 \log l$
C ♀ only	97	0.962	$\log w = -1.782 + 3.072 \log l$
R combined	159	0.985	$\log w = -1.899 + 3.120 \log l$
R ♂ only	62	0.959	$\log w = -1.955 + 3.239 \log l$
R ♀ only	94	0.992	$\log w = -1.885 + 3.093 \log l$

Disc Gel Acrylamide Electrophoresis Recipes1 N HCl

21.55 ml conc. reagent (36.0%) to 250 ml

Stock A

pH 8.9

48 ml 1 N HCl

36.6 g THAM (TRIS)

0.23 ml TEMED

to 100 ml H<sub>2</sub>O

Stock C

70 g Acrylamide

1.84 g Bis

to 250 ml (H<sub>2</sub>O)

-Catalyst

2.8 g to 1000 ml

Ammonium Persulfate

Small Pore Gel

5 ml A      10 ml C      5 ml H<sub>2</sub>O      20 ml catalyst

gel stock good for 4000 runs.

Buffer

6.0 g THAM (TRIS)

- dilute to 1/10 strength for run

28.8 g Glycine

- ignore for various enzyme systems

to 1000 ml (H<sub>2</sub>O)

pH 8.3

Destain

50:50:10 methanol:H<sub>2</sub>O:glacial acetic acid

Storage

7% Acetic Acid

Tracking Dye

0.1 g Bromophenyl Blue to 1000 ml (H<sub>2</sub>O)