
CSIRO PUBLISHING

Marine & Freshwater Research

Volume 51, 2000

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A journal for the publication of original contributions
in physical oceanography, marine chemistry,
marine and estuarine biology and limnology

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Marine and Freshwater Research

CSIRO PUBLISHING

PO Box 1139 (150 Oxford St)

Collingwood

Vic. 3066

Australia

Telephone: 61 3 9662 7618

Facsimile: 61 3 9662 7611

Email: ann.grant@publish.csiro.au



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the Australian Academy of Science



Species boundaries in carp gudgeons (Eleotrididae: *Hypseleotris*) from the River Murray, South Australia: evidence for multiple species and extensive hybridization.

Terry Bertozzi^A, Mark Adams^{B*} and Keith F. Walker^{A,C}

Allozyme profiles at 20 variable loci for the 55 *Hypseleotris* in the overview study.

Each individual has a binomial code including its site (1-5, Fig. 1) plus a unique number.

Individuals are grouped according to Fig. 2. Code for loci: 1=*Ada*, 2=*Ca*, 3=*Enol-1*,

4=*Enol-2*, 5=*Fdp-2*, 6=*Fum*, 7=*Got-1*, 8=*Got-2*, 9=*Gp-3*, 10=*Gpi-1*, 11=*Gpi-2*, 12=*Hbdh*,

13=*Idh-2*, 14=*Mdh-1*, 15=*Me*, 16=*Pepd-1*, 17=*Pepd-2*, 18=*Pgam* 19=*Pgm*, 20=*Tpi*.

Fish	Group	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
3.2	HA	b	ad	b	a	b	a	b	c	a	c	b	a	b	c	b	c	b	b	bc	a
3.3	HA	b	bd	b	a	b	a	b	cd	a	cd	b	a	b	c	b	c	b	b	b	a
3.12	HA	b	bd	b	a	b	a	b	c	a	cd	b	a	b	c	b	c	b	b	bc	ab
3.13	HA	b	bd	b	a	b	a	b	c	a	d	b	a	b	c	b	c	b	b	c	a
3.15	HA	b	bd	b	ab	b	a	b	c	a	c	b	a	b	c	b	c	ab	b	bc	a
2.2	HB	e	d	b	a	b	b	b	c	a	bd	d	a	b	c	b	b	d	b	c	a
2.3	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	b	d	b	c	a
2.5	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	b	d	-	c	a
2.7	HB	e	d	b	a	b	b	b	c	a	bd	d	a	b	c	b	b	d	b	bc	a
2.8	HB	e	d	b	a	b	b	b	c	a	c	d	a	b	c	b	b	d	b	c	a
2.9	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	b	d	b	bc	a
2.10	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	ab	d	b	c	a
2.11	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	b	d	b	c	a
3.8	HB	e	d	b	a	b	b	b	c	a	cd	d	a	b	ac	b	b	d	b	bc	a
4.5	HB	e	d	ab	a	b	b	b	c	a	d	d	a	b	c	ab	b	d	b	c	a
4.7	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	b	d	b	bc	a
4.8	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	b	d	b	c	a
4.9	HB	e	d	ab	a	b	b	b	c	a	cd	d	a	b	c	b	b	d	b	bc	a
4.10	HB	e	d	ab	a	b	b	b	c	a	cd	d	a	b	c	b	b	d	b	c	a
4.11	HB	e	d	ac	a	b	b	b	c	a	cd	d	a	b	c	b	b	d	b	c	a
4.13	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	b	d	b	c	a
4.14	HB	e	d	b	a	b	b	b	c	a	d	d	a	b	c	b	b	d	b	c	a
5.4	HB	e	d	ab	a	b	b	b	c	a	bd	d	a	b	c	b	b	d	b	c	a
5.8	HB	e	d	ab	a	b	b	b	c	a	d	d	a	b	c	b	b	de	b	c	a

Fish	Group	1	2	3	4	5	6	7	L	o	c	u	s	13	14	15	16	17	18	19	20
3.4	HC	dg	ab	b	a	b	a	b	bc	a	bc	c	ab	a	ac	b	b	ac	b	b	a
3.5	HC	d	ab	b	a	b	ab	b	bc	ab	b	c	ab	ab	ac	b	b	ab	b	b	a
3.6	HC	ad	be	b	a	b	b	b	ab	a	bc	c	ab	ab	bc	b	b	b	b	ab	-
3.9	HC	af	bc	ab	a	b	b	b	bc	a	b	c	ab	a	ac	b	b	ac	b	b	a
3.10	HC	d	be	b	a	b	ab	b	bc	a	b	c	b	b	c	b	b	bc	b	b	a
3.11	HC	d	e	b	ab	b	a	b	bc	a	b	c	ab	b	c	b	b	b	b	b	a
3.14	HC	ch	b	cd	-	b	ab	b	bc	a	b	c	b	b	ac	b	b	ac	b	b	a
4.1	HC	cd	ac	b	a	b	ab	b	b	a	b	c	b	ac	ac	b	b	ab	b	b	a
4.2	HC	d	ab	bd	a	b	ab	b	b	a	b	ce	ab	b	ac	b	b	ab	b	b	a
4.3	HC	ad	ab	bd	a	b	ab	b	ab	a	bc	c	ab	a	c	b	b	ac	b	b	a
4.4	HC	d	a	bd	a	b	ab	b	b	a	b	c	ab	ab	ac	b	ab	ab	b	b	a
4.12	HC	dh	be	b	a	b	a	b	c	a	bc	c	ab	a	ac	b	b	a	b	ab	a
5.9	HC	d	a	b	a	b	a	b	b	a	ac	c	ab	a	ac	b	b	ab	b	b	a
1.1	HA×HB	be	d	b	a	b	ab	b	c	a	cd	bd	a	b	c	b	bc	bd	b	bc	a
1.2	HA×HB	be	d	b	a	b	ab	b	c	a	cd	bd	a	b	c	b	bc	bd	b	bc	a
1.3	HA×HB	be	d	b	a	b	ab	b	cd	a	cd	bd	a	b	c	b	bc	bd	b	bc	a
1.4	HA×HB	be	d	b	a	b	ab	b	c	a	cd	bd	a	b	c	b	bc	ad	b	bc	a
1.5	HA×HB	be	d	b	a	b	ab	b	cd	a	cd	bd	a	b	c	b	bc	ad	b	bc	a
3.1	HA×HX	bc	a	b	a	bc	ab	ab	c	a	c	bd	a	b	c	ab	bc	bd	b	bc	a
3.7	HA×HX	bc	a	b	a	bc	ab	ab	c	a	c	bd	a	b	c	ab	b	bd	b	bc	-
2.1	HB×HX	ce	ad	b	a	bc	b	ab	c	a	cd	d	a	b	c	ab	ab	d	ab	cd	a
2.4	HB×HX	ce	ad	b	a	bc	b	ab	c	a	bc	d	a	b	c	ab	ab	d	ab	cd	a
2.6	HB×HX	ce	ad	-	a	bc	b	ab	c	a	cd	d	a	b	c	ab	ab	d	ab	cd	a
4.6	HB×HX	ce	ad	b	a	bc	b	ab	c	a	bc	d	a	b	c	ab	ab	d	ab	cd	a
5.1	HB×HX	ce	ad	b	a	bc	b	ab	c	a	cd	d	a	b	c	ab	ab	d	b	c	a
5.2	HB×HX	ce	ad	b	a	bc	b	ab	c	a	cd	d	a	b	cd	ab	ab	de	b	bc	a
5.3	HB×HX	ce	ad	b	a	bc	b	ab	c	a	c	d	a	b	c	ab	ab	d	b	c	a
5.5	HB×HX	ce	ad	b	a	bc	b	ab	c	a	cd	d	a	b	c	ab	ab	de	b	c	a
5.6	HB×HX	ce	ad	ab	a	bc	b	ab	c	a	cd	d	a	b	c	ab	ab	d	b	c	a
5.7	HB×HX	ce	ad	b	a	bc	b	ab	c	a	bc	d	a	b	c	ab	ab	d	b	c	a
5.10	HB×HX	ce	ad	b	a	bc	b	ab	c	a	cd	d	a	b	c	ab	ab	d	ab	bd	a