

Accessory publication

Allozyme profiles at the 37 variable loci for the 122 *Retropinnna* surveyed in the overview study. Individuals are identified by site code plus a unique number (eg 1-1 = fish #1 from site 1). A dash indicates no genotype was assignable and reflects either poor activity or mobility overlap with another locus in this taxon.

Fish #	OTU	<i>Acn1</i>	<i>Acn2</i>	<i>Acyce</i>	<i>Aldaa</i>	<i>Alkl</i>	<i>Argk</i>	<i>Ca</i>	<i>Enl1</i>	<i>Enl2</i>	<i>Est1</i>	<i>Est2</i>	<i>Fdp</i>	<i>Fum</i>	<i>G6pd</i>	<i>Glo</i>	<i>Got1</i>	<i>Goti</i>	<i>Gpi2</i>	<i>Gsr</i>	<i>Idh1</i>	<i>Ldh1</i>	<i>Ldh2</i>	<i>Mdh</i>	<i>Me1</i>	<i>Me2</i>	<i>Mpt</i>	<i>PepA1</i>	<i>PepA2</i>	<i>PepB</i>	<i>PepD</i>	<i>Pgkm</i>	<i>Pgk</i>	<i>Prl2</i>	<i>Tpi2</i>	<i>Ugap</i>
1-1	CEQ	c	b	f	d	b	c	d	a	c	a	c	a	b	c	b	e	b	c	d	e	b	d	c	b	b	a	d	c							
1-2	"	c	b	f	d	b	c	d	a	c	a	c	a	b	bc	eg	bf	e	b	d	c	d	e	b	b	a	d	c								
1-3	"	c	b	f	d	b	c	d	a	c	a	c	a	b	bc	ce	b	e	b	d	c	d	e	b	b	a	d	c								
1-4	"	c	b	f	d	b	c	d	a	c	a	c	a	b	bc	ce	b	e	b	d	c	d	e	b	b	a	d	c								
3-1	"	c	b	f	d	b	c	d	a	c	a	c	a	b	bd	c	d	c	cd	c	d	e	b	d	c	b	b	a	d	c						
3-2	"	c	b	f	d	b	c	d	a	c	a	c	a	b	bd	c	d	c	cd	c	d	e	b	d	c	b	b	a	d	c						
3-3	"	-	b	f	d	b	c	d	a	c	a	c	a	b	bd	c	d	c	cd	c	d	e	b	d	c	b	b	a	d	c						
4-1	"	c	b	f	d	b	c	d	a	c	a	c	a	b	bc	ag	b	e	b	d	c	d	e	b	d	c	b	b	a	d	c					
4-2	"	c	b	f	d	b	c	d	a	c	a	c	a	b	bg	b	be	b	d	c	d	e	b	d	c	b	b	a	d	b						
4-3	"	c	b	f	d	b	c	d	a	c	a	c	a	b	bd	gh	b	e	b	d	c	d	e	b	d	c	b	b	a	d	c					
2-1	SEQ	-	b	ab	f	b	c	d	a	c	a	c	a	b	bc	ab	a	d	a	b	c	d	e	b	d	c	b	b	a	d	c					
2-2	"	-	b	ab	f	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
2-3	"	-	b	ab	fg	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
2-4	"	-	b	b	f	b	c	d	a	c	a	c	a	b	bf	e	a	d	a	b	c	d	e	b	d	c	b	b	a	d	c					
4-4	"	-	b	b	df	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
4-5	"	-	b	b	f	b	c	d	a	c	a	c	a	b	bf	e	a	d	a	b	c	d	e	b	d	c	b	b	a	d	c					
4-6	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
5-1	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
5-2	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
5-3	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
5-4	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
6-1	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
6-2	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
7-1	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					
7-2	"	-	b	b	e	b	c	d	a	c	a	c	a	b	bc	cc	a	c	a	b	c	d	e	b	d	c	b	b	a	d	c					

Fish #	OTU	<i>Accon1</i>	<i>Accon2</i>	<i>Acyc</i>	<i>Ada</i>	<i>Akt1</i>	<i>Argk</i>	<i>Ca</i>	<i>Ehol1</i>	<i>Ehol2</i>	<i>Est1</i>	<i>Est2</i>	<i>Fdp</i>	<i>Fum</i>	<i>G6pd</i>	<i>Glo</i>	<i>Got1</i>	<i>Gpi1</i>	<i>Gsr</i>	<i>Idh1</i>	<i>Ldhl1</i>	<i>Ldhl2</i>	<i>Mdh</i>	<i>Me1</i>	<i>Me2</i>	<i>Mpi</i>	<i>PepA1</i>	<i>PepA2</i>	<i>PepB</i>	<i>PepD</i>	<i>Pgk</i>	<i>Pgmd</i>	<i>Pkr2</i>	<i>Tpi2</i>	<i>Ugapd</i>				
61-1	"	d	c	g	d	c	b	c	b	-	c	bd	d	i	b	c	d	c	c	d	c	a	cd	c	c	b	a	b	ab	e	b	d	b						
61-2	"	d	c	g	d	c	b	cd	d	a	e	a	c	d	d	i	b	cf	d	b	bc	c	a	c	c	c	d	b	a	e	b	d	b						
62-1	"	bd	c	g	d	c	b	c	d	a	ce	a	c	b	a	c	d	d	i	b	c	d	b	bc	c	a	c	c	d	be	a	b	b	be	b	d	ab		
62-2	"	d	c	g	d	c	b	cd	d	a	e	a	c	b	a	c	d	i	b	cd	d	i	b	c	d	b	a	c	c	bc	d	bg	a	b	b	be	b	d	b