

big skin of Dolly Varden fr. Mont.
- didn't know where - made plea to
foreign locality fr. Bell

Bean, T. H. 1888. Eastern limit of Dolly Varden.
Cem. Angler 13(3): 44 - quote fr. letter (Mr. Boring to
Mr. M²g^o-u) "I caught it in a stream which
joins the upper & lower St. Marys Lakes, ^{Thinks} ~~is~~ Belly R.
Last year caught two others in same place one of 6 lb.
never seen them in any other stream. -

Partello, S. M. T. 1888. Some fishes of the
Yellowstone. ibid 13(4): 57-58.

Yellowstone full of trout to junction of
Big Horn - few after - Tongue R. up 100 miles in Mtn.
many trout.

- Bean, T. H. 1888. 13(4):⁵⁹⁻⁶⁰ Distribution of the lake trout
- specimens. - last Nov. Mrs. Gilman Sawtell sent 3 specimens
of lake trout fr. Henry's Lake Idaho -
- 13(5): 77 - lake trout in St. Marys Lake,
Mont. - received salted skin and excellent photographs
of a lake trout fr. St. Marys L. + 2 other fish which
seem to be Elviki trout.

● よろこび/KRR札幌へ。ここは道庁北門前。

1. Manuscript for proceeding of the symposium should be send not later than 31 Oct.
2. Editor's Commission will send a copy of the manuscript to chairman of each session
3. Each chairman ^(must) read the manuscript as a ~~ref~~ 1st referee. If necessary, send it to adequate 2nd referee.

4. As Japanese participants are usually unskillful to write English. Please, rewrite sentence, if necessary
5. ~~Style of paper~~ About the style of paper refer editorial memorandum of Japan.
- J. Ichthyology, please



Cavender & Kimura (1989) - Cytotaxonomy and interrelationships of Pacific basin Salvelinus

H. perryi 2N = 62 NF 104

42 meta, submetacentrics 4 subtelocent. 16 ^{acrocentric} telocent.

Hensel & Hočik (83) - H. leuco 2N = 82

24-32 msm + 52-58 t

Dorofeev (cited Viktorovsky) - tsimen 2N = 84 18 + 66 = 102^{NZ}

O. masou 2N = 66 NF 102

36 msm 22 ST 8 acrocent.

rhodurus identical except for NOR position

S. leucomensis leucomensis 2N 84 NF 100

same for imbricus

96 msm 68 ST + T

" " pluvius

* only 8 biarmed-metacent. - also only

8 metz. - in southern malma, fontinalis & maximus = primitive

confluentus 2N = 78 NF 100 22 + 56

same as Kronocius - but Kronocius gillrakers, derethmoid & ethmoid fontanelles all similar to malma.

S. malma krascheninnikovi - Hokkaido - 2N = 82 NF 98

16 sms + 66 ST + T

lordi

same

w/ 'big' very long acrocentric marker acrocentric

according to Viktorovsky's photos - some 'marker' in malma malma.

(?) - malma malma 18-20 sms, 56-60 ST + T

2N = 76-78 NF 96

NOR's AK malma differ from Hokkaido malma

miyabe 16 + 66 2N = 82 NF 98 but differs in NOR

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Hennig, W. 1966. Phylogenetic systematics. Univ. Illinois Press, Urbana, 263 pp.

Hubbs, C. L. 1960. The spiny-rayed cyprinid fishes (Plagopterini) of the Colorado River system. Misc. Publ. Mus. Zool., Univ. Mich., (115): 1-39, pls. 1-3.

Farris, J. S. 1981. Distance data in phylogenetic analysis. Pages 3-23 in V. A. Funk and D. R. Brooks, eds. Advances in cladistics: Proceedings of the first meeting of the Willi Hennig Society. New York Botanical Garden, New York.
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SYNTHESIS OF INFORMATION
FOR A PHYLOGENETIC INTERPRETATION OF SALVELINUS

Robert J. Behnke
Colorado State University
Fort Collins, Colorado

Since my 1980 and 1984 papers on charr systematics, considerable new information has been compiled, particularly from the areas of karyology and biochemical genetics. In view of the latest information, I have modified my former concepts of relationships in the subgenus Salvelinus.

It is now apparent that organismal (morphological) and molecular evolution may proceed at very different rates, particularly among sympatric populations selected for different niches. Problems associated with a best interpretation of available information concern decisions on primitive vs. derived character states and convergent evolution. Ongoing studies of mitochondrial and ribosomal DNA and chromosomal banding, particularly of nuclear organizing regions, imply that much greater resolution and refinement for phylogenetic analysis will be possible.

Some controversial aspects of Salvelinus systematics, such as species criteria, are not amenable to resolution by quantification. A realm for qualitative speculation will always remain for future symposia.

Thingvallavatn

Magnusson, K.P. and M.M. Ferguson. 1987. Genetic analysis of four sympatric morphs of Arctic char, *Salvelinus alpinus*, from Thingvallavatn, Iceland. *Environmental Biology of Fishes* 20(1): 67-73.

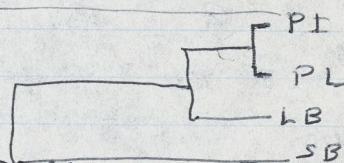
- small ^{SB} benthivore, large ^{LB} benthivore, planktivorous, and ^{PI} piscivorous "morphs" - 5 of 36 enzyme loci were polymorphic Est 2, Gpi 3, Ldh 4, Mdh 4, 5 and Pgm 2. all very closely related Nei's D values from .00004 to .00126 - small benthivore most divergent.

- Benthic forms feed mainly on snails (*Lymnaea peregrina*) and to lesser extent on chironomids.

planktivore = percivore more pelagic - planktivore eat mainly *Daphnia longispina* and *Cyclops abyssorum*, some chironomids. Piscivore mainly on 3-spine stickleback

- Benthic char dark color, rounded snout, subterminal mouth. - Pelagic char - slender, lighter ^{silver} color, pointed snouts, terminal mouth.

	SB mature 3-5 yrs. (27-20cm)	PL - 5-6 yrs. (18-22cm)	LB & PI 6-10 yrs. (25-30cm)
spawn	July-Sept. - north	Sept-Oct	July-Aug. Sept-Nov.
	Sept-Nov. - south	July-Aug	
Mdh 4,5	100 .42	.33 .25 .28	
	130 .58	.67 .75 .72	



Sundlund, O.T., B. Jonsson, H. B. Malmquist, R. Gydemo, T. Lindem, S. Skulason, S. S. Snorrason, and P. M. Jonsson, 1987. *Ibid* 20(4): 263-274. Habitat use of Arctic char, *Salvelinus alpinus*, in Thingvallavatn, Iceland, I.

last glacial retreat 11,500 yrs. B.P.

84km² max depth 114m - only few brown trout &

3-spine stickleback - Great niche diversity - selection to divide up resources -

Loch Rannoch

Walker, A.F., R.B. Greer, and A.S. Gardner, 1988. Two ecologically distinct forms of Arctic char, *Salvelinus alpinus* (L.), in Loch Rannoch, Scotland. *Biological Conservation* 43(1): 43-61.

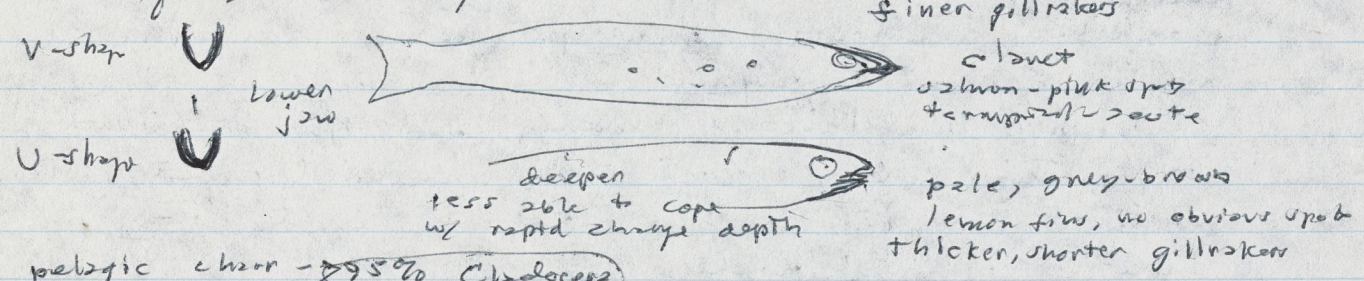
- small headed, claret colored (described by Maitland 1851) - other large headed, pale color.

* - 2 'morph' - diet more distinct non-overlapping than genera - same - *S. salar* et

L. Rannoch, Tayside, Scotland.
 small-headed form is pelagic - mainly zooplankton
 large-headed form benthic - mainly benthic insect
 pelagic form spawns sublittorally in Oct. early Nov.
 benthic form found ripe in Nov. at head of Loch
 in dead inflow zone of River Gaur - main inlet

Qualitative
 Iwama
 form more
 1.2m plexus
 only 500m
 3.11
 why?

S. struanensis Loch Rannoch - Maclean 1951
 Loch Rannoch - long, deep (max 134m) - 16.7 km. long
 el. 204m. 1902 ha., of which about 25% < 15m. deep.
 + pike, perch, eels, 3 spine stickleback, *Phoxinus p.*, *S. salar*,
 & *S. trutta*. - *Trutta* average < 300g. but occasional
 'ferox' to 10 kg.



pelagic charr - 95% *Chirocentrus*
 4-5% insects
 benthic charr *C. edulis* stoneflies - 38% ~~stoneflies~~ 20% debris, 13% clams *Pisidium*,
 6% *Lochimura* 4% fish. 1% *Chirocentrus* some *Gymnarus*

4 of largest fish had eaten small charr,
 max changed 202mm. st eye 8 (1970) largest fish of 375mm. contained 180mm. charr (48% of its length)
 260mm (1987) max size - pelagic at 250-260mm age, 2-8 (4.3)
 benthic - 375mm - 2-12 (6.1)
 only 1 of 243 fish > 240mm. (= ^{max} 375mm)

- differ in MDH allele freq.
 - Loch Ericht (which now enters L. Rannoch from hydro plant) - reported to have large (1134g) benthic charr by Malloch 1910 - gillnetted L. Ericht found charr closely resembling Rannoch benthic form - preliminary data - Loch Ness also seems to have benthic & pelagic forms, but less distinct. also L. Awe.

Sparholt 1985. J. Fish Biol. 26: 313-34
 Greenland lake charr - 2 forms by age growth but no sig. dif. est. so concludes all one pop

Hindar, K., N. Ryman, and G. Ståhl. 1986.
Genetic differentiation among local populations and
morphotypes of arctic charr, Salvelinus alpinus.
Biol. Jour. Linnean Soc. 27(3): 269-285.

drawf, normal, = anadromous pop. compared
EST alleles - no consistent diff. - not monophyletic
3 sp. - all from recent common ancestor.

Some sympatric drawf-normal have signif. dif.
of alleles, some do not - Char introduced
into barren lakes 1910 - now have drawf-normal
in some lakes w/o ^{detectable} genetic dif. - suggest -
ecological - trophic polymorphism - like Cuatro Ciénegas cichlids.

Hindar, K., N. Ryman, and G. Ståhl. 1986.
Genetic differentiation among local populations and
morphotypes of Arctic char, *Salvelinus alpinus*. Biol. Jour.
Linnean Soc. 27(3): 269-285

ecological
polymorphs
- Quers
- Cienega
disruptive
selection

some sympatric dwarf-normal signif. diff. - others not -
some lakes introduced char developed normal-dwarf forms.

no evidence of evol. lines - common ancestor in 1, ancestral

2-dwarfs & normal char no relation between

1-2-3 & EST - allele freq. - all Scandinavian

char fm. on recent common ancestor. - lake spawning

promotes sympatric speciation - lake-stream.

+ same issue "sympatric speciation: when is it
possible?" 201-223 - *Alexeev 1983 - Lenok of

Amur R. - Uda R. - Zool. Zhurn. 62: 1057-68.

Wilson, A. C., et al. 1985 - Mitochondrial DNA
and two perspectives on evol. genetics. Ibid.
26(4): 375-400.

* Zool. Zhur. 61(9):1372 DMT

QH
431
A154

* Genetics 1983 19(4):584-73

* ~~Ecologia~~ 61:319-25 opt. reprod. zone

~~Genetics~~ 111:905-15

QH
506
M 6612

* Mol. Biol. Evol. 1:183-94

Proc. Nat. Acad. Sci. 86:1397-00

~~J. Fish. Ecol~~

* Environ. Biol. Fish. Alley & Reed microhab. segregation

Joley, P.L. (ed.) The Evolving Biosphere.

Conservation Biology - Frankel, O.H. & M.E. Soule. 1981

Darlington, C.S. Conservation & Evolution

Soule & Wilcox. 1980

Conserv. Biol.

QL
186 Zool. Res. 4(3):227-34

+ 27(4):433-40

~~Can. J. Genetics & Cytology - June 85 - RB&CT - Myra Ferguson~~

Mont. 16(4) Outdoors - 1985! - cult. broodstock genetics.

* Critique IFG - 42(4):825-31 + 785-90 - visual pmv - + 42(3):449-

M^cChimmon & Gots - 1979 36:422-57 42(2):791.

Biochem. Genet. 23(7-8):517-70 RB&CT - no incompatibility

QH
C143

QL 614
585

Eisenberg QL J. Fish Biol. 27(1):73-79 + 26(6):691-99

708.5 * Aquaculture 46(4):341-51 + 47(2-3):105-11

E 57

43(1-3):315, 323; 44(2):83

Sindermann Q 175

5569

* J. Fish. Biol. 26(2):318-cha.

QH
56 J

* Nat. Hist. 18(6):327-58 Oman cichlid

* Bamid geth 37(4):27-31 cichl. Gallina

Sci. Rep. Hokkaido Fish Hatchn no 39 Dolly varden

Dronningholm Rep.

16:497- Reif

Cyprinid phylog.

romer p. 6 N.

Brook trout - ¹²⁵² Kerguelen Islands ca. Indian Ocean

- Subsp. - type N.Y. ^{Sumner} Mitchill 1814 -

← west limit Manitoba to Seal R. (Hudson Bay)

Churchill
res.

→ Severn

- Knife

- Churchill rivers

- N.E. Brazil, Savannah, Chokchoochee, Catwaba

- Georgia

↑

Thunder Bay
Huron

Grand Traverse Bay

on the
grays

Mich

C. J. F. A. S. 1986. R. Jørgen, Nygaard,
& Christensen. S. alpinus Greenland - 3
groups exhibiting incipient sympatric
speciation. 43(4): 985-92.

- Interpretation HRM - water

Rigani.

- Biol.

QH1
C143

42(12):1946
43(1):243, 2-11

FAVRO 43(4):896

C. J. F. A. S.

- 211 + 86 -

brook trout: 1969 26:1699
1971 28:452

43(4), (5), (7)

SRP - 43:1093 Matha

~~43:1092-93~~

QH98
ISS

Int. Ver. Theor. Angew. Limnol. Verh.

20(3):2040
2070
2065

22(4):2516
2509
2631

SH 328

N6 Am. S. Fish Mgt. 6(2):296

brown trout col.

Drothryphids

SH 287
D73

SRP: Ecology - 67(4):898

cott. invent. drift

- Environ. Biol. Fishes - 16(4):220-3.

SRP - S.W. Nat. 30:129-
March - 1985

USFWS E.S. Bull. 11(10):3

Loach minnow - proposed Jew

threatened

- Critical Habitat (proposed till June 87)

final rule Oct. 28, 1986

- only 1985 found 5 2k white R. -

- Nehring Stream Fish. Invest. 1986 - flow investigation
2nd Ad Proj. 7-51-R

Altemetsen, A., P. E. Grotner, H. Holthe, &
K. Kristoffersen. 1955. Bear Island Char.

Rep. Inst. FW Res. Drottningholm 62: 98-119.

- several lakes - only few sp. S. G. -

2 morphs - small, spawn younger, shorter life
darker coloration ^{less bright} - Esterase - ~~no dif.~~ ^{between morphs.} ~~between~~

morphs - dif. lake 7 allele .75 - 1.00 -

- but caeca: small char - 25 large char

" may have genetic
basis "

Ellsjoen 36 " 45

Stevata 35 Ojungen 46

Small char 23-24 lakes, large 24-25

- ~~Stevata~~ ~~2 morphs~~ dif. esterase freq. - isolated
pop. → .

QH431 Kortavtsev, Yu. P., M.K. Glubokovskiy, &
A164 I.A. Chevershnev. 1983 Genetic differentiation and
level of variability of two sympatric char species
(Salvelinus, Salmonidae. Genetika 19(4):584-593
- 28 protein loci - S. malma, S. taranetzki
Chukotka - Dif. freq. of alleles at Idh &
Est-2 loci - and unique fixed alleles at AcpH-1 locus
- Thus no doubt distinct sp. $D = .08$

Ben Voyago
493-8571

Larry Reno
534-1700

BOZEMAN

① Gresswell - coming CSU 12/1

② Field Trip - Yellowstone
Mont. - - - - - Glacier

overnite camping

PBS-TV cutthroat Marty Stoffer on Yellowstone

Bozeman transportation -

Jerry Wells - 586-5410

Mont.
5x7

994-3551

Rob Lezny

Dolly Varden, Arctic Char

(Dolly Varden L.) - ca 6 of each in 3 dif. waters

ca 40 loci - only few allels at 2 loci

show any real dif,

Bull trout - 2 loci

~~III~~

585 malma	7 pop.	756 taroneta
cœca 17-36	(26.9)	31-64 (45.4)
range \bar{x} 26.1	- 27.8	431 - 48.7

of 756 taronetai spec. 12 (1.5%) have less than 35 (31-34) cœca.

of 585 malma 7 (1.2%) had 35 or 36 cœca.

but specimen readily classified on basis of spot size, body shape \bigcirc lat. compressed \bigcirc taronetai, fin & mouth coloration, malma rounded

gill rakers --- no intermediates --- no hybrids

range of overlap + even 31-36 - clearly identical as malma or taronetai

19 of 1314 specimens (1.5%) separate on cœca -

13.1 $\frac{1.4}{19.0}$
1314
5860
525.6

verbal legend main

- Volokov et al. - form I from a "hybrid"

ignore Chershev's diagnosis - < 30 cœca > 35 cœca 30-40
almost desperate but feeble, naive attempt to conclude no reprod. isol. - single polymorphic sp.

Chershev skillfully hopefully

dissected arguments & bits of Volokov et al. demolished the hybrid's

and hopefully put to rest once and for all the "polymorphic sp."

concept to explain all diversity of Fin. Farther charr

- all over Chershev's Pen native people no trouble

recognize the sp. Polly Varden is "Kingyn" and "Taronetai" charr -

"phenotypic diff" "lyginnen"

- Chershev correctly emphasize that the "polymorphic sp." concept is a "very negative influence on char. fisheries" by ignoring little hereditary basis for dit. malma, lipkin - igry "stock" concept of ^{problem} fish mgt. -

SH 287
D 73

Nyman, et. al. 1980⁸¹ (¹²) *Nottingham Rep. 59: 128-141*

alpinus complex extreme low level of polymorphism - seems to lack genetic variability - heterozygosity but ecol. plasticity -

3 species based on esterase allelic freq. - Beluk (1980)

asked what 3 sp. -

alpinus

salvelinus

stageros

Alpine in Austria

Greenland

no data type locality - all "3 spec" in N. Europe.

electroph. limitation - clonalistic - tax.
 - Ferguson - 5 loci - 12k trout
 - STOC - 5 loci - 12k trout

* markers primers
type

- Nyman - no attempt correl. est. w/ other tax. chem
 (note: "species" assigned by other fu - but esterase
 labile - red shiner - Kornfeld - if ~~ex~~ Nyman intelligent
 - but suppose - track over fish sp. data - see ~~particular~~ ~~data~~ on
 polymorphism - how utterly foolish to assign "species" to
 arbitrarily allocl. freq. - complete neg. - what if?
 "adds new dimension to view of treating all Mendelian
 polymorphisms as simple pop. markers" - right! - trust as sp. markers

ment: n

* STOC Symposium -
 Stockholm - Gene pools -

Ryman & Stah (81) - STOC 3f(12): 1562
 37 loci 9 Swedish lakes - 5 showed no polymorphism
 (no heterozygosity detected) - EST-2 - 4 w/ polymorphisms only on
 w/ MDH-4 polymorphism - correlation - phyletic implications

Ferguson (81) - only EST
 EST-1^{low} (100) = "S" of Nyman -

EST-1^{fast} (115) - variable .9 lakes
 8 Irish lakes
 + 2 pop. Windemere

markers

7 Irish lakes - 22.5 - 24.7
 Coomasaharn 29.1 N=15
 Windemere sutor 21.15 - 19
 Spring 25.33 - 12

* > 60 loci ?

data 39 loci
 Gen. distance (Nei) sin.
 brook-lake = .353 (647)
 brook-bull = .373 (627)
 bull-lake = .267 (923)
 Stoneking et al (81) p. 810
 Leary
 6 no alleles in
 4

Exp. back

Kornfeld et al.
 polymorph. only (>2%)
 EST-4
 except LDH "5" only Arctic clade
 + PGI - 105 unique -

211 monomorphic in alpinis	confluentus	fontinalis
Aat-1	100	54, 191
CK-1	85	100
Idh-4	106, 135	135, 82
Ldh-1	54	0
Ldh-4	76	28
Mdh-1	144	110
Me-1	110	56, 110
Me-4	95, 97	97
Sdh	120, 125	190
Sod	177	97

Clayton & Ihssen (80)

* LDH "E" - Nettling L., Baffin Is.

AA	AB	BB
5	5	2

Cambridge Bay (56%)

Table 1

LDH E

(A allele incl. freq = 3% to 7%)

6 samples N.W. territory

but Montreal L. (Quebec) - inter-ogone-ti - 0 freq.

Globokovsky and Chereshevnev 81

Unresolved problem.

Barsukov (1960) - suggested one polytypic sp. - touch pulled in and championed by Savitskaya and colleagues.

- Numerical Tax (small phenotypic unit) 60 characters of skull bones

malma malma 18-25 (21-23) long, thin
65-69 (32-36 ab., 30-35 caudal)
18-36 (25-28) thin, rel. long

truncate

23-26 (25) - short, thick

65-68 (34-36; 31-32)

39-52 (46) rel. short, thick

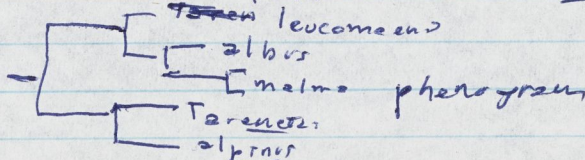
forked

Anguema & Erguveem river basins → malma, truncate, albus

alpinus sympatric

leucomaeus sympatric

Kamchatka R. - malma albus



discrete or phenogram

Cladistic interpretation of charac. → concludes alpinus, truncate +

leucomaeus more primitive malma albus more derived character - but:

characters - bone shape - weak basis for prim. der. - subject to convergence &

independent der. → no. + phylog. stamp.

selective use -

origins too recent for

'type' - south mt low local vari. rel.

trenchant osteological dif. sufficient

valid basis cladistic interunit

"Liver esterases"

McCant (1980) Can. Arctic - all eastern char (S. 21-50, 22-50, 23-50, 24-50, 25-50, 26-50, 27-50, 28-50, 29-50, 30-50) show "fast"

pattern (drafts of Nyman) - but largest # all alpinus - would second - but 98.6% of west.

2 malma have "slow" esterase

- but to 11 yr, s.a.c.

04

221 Thesis - Mayr & Speed 20

Mc Cort (cont)

Liver esterase" (EST-1)" Nyma?

East Arctic (colpina)

78 spec, 5 pop. = 100% "fast" = alpinus

West Arctic (malina)

212 spec. 13 pop. 209 "slow" (98.6%) stagnalis

3 "fast" (1.4%)

Table 6 meristic data range \bar{x}

13 pop. N=617	fishers	\bar{x} 21.5	range 19.4 - 23.4	range
N=618	gacez	\bar{x} 29.6	(18-40) 27-33	18-45
N=201	venti	168.0	67.1-68.9	60-70

-Trend for small isol. resident pop. for lower meristics

Fraser & MS. Thesis - The interactive segregation Arctic lake food chain, southern Ungava, Quebec.

No Name L. 211 3 sp. - alpinus: gastropods, Gammarus, 378h

Ducreeux L. - only alpinus - piscivorous \bar{x} depth 10 - max 32

chain + 3 spine stickleback

3 sp. charr, sticklebacks, + few burbot x depth 2m 407h

alpinus predaceous - 20-30cm > 40cm ~100% pred 95% fish vs. resident -

but overwhelmingly cannibalize own young in pref. to stickleback.

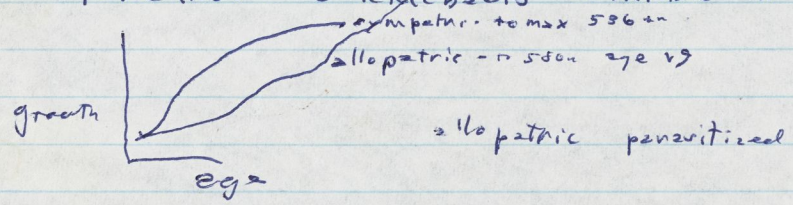
L. Nonano - alpinus all sizes (to >50cm) virtually no fish in diet - zooplankton to 30cm then snails, Gammarus.

Snails & Gammarus only few% in Ducreeux - main inverte. Trichoptera.

- Table 5: Nemayech - predominantly fish from smallest size (20cm)

fontinalis - ^{benthic} mainly insects, caddis, larger more snails

lake trout - sticklebacks dominate to 50cm then charr (+ burbot)



allopatric mature at 4+ 5+ (15-20cm)

symp. 8+ (>30cm)

Covender (this vol.) : S. malina "londi" 2N = 82, 98 snms - identical to terrestris

contaminant: 2N = 78 102 snms

alpinus 80 100 snms

4 AM)
W 205

Hinder and Jonsson (1982) - Habitat and food segregation of dwarf and normal Arctic char (S. a.) from Vangsvatnet Lake, western Norway. Can. J. Fish. Aquat. Sci. 39(7): 1030-1045.

Nordeng this symp. "solution to char problem" -

dwarfs 25.2 \bar{x} rakers + no spawning colors

normal 28.5 \bar{x} - bright colors

change from pair \rightarrow normal char corresponds to pair \rightarrow small transformation in anal. char.

sharp niche segregation (dwarf + pair) + (normal)

called "morphs" implied but not explicitly stated that dwarf & normal are polymorphism ^{non-lex} in single pop (no data given \rightarrow spawning - do ~~dwarfs~~ ^{only} spawn only w/ dwarfs many or no? -)

influenced by Nordeng (this probably wrong)

- May be genetic polymorph as w/ Arctic char ^{max.} ciakha C. W. (Kornfield 83).

Kornfield, et. al. 1981

samples of blueback char 5 pops.

Sanctuary (Florida Fl.) 1

Quebec red char 1

Arctic char, Cambridge Bay N.W. 1

genetic sim. \rightarrow tax. \rightarrow see Hinder - sturgeon -

see Gen. Stock

G.S. Homo sapiens chimpanzee

Eastern red stream Br2205 R.

else pattern

enzymes p₁ p₂ gene loci examined

2 esterase enzymes - EST-4 variable

Wadley	Quebec
11 alleles 100 - 84	98.50
95 96%	1.50

*LDH-5 all monomorphic for 100 allele

Selectivity, interpop needed: except Arctic only 40%, 100 50% 92

only 7 fish Arctic sample

+PGI -3 80% occurrence of 105 allele, not found at all in others - 6 of 7 here

"(Hinder when all data ^{set loci} "homogenized" the overall ^{genetic} similarity ^{of} monomorphic loci"

masks dif - and Arctic char - not stand a

"numerical" leading to erroneous ~~conclusion~~ ^{conclusion} that separate subspecies recognize for "Arctic" as ^{may} not be biol. reality; (Hinder recognizes "S. a. alpinus" by authors)

consistent by 6-7 mm rakers; 4 rest - no way this dif + explicit in genetic info.

UW

Russian book

Cherekhov - size Pzc. salmonids
for IGA* malma to 830mm. (33 in) 5700g. (12.5 lb.)
taranteli to 680. (27 in) " (6.6 lb.)

E. Sakhalin charr 21-25 cases malma?
21 taken
60-62 mt.

Viktorovsky et al. Charrs of the genus Salvelinus from Lake Elgygytshyn (Central Chukotka)

Mountain L. El. located "border" Anadyr R and Chaunsk Bay drainages - 489m el. - never submerged by late Pleistocene marine transgression - 12 km. diameter, 169m

if refuse
p. 456
extinct
convergence

max. depth - summer temp. hypolimnion 2.5-3°C
besides Salvelinus only Cottus cognatus and Thymallus arcticus which both occur in Anadyr. and Enmyraem R.

July - Sept. 79 115 charr collected from 17-70 cm.
2 forms well represented -

S. bogenidae
N = 38

S. elgyticus
N = 67 D S A T

caeca 42-66
taken 23-31 (27.32)

25-58
39-51 (44.03)

fig. 1 - small mouth - max mt. to ant. eye sparse, large spots.
fig 2 "S. bogenidae" - sparse, large

- one of forms appears close to S. bogenidae Berg. 1926, 48, but with slightly more taken (27)

fig 3 skull-chondrocranium S. e., S. b.

S. b. - prob > 1m. in L. E.

S. elgyticus Viktorovsky et Glubokovskiy - "Malorotsya" charr (smallmouth charr)

Holotype 3414 45397 large spots, no need spots,

feeds only on plankton

L. E. - reposit - meteorite crater lake of late Pliocene

B. (1980) - 2000s Sartan Siluria interior Lena - Indigirka
isolated lake ca. 29-36 maa erethroni

p. 456 - state Chukotka charr long thin taken = alpin
wrong!! short, stubby " = malma

Jenkinson (80)

EST 1 (100) allele = slow or "S" type

EST 1 (115) " = fast "F" "

Myrica senon estensis

S₂ alpinus - Linn. Swedish Lapland

"F" char. F (in 115 allele) = .9 - 1.0

least competitive, stunted sympatric, ^{may be} large to adapt

So. Swede
L. Västern =

- S₂ saluelinus Linn. Lake Austria.

large, predator

N - normal

.4 - .7 at 115 allele - but may be fixed = ?

1.0! (how diff. to alpinus |

S₂ stagnalis - Greenland.

115 allele .0 - .2

"- introgression?"

January 5, 1982

Professor Y. Yasue
Nakamachi 5-17-0
Setagaya-ku
Tokyo 158, Japan

Dear Professor Yasue:

I received the special publication on Salvelinus. It is a beautiful work and I send my sincere thanks. I received another copy of this same work from the "Freshwater Fishes Preservation Society", Osaka, but there was no letter revealing the name of the person who sent it to me. If you know who sent me the additional copy, please give him my thanks for his thoughtfulness. I will give the extra copy to Dr. Ted Wavender who also has an interest in the taxonomy of Far Eastern Salvelinus.

I also received your abstract on the comparison between Formosan and Japanese O. masou. There is much additional data supplementing my publication of 20 years ago. I would suggest that you publish an English translation of this paper in the Japanese Journal of Ichthyology. If you can make a "rough" English translation and send it to me I can edit it for you for publication.

Sincerely,

Robert Behnke

C
O
P
Y

3 forms Arctic char

L. Greenland

- may change into each other
- transitional stage for speciation and isolation

1986
C J F A S 43(5): 985-92

- visa - R...
- D...
- passport
- tickets if not
O...?

Review -

Syst. - Tax.

- study of diversity actual alborit. limits modern techs

- 1980 - plus...
- main change

- electrop gene loci
Ferguson et al -
Creative Sol. of Solun

sympatric
Iceland Drottninghol
Sweden
Cyprus
Thiruvalluvar
1984
1988

Dolly Undergo
78 South
bull, time
82 dm

Theoretical
but the
good
review
observed data

R.B. - cuts
- subsp
overall phenom
- not cladistic

new pap
have may do
- Greenland
- Scotland
morphol - vs - genetic

but alleles
mains - electrop
confluent
* new Cladistic

1980 books - character states
- no firm knowledge of phylo-sp.
+ best basis of playing genes - ex. K...
chromosomes - homologous -
- playing genes - ex. K...
- homologous -
- playing genes - ex. K...
- homologous -

- Karyotypes
--- no m2
high - low

welcome comments

why study - intern
- smart - met -
- 2 the - quest
- culture quest
- remain unknown

but indelent -
some charac
near
2nd spec
ex - 78 - 2 spec
8 - confl
convergence
- as of volck
no. vent
costs
preserv
above
- voter tests

review changes open 1980 books
- 1981 - (54)

may we look
for major contribution
- electrop
- genetic
- genetic

hope Dolly char compares part to rest to all D
(V.I. - - Am - Kauch... - 2 spec
- 2 spec

L. Elgygys - small words
by next symp. - perhaps phylogenetic units
small words - under-established?

profundus - elgygys
- material work
with
- don't mix spec
- electrop, ch... bud...
- DNA hyd...
- elgygys - prim... - det... - G.D. 1.0 - 740...
state - -
- isthmus Panama
- 2 spec

My major change - leucomensis
ball - stone

ughy - how so adapt
ecol. polymorph
- mensuraphic loci
- structural: vs. regulatory gene
- ne. Cooper. Biol - 'subbreeding'
- danger. S.R. cuts - knill.
- quest. wgt. - - quest. - like birds

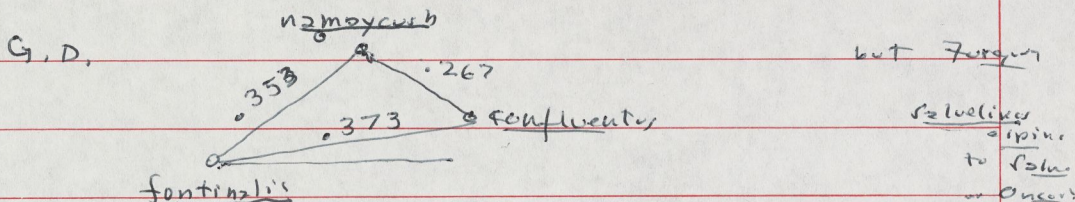
but new genus
- Conf
- 3

Stages of sympat. divergence
subgenus

1 spec
- bull
- 2 spec
- with
- longhead
- profundus

Stoneking et al 1981: 810

Leary - Allendorf - Syst. Zool.



need electron = 40 in one

leucomensis

straw silva

malma = north

profundus

* Elgygytyn

L, Elgygytyn -

tax, not autonomy S. solvayensis

malma all 98% 100

EST-1-100

Arctic silva profundus

all part EST-1-115

-P01-

h. Elgygytyn - why no Conegoua??

- Katsvtsjev, Glubokovskiy, Cheresknev 83 -

28 loci - malma - taranetz: Chukotka

Idh - EST-2 freq. dif. - but fixed dif.

of Aeph-1 - D. .08

60 64 64
58 - 64

60 2x indopina
64 + 64 orth.

sub genus Cristivona 84 84 84

leucocentris 84 NR 100

Czender (1) (100) ?

confluents 78 - 102
Phillips - Arrow L. B.C. 12th metacentric
of Czender actually large acrocentric
4 NOR on short arm

alpinus
78
25 - 98

all west. Pacific Isl.

1 pr. NOR.

malma southern 82 NR 98

Phillips

- Although all sp. Salmo & Oncorhynchus have NOR in only 1 pr.
NORs on several chromosome pairs of mamajacob, fontinalis
alpinus -

malma - Juneau AK

total no. NORs / genome 4-10 fontinalis

4-12 mamajacob 2-6 alpinus

variation on NOR placement in chromosome between individ.

ie. - "constant" sites & "variable" sites

+ leucocentris ?

- confluents similar NOR to fontinalis + mamajacob polymorph

- malma + alpinus have synapomorph - NOR on largest submetacent

malma (Cronholm) & alpinus share derived acrocentric which
is 2x longer than any acrocentric in other Salvelinus.

★ - malma → alpinus by electroph. - karyology

rather than leucocentris - canis

- NOR location
transloc. on sub
synapomorph.

* NORs - alpinus 2 in N.W. Terr. - 6 in Scotland

- miyabe - great dif. NOR placement long-short.

enzymes at

Henry - 40 loci - no real dif. so: AK malma - AK alpinus Tarnetz

2 loci distinct w/ bull trout.

- Conserv. Biol.

why such
ecol. - evolution

plasticity or not

low levels heterozygosity

- inbreeding to

what's overrepresented

genetic genetics

Sparholt 1985 J. Zool Biol. 26: 313-85

lake in Greenland - 2 forms differ size-growth,

no sig. dif. & conclude one pop. -

1. Rapid morphol. dit
- little genetic change

Walker, Green, & Gardner 1988 - Biol. Conserv. 43(1): 43-61

2 forms - Loch Rannoch -

- Norway
- introduction
divide into
1000s
A 2 types
schol. 2
17 - Greenland
1000 yrs
1 - Iceland
4 - Thin

benthic dull color - pale large head 1% cladocera
pelagic clear color small head 95% cladocera

- rarer
niche separation
sp. feeding
reproduction

- Nyman: 2 sp.
esters:

divide
- redivide

Magnusson & Ferguson 1987 - Gen. Analysis sympatric
morphs Thingvallavatn - Environ. Biol. Fishes 20(1): 67-77

Small benthivore, large benthivore, planktivore, piscivore

D values ~ .00004 = .00126 re. Loch Malin trout - much greater
longer? or Salvelinus

Sandlund et al (Gydebo) 1987 ibid. 20(4): 262.

habitat use - post glacial retreat 11,500 yrs B.P.

polymorphism
Cuz-to
change
one system
in one sp.

why? increase abund. sp. - but reprod. isol.

Netun - unture

mgt. vs. sympatric
obscurely many species

Condens. Biol.
presumably distinct sp. diverg. in
for 1000 years
Pyramid
ecol. potys

Riget et al. 1986 C. J. Zool. 43(4) : 98
S. alpinus Greenland 3 groups exhibiting
incipient, sympatric speciation.

(Emerald L.)
CO.
100 yrs

Klemetsen, et. al. 1985. Drottningholm Rep. 62 98-119

loch
Rundeech

Boer Is. chain - several lakes S. alpinus only
fish sp. - find 2 'morphs' - small, young spawning,
short life - vs. larger - + coloration

esterase - no signif. dif. - all predom. 7-11

but - caeca	L. Ellsjoen	36	small 22-25	large 45
	L. Stevns	35	23-24	24-25
				46

re. loach
Melvin
3 brown trout

Hindor, Ryman Stahl et. al. 1986 - Biol. J. Linn. Soc. 27(3) : 267-85

- ex. of known introductions 1910 some lakes - normal and dwarf appear ... (schooling) - Review evidence 3 sp. ancestral sp. Sci

1. Anadromous 2 dwarf 3 normal - no relation in EST freq. allele

- All Scandinavian chain - one, recent common ancestor - lake spawning promotes symp. spec. - (see same issue - re. when symp. spec. possible : 201-223) - Cuzto -

chain L. Elgryn
- similitude - change by host
+ chain 57 m
better phylog. pos.
1984 - hopefully the
erroneous - strain
that make - a lot
one sp. - put to rest for all m.
- V.I. - Swazis
- in Mison
- 1/2

- Gydeno 1984 1981 Thingvallavatn Iceland
species from Scandinavia as defined by Nyman et al. (1981) are present in Iceland also. "The landlocked chain of Iceland were already separated into species when immigration occurred." - 1987 - Enviro. Biol. Fish. The Arctic chain of in

Thingvallavatn. is polymorphic. Genetically the four morphs are very closely related. The morphs are conspecific and do not represent different evolutionary lineages. "The morphs, in Thingvallavatn have probably developed within the lake, and the morphs are locally adapted to different niches."

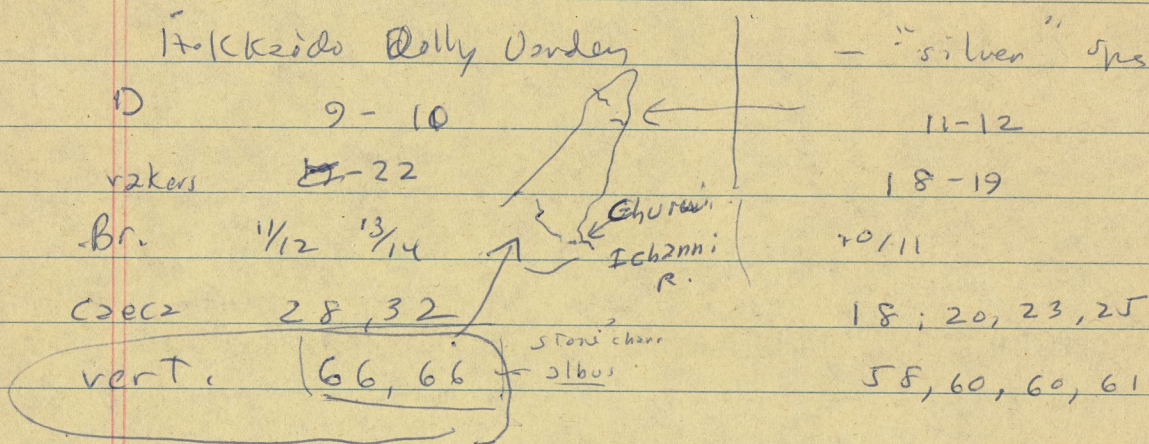
compare
L. Elgryn
Convergence
benthic - pelagic
growth, stream, mtn
low mtn
- head, river, lake
- convergence
- but Dolly
- only stage

Hindor, Jonsson 1982 - Vangvatnet - dwarf, pale morph, large, colored m.
- same gillraker no. - niche segregation - local phenomenon
in relation to lake morphometry, spec. acc., niche availability -
- dip. color, growth, eye melanin, - not gillrakers, (caeca?), etc.

Andersson, Ryman, Stahl 83 - Nyman wrong.

2N: pluvius - leucomaculatus → curilus D.
curileus D.

see Mj -
 Sex-run silvery specimens Oketchiushi R. a
 Shoji R. - Shiretoko Pen. Hokkaido



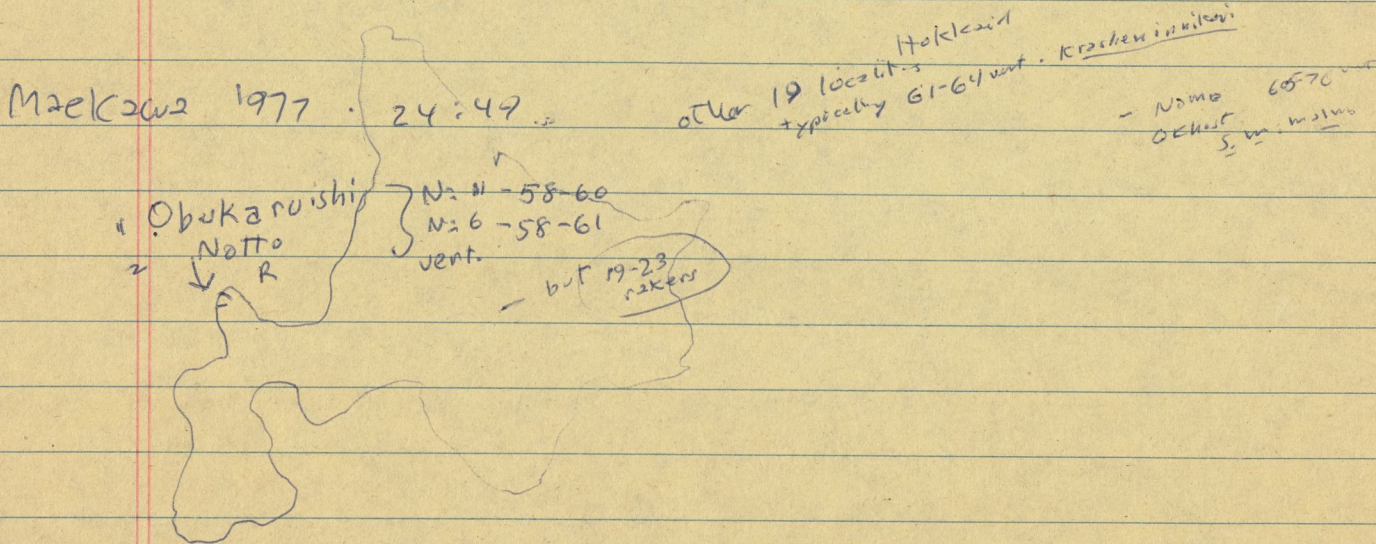
Sm. Krashennikovs
 19-23 2N = 52-84 982m
 61-63 vert. 76-78 965m malms
 25-30 cae

84-86 = leucomaculatus + curilus
 100 arms

↑ 1m
 ← expect Hokkaido
 Hokkaido

19-22

61-63



- Impact assessment 400 cfs depletion.

- Squawfish, humpback chub, ^{De Beque} razorback, channel catfish
= Mitigation-enhancement measures

- flow no way solid flow requirements

- 2 of 3 yrs. ... Aug. ... temp more in fl flow abt 2000-3000

- Need actual study ... :

^{FWS}
^{HEP}
^{IFG} - Use FWS on L - miller = 75% exceedence flow. - not to drop below

- go over steps 1-2-3 best success - 1980 : IFG (depletion work) -

75% exceedence flow ...

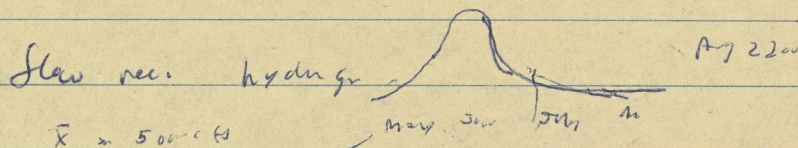
- flush flow)

- Status of squawfish "Archer" - not valid. - no Colo. R.

- actually more adults found recat ya - no data < 1975

rationale assessment - background - in-depth info available to GCC

- actually humpback
CFT on
humpback
razorback - rsc
squawfish
bottom
the



$\bar{x} = 500 \text{ cfs}$
- normal yrs. - wet - low - very low

- err on side of endsp. - no. 1000 run 10-20 years ago may still be declining
- play; need more than present conditions - Archer = 66% decline juv. & adult - 1960-80

94% yoy 1960-1980

Squawfish - 30 days 20+ spawning larvae 240 d on
humpback - 16-18 spawning 20-26 egg 120d develop. 24-26 growth

- regulated river reduce peak - increase some beneficial to aquatic life

- Bolon's Reprod. Guilds

I

pluvius = leucomænis - convinced by Japanese ichth. + caecid. (Tri) gillnetter may 1925

②: For East Dolly - southern - sub sp. Hokkaido
19-23 rakers - 61-63 vert. (Krascheninikovi) - also 2N chromosome
-- Komiyama et. 21. 1921 - "Dolly Varden"
curilus 2N: Shiretoko Pen. E. Hokkaido - color photo
plo. "pluvius-lila" 58-61 vert., 18-25 caeca,

	rakers	caeca
Viktorovsky - <u>Krascheninikovi</u>	15-21	16-31
<u>leucomænis</u>	12-20	11-33

small spots
feverish
Dolly - low caeca
 $\bar{x} < 35$
= 1 pr.
> 35 (except Labrador)
gill net $\bar{x} < 9$

2N Viktorovsky (78) - but aware of
caeca
5. gillnet
5. gillnet
40 subsp
morphe
prim 5f
-2- →

<u>madra madra</u>	76-78	- 96	ca
<u>Krascheninikovi</u>	82-84	- 98	lonki
<u>curilus</u>	84-86	- 100	
<u>schmidti</u>	78-82	- 100	
<u>Kronocius</u>	78-82	- 100	- 102 - 78 confusio
<u>leucomænis</u>	84-86	- 100	

80 - 100 2/3 pr

3rd group predator
leucomænis - bull, ston, kronocius

Kronocius 63-68 vert. (66)
18-26 (24) rakers
24-43 caeca

schmidti 75-24 (29) rakers
63-68 (66) vert.
23-37 caeca

③ Volobuyev's OT 21 (78) cast subsp Hokkaido, K. - ca
- S. elegans

III

Chereshnev (2) transferred ch=rr

Dolly Vant

light dark mouth coloration mixture of or kype brown jaw

light mantle color - no kype

Mackwa 1977

Hokkaido vent.

caeca personal comm. c. 30

2 on west coast 58-61

others \bar{x} : 61-63

rakers - 19-23 \bar{x} : 21

others \bar{x} : 20-22 miyake 25+

1983 MS Dempson & Misra

samples from

7 samples Labrador charr

vent. 64.8 - 65.5 62-72

rakers 23.9 - 25.7 20-30

no caeca curbs 20-21 27-28

Chereshnev 1982 V.R. 22(6): 922-96

S. taranetzi Kaganovsky, A.G. 1955.

To Golets iz basseina Beringova Morya. V.R. (3): 54-56.

illustration - few, small spots, long jaw - prob. malma

rakers = 23-25 caeca ^{mal=} 32-42 - certainly also

morphomet. data 10 spec. 165-257 mm T.L. - young = syntypes found

S. taranetzi coll. 12 Aug. 1948 by P.G. Nikulin

L. Achchen

but meristic data - Table 2 - no. specimens not given

no info given on fate specimen - museum? museum w. (Watanabe)

+ T UhtPO - Pacific Ocean. Fish. Res. Inst.

Compare of morph. data

taranetzi lectotype redescribed

Borsukov (1980) - Bengi type # ZI 31107 of andriashovi - but only 14

rakers, but resembles of this spec. - when 19 rakers on right,

21 left side -

- Chereshnev (cont.)

584 malma 7 pop.

18-26 (22.4) rakers

2.5 - 21.8 - 22.5

L. Achchen 20-26 (22.5)

A fig. 1 rakers malma

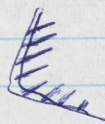
756 taranetzi

21-30 (25.4)

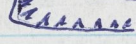
25.0 - 25.75

22-30 (25.75)

taranetzi



denticles



as leucogrammus

Kozlovskiy

Unanswered Questions

Hokkaido

- Euzona -

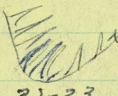
Komiyama et al. 84(?)

Sessun silvery specimens Oketchiushi R. & Shoju R., Shiretoko Penr., Hokkaido
Hokkaido Dolly Varden silver specimens

D	9-10	11-12	
g.r.	22	18-19	leucocarinis
caeca	28, 32	18, 20, 23, 25?	
Branch.	11/12 <u>13/14</u>	10/11	
Vent.	<u>66, 66</u> ?	58, 60, 60, 61	

Maekawa 1977 : 19 Hokkaido localities, Dolly Varden
 61-64 vent. - but Obukaruishi k., Noto k. -

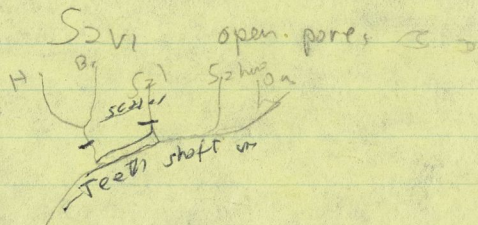
N=11 58-60 N=6 58-61 typical leucocarinis
 rakers 19-23 typical malma

Chereshnev - Chukotsk. - malma  21-23 Tsrenetzi Alaska type Andriashvili 26-28 Andriashvili
 E. Arctic form? N. slope AK Andriashvili 23-26 Andriashvili 26-28 Andriashvili

N-S Dollies AK 1957 -
 AK willow Crk - near Anchorage - south-north D.V. sympathy ???
 simply count gillrakers. !!! no one took since ???

Kensi R. - 9-10 lb - Tsrenetzi - Andriashvili AK ??

Savsitava - Hass - confluent - ecol. equivalent of stone charr (river predator)
 white, longnose, snout charr Kurotsuka - not stone charr. albus but stone charr



(1)

Welcome comments > 35 y.o. Sanders - Okunskas in R.I.R.
Iwona, Yamae ... before formal higher ed. - but set direction

This Int. Symp. - Proc. - 3rd major vol. devoted to Salvelinus - Note range info - ecology, physiology, culture, disease, conservation - but our Sect. B

Taxonomy & Systematics - the heart of the matter -
The PROBLEMS ^{then} answered questions, disagreements, controversies - w/o this controversy - probably not sustain interest to get all people together for another meeting of chair fanatics - so let us agree to disagree in ^{friendly} _{cooperation}
- Thus in anticipation for next symposium

benjo for benjo
zealot
hass reason
- even spelling
r vs rr

I do not want to do away w/ the stimulus of controversy by telling you that I have now indisputably resolved all issues - and all will surely agree! - Chernshnevi's new sp. 'genus' let's us know what is yet to be discovered
- BUT re. PROBLEMS - re.

Patterns & Processes

Taxonomy & Systematics - perhaps helpful if attempt to separate - Taxonomy - actual classification - ordering of patterns observed into phylogenetic framework - Systematics - attempt to understand the processes of evolution.

polyphyletic
not
monophyletic

- Re. sibling sp. - sympatric pop. - N. Europe matter at systematics - not taxonomy disagreement
Nyman - ^{pattern of test alleles} Not 3 ancient phylogenies - but if 3 coexist = 3 "biol. sp." - 3 niches for mgmt. - for conservation

Re. How can chair do this?? - How can such ecol. polymorphic sp. be so genetically monomorphic?

- Questions of systematics - the processes of evol.

→ When phylogenies separate for sufficient time or 100,000 - mil years - identifi. & describe of them

W

= taxonomy - Thus trilma - alpinus matter of tax. and species recognition -

- Don't expect Nyman's system - you're correct - we're wrong - Belief system -

Personally - open minded, flexible, changeable when new info available - wishy-washy - non-fanatic

- Lot's Review my earlier position 1980 book

1981 symp. (1984 pub)

(11)

Assess 'Progress' - or Hindsight more accurate than foresight
1980 book - (written 10 years ago). Re. interpretation of
phylogenies - assessing evidence for classification -
"Without a sound basis for ~~determining~~ ^{establishing} plesiomorphic
and apomorphic character states (primarily-derived condition) for
determining phylog. branching sequences and 'sister
groups', we are only playing games."

~~1981~~ Nyman (Drothingolun school - Svendsen - Mayr) -
esterase allelic freq. for phylogenetic markers (OIL for 10,000 y
not 1 mil) - 81-3 sp. paper - Taxonomy alpinus, salvelinus,
stagnalis - Gydemo 1984 book - Iceland - Thingvallavatn

"It is clear that all of the species from
Scandinavia as defined by Nyman et al. 1981. are
present in Iceland also" - "The landlocked charms
of Iceland were already separated into species
when immigration occurred."

- Gydemo (1987 as coauthor) - Environ. Biol. Fishes. -

"The arctic charm in Thingvallavatn is polymorphic.
Genetically the four morphs are very closely
related (G.D. 00004). The morphs are conspecific
and do not represent different evolutionary lineages.
The morphs in Thingvallavatn have probably developed
within the lake and the morphs are locally
adapted to different niches." (- As Thingvallavatn ≈ 11,000 years
old - sp. flock-rival cichlids of L. Malawi) -

- * Major change outlook sep. syst. - tax. - not "species" - but
symp. reprod. isol. pop. real. - not single 'polymorphic' pop.
term "morphs" - not good - such as Cuzco Cuzco cichlids.

FC - Symp.
- 1984 book "Any reasonable standard for sp. recognition
malma = alpinus are separate sp. - "Hopefully the erroneous
notion that Dolly Varden = Arctic charm are only polymorphic
forms of single sp. put to rest once & for all" -
- then V. I. - Moscow school - S. alpinus Komchuk - Amer - Sokolov.
(Sov. 1988 need for Int. Coop.) -

- My changes 1980-84 - pluvius for malma - leucocomensis
re. 1984-1988 - slide - new subgenus Salvelinus

malma + alpinus (super sp. - complex) -

leucomænis - ~~confluens~~ - albus - Why

- Karyotypes ^{RUT 011190} NOR ^{Covered} - N7 - 100 w. 99
electroph. - 40 loci Univ. Mont. malma - alpinus - now
confluens - 2 of 40

Predicting ahead

- Needs - ^{limited but} electroph. profiles - leucomænis, L. Kronotskoe
Kamchatka - 3 - longhead, white, snout - stone char

Elgygyta - profiler - karyotypes - but so remote - Europe
- profundus Tifferssibling Bodensee - no data !

- mtDNA mtRAA -

But limitations - overall phenotypic sim. = numerical -
non-phylogenetic tax. of prim-derived states -

- Recognize not constant rate divergence between
any lines any characters - Morphol. - ecol. (niche
filling) - can be very rapid - electroph. - mtDNA - variable
- Isthmus Panama - Ann. Rev. Ecol. Syst. (1957) - Hillis

- Karyotypes - NOR - banding - ^{best} hope for
cladistic arrangements - but NOR. stable vs. variable
(mixes ^{+ local variation} - quite distinct)

Cont. to line
Rev. since 81

Adds

Strong
ANTI
sympat.
spec.

Myr -
revisionist

- Anatomy of
Controversy - Mayr 1940s sympatry, reprod. isol. = sp. "biol. sp.
concept" - to taxonomy - Drottingholm school - belief system
Sweden - Coregonus - Outstanding systematic - wrong
taxonomy. - non ancient phylogenies = polyphyletic sp.

- Some Pleistocene period - 100,000 - 1 mil. yrs = ancient
lineages - - - 3 sp. charn alpinus complex - -

Rev. ^{2/11} - Early stages: ① Hindor et al. 1986 (Biol. J. Linn. Soc.)

Norway - 1910 some lakes stocked w/ common trout -
now drafts - normal pop. (spawning - isol.)? -

② Hindor - Jensen
1982-84 -
Vaggevatnet
Norway - no
tax. dif.

EST freq. a quarter of sp. - "All Scandinavian charn
from one recent common ancestor" - No 3 sp.

Emerald
L.

Convergence
at divergence

③ Klometson et al. (1985) Drottingholm Rep. - Bear Is.
sympat. pop. - no EST dif. (if introgressed to common gene
pool - then no 2 pop) - dif. markers (12) - (2002 (10) draft -
normal.

④ Riget et al. (1986) C57AS - 3 forms charn ^{12 ker} Greenland -
early stage of incipient sympat. speciation

IV

- more advanced - Thingvallavatn = 11,000 years
benthic - pelagic Then again = 4 w/ little genetic - good morph.
"niche" filling - low overlap.

Loach
Melvin
Trotter

(5) Loch Rannoch Scotland - Walker et al. 1988. (Biol. Conserv.)
- sharp morphol. divergence - benthic - pelagic -
- Alpinus - polyphyletic, rapid divergence
but convergence

conclude

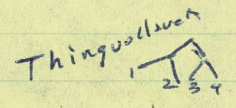
benthic	-	pelagic
size	smaller, slower	larger, faster
growing		
age	shorter life span	
	earlier reproduction	
color	dull, pale	bright coloration
rostrum	shorter, fewer	exp. spawning
caeca	fewer	
head morphol.	blunt snout	
	subterminal jaws,	
	short, thick maxillary	

conclude - slight sp
reprod. - but - not
to taxonomic sp.
wt.

Conservation

Ex. morph. rapid change
wo - Silva robusta
genetic
pointing
Cyprid
elegans
Gasterosteus aculeatus

Why only Alpinus? - fontinalis - malinus - leucodanus



- Message protect Biodiversity (book)
course - Society - - Danger quantitative
(pop.) genetics - if no dif. not gen. dif. (nature - nurture) - what highlight protection
ecol. distinct, dif. life history types -
Do not manage on 70s allele - can lose great diversity -

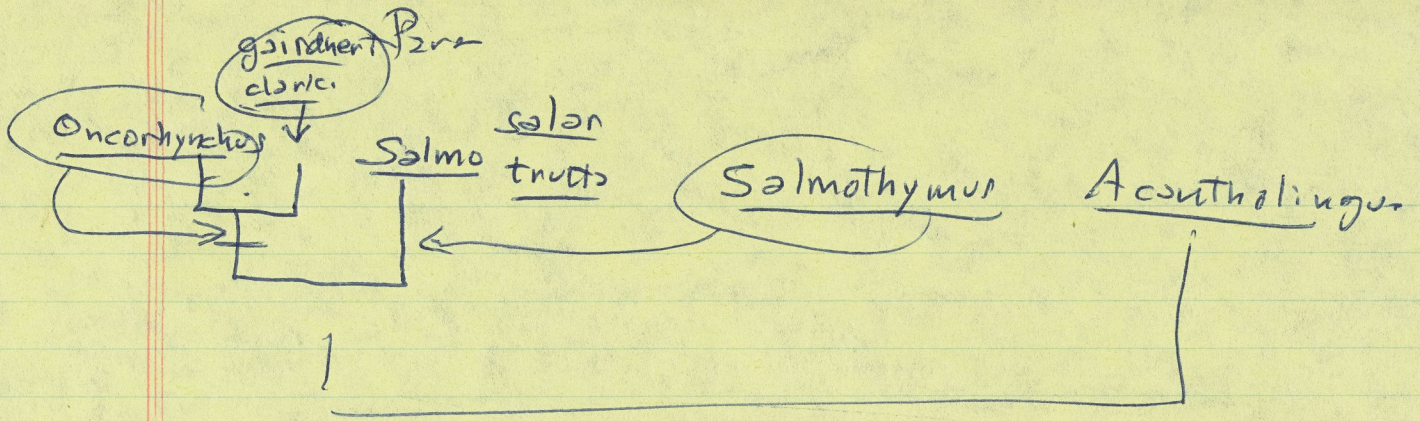
return
to look
ahead
to prevent
my change

Genetic
mistakes
Biodiversity

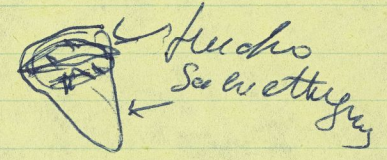
Human mind seeks order - Illusion techniques
order from chaos. - Quant. vs Qual. - lesson to
live w/ and appreciate chaos, on Polytomy -
Vitus vinifera - can etymol. profile - DNA character -
tell us how to make all Chateaux de Rothschild -
both nature - nurture - involved.

slides

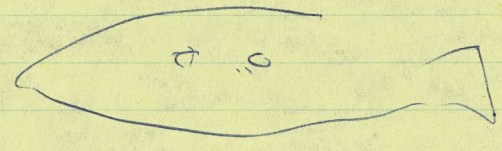
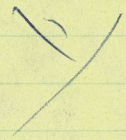
- if live to 100
- look forward.
future met of change function



Parasalms in
genus Oncorhynchus



1984. Kendall and Behnken



Pravda

10:30

Biodiversity - - so good to see again ^{Tovarich Barroly}

- 23 years: - but few hrs. of arrival - first bite of chummy kleb - as if 23 years but 2 day
- Editor for English V, L, G, K. - stay up to date Soviet fisheries

My topic - Preservation biodiversity - Conservation Biology society - 2000 members x journal - intraspecific diversity

weina grape vitis ^{vinifera} w/in sp. - let me demonstrate w/ slides

example Species - cutthroat trout S. elarki -

"Clark's forest" - even only Donofeev - Barsukov - even Amer...

- but - once great distrib. --- Rich - Colo. - only trout -

- big brown, 100 yrs ago - only cutthroat - → 95% gone

- What look like - Great Diversity - 14 subsp. - many, 1200

30 years many believed extinct - discovered - S. c. utah -

- Greenback - Colo. - E. Cont. Divide -

- Why - what benefits - beauty - spiritual ^{to us} ~~recreation~~

* L. Boical - ~~Techno-illustrate~~ illustrate similar ~~not conservation~~

Problems - Technocrats - ^{just} forest-wood → paper - horrendous pollution - stink. Nation snored - - blight on ~~waterfalls~~

- first great victory -

- Practical values - unique genetic resources

Pyramid L. cutthroat 1938 - why save? - subsp. not extant

- Redband - 28-29°C -

Chukotsk - Kamchatka - czar - golets, polix -

how many sp. - Moscow school. - - Do not let

classif. controversy block preservation - S. t. caspius -

same Kurz in ^{winter} pop. - (stock L. Michigan when ^{clupeid} ~~lewis~~)

* Syn. 1st spawning - Genetic resources ~~don't~~ saved -

- Educational - Lit. - St. - Fed. WFA - Public -

Progress Conservation of Nat. Resour. - Evolutionary - slow grad

not revol. - - change ^{2w} ~~attitudes~~ ^{attitudes} -

Gen Kstuzov - ~~napoleon~~

costly for \$ Pechor 15-20 kg ^{salmon} - 1000 R ^{fish} - age vs. command - - tags ^{word} ~~hydropony~~

Caspian
- Kurz R.
51 kg. -
save

Karluk R. —

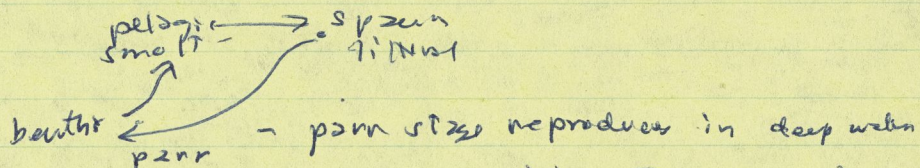
(Liver) Aat (perhaps?) allele 100 Dolly Udder allele 30 char
AST 9 (pinks - chum)

~~PGF~~ 1-2 - mainly pattern dif. - mobility similar
but intensify size of band.

Leucyl - leucyl peptidase - LLP - dominant char
is very rare in Dolly Udder (may essentially
be 0 in Dolly) — total loci: 27

— Wilmot - No Slope - stock repetition
US FWS Anchorage

— Sympatric spec.



— Play free & loose w/ taxonomy - if taxonomists
ambiguously played so free - unsubstantiated - speculation on
ecol. subjects - ex. - look radical -

- ex. type locality - EST Junc. fr. - type locality salvelinus stagnalis

- tax. - dollar plus
- p blood thick as thin blood ecology - fat rich coarse ground habitat

*bot
salvelinus
valid sp.
second name
Alice

No

- Chenashvili - general (problem)

- Resolve problems - not all but up to date 1980-84-5 - helpful

I Taxonomy & Systematics Patterns & Processes

patterns - processes - taxonomic disagreement not necessarily systematic disagreement or 'problem' - Ex. 3 sp. ^{Scand. N. Europe} - Iceland -

S. a. alpinus

flexible - open minded - to assimilate new information - new evidence

Review: 10 yrs. ago 1980 I wrote " " " more specifically

EST 7-5 not - type of plasio-eps - character state for identifier ancient evol. lines - to date all lit. agree

- ex. Thingvellir Iceland

- sympatric speciation - - - - - rapid -

varying degrees of convergence =

re. taxonomy S. a. alpinus 1 tax vol = 2-3-4 ^{symp.} rapid

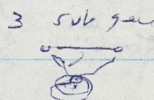
notes - mgt. - - -

alpinus - malina - saussurea not here - etc -
- - - - - 2+2=4 - insist = 5 =

tax. S. salvelinus taxonomy

II present - look ahead - albus subgenus salvelinus confusus change - malina - alpinus leucomensis → malina - alpinus

- N 7 100 - NOR... + Leary, Allendorf 40 loci.



- Unanswered questions.

- mtDNA rDNA

- electroph. profile - leucomensis (40 loci) -

* Kirotskoe - 3 ^{longhead} ^{snout} white

- profundus - Elgygytn.

human mind

deluded by

Hardy Weismann Technique

quant - qual. -

objective, quantitative vs correct, biol. reality

- Recognize limitations - plasio-eps - vs - overall phenotypic - numerical tax

- Ponnans - 1987 Ann. Rev. Ecol. Syst.

NOR: ? - miyabei

- Conclude - preserve biodiversity - Vitus vinifera Chater R - some EST

- Slides

- future - 100 - still going - still problems.

by and mouth
send M.S

Swedish postage stamp Salvelinus salvelinus

- Canadian - he Gaudin

evidence - 'Hold up in court' 3 ancient lines
Polly U

- G fanatic - unreason

- 10 - 14
- 17 type 2# send by
24' text - 2#
31

- Int Coop.

- farms L. Elgy ^{glacial} - headlight
L. Krenatskae Siberian mts

510, wDNA
- but hDNA
leucumorus
- malms - M.S.

profundus - 21pin 1#

Arctic Ch. ^{Malms} - Bidouise

Sietoniden - genui? - 504 ca

- how pr 20 84 = 100 -

14vch B - 2 -

penry - 52

- Want to play the game - first learn the rules

- violation of any standard of

1. Type-topotype - Eri
- if possible

on what basis

2. specular A

- 55 - no! 27-28

character

sub gen

Salvelinus
A leucos
confund
21m-
B elpin
malms -
profundus
- 504
- 51m
- 52

Km

levelo ^{alms} ^{coml}

why symp.

esp. 21

esp. 22

Pacific

bill - 21m

Dolly Veen

- recent 50,000 yr
N-S - 100,000
m - 2 - 50,000
- 2 - 100,000
- 2 - 100,000
- 2 - 100,000
- 2 - 100,000

- C.R. map - 20-25

- stock other water

Farmland - 20-25

Drs Hattuglund school
belief system
- sympet. pr = anarchy
Mayr - Sweden - Nyman
admits = fanatics

- Systematics - Big Picture - Subgenus 120
 - Cristivomer - Bioner - Salvelinus - Chavarchner

- electroph. - bull trout - homocyst. - festinus
 slight dif.

alpinus - malus - no dif. 40 loci
 EST-1 - mlt 100
 216m 117 nodes

- Chavarchner - Acph - 1 - Transloc
 malus

Need leucosensis - stone - albus

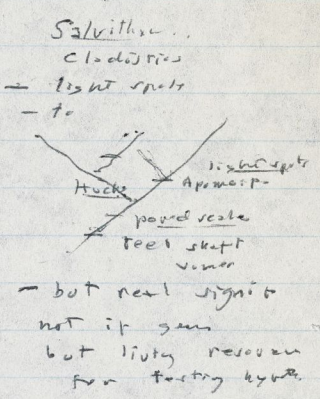
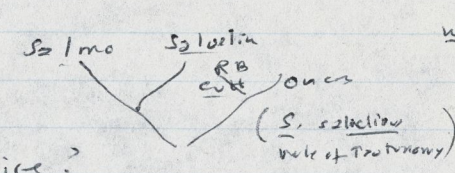
- Kronotus - white, longhead, bare char

- elgygytn - new sp.

- But understand limitations - dif.

numerical overall phenotypic sim. - between G.D. or G.E. scores & cladistic - branch.

Ferguson -



DNA hybrid. - since 1950 - noise?

mt. DNA rb DNA - some limitations - elect - more correct - support, supn

1987 Anni Rev. Ecol. Syst. Hillis

- Review evidence uneven rates
 3.5 mil. Isthmus Panama - Box di
 calibrate G.D. 1.0 = 1 - 40 mil. yr. -
 Hillis -

Karyotypes - NORs - w/ banding - great hope for tracing phylogenetic - branch. changes

- Bles - micro-macro - big picture - provincialism forest / trees
 profile - leucosensis, Kronotus L. - L. Elgygytn -
 - Bodensee profundus - ? After 21 years -

- Karyotypes -- NF 100 - all 3 subgenera present

98 - sporophyte malus alpinus

Convergence no. 84 pls. - 82-78 - 6.73 north. malus
 28 confluentis convergent? alpinus
 Stone 04

1 pr. NOR - Salmonidae - but homocyst. festinus, alpinus

NF 98 - electroph.

malus - alpinus synapomorph.

" NOR - translocation of NOR on long acrocentric 2x long

- miyabei - distinct ty. NOR

how constant character

- phyl. groups indicate

- new spec.?

post. elect. inbreeding
 genetic diversity because

Because of one in 2000 in one -
 Illumin. Technique

Frish of Conservation message - 1984 - char. watch - Conserv. Biol.
 - 10 yrs old - attend char. conf.

Yappie Speke (winnin) tight-mild)

Schedule
1/2 hr. 1/2
if fin. ok 3/4
5 min ok 1/4

happy as
- special god - Ebisu for fishermen

After so many years - Here ^{JAPAN} first began obsession with study of salmonid fishes - Sendai - Okunikawa
IWANA - YAMAME (books) - what are these beautiful trout?

This Proceedings of this symp. = 3rd major vol. on Salvelinus - Note range of interests ecology, physiology, culture, disease, conservation but - Sect. B Taxonomy & Systematics is heart of matter - 'Problems' & 'Controversies', ^{Unanswered} - That stimulate & sustain interest & enthusiasm necessary to get people from Holarctic region together for another symp. - Exp. 5. Ichthyol. --- "Arctic char fanatics" - "Fanatics" - "strong" word - zealot - emotional based belief system immune to reason - closed mind - - - Perhaps good term.

* Advice
Open mind
wacky-wacky

Perhaps best, even if possible, to answer all ~~unanswered~~ questions, solve all problems that we complete this symp. with sufficient ^{unresolved} problems for next int. symp. - - expect it live to 100 - still be char problems for future -

Ex. Chersherner - dramatic, exciting discovery new "genus" - but in perspective - all other represent. - 3 sub genera - Salvelinus (alpinus)
Cristivomer (lake char) Baione (fontinalis) - last up.
described 1814 - 175 years ago -

T.O. offer trout ^{char} - Suspect Dolly Urvan, Brook trout, lake trout

Pop. genetics
or
evol. genetics

Smith & Todd & Gilg clubs

intro/culture spec.
- neutral allele - not exposed + change

with DNA - O. - S. masou - C. J. 2. 1 mil yr for 10,000

I

Taxonomy & Systematics & To see what ^{problems} we might resolve from ^{part} 1980 (Bolon) 1981 Symp (1984 pub) - I review problems & some work to present - Helpful define tax. & syst. terms & patterns & processes ^{of evolution}
tax. = assessment of patterns of evol. ^{we observe} differentiation for classification - w/o reference to the processes causing the differentiation (Can have good tax. w/o ref. to evol. - tide tables - sun rise-set - calendars - flat earth)
Syst. = processes that produce the patterns.
- understanding phylogeny - the evol. & zoogeogr. processes - the evidence used to interpret.
- Thus If there still disagreement - more precisely focus on if a taxonomic problem (of classif.) or systematic prob. of origin - cause of differentiation -

1 Problem Ex. Sympatric sibling sp. problem - I said ~~the~~ Scandinavia, Iceland, Great Britain

1 sp - subsp. S. a. a. - i.e. one common ancestor form char established during & since last glacier - i.e. all for monophyletic to ancestor ~ 50,000 yrs. - most 10-20,000

Nyman (81) - 3 ancient sp. - alpinus, svalveticus, stagnalis - ~~before~~ 3 sp. estab. < last glacier separate invasions - my evidence

— Hindsight more accurate than foresight. but - 1980 book - "Without sound basis for establishing plesiomorphic & apomorphic character states (prim. - derived). for determining

pop. genetics
vs.
evol. genetics

should be
basis in
present.

II

phylogenetic branching sequences and sister groups, we are only playing games."

- Re. tax.-syst. sympst. sibling sp. - more precise
- the ~~Allelic~~ ^{gene} freq. at one esterase locus is not sufficient basis for plesio-spo. state and should not be used as basis for I.D. ancient phytoevol. lines -

Changing vicupoint

Ex. → 1981 Symp. (1984 book) Gydemo chair of Iceland (Thingvallavatn symp. try) -

"It is clear that all of the species from Scandinavia as defined by Nyman et al. (1981) are present in Iceland also. The landlocked chair of Iceland were already separated into species when immigration occurred."

clear statement of 3 sp. theory

— Gydemo (et al.) 1987 - Environ. Biol. Fishes
 "The Arctic chair in Thingvallavatn is polymorphic. Genetically the 4 morphs are very closely related. The morphs are conspecific (all one sp.) and do not represent different evolutionary lineages. The morphs in Thingvallavatn have probably developed within the lake and the morphs are locally adapted to different niches" - Thingv. 11,000 yrs
 (L. Malawi, 4 pop. symp. reprod. isol. - L. Malawi 300 sp. cichlids) -

* sympatric speciation

* Biol sp.

sy. st. tax.

- i.e. for mgt. - different niches - different functional roles in ecosystems - mgt. as species - tax. - NO!

— Studies influencing new acceptance rejection of 3 sp. theory:

1. Hindar et al. 1986 (Biol. J. Linn. Soc.) Norway - lakes stocked 1910, common ancestor

III

-75 yrs. - some lakes w/ dwarf & normal (reprod. isol. ?) - rec. EST allele freq. -

"All Scandinavian obsorr are derived from one recent common ancestor" - No 3 sp. involved,

Winder Jensen 1981-84

3. Klemetsen et al. 1985 (Drottningholm Rep) Bear Is. Arctic Ocean - several lakes, sympat. clon No dif. EST. but $\frac{c_{1-1.2}}{c_{10}}$ (time, convergence (introgression of 2-3) or divergence (1 common ancestor) - recent - if EST homogeneous - no dif. ratios & such.

4. Rigot et al (1986) - C.S.F.A.S. - 3 forms Greenland lake "early stages of ^{incipient} sympatric speciation".

Advanced stage

5. Walker et al. (1988) - Biol. Conserv. - Loch Rannoch Scotland. - sharp morph. divergence benthic-pelagic head, jaws. - Loch Melvin S. trutta 3 sp.

- Summary of sibling sp. problem

- S. alpinus strong tendency in lakes (esp. ^{strong selection to diverge} sole or dominant fish) - to become more efficient resource (-selection increase sp.) divide into benthic-special. & pelagic.

utiliz.

consistent, convergent trends -

benthic - pelagic

size smaller
maturation - slower growing - earlier, smaller

age - shorter lifespan

color - dark, dull, pale

rakers - shorter, fewer

Coeca - fewer

head morph - blunt, snout short jaw thick max.

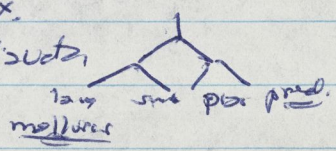
exception or Windermere

Sympatric speciation at Windermere

looks like 2-3 ancest.

sp. - but not - 10,000 yrs. -

- Thingvallavatn



IV

Why only alpinus - why only Europ subsp?
lakes - lake spawning - few other sp. - where malma - no

not leucos fontin - fontin

Present - Future: electroph. G.P. limitations

- mt DNA rbDNA - some limit. - pleistocene - spirograph

- numerical tax. - overall phenotypic sim. - additional evidence - Karyotypes - NORs - Rsa

hopeful for phylogenetic esp. bonding - tetras more about - but led me to change

Since sl - malma → alpinus - leucosensis confluens
how can 40 loci be so monomorphic? 3 subgenus albus

mol. clock
Isthmus
Panama
10-30 x diff.
1987 Ann Rev
Ecol. Syst.

how can
polymorphic be so
monomorphic?

→ Plea for broader scope - characterize

Vegetation of Hokkaido - limit counting sp.

in garden or park of limited value. - -

ex. Europe Tiefsee sibling - what is it? who is

closest relative? - electroph. - karyotypes - mt DNA nb

- N. Am. - S. - n. Dolly Varden - sympatric?

1957 Susitna - northern - count gill rakers!

→ Kamchatka - L. Kronotskoe - longhead, white, snout? - what are they??

- electroph. profile - mtDNA etc. - leucosensis link

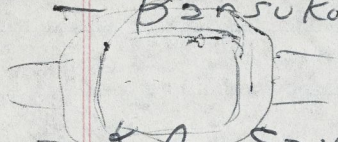
→

slides

→ Sticklebacks

- Gila chub - L. ventra
- Colo. R.

- Barsukov - get batteries for CASIO digital watch



- K.A. Savvaitova, Dept. Ichthyology, Moscow St. Univ.

Moscow 119899, USSR - copies abstracts - reprints

documentary distrib. of 2 sp. *Brachymystax*.

- Mornell et al.

* Lavrosky? - C.S.U. stuff

- Dr. Norihiko Okada

- phylogeny papers

Inst. Biol. Sciences, Univ. of Tsukuba

Tsukuba City 305, Japan.

- Herbst (pipes)

T.U. Japan - Delwa card (chann books)

offer send Trout r. brook, lake, Sunapee, Dolly Varden

- Cortland Pro Shop

And Kirchoff T.V. souvenirs

- Schollery

- Martin Bohl - after Nov. - slide of shorts - steelhead fly

- article shorts RB (written on European -

1697 Hitsu-dai. Hitomi. Honty = Shokkun

(Encyclopedia) - Recently fly fishing discovered at River

Ōi (Shizuoka Prefecture) for 2 yr - is independent

- Kawasabe origin

chann system electroph. - Tony Chernet

- Dick Wilmot, USFWS, AK Field Station, ~~1011 Foster~~

1011 E. Tudor Rd. Anchorage. 99503

Nymen 84 " Taxonomic arrangements - a
the result of human reasoning" - ^{this is only}
"approximate translations of real truth!"

84.8 " Taxonomy is not a matter of whether osteology holds
the key to sound systematics - but rather the drawing
together of conclusions from whatever methods provide convergent
results."

- re weighting ^{character} EST - , rays, karyotyp, etc -
total no. fins more weight than
no. rays in any fin -
- then abandon all reasoning = total weights = EST k
- non showing of allels at loci - was in pt. a su.
- et al (81) "new dimensions added to view of trestles
= mutation polymorphism & p.p. marker"

divine power

Before moving into interspecific problem it
would useful to clarify ^{elucidate} differences between population marks
and phylogenetic markers that ^{define = karyotyp, etc =} are used as sp. criteria.
- it is to quote ^{Dros} Nyman views.

By what divine power are we endowed with the ability
to say that all polymorphism exposed by electrophoresis
have equal resolving power as population and
species discriminating? - Cont Dros

reflective judgement
if D.V. or 100%
consistent - limitations - Inter -
- S. Zool. - methods -

① - No taxonomist would give equal taxonomic
weight to the number of rays found in a particular
fin compared to the total number of fins when
separating two species of fish.

- precisely! - if the divine power that we are endowed with
is the complex contraption of neural development called brain
particularly the cerebrum or gray matter, developed for fish etc

Mendelian polymorphism w/ corroborating evidence
not valid marker

Nyman et al. (1981) EST-1 (115) allele $\frac{1}{115}$ = alpinus .9 - 1.0 smallest
largest - 'N' svalvinius .4 - .7
'S' stagnalis .0 - .2 large

type localities -
redescription of topotype specimens

- all 3 "sp" in New Zealand ^{wh. co.} ogusarra

transmission eastern Arctic = 115 1.0 f =

if same 66-70 vent 25-30 notum

male 2: male 2: 201 = stagnalis - M

sex dim type = slab body - m - - -

ex. how morphol. test ops

observe? - ex. Australian ^{native - 4 persons} barb. ash. pers. co.

distinct morphol. - if colony

exists Sweden - -

* taxon. recognition - sympt.

key - - -

"gen. model - - -"

- EST frag. -

caution
electroph
not cladistic
but.

POP. genetics

quote Nyman 6/5 - "By what divine power do we
we to say that all polymorphisms exposed by electrophoresis
have equal resolving power as population and species
discriminants?"

allows for professional judgment = common sense. - when
seen that EST-1 is pop. marker - many individuals

and diverse subsp. alpinus, in malmo - -

If indeed, EST 100 / 215 - only -

pop. not 1:2!

Can alpinus sympatric - pop.

beginning host
"alpinus"
no. 1st year
23-25 vent
63-65 vent

p. 71 fig. 2
Remetsu - No Norway fixed in (115)
.2 in South

Fennoscandian clades

may be descended from one ancestor from east and one
from southwest -

White Sea
low water
18-21

Scm
25-30

Patterns of Chromosomal Nucleolar Organizer Region (NOR) Variation in Fishes of the Genus *Salvelinus*

RUTH B. PHILLIPS, KAY A. PLEYTE AND PETER E. IHSSSEN

The chromosomal locations of the nucleolar organizer regions (NORs) in two North American species of *Salvelinus*, *S. confluentus* and *S. malma*, are reported. This brings the total number of species of *Salvelinus* examined for chromosomal NORs to six, and the total number of species of salmonid fishes examined for NORs to 17. Arrangement of the NOR differences into a phylogenetic hypothesis supports morphological, biochemical, and cytogenetic data which group *S. malma* with *S. alpinus*. The relationship between these species and the other species of *Salvelinus* remains unresolved.

THE chromosomal locations of the nucleolar organizer regions (NORs) of three of the five North American species of *Salvelinus* were determined previously by using chromomycin A3 (CMA3) staining and silver staining (Phillips and Ihssen, 1985). The fluorochrome chromomycin A3 (CMA3) stains active and inactive NORs in amphibians and fishes (Schmid, 1982; Amemiya and Gold, 1986). Although all the species in the related genera of *Salmo* and *Oncorhynchus* have only one chromosome pair with NORs (Phillips and Ihssen, 1985; Phillips et al., 1986), the NORs were found on several different chromosome pairs in these three species of *Salvelinus*: *S. namaycush* (lake trout), *S. fontinalis* (brook trout) and *S. alpinus* (arctic char). In this paper the chromosomal location of the NORs in the other two species, *S. malma* (Dolly Varden char) and *S. confluentus* (bull trout), is described and a phylogenetic hypothesis for relationships among these species using these data is presented.

The five North American species of the genus *Salvelinus* have been assigned to three subgenera by Behnke (1965, 1980). These are the subgenus *Cristovomer* including *S. namaycush*, the subgenus *Baione* including *S. fontinalis*, and the subgenus *Salvelinus* which includes the species in the arctic char complex. This latter group includes *S. alpinus*, which has a circumpolar distribution in the arctic, *S. malma*, which occurs in sympatry with *S. alpinus* in the north Pacific, and *S. confluentus*, which is found in the Rocky Mountains (Cavender, 1980). In the Far East another species, *S. leucomaenis* ($2n = 84$) is considered to be more closely related to *S. namaycush* by Savvaitova (1980) and Viktorovsky (1978), but is placed in the subgenus *Salvelinus* by Behnke (pers. comm.). Behnke divides the subgenus *Salvelinus* into the *S. alpinus*—*S. malma* complex and a group comprising *S. confluentus*, *S. leucomaenis* and *S. albus*, which is found in the Kamchatka peninsula.

The possible phylogenetic relationships

among the North American species of *Salvelinus* were discussed in a paper by Cavender (1984), in which he summarizes his work and that of others in the cytotaxonomy of *Salvelinus*. He presented two alternative cladograms based on chromosome number ($2n$) and chromosome arm number (NF). A major point of uncertainty is whether *S. confluentus* has a sister-group relationship to *S. malma* and *S. alpinus*, or is more closely related to *S. fontinalis*, *S. namaycush* and *S. leucomaenis*. The data on NORs provide some additional evidence relevant to this problem, which remains unresolved.

MATERIALS AND METHODS

Eggs of *S. confluentus* were obtained from fish in Arrow Lake, British Columbia and eggs of *S. malma* were obtained from fish collected near Juneau, Alaska. Chromosome preparations were made from embryos from fertilized eggs using methods described previously (Phillips and Zajicek, 1982). Embryos of approx. 180° d were dissected from the eggs, incubated in culture media with 25 $\mu\text{g}/\text{ml}$ colchicine and fixed after 4 h with 3:1 methanol:acetic acid fixative. Chromosome slides were made according to the methods of Kligerman and Bloom (1977), in which fixed material was pulverized in 45% acetic acid and the material dropped onto heated slides. Slides were stained with the silver staining procedure of Howell and Black (1980) to visualize the Ag-NORs, and with chromomycin A3 followed by counterstaining with distamycin A as described in Phillips and Hartley (1988). Slides were viewed with a Zeiss Universal microscope on bright field for the Giemsa-stained and silver-stained slides and with a HB 200 mercury lamp with BG 12 excitation filter and 47 or 50 barrier filter for the CMA3 fluorescent stained slides. Six figures were examined for each individual. Photographs were made using Kodak Technical Pan film, developed with D-19 and 8 \times 10 prints were made for karyotyping.

RESULTS

Location of NORs.—The locations of the NORs in *S. confluentus* and *S. malma* were identified by silver staining and CMA3 staining. In these two species, the NORs were found at one location on one chromosome pair in all individuals examined. There was a complete correspondence between NORs as identified by CMA3 staining

and silver staining. In *S. malma*, the NORs were found at a telomeric location on the short arm of the largest submetacentric chromosome and, in *S. confluentus*, the NORs comprised the entire short arm of a large acrocentric chromosome (Fig. 1).

The locations of the NORs in five species of *Salvelinus* are illustrated in Figure 1 and summarized in Table 1. In the other three species the location of the NORs was multi-chromosomal (Phillips and Ihssen, 1985). Considerable variation among individuals in the number of chromosomes with NORs was found in all three of these species, although the location of the NORs was constant for all of the cells of a given individual. The total number of NORs per genome varied from 4–10 in *S. fontinalis*, 4–12 in *S. namaycush*, and 2–6 in *S. alpinus*.

In each species with multichromosomal NORs some of the NORs were found at the same location in almost every individual, while other NORs were quite variable between individuals (Phillips et al., 1988a, 1988b). The constant NOR sites were present on both members of the putatively homologous chromosomes, while the variable sites were often present on only one of the two homologous chromosomes. (Exact identification of homologous chromosomes would require G-banding or meiotic studies which were not done. The chromosomes were grouped according to size and centromere position.) The constant NOR site in *S. alpinus* was at the telomeres of the short arms of the largest submetacentric chromosome pair in the genome. The constant NOR sites in *S. fontinalis* were on the short arms of two acrocentric chromosome pairs. In *S. namaycush*, they were on the short arms of one acrocentric chromosome pair and the telomeres of the long arms of another acrocentric chromosome pair.

Phylogenetic analysis of NOR character states.—Differences have been found between related fish species in the haploid number of chromosomal NORs, the specific chromosome(s) on which the NORs are located, and the precise chromosomal location in the Salmoninae (Phillips and Ihssen, 1985; Phillips et al., 1986) and Cyprinidae (Gold, 1984; Gold and Amemiya, 1985).

In order to arrange NOR character states into a phylogenetic hypothesis, one must identify homology of character states between species and determine character state polarity. In order to determine character state polarity, the character states of species in related taxa must be examined so that the plesiomorphic character

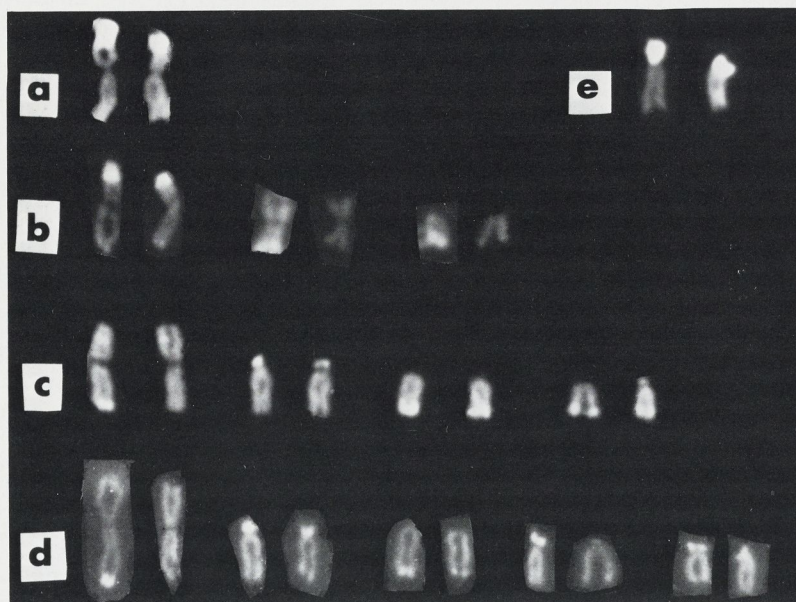


Fig. 1. Partial CMA3 stained karyotypes showing the location of NORs in the five species of *Salvelinus*. In the species with multi-chromosomal location of NORs, intraspecific differences occur in the number and location of the variable NORs which are found on only one of the two homologous chromosomes, so that the partial karyotype shown is only one of several found for that species. a) *S. malma*, b) *S. alpinus*, c) *S. namaycush*, d) *S. fontinalis*, e) *S. confluentus*.

state can be determined. The NOR phenotypes of 17 salmonid species analyzed to date are presented in Table 2. Fourteen of the 17 species have the NOR on one chromosome pair, and nine including at least one species in each genus have the NOR on the short arms of an acrocentric chromosome (Type A in Table 2). An additional four species have the NOR on the short arms of a submetacentric chromosome (Type A'). This is considered a minor change, because it is often simply the result of additional heterochromatin added to the region flanking the NOR on the short arm (Phillips and Hartley, 1988). Because NOR phenotype A is found in

each genus including one species in the related Coregoninae (Phillips, unpubl.), we have assumed that this phenotype is plesiomorphic for the Salmoninae. Thus *Coregonus clupeaformis* could be considered an outgroup for the Salmoninae.

NOR phenotype A is also found in the majority of fishes (Gold and Amemiya, 1985) and this type is also thought to be primitive for most vertebrates (Hsu et al., 1975; Schmid, 1978). If this is the case, then the initial ancestral tetraploid salmonid fish would have had two pairs of acrocentric chromosomes with the NORs on the short arms, and the NORs must have been

TABLE 1. NUMBER OF NORs PER DIPLOID GENOME IN DIFFERENT SPECIES OF *Salvelinus* FROM NORTH AMERICA. Numbers indicate the range of NORs found in different individuals of a given species.

Species	Number of specimens examined	Chromosomal location of NORs				Total NORs
		Acrocentric short arms	Acrocentric telomeres	Submetacentric telomeres	Metacentric telomeres	
<i>S. confluentus</i>	12	2	—	—	—	2
<i>S. fontinalis</i>	20	2-8*	0-2	—	0-1	4-10
<i>S. namaycush</i>	91	2-6*	2-4*	—	0-2	4-12
<i>S. alpinus</i>	60	—	0-2	2*	0-3	2-6
<i>S. malma</i>	12	—	—	2	—	2

* Indicates major NOR sites in species with NORs at more than one chromosomal location.

TABLE 2. NUMBER AND CHROMOSOMAL LOCATION OF NORs IN DIFFERENT SALMONID SPECIES.

Species	Major NORs		Minor NORs	
	Num-ber/2n	Loca-tion*	Num-ber/2n	Loca-tion
Coregoninae:				
<i>Coregonus:</i>				
<i>C. clupeaformis</i>	2	A		
Salmoninae:				
<i>Salvelinus:</i>				
<i>S. alpinus</i>	2	C'	0-6	A, B, C
<i>S. confluentus</i>	2	A		
<i>S. fontinalis</i>	2-4	A	2-6	A, B, C
<i>S. leucomaenis</i>	2	A		
<i>S. malma</i>	2	C'		
<i>S. namaycush</i>	2-4	A, B	2-8	A, B, C, C'
<i>Salmo:</i>				
<i>S. clarki</i>	2	A'		
<i>S. gairdneri</i>	2	A'		
<i>S. salar</i>	2	A'		
<i>S. trutta</i>	2	A		
<i>Oncorhynchus:</i>				
<i>O. gorbuscha</i>	2	D		
<i>O. keta</i>	2	B		
<i>O. kisutch</i>	2	A		
<i>O. masu</i>	2	A		
<i>O. nerka</i>	2	A'		
<i>O. tshawytscha</i>	2	A		

* A = acrocentric short arms, A' = submetacentric short arms, B = acrocentric telomeres, C = metacentric telomeres, C' = large submetacentric telomeres, D = metacentric adjacent to centromere.

consolidated to the short arms of one acrocentric chromosome pair or lost altogether from one pair shortly after tetraploidization. Studies of several tetraploid cyprinid fishes (Takai and Ojima, 1982) have revealed only one chromosome pair with NORs, suggesting that consolidation of NORs to one chromosome pair may be a common event in tetraploids.

If we consider NOR type A as the plesiomorphic NOR character state for the Salmoninae, then *S. confluentus* and *S. leucomaenis* (Ueda and Ojima, 1983) would have the plesiomorphic NOR character state, and the other species in the genus *Salvelinus* would be placed into two groups in which the members of at least one group share derived character states (synapomorphies).

In order to define the synapomorphies precisely, a sequential method involving G-banding for identification of chromosomes followed by

CMA3 staining or CMA3 staining of meiotic chromosomes in hybrids would be required. Because a reliable G-banding method was not available, chromosomes were divided into groups on the basis of size and centromere position.

Salvelinus fontinalis and *S. namaycush* have a very similar pattern with a constant NOR site on the short arms of the largest acrocentric chromosome pair and additional NORs on 5-12 different chromosome pairs in three different types of locations: short arms of acrocentric pairs, telomeres of acrocentric pairs, and telomeres of metacentric pairs. The only NOR in *S. confluentus* is on the largest acrocentric pair, which is similar in size to the acrocentric chromosome with the constant NOR found in *S. fontinalis* and *S. namaycush*. *Salvelinus malma* and *S. alpinus* share a synapomorphy because the only NOR chromosome pair in *S. malma*, and the constant NOR chromosome pair in the North American *S. alpinus*, is the largest submetacentric chromosome in the karyotype. In order to be completely certain of this synapomorphy, homology of the two chromosomes would have to be shown using G-banding, which has not yet been accomplished. However, the gross karyotypes of these two species are quite similar to one another (Cavender, 1984), so this is a plausible assumption. G-banding might also reveal shared derived character states between *S. fontinalis* and *S. namaycush* because several of the minor NOR sites may be identical between these two species. However, without G-banding, the minor sites must be considered autapomorphic in both species. One possible cladogram summarizing all of the chromosome data is shown in Figure 2.

DISCUSSION

Examination of the NOR character states in the species of *Salvelinus* supports the other chromosome data which indicate a major split between *S. namaycush* and *S. fontinalis* on the one hand and *S. alpinus* and *S. malma* on the other. Analysis of the other karyotype data suggests that *S. leucomaenis* from Japan should be grouped with *S. fontinalis* and *S. namaycush*, because all three species have $2n = 84$, $NF = 100$, and eight metacentric chromosome pairs which are all larger than the 34 acrocentric pairs, a karyotype considered primitive by Cavender. However, if we are correct in assuming that *S. leucomaenis* also has the primitive NOR character state, its

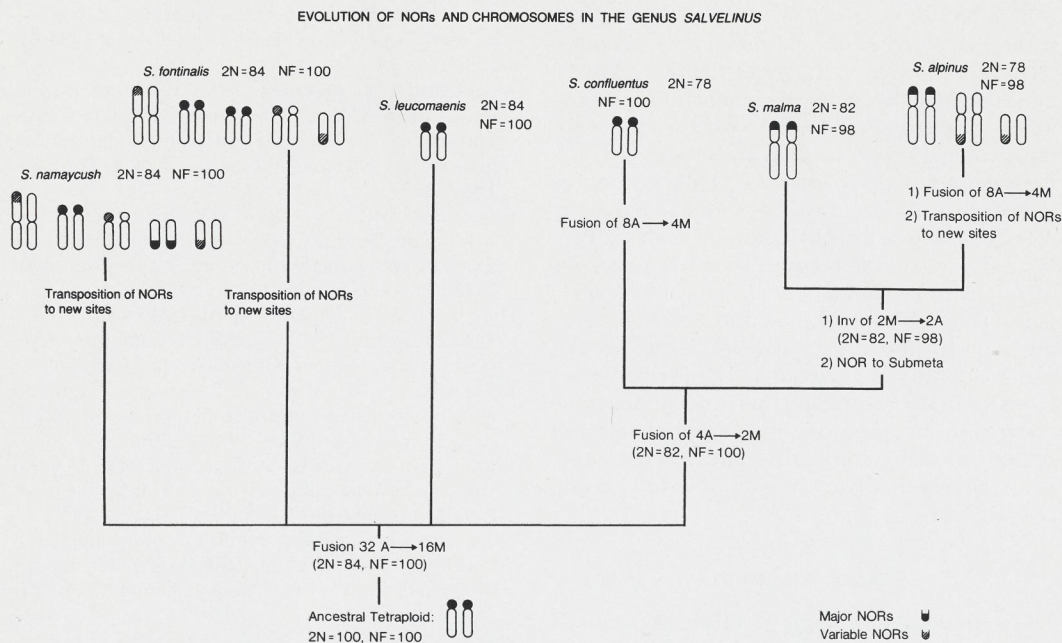


Fig. 2. One possible cladogram showing relationships between species of *Salvelinus* based on chromosome data. The locations of the major NORs which are always found on both members of a homologous pair in all members of the species are shown in black. The locations of the variable NORs which show intraspecific variation in number and chromosomal location and are usually found on only one member of a chromosome pair are shown in stippling. The chromosome data for all of the species except *S. leucomaenis* are based on karyotypes prepared from representatives of the five North American species of *Salvelinus*.

placement with respect to the other species cannot be resolved using NOR data alone. There is some question about this, because the location of NORs in the karyotype of *S. leucomaenis* was deduced from the description by Ueda and Ojima (1983) of satellites identical to those found by us on the large acrocentric chromosome pair with NORs in *S. confluentus*, and from the report of positive N-banding of these sites (Ueda, 1987). Because N-banding is usually considered equivalent to silver staining, this assumption is probably valid. The three other species have derived karyotypes, with $2n = 78$, $NF = 102$, and 12 metacentric pairs for *S. confluentus*, $2n = 78$, $NF = 98$ and 10 metacentric pairs for *S. alpinus*, and $2n = 82$, $NF = 98$ and eight metacentric pairs for the southern form of *S. malma* in North America. The northern form of *S. malma* in North America has not been karyotyped. A close relationship between *S. alpinus* and *S. malma* is suggested because they share a derived large acrocentric chromosome, which is twice as large as the others or any of the acrocentrics in the other species of *Salvelinus*.

Cavender (1984) suggested that the closest relative to *S. confluentus* may be *S. kronicus* (also called *S. albus*) from the Kamchatka River basin, which was reported to have $2n = 78$ and $NF = 100$ with 11 metacentric chromosome pairs. Our results from the karyotyping of *S. confluentus* from Arrow Lake, British Columbia, suggest that *S. confluentus* and *S. kronicus* may have identical karyotypes. Silver staining reveals that the 12th metacentric pair in the karyotype of *S. confluentus* prepared by Cavender is actually a large acrocentric chromosome pair with the NOR comprising its entire short arm.

The chromosome data support the morphological data, which suggest that there are five distinct species in North America subdivided into three groups: 1) *S. fontinalis* and *S. namaycush*; 2) *S. malma* and *S. alpinus*; and 3) *S. confluentus*. A recent electrophoretic survey of protein loci in the five species (Leary et al., pers. comm.) found that *S. malma* and *S. alpinus* shared alleles at all loci, while the other three species were distinct from each other and the *S. malma*—*S. alpinus* complex.

The NORs are the sites of the 18S and 28S ribosomal RNA genes in animal chromosomes (Hsu et al., 1975). The genes for the 18S and 28S ribosomal RNA are present in multiple copies of a repeating unit which contains both rapidly evolving spacer regions and conserved coding regions. Comparison of DNA sequences between the different species should yield many more characters for phylogenetic analysis. Preliminary restriction maps of the rRNA cistrons in *S. namaycush* and *S. fontinalis* have been prepared (Popodi et al., 1985), and we are preparing maps for the other species of *Salvelinus*. We plan to identify phylogenetically informative regions in these cistrons and then do fine structure restriction mapping or sequencing of selected regions in order to obtain multiple characters for a phylogenetic analysis of *Salvelinus* using rDNA.

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A New Arctic Char, *Salvelinus alpinus* sp., from Western Alaska

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Abstract.- A previously unknown char, *Salvelinus* sp., was discovered in certain lakes in the Kigluaik Mountains on the Seward Peninsula of western Alaska during routine inventories of fish and their habitats during summer, 1987. The fish from two of these lakes were judged to belong to the Arctic char complex, *S. alpinus*, based on morphometric and meristic features attributed to this group. Until the discovery of these populations only Dolly Varden, *S. malma*, were known from this area of western Alaska. Additional examination of physical features have revealed some apparently unique characteristics compared to other populations of the *alpinus* group in Alaska and perhaps elsewhere. Also, the populations from the two lakes are different from each other in that the sexually mature individuals from one lake have neotenic features (i.e., parr marks) and the sexually mature adults in the other lake do not exhibit neoteny. Subsequent examination of these fish in which the sequence of the ribosomal DNA first internal spacer region (rDNA ITS1) was determined, indicated that these fish were different from other Arctic char that were available for comparison which included *S. alpinus* sp. from the Kenai Peninsula in southcentral Alaska and *S. alpinus erythrinus* from Nayuk Lake in the Northwest Territories, Canada. Further genetic analysis using mitochondrial DNA restriction site fragment length polymorphisms (RFLPs) indicates that the populations from the two lakes are also different from each other. This study provides further insight into the ongoing effort by char specialists to explain the postglacial dispersal and distribution of the Arctic char complex in North America and Asia.

Morphological and meristic characteristics that are used to identify and discriminate among the different subspecies of other fish species often are unsatisfactory when trying to determine the subspecies status of different Arctic char populations. Evidence of this difficulty is demonstrated by comparing the opinions of several char specialists (Benhke, 1984, 1989; Mcphail, 1961; and Wilson et al., 1996) when discussing the subspecies status of Arctic char from the Kenai Peninsula, Alaska. Recent taxonomic studies have increasingly relied on employing techniques and methods of molecular systematics to determine the subspecies status of different Arctic char populations. Biologists with the U. S. Bureau of Land Management discovered a char, *Salvelinus* sp., inhabiting several of the glacially formed lakes of the Kigluaik Mountains on the Seward Peninsula of western Alaska in 1987. Additional char were collected from Crater Lake and Fall Creek Lake in 1992, 1995, 1996 and 1997.

Methods

Fish were captured using variable mesh monofilament gill nets and hook and line. Fish were identified to the species level using standard taxonomic keys (Scott and Crossman, 1973; Morrow, 1980). The physical characters used to determine the species to which these populations belonged were the numbers of pyloric caeca and gill rakers, the relative abundance and size of lateral spots, and whether the caudal fin was strongly or shallowly forked.

Subspecies determination was accomplished by using molecular systematic techniques in which the DNA is compared to the DNA from previously known and identified populations. Kigluaik Arctic char DNA was compared to DNA from tissues available from specimens taken from populations endemic to the following lakes: Storvatn, Norway (*S. a. alpinus*); Nauyuk Lake, Northwest Territories (*S. a. erythrinus*) and East Finger Lake, Kenai Peninsula, Alaska (*S. alpinus* sp.). In addition, the Kigluaik char were compared to Dolly Varden char (*S. malma lordi*) from the Fox River, Kenai Peninsula, Alaska.

Two genetic markers were examined in the Kigluaik char. The ITS1 sequence was selected because previous work showed that the three subspecies of Arctic char have different sequences for this region (Phillips et al., 1995). The mitochondrial RFLPs method was selected because of variation in several restriction enzyme sites among the subspecies examined previously (Grewe et al., 1990 and Wilson et al., 1996). For nuclear ribosomal analysis the DNA was extracted from three fish from Falls Creek Lake and two fish from Crater Lake using standard methods. The ITS1 of the ribosomal DNA was amplified from each fish. The products were cloned into pGEM and sequenced. The sequences were aligned and phylogenetic trees were constructed using maximum parsimony of the PAUP computer program (Swofford, 1997) and the neighbor joining method of the MEGA program (Kumar and Nei, 1993). For mitochondrial analysis the DNA was extracted from six fish from Fall Creek Lake and five fish from Crater Lake using standard methods. Approximately 50% of the mtDNA molecule was amplified using the Expand Template PCR System (Boehringer Mannheim). The products were subjected to single and double digests with four enzymes: BamH1, HindIII, Nco1 and Sma1.

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Rise R. ♀ bull x sneaker ♂ DV

upper col. chinook
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Symp 17: 149-68

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This winter capture record evidently biased the authors to assume all of the "anomalous" fish spent their winter in the sea. Considering the limited sites sampled, the recapture data showing considerable movement in the sea, and the fact that some northern Dolly Varden of Alaskan origin overwinter in Russian rivers, it should be assumed that in a non-spawning year, many southern Dolly Varden may overwinter in distant non-natal freshwater sites.

The statement on p. 14 that failure to return every winter to freshwater is a trait of coastal cutthroat is not correct. I know of no marine winter record for coastal cutthroat. They may not all return to the sampling sites but not to return to freshwater for overwintering is certainly not a "trait" of coastal cutthroat.

TRANSACTIONS of the AMERICAN FISHERIES SOCIETY

MANUSCRIPT REVIEW

Manuscript Number: T 94-162-1

Author(s): Bernard et al.

Title: Some Tests of the "Migration Hypothesis"...

Summary Recommendation:

Publish with minor revision

Publish with major revision _____

Unacceptable _____

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TRANSACTIONS

OF THE AMERICAN FISHERIES SOCIETY

GUIDE FOR REVIEWERS

The accompanying manuscript has been submitted for publication in the *Transactions of the American Fisheries Society*. We ask your assistance in judging its technical competence. Please address yourself to the following basic concerns.

- (1) Is the paper understandable, scientifically sound, and technically reliable?
- (2) Are the statistical tests, if any, appropriate for the data and correctly applied?
- (3) Are the conclusions adequately supported by the data?
- (4) Is the contribution sufficiently integrated with existing knowledge?
- (5) Does the paper represent a substantial, a marginal, or a trivial advance beyond existing knowledge? Is it relevant to its field?
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You may range as widely in your commentary as you wish, but you need not be concerned about journal style or format, which will be checked by the Editor. Additional remarks may be placed directly on the manuscript. Whenever possible, please make your criticisms constructive. It will be helpful to both author and Editor if you will point out the paper's strong points as well as its weak ones.

Please begin your review on the reverse side of this sheet, and continue on additional sheets as needed. Remember to check your summary recommendation. We would like two copies of the review, one of which will be forwarded to the author. We encourage reviewers to sign their critiques, though this is not a requirement, and urge a civil tone even for unfavorable assessments.

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Please return both copies of your review, *together with the manuscript*, to the Associate Editor who contacted you. Your assistance is greatly appreciated. *Thank you.*

Salvethymus svetovidovi gen. et sp. nova - A New Endemic
Fish ~~from~~ the Subfamily Salmoninae from Lake El'gygytgyn
(Central Chukotka) *

I. A. Chereshnev and M. B. Skopets

Institute for the Biological Problems of the North,
Far East Division, Academy of Sciences of the USSR, Magadan

A description is given of a new ~~endemic~~ genus and species of salmonid, Salvethymus svetovidovi - Svetovidov's long-finned char, ~~caught in the~~ ^{from} ancient lake El'gygytgyn. This species differs from all other salmonid genera in the coloration of the fins, in the shape and structure of the ^{chondocranium} cartilaginous skull and many of the bones of the head, in the reduction of the orbitosphenoid (to the point of complete disappearance) and of the basisphenoid, of the teeth ^{and} the vomer, and in reduction^s in the number of predorsals, ⁱⁿ of the axillary lobes and ^{process} of a number of dermal bones of the head. The new species is morphologically more similar to chars of the genus Salvelinus and it is apparently ~~from~~ phylogenetically ^{close} similar to the ancestors of that group. It is suggested that the chars should be ^{revised} ~~erected~~ as a separate tribe of the subfamily with four genera: Salvelinus Richardson, Baione Dekay, Cristivomer Gill et Jordan, Salvethymus Chereshnev et Skopetz.

ev/

The systematics, fauna and geographic range of salmonids of the subfamily Salmoninae have ~~apparently now~~ been almost completely investigated at the generic level. There is not usually any disagreement concerning the generic status of such taxa as the Pacific salmon, Oncorhynchus Suckley, the Atlantic salmon, Salmo Linnaeus, the char, Salvelinus (Nilsson) Richardson, the ~~sea-trouts~~ taimen^f, Hucho Günther, the ^{lake} genus Brachymystax Günther, and the endemic ~~salmons of the Balkans~~ ^{trout} Salmothymus Berg (Chernavin, 1923; Berg, 1948; Svetovidov, 1975; Dorofeeva et al., 1980; Viktorovskiy, 1978; Glubokovskiy, 1983; Vladykov, 1963) ~~the list is not exhaustive~~. Current views diverge only as regards the rank of some groups of salmonids that are fairly clearly segregated with respect to a number of characters. In particular, doubt is cast on the generic status of Salmothymus (Norden, 1961), but separate genera ^{have been recognized} ~~are erected~~ for the North American lake char^f,

* Originally published in Voprosy Ikhtiologii, No. 2, 1990, pp. 201-213.

4/18/90

Hi Bob,

Thanks for the fishery notes, especially the 1936 PFC reference. Enclosed is another bull trout paper by Fralay & Shepard. I reviewed it for NW Sci. about a year ago. Fralay, incidently, was one of the reviewers for our CTFAS paper.

See note about Fred's fish getting croaked - Jesus!

Any chance you'll make it up to GVP this summer? I hear our next Sci. Council meeting will be in March of 91.

Leo

Chlorine ruins UM trout lab

Weekend water treatment destroys 238 fish from 8-year genetic study

By PATRICIA SULLIVAN
of the Missoulian

Missoula's episode with tainted water has claimed the lives of more than 200 trout, virtually wiping out the basis of an 8-year-old federally funded research project at the University of Montana.

Six families of trout died of chlorine poisoning after Mountain Water Co. put the chemical into the coliform bacteria-contaminated water system Saturday afternoon. Only two of the approximately 240 trout survived.

"I was livid," said Kathy Knudsen, the research specialist who's spent eight years on the project with one of UM's leading researchers, Fred Allendorf. "I lost several years from my life Sunday afternoon."

"It's a tragic thing and it's a real mess," said Ray Murray, associate vice president for research.

ing whether or why the proper authorities at UM were not notified in time to prevent the accident.

"There was a tremendous amount of concern" for mammals that make up most of UM's research population, Murray said. They are susceptible to diseases carried by coliform bacteria, as are humans, but are little affected by chlorination of the water.

The fish are part of an evolutionary genetics project in which six families of trout were bred and cross-bred for specific genes. Male trout mature at two years and females at three years, so the loss of adult members of the basic stock

UM officials still aren't sure exactly how the fish kill occurred. Knudsen was in Utah during the weekend, so she didn't know about the water contamination and subsequent decision to chlorinate the water Saturday. A graduate student who works on the project also was out of town.

Phil Bowman, manager of UM's animal experiment laboratory, said he didn't know the water had been chlorinated until he got a phone call Sunday morning from a UM maintenance employee.

"Obviously, by that time it was too late," Bowman said. "I don't think it reflects how we care about or for the animals under our care. ... It was an obvious miscommunication with some pretty tragic results."

Murray said he's still investigat-

puts the project at least three years behind, Knudsen said.

The two surviving trout, circling in a small tank Wednesday afternoon, looked sluggish.

"I don't know if they'll make it," Knudsen said. "Chlorine damages the gill tissue, so they can't respire. They suffocate. Their tissue is still damaged. I was really amazed that anything was alive, because chlorine is really toxic."

UM's fish tanks do not have a dechlorinator, no separate water supply and no backup generator, Knudsen said.

"We need a probe in each tank to monitor the temperature, water levels ... but, dream on," she said. "It's always been in the back of my mind that we need a generator. If we lose power for more

than a couple hours, we lose the trout."

Knudsen said research findings based on the trout have been published in at least 30 articles. Some fish have died before, but nothing of this magnitude, she said.

"This was a setback, but it didn't totally trash" the last eight years of work, she said. In addition to the articles, the research has been worth "several hundred thousand dollars" of federal grant money to UM. Murray confirmed that the trout research represents "one of our biggest projects" at the university.

Knudsen has started to review records to begin figuring out how to continue the research. But she has not yet reached Allendorf, who is on a sabbatical leave to the National Science Foundation, and who still does not know about the fish kill.

Missoulian, Thursday, March 29, 1990

Egad!

Life History, Ecology and Population Status of Migratory Bull Trout (*Salvelinus confluentus*) in the Flathead Lake and River System, Montana

Abstract

Life history, ecology, and population trends of migratory bull trout (*Salvelinus confluentus*) were investigated in the Flathead Lake and River system of northwest Montana and southeast British Columbia. We conducted these studies to obtain information to manage the species in light of threats posed by timber harvest, hydropower development, and a proposed coal mine. We estimated that about half the adult bull trout in Flathead Lake embarked on a spawning migration from May through July, swimming 88-250 km to reach tributaries of the North and Middle Forks of the Flathead River. Bull trout entered the tributaries when water temperatures dropped below 12°C, and spawned from late August through early October after water temperatures were below 9°C. They spawned in areas of tributaries with low gradient, loosely compacted gravel, groundwater influence, and cover. After spawning, females left the tributaries and returned to the lake sooner than males. Most spawners were six or seven years old and they averaged 628 mm in length. Juveniles were found close to the substrate in streams with summer maximum temperatures less than 15°C. Juveniles migrated out of the tributaries to the river system from June through August, at age I (18%), II (49%), III (32%), and IV (1%). Population status was monitored through redd counts and estimates of juvenile abundance in natal tributaries. The population may be limited by quantity and quality of rearing and spawning habitat, and spawning escapement. Specific requirements for spawning and rearing habitat, and general sensitivity of each life stage, make the bull trout an excellent indicator of environmental disturbance.

Introduction

The bull trout (*Salvelinus confluentus*) is the largest species of fish native to the Flathead drainage, attaining a length of nearly one meter and a weight of 10 kg. The bull trout inhabiting the inland waters of northwestern North America is considered a separate species from the smaller, coastal Dolly Varden (*Salvelinus malma*) (Cavender 1978). The bull trout population in the Flathead system is largely migratory, growing to maturity in lakes and migrating through the river system and into the tributaries to spawn. Juveniles live in tributary streams from one to four years before migrating to the lakes.

Much information has been published concerning the life history of coastal Dolly Varden (e.g., Blackett 1968, Afmstrong and Morton 1969, Armstrong and Morrow 1980, Balon 1980). Published information on the bull trout is limited. McPhail and Murray (1979), Leggett (1969), and Allan (1980) studied various aspects of the life history of bull trout in British Columbia and Alberta. Gould (1987) described the characteristics of larval bull trout.

The Montana Department of Fish, Wildlife and Parks has studied the bull trout population in the Flathead drainage since 1953 (Block 1955, Hanzel 1976). More intensive work was undertaken from 1979-1984 during the EPA-sponsored

Flathead River Basin Studies (Graham *et al.* 1980, Fraley *et al.* 1981, Shepard *et al.* 1982, 1984b, Graham *et al.* 1982, Fraley and Graham 1982, Graham and Fredenberg 1982, Leathe and Graham 1982). We studied bull trout age and growth both in the lake and in the river system, harvest by anglers, the adult spawning migration, spawning site selection and use, and the densities, habitat selection, and emigration of juveniles growing in tributaries. Methods included tagging, gillnetting, stream trapping and electrofishing, creel survey, otolith and scale analysis, redd counts, and substrate analysis (Graham and Fredenberg 1982, Shepard and Graham 1983).

In this paper we summarize our findings on the life history, ecology, and population status of adfluvial bull trout in the Flathead Lake and inlet river system and compare our information to the results of other investigators.

Study Area

The Flathead Lake and River system is a head-water drainage of the Columbia River Basin (Figure 1). Flathead Lake is a large oligomesotrophic lake with a surface area of 476 km² and a mean depth of 32.5 m (Potter 1978). The upper 3 m of Flathead Lake is regulated by Kerr Dam, constructed on the outlet in 1938. The Flathead River enters the north end of the lake. This study

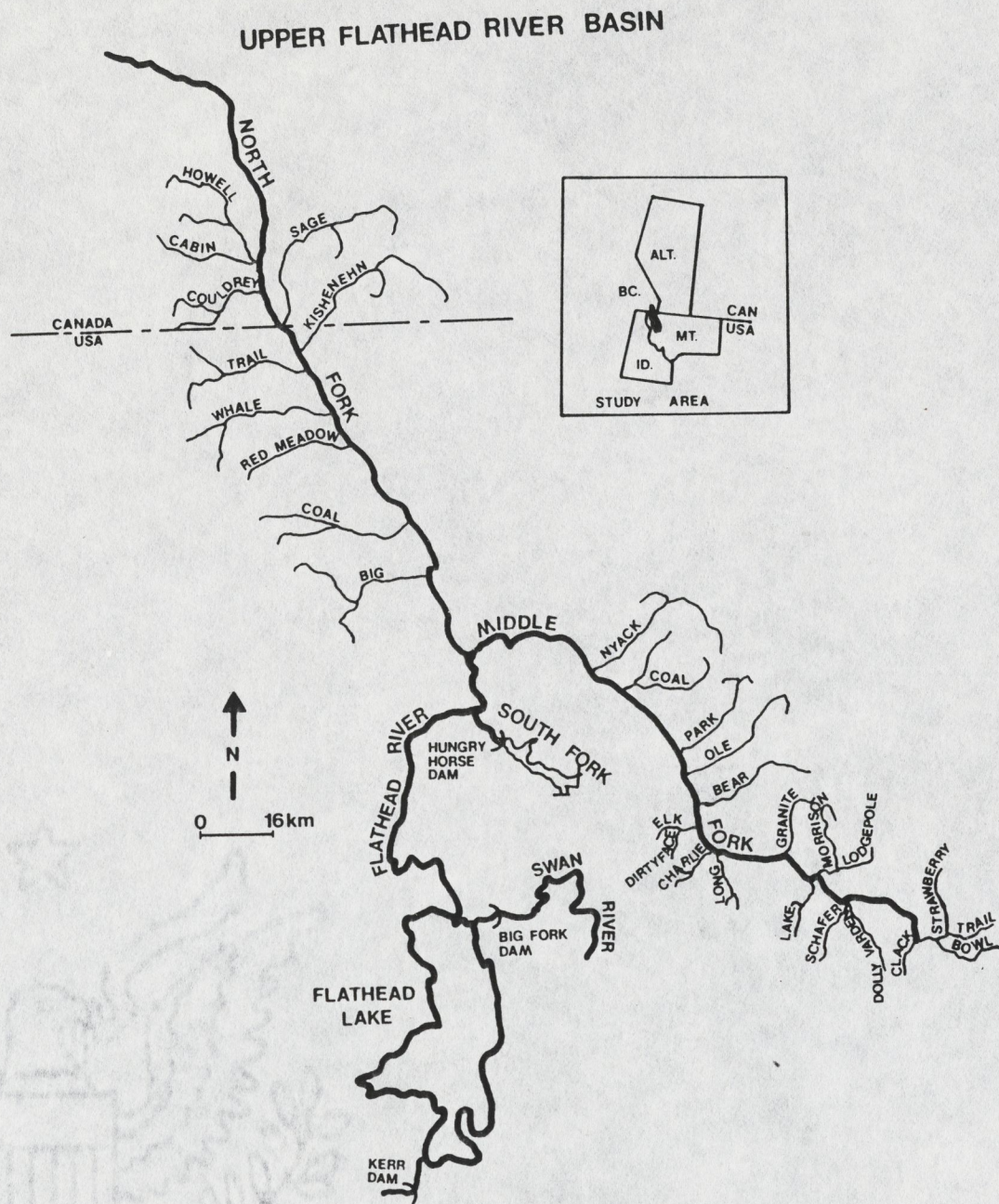


Figure 1. The upper Flathead River Basin. The 28 tributaries shown were used by spawning bull trout.

was conducted in the upper Flathead Basin which includes Flathead Lake and the river system upstream from Flathead Lake.

The South, Middle and North forks drain areas of approximately equal size in portions of the Great Bear and Bob Marshall wildernesses, Glacier National Park and the Flathead National Forest. The upper North Fork drains southern British Columbia. The South Fork is regulated by Hungry Horse Dam, located 8 km above its mouth. The Swan River enters Flathead Lake near the mouth of the Flathead River. Bull trout coexist with 23 other species of fish in the Flathead Lake and River system (Leathe and Graham 1982).

Most bull trout that spawn in the North and Middle Fork drainages mature in Flathead Lake, but fish maturing in large lakes of Glacier National Park may spawn in some tributaries. There are a few populations of bull trout in tributaries of the North Fork that spend their entire lives in the streams.

Bull trout originally used the tributaries of all forks of the Flathead and the Swan rivers. The construction of Bigfork Dam in 1902 blocked bull trout migrations into the Swan River. Limited numbers of bull trout move downstream from the Swan drainage via a marginal fish ladder, as evidenced by tag returns. Hungry Horse Dam, a 164.6-m structure which was closed in 1951, blocked all movements of bull trout into the South Fork drainage and probably resulted in a substantial reduction of the population in Flathead Lake.

The 28 tributaries used by spawning bull trout in the North and Middle Fork drainages (Figure 1) are characterized by gravel-rubble substrate, low flows of 0.057-1.70 m³/sec, and maximum summer water temperatures less than 15°C.

Results and Discussion

Life History

Lake Residence

Bull trout populations residing in Flathead Lake were found to include recently arrived juveniles from the Flathead River system, subadult fish less than about 450 mm in length, and mature fish five to six years or more in age. Most bull trout in Flathead Lake matured at age VI. A similar

age of maturity was reported for bull trout in Arrow Lakes, British Columbia (McPhail and Murray 1979).

The diet of bull trout in the lake consisted almost exclusively of fish. Whitefish species and yellow perch (*Perca flavescens*) were the most important food items, followed by kokanee (*Oncorhynchus nerka*) and nongame species (Table 1). Small bull trout have been found to feed incidentally on *Mysis* in Flathead Lake. *Mysis relicta* was discovered in Flathead Lake in 1981 and densities increased dramatically through 1986. Kokanee were the major food item for bull trout in Pend Oreille Lake, Idaho (Jeppson and Platts 1959), while whitefish were the major food in Upper Priest Lake (Bjornn 1961).

The annual growth increment for bull trout in Flathead Lake, based on analysis of scales, ranged from 60-132 mm (Table 2). Back calculations of length at annulus formation were made from 1,813 scale samples. Aging was checked with otoliths from 451 of the fish. Agreement of aging between otoliths and scales ranged from 100 percent for fish zero to three years of age, to 52 percent for older, mature fish. Growth of lake-resident fish was relatively constant after age IV. Growth rates of bull trout in Flathead Lake were similar to those reported for Priest and Upper Priest Lakes, Idaho (Bjornn 1961).

Not all mature bull trout spawned annually. Adult-size fish were relatively less abundant in the lake during the summer and fall, as compared to the spring. It appeared that 38 to 69 percent (average 57%) left the lake each spring and summer to spawn. The frequency of successive year spawning varied by age and sex (Leathe and Graham 1982). Alternate year spawning has been reported for inland Dolly Varden char (Armstrong and Morrow 1980).

Upstream Migration

Bull trout maturing in Flathead Lake began their spawning migration into the river system during April and moved slowly upstream, arriving in the North and Middle forks during late June and July. They traveled more than 250 km to spawn in some North Fork tributaries in British Columbia. The shortest distance traveled from Flathead Lake was 88 km to the mouth of Canyon Creek in the North Fork drainage. Observations and tag returns from 1974-1982 indicated that adult bull

TABLE 1. Composition by number, weight, and frequency of occurrence and calculated index of relative importance (IRI, George and Hadley 1979) for major food items in the stomachs of 95 bull trout collected between November and January, 1979, 1980 and 1981 in Flathead Lake.

Item	Number	(%)	Wet weight -g.	(%)	Index of Relative Importance (IRI)
Pygmy whitefish	5	(2.4)	37.0	(4.0)	3.2
Lake whitefish	1	(0.5)	104.1	(11.2)	4.3
Mountain whitefish	1	(0.5)	24.3	(2.6)	4.4
Unidentified whitefish	11	(5.3)	281.2	(30.3)	15.0
Total whitefish	18	(8.7)	446.6	(48.1)	23.5
Kokanee	2	(1.0)	82.8	(8.9)	4.0
Unidentified trout/salmon	2	(1.0)	13.2	(1.4)	1.5
Total trout/salmon	4	(1.9)	96.0	(10.3)	5.1
Sculpin	3	(1.5)	7.6	(0.8)	1.8
Redside shiner	5	(2.4)	15.0	(1.6)	2.0
Peamouth	1	(0.5)	3.6	(0.4)	0.7
Sucker	2	(1.0)	74.4	(8.0)	3.7
Yellow perch	83	(40.3)	105.1	(11.3)	24.6
Total nongame	94	(45.6)	205.7	(22.1)	31.0
Unidentified fish	90	(43.7)	181.1	(19.5)	41.4

TABLE 2. Back calculated lengths at annulus formation of bull trout in the upper Flathead Basin (n in parentheses). Calculations were made based on methods in Hesse (1977).

Drainage	Total length (mm) at annulus								
	I	II	III	IV	V	VI	VII	VIII	IX
<i>Adults and Juveniles</i>									
Upper Flathead (1968-81)	66 (1,813)	121 (1,538)	196 (1,161)	292 (927)	385 (669)	475 (349)	566 (129)	657 (32)	731 (4)
Flathead Lake (1968-81)	68 (931)	129 (931)	204 (928)	291 (853)	384 (603)	472 (291)	566 (102)	658 (28)	731 (4)
North Fork of the Flathead River drainage (1975-81)	73 (533)	117 (306)	165 (60)	301 (12)	440 (8)	538 (7)	574 (3)	—	—
Middle Fork of the Flathead River drainage (1980-81)	52 (349)	100 (300)	165 (172)	297 (61)	399 (57)	488 (50)	567 (24)	655 (4)	—
<i>Juveniles Only</i>									
North Fork drainage	73 (525)	117 (298)	155 (52)	228 (4)	—	—	—	—	—
Coal Creek	75 (145)	124 (62)	202 (23)	323 (14)	—	—	—	—	—
Red Meadow Creek	65 (145)	113 (113)	168 (29)	360 (7)	—	—	—	—	—
Trail Creek	74 (473)	119 (264)	158 (46)	228 (4)	—	—	—	—	—
Whale Creek	56 (52)	98 (34)	139 (6)	—	—	—	—	—	—

trout remained at the mouths of spawning tributaries for two to four weeks during which time feeding was thought to be limited.

Based on observations at stream trapping sites, adult bull trout entered tributary streams at night from July through September; the majority entered in August. Because most bull trout moved through the traps in pairs, we believe bull trout formed pairs near the mouth of the spawning tributary. Bull trout which entered the spawning tributaries were generally not in final spawning condition, but held in the tributaries for up to a month or more in deeper holes or near log or debris cover before spawning. Similar pre-spawning behavior and spawning timing was reported for bull trout in Mackenzie Creek (McPhail and Murray 1979) and John Creek (Leggett 1969) in British Columbia.

Most bull trout spawners in the North and Middle Forks were six or seven years of age (Table 3) whereas most spawners in the Swan system were five or six years old (Leathe and Enk 1985).

TABLE 3. Age of bull trout spawners in the Flathead system.

Stream	Percent by Age				
	5	6	7	8	9
North Fork Flathead River 1954 (N = 41)	24	39	34	3	0
Middle Fork Flathead River 1981 (N = 31)	10	48	35	10	0
Swan River 1983 (N = 57)	33	35	23	9	<1
Swan River 1984 (N = 76)	43	37	17	3	0

Spawning

Most bull trout spawned during September and early October in the Flathead River system, as did adfluvial bull trout in Idaho (Heimer 1965) and British Columbia (McPhail and Murray 1979). Initiation of spawning in the Flathead appeared to be related largely to water temperature, although photoperiod and streamflow probably also played a part. Spawning began when water temperatures dropped below 9-10°C. McPhail and Murray (1979) reported that 9°C was the threshold temperature for the initiation of spawning in Mackenzie Creek, British Columbia.

Bull trout spawners selected areas in the stream channel characterized by gravel substrates, low compaction and low gradient (Table 4). Groundwater influence and proximity to cover also were important factors influencing spawning site selection. These relatively specific requirements resulted in a restricted distribution of spawning in the Flathead drainage. Bull trout from Flathead Lake spawned in only 28 percent of the 750 km of available stream habitat according to basin-wide surveys from 1980-1982.

TABLE 4. Mean measurements of physical habitat variables in 34 stream reaches where no redds were located, 29 reaches where redd frequency averaged 1.2 redds/km (low), and 31 reaches where redd frequency averaged 6.9 redds/km (high).

Parameter	Redd frequency categories		
	None	Low	High
Stream order	3.0	3.1	3.6
D-90 (cm; the size of material larger than 90% of the substrate)	51	37	33
Gradient (percent)	3.2	1.8	1.5
Boulder (percent of substrate)	16	12	10
Gravel-Cobble (combined percent of substrate)	54	62	62
High quality pool (percent of stream)	5	7	8
Overhang cover (percent)	14	10	11
Total cover (percent)	16	15	13

Average length of adult spawners in the Flathead River system was 628 mm (Table 5). The female chose a spawning site and constructed the redd, while the male defended the area. Male bull trout in Trail Creek, a North Fork tributary,

TABLE 5. Average total lengths of adult bull trout spawners in the Flathead drainage.

Stream	Year	Average Length (mm)	Number of Fish
North Fork	1979	638	36
	1977	645	32
	1953	617	165
Middle Fork	1980	618	35
	1957	622	87
Both Forks	1975	628	46

remained near the redd an average of two weeks after spawning. Bull trout redds in the Flathead drainage averaged 2.0 m long x 1.0 m wide, and sometimes overlapped. Block (1955) observed one male spawn with three females in succession; the size of the redd expanded each time. McPhail and Murray (1979), Leggett (1969), and Block (1955) provided detailed descriptions of spawning behavior and spawning site activities. After spawning, the spent adults moved out of the tributaries and downstream to the lake, possibly feeding on mountain whitefish (*Prosopium williamsoni*) during the journey.

Fecundity varied with fish size, averaging 5,482 eggs per female for a sample of 32 adults averaging 645 mm. One female bull trout weighed 15 pounds and contained 12,000 eggs. Bull trout in Arrow Lakes, British Columbia, were smaller and contained fewer than 2,000 eggs per female (McPhail and Murray 1979). The sex ratio of bull trout spawners averaged 1.4 females per male in Trail Creek in the North Fork drainage, and 1.37 females per male in the Swan drainage.

Incubation and Emergence

After deposition by early October, bull trout embryos incubated in the redd for several months before hatching in January. The alevins then remained in the gravel and absorbed the yolk sac, with the first fry appearing in electrofishing samples in mid April. Emergence occurred approximately 200 days after egg deposition. Newly emerged fry averaged 23-28 mm and more than doubled their length during the first summer of growth (see Table 2).

Weaver and White (1985) found that incubation time was dependent on temperature. Bull trout eggs required 113 days (340 temperature units) to 50 percent hatch in Coal Creek, a tributary of the North Fork of the Flathead River. The fry emerged from the gravel 223 days (635 temperature units) after egg deposition. Intergravel temperatures during the incubation period (October-March) in Coal Creek ranged from 1.2-5.4°C. Survival to emergence in Coal Creek averaged 53 percent. McPhail and Murray (1979) reported the best survival of bull trout embryos at 2-4°C.

Juvenile Occurrence and Emigration

Juvenile bull trout were present in about half of the stream reaches surveyed during studies in the

upper Flathead River Basin. Juveniles were present in many reaches that were not used by adult spawners; they apparently swam upstream to these sections to grow. Distribution also was influenced by water temperature as juvenile bull trout were rarely observed in streams with summer maximum temperatures exceeding 15°C. Oliver (1979), Allan (1980) and Pratt (1984) also reported that bull trout distribution was affected by temperature.

Young-of-the-year bull trout were generally found in side channel areas and along the stream margins in Flathead tributaries. Blackett (1968) reported a similar habitat preference for juvenile Dolly Varden char in southeast Alaskan streams. McPhail and Murray (1979) found young-of-the-year bull trout in areas of low velocity near stream edges.

Densities of bull trout juveniles in Flathead tributaries were greatest in pools, and lower but generally similar in runs, riffles and pocketwater habitat. Juvenile bull trout were found closely associated with stream substrate. Pratt (1984) studied microhabitat preferences in the Flathead drainage and reported that juvenile bull trout (less than 100 mm) usually remained near the stream bottom, close to streambed materials and submerged fine debris. Juveniles 100 mm or longer also remained near cover, including larger instream debris. As the juvenile bull trout grew, they became less associated with the streambed.

During stream residence, juvenile bull trout were opportunistic feeders, mainly ingesting aquatic invertebrates (especially Diptera and Ephemeroptera) in similar percentages as they were available in the stream (Fraley *et al.* 1981). Bull trout larger than 110 mm also ate small trout and sculpin.

Snorkeling estimates of juvenile bull trout densities in Flathead drainage tributaries averaged 1.5 fish/100 m² of stream surface area (range: 0.1-7.1). Juvenile bull trout are difficult to observe because of their close association with the stream bottom, so these numbers are probably underestimates. Electrofishing estimates ranged as high as 15.5 fish/100 m² in certain streams.

Most juvenile bull trout in the Flathead drainage remained in the tributaries for one to three years before emigrating to the river system. Of 246 juvenile bull trout captured in downstream

migrant traps placed in three tributaries to the North and Middle forks, about half (49%) were age II, a third (32%) age III, and 18 percent age I (Table 6). Only 1 percent of the emigrants were age IV. The ages of emigrating juveniles were similar in Idaho and British Columbia (Bjornn 1961, Oliver 1979, McPhail and Murray 1979). The average lengths at annulus formation of Age I, II, and III juvenile bull trout in tributaries of the North Fork Flathead were 73, 117 and 155 mm, respectively (Table 2).

TABLE 6. Percent and number of age I, II, III and IV bull trout emigrating from tributary streams.

Location	Years of migration sampling	Age Classes			
		I	II	III	IV
Red Meadow Cr.	1973, 79	6 (3)	76 (42)	18 (10)	0 (0)
Trail Creek	1977, 79	34 (41)	43 (52)	19 (23)	3 (4)
Geifer Creek	1981	0 (0)	37 (26)	63 (45)	0 (0)
All Sites	(%) (number)	18 (44)	49 (120)	32 (78)	1 (4)

Emigration of juveniles from the tributaries into the Flathead River system took place largely from June through August (Table 7), similar to the emigration period reported for the Wigwam drainage, British Columbia (Oliver 1979). After juvenile bull trout entered the river system they appeared to move rapidly downstream into the main stem Flathead River, arriving below the South Fork during August and September. Although juvenile bull trout were captured by electrofishing in the main stem throughout the year, their numbers peaked during the fall months (McMullin and Graham 1981). Snorkel observations indicated that some juveniles lived along the shallow margins of the Middle and North forks. Residence in the lower Flathead River before entry into Flathead Lake has not been well documented.

Trends in Spawner Abundance

Drainage-wide counts of bull trout redds in 1980 (568), 1981 (714), 1982 (1138), and 1986 (814) were used to index the number of adfluvial bull trout

which successfully spawned in the river-tributary system. We converted the redd counts to approximate fish numbers by making the following assumptions: 1) 75 percent of all redds were located, and 2) an average of 3.2 spawners entered the tributary for each completed redd. From partial trapping results on several tributaries in 1977-1981, we estimated a spawner:red ratio of 3.2:1. In 1953, 55 bull trout entered Trail Creek and constructed 18 redds for a spawner:red ratio of 3.2:1 (Block 1955). During 1954, 160 bull trout constructed 48 redds in Trail Creek, yielding a ratio of 3.3:1. Based on these assumptions, we calculated that an average of 3,450 bull trout successfully spawned annually in the Flathead drainage during our period of record.

Bull trout spawned in 28 tributaries to the North and Middle forks (see Figure 1), but only a small percentage of the stream reaches were used for spawning. Important spawning tributaries in the North Fork were Howell, Trail, Whale, Big and Coal creeks. Major spawning tributaries in the Middle Fork were Morrison-Lodgepole, Granite, Ole, Trail and Dolly Varden creeks. The portion of the drainage in Canada supported 23-31 percent (mean 29%) of the spawning in the North Fork drainage during the 1980-82 period. Howell Creek supported 13-19 percent (mean 16%) of all North Fork spawning.

Monitoring of bull trout spawning at selected sites indicated that escapement was highest in 1982 (Table 8). These sites are considered representative of the drainage, and comprised 32, 30, 31, and 43 percent of the total drainage-wide counts in 1980, 1981, 1982, and 1986, respectively. Monitoring areas reflected drainage-wide trends.

Juvenile bull trout densities have been used as an index of population status. Juvenile bull trout populations in sections of Coal and Morrison creeks have been monitored for a six-year period (Table 9). Numbers of juvenile bull trout in these sections were highest in 1987 for both streams. Continued population estimates in these streams will provide valuable baseline information for future monitoring.

Sampling for bull trout in Flathead Lake indicated that the population had been relatively stable through 1981. Average catches of bull trout in sinking nets were 1.2 to 2.1 fish per net

TABLE 7. Number of stream trapping days, number of juvenile bull trout passed downstream through traps, and number of trapped juvenile bull trout per trap day by month from North Fork tributaries during 1976 to 1980 and Middle Fork tributaries during 1981.

	June	July	August	September	October
<i>North Fork tributaries (1976-1980)</i>					
Trap days	42	443	424	264	131
Number of fish	42	709	340	116	6
Fish/trap day	1.00	1.60	0.80	0.44	0.04
<i>Middle Fork tributaries (1981)</i>					
Trap days	43	74	62	14	—
Number of fish	60	28	19	8	—
Fish/trap day	1.40	0.38	0.26	0.57	—

TABLE 8. Bull trout redd counts for selected areas of tributaries chosen for monitoring in the Flathead Drainage.

	1979	1980	1981	1982	1983	1984	1985	1986	1987	1988
North Fork:										
Big	10	20	18	41	22	9	9	12	22	19
Coal	38	34	23	60	73	61	40	13	48	52
Whale	35	45	98	211	141	133	94	90	143	136
Trail	34 ^a	31 ^a	78	94	56	32	25	69	64	62
Total North Fork	117	130	217	406	292	235	168^b	184	277	269
Middle Fork:										
Morrison	25 ^a	75	32 ^a	86	67	38	99	52	49	50
Granite	14	34	14 ^a	34	31	47	24	37	34	32
Lodgepole	32	14	18	23	23	23	20	42	21	19
Ole		19	19	51	35	26	30	36	45	59
Total Middle Fork	71	142	83	194	156	134	173^b	167	149	160
Total Drainage Monitoring Areas	188	272	300	600	448	369	341	351	426	429

^aCounts may be underestimated due to incomplete survey.

^bHigh flows may have obliterated some of the redds.

TABLE 9. Juvenile bull trout densities in sections of a North Fork tributary (Coal Creek) and a Middle Fork tributary (Morrison Creek) from 1980-1985.

	Date	Population Estimate (Number/150 m section)	95% Confidence Interval
Coal Creek (at Deadhorse Bridge)	08/05/82	130	± 36
	03/23/83	99	± 33
	08/31/84	89	± 27
	08/26/85	167	± 66
	08/12/86	149	± 45
	09/01/87	179	± 55
Morrison Creek	09/23/80	91	± 48
	09/01/82	93	± 5
	08/18/83	62	± 11
	09/25/85	93	± 27
	08/27/86	114	± 15
	08/25/87	138	± 10
	08/30/88	126	± 23

in 1967-1970, 2.2 to 2.9 fish per net in 1980-81 (Leathe and Graham 1982). Average length of bull trout sampled in Flathead Lake increased by 24 mm from 1967 to 1980. A larger percentage of the fish were greater than 500 mm in the 1980-81 sampling period. The percentage of trophy fish (greater than 634 mm) was similar in both sampling periods. Migrating spawners, captured in the river system, were similar in size from 1953 through 1981 (see Table 5).

Sensitivity to Environmental Disturbance

All bull trout life stages are sensitive to environmental disturbances. The population in the Flathead system is threatened by several major forms of resource development. The proposed Cabin Creek coal mine in the North Fork drainage in British Columbia received preliminary approval by the Canadian government and was referred by the U.S. and Canadian governments to an International Joint Commission for review (Flathead International Study Board 1988). This mining activity could harm bull trout spawning and rearing habitat in the upper North Fork and in Howell Creek, the major spawning tributary in the Canadian portion of the drainage. The major concerns are increased sedimentation, alteration of flow and water quality degradation (Biological Resources Committee 1987). In addition, timber harvest and road construction in both the North and Middle Fork drainages are potential threats to bull trout spawning and rearing habitat.

Increased fishing pressure is often associated with resource development. Because of the restricted distribution of bull trout spawning in the basin and the limited size of the known annual escapement (3,000-4,000 individuals), harvest of fish by anglers could reduce the population. Any increase in harvest by anglers in a particular area or subbasin could result in a loss of recruitment from that site, in turn reducing the overall population in Flathead Lake.

The long overwinter incubation and development phase for bull trout embryos and alevins (223 days in Coal Creek) leaves them particularly vulnerable to increases in fine sediments and degradation of water quality. In laboratory experiments, survival was shown to be inversely

related to the percent fine material (<6.35 mm) in the gravels (Weaver and White 1985). Survival to emergence ranged from nearly 50 percent in substrates which contained 10 percent fines, to zero percent in mixtures which contained 50 percent fines. Juvenile bull trout could be affected by streambed changes because of their close association with the substrate. Shepard *et al.* (1984a) found a significant relationship ($r^2 = 0.40$, $P < .01$) between substrate score (a measure of unimbedded instream rock cover) and juvenile bull trout densities in tributaries of the Swan River.

As our studies of bull trout in the Flathead River system continue, we hope to define more precisely the factors which negatively affect the population. It is not clear whether the tributaries are at carrying capacity for juvenile bull trout, nor whether juvenile densities are limited by spawner escapement levels. The answer to these questions will require monitoring of the escapement levels and resulting juvenile densities in the tributaries over a longer period of time. McPhail and Murray (1979) suggested that limitations in juvenile rearing habitat may form an "ecological bottleneck," greatly affecting overall population levels of bull trout.

Bull trout in the Flathead River system are dependent on habitat quality and management of the interconnected river, lake, and tributaries. Cumulative losses of spawning and rearing habitat would reduce the bull trout population in Flathead Lake.

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