

Descriptors of Sun Catfish, *Horabagrus brachysoma* (Teleostei: Bagridae) and Genetic Stocks

National Bureau of Fish Genetic Resources

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ACCESSION CODE.

A. SPECIES : INDIA_FINISH_HORABAGRUS_31602

B. GENETIC STOCKS

1. MEENACHIL : INDIA_FINISH_HORABAGRUS_31602_01

2. CHALAKKUDY : INDIA_FINISH_HORABAGRUS_31602_02

3. NETHRAVATI : INDIA_FINISH_HORABAGRUS_31602_03

I. GENERAL DESCRIPTION

1. Name of the Finfish Species (Scientific Name) *Horabagrus brachysoma* (Gunther)
2. Name of the Variant/ Genetic Stocks *Horabagrus brachysoma* (Gunther), Three genetic Stocks: Meenachil, Chalakkudy, Nethravathi.
3. Local Name & Language Manja Koori, yellow catfish
4. Background of the local name Bright yellow coloration over the body.
5. Close related common species/variant *Horabagrus brachysoma* (Gunther)
6. Max. Size Reported 485 mm TL/ 1.5 kg
7. Common Habitat Rivers, Freshwater Ponds, Acclimatized to Farm Conditions.
8. Native Distribution Endemic to West flowing rivers of the Southern Western Ghats
9. River basin/ Major River West flowing rivers of the Southern Western Ghats
10. Reservoir/ Any other water body Not Known Yet
11. Local region of High Abundance (if any) West flowing rivers of the Southern Western Ghats. 1. Meenachil River
2. Chalakkudy River
3. Nethravathi River
12. Collection site (Name & Lat. - Long., Altitude) 1) Meenachil River at Kumarakom, Kottayam, (09°33'N; 76°25'E);
2) Chalakkudy River at Kanakkankadavu (10°08'N; 76°07'E)

		3) Nethravathi River at Kankanadi, Mangalore (12°52'N; 74°54'E)
13.	Nearest Railway Station	Kottayam, Chalakkudy, Mangalore
14.	Specific Gear Used	Cast net
15.	Known Economic Importance	<p>a.The sun catfish as a food fish due its size and local acceptance</p> <p>b.The sun catfish can also be used as ornamental fish due to presence of attractive coloration</p> <p>c.The fish is potential aquaculture species.</p> <p>d.Captive breeding was successful through induced spawning and upto F₂ generations were produced. Captive bred individuals are being maintained at RARS, Kumarakom.</p> <p>The distinct genetic stock information indicate the need for stock specific breeding programme for stock enhancement for natural fishery and conservation.</p>
16.	Local Importance	<p>Table Fish,</p> <p>Fisher folk expect antimicrobial potency in the body mucus and use on small cuts and injuries.</p>
17.	Any specific use such as Medicinal / Local Dish & Recipe/Special occasions/Tribal	Not known
18.	Traditional knowledge (Give Details): Ref. In Local/Community/tribal mythology:	Not known
19.	Restrictions/Protection/ Conservation / under any local Community/ sentiments.	Not known

i.	Morphological and Meristic Characters	<ul style="list-style-type: none"> ★ Body is moderately elongated, compressed body with a large head and wide sub terminal mouth. ★ Teeth in villiform bands on jaws; occipital process exposed, extending to predorsal plate; eyes are large, inferior and ventro-lateral in position, visible from under side of head. ★ The dorsal fin, consisting of the rayed fin with 5-7 rays, possesses a hard spine and is serrated from the softer smaller adipose dorsal fin. Adipose fin short, commencing over the last fourth of the anal. Pectoral just reaching or not reaching pelvic; its spine stronger than that of dorsal and as long as the head, excluding snout, serrated externally and with 16-18 strong teeth internally. Ventral fin about half as long as the pectoral and reaches the anal; the base of the anal fin almost equals the length of the head. Caudal fin slightly or deeply lunate or even forked. ★ It has 4 pairs of barbels: one nasal, two mandibulars and one maxillary. Maxillary barbels extend posterior to pectoral fin base, outer mandibulars an eye diameter shorter than maxillaries; inner mandibulars, 1/4 shorter than outer mandibulars, nasals nearly 1/2 as long as head. ★ Fin formula: D I 6-7; A iii 20-25; P I 8-9; V I 5.
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ii.	Coloration	Colour: In life, greenish yellow above, the flanks brilliant golden, belly white, with a large round black mark on shoulder surrounded by a light yellow ring. Dorsal and anal fins yellowish orange stained darker at their margins. Caudal fin yellow with black base and dark edges.
iii.	Ref. Taxonomic Key	Jayaram, K .C. (1999) <i>The freshwater fishes of the Indian region</i> . Narendra Publ. House, New Delhi, pp. 551. (For yellow catfish)
iv	Source/ Reference	Three distinct genetic stocks of <i>Horabagrus brachysoma</i> (Gunther) were identified based on 25 allozyme and 8 microsatellite loci under National Agricultural Technology Project, Indian Council of Agricultural Research, New Delhi entitled " Germplasm inventory, evaluation and gene banking of freshwater fishes ". Abdul Muneer, P. M., A. Gopalakrishnan, K. K. Musammilu, Vindhya Mohindra, K. K. Lal, V. S. Basheer, W. S. Lakra (2009) Genetic variation and population structure of endemic yellow catfish, <i>Horabagrus brachysoma</i> (Bagridae) among three populations of Western Ghats region using RAPD and microsatellite markers. Molecular Biology Reports , DOI 10.1007/s11033-008-9381-6.
v	Collected by Genetic Stock Identification	Dr. A. Gopalakrishnan & V.S. Basheer, National Bureau of Fish Genetic Resources, Cochin Unit, Kochi 682 018. Dr. A. Gopalakrishnan Abdul Muneer, P. M., Vindhya Mohindra, K. K. Lal, K. K. Musammilu, V. S. Basheer, W. S. Lakra

II. DIAGNOSTIC TAXONOMIC CHARACTER (Description)

iv. Morphometric characters and measurements of sun catfish, *Horabagrus brachysoma*.

Total length (mm.)	320 (Maximum observed: 480mm)
Total body weight (g)	900 (Maximum observed: 1500g)
Standard length (mm.)	270
Head Length (mm.)	86
Lateral transverse rows	--
Lateral line scale	--
Pre-dorsal scale	--
Insertion of 1 st dorsal fin	Closer to tip of snout than to the caudal base
Barbels	4 pairs (1 nasal, 2 mandibular and 1 maxillary).
	In relation to % of standard length (SL)
Head Length	34
Pre-dorsal length	41
Snout to pelvic base	59
Snout to anal origin	70
Length of caudal peduncle	14
Adipose to caudal fin	19
Depth of caudal fin	12
Dorsal to adipose	37
Anal fin base	30
Length of dorsal fin	25
Dorsal spine length	20
Length of pectoral fin	Does not reach pelvic fin base
Length of caudal fin	16.7
	In relation to % of head length (HL)

III. Molecular Descriptors for the Genetic Stocks identified in sun catfish, *Horabagrus brachysoma*

In *Horabagrus brachysoma*, three distinct genetic stocks were identified inhabiting three river systems in Western Ghats, native distribution range of the species. Two types of molecular markers, allozyme and microsatellite DNA were used in analysis for the samples collected from wild populations. The detailed information on molecular descriptors and parameters leading to identification of three distinct genetic stocks is given below.

1. General Information

1.1 Sample collection details and sample size analysed

Sampling Location and Sample Size	Region	Western Ghats, Kerala, India		
	Rivers	Meenachil	Chalakkudy	Nethravathi
	Locality	Kumarakom	Kanakkankadavu	Kankanadi
	(Lat. & Long.)	09_33°N; 76_25° E	10_08° N; 76_07° E	12_52° N; 74_54° E
	Sample Size	70	70	70
	Total Sample Size	210		
For Both the Markers, Allozyme and Microsatellite DNA, same sample size from the above locations was analysed.				

1.2 Overview of the two type of markers used in analysis

	Allozyme,	Microsatellite
Total Sample Size	210	210
Total Loci Examined	25	8
Polymorphic Loci	13 (56%)	8
Significant Loci Over all the Three Populations. (P < 0.0001)	10 AAT-2* EST-1* EST-3* G ₃ PDH* G ₆ PDH* GLDH* LDH-2* ODH-2* SOD* XDH-1*	7 Phy01 Phy05 Cma3 Cma4-2 Cga06-1 D33-2 D38-1
Coefficient of Genetic Differentiation(Fst) Over all the Three Populations. (P < 0.0001)	0.1537	0.1055

No significant linkage disequilibrium was detected between any pair of loci in each population sample or over all the populations		

2. Molecular Descriptors:

2.1. Allele Frequencies of Thirteen Polymorphic Allozyme Loci, Private Alleles and Parameter of Genetic Variation for *Horabagrus brachysoma* from Three Rivers.

1. Allele Frequencies					
	Locus	Allele s	Meenachil	Chalakkudy	Nethravathi
<i>i.</i>	<i>AAT-2*</i>	100	0.7571	0.4500	0.6214
		117	0.2429	0.5500	0.2929
		126	-----	----	0.0857
<i>ii.</i>	<i>EST-1*</i>	083	----	-----	0.8429
		100	1.0000	1.0000	0.1571
<i>iii.</i>	<i>EST-2*</i>	100	0.7214	0.8500	0.9143
		106	0.2786	0.1500	0.0857
<i>iv.</i>	<i>EST-3*</i>	095	----	0.1857	----
		100	1.0000	0.8143	1.0000
<i>v.</i>	<i>G₃PDH*</i>	088	0.5000	0.5929	0.2571
		100	0.5000	0.4071	0.7429
<i>vi.</i>	<i>G₆PDH*</i>	086	0.7071	0.2786	0.4786
		100	0.2929	0.7214	0.5214
<i>vii</i>	<i>GLDH*</i>	080	----	0.0571	-----
		089	0.5357	0.7214	-----
		100	0.4643	0.2214	0.4071
		117	----	----	0.5929
<i>viii.</i>	<i>GPI-2*</i>	096	0.1714	0.2571	0.2143
		100	0.8286	0.7429	0.7857
<i>ix.</i>	<i>LDH-2*</i>	100	0.8286	0.6643	0.9429

		112 134	0.1714 -----	0.2857 0.0500	0.0571 -----
x.	<i>MDH*</i>	086	0.3571	0.4000	0.2357
		100	0.6429	0.6000	0.7643
xi.	<i>ODH-2*</i>	091	0.2571	0.3714	0.7857
		100	0.7429	0.6286	0.2143
	<i>PGM*</i>	093	0.3143	0.2429	0.2786
		100	0.6857	0.7571	0.7214
xii.	<i>SOD*</i>	093	0.1929	0.5786	0.7286
		100	0.8071	0.4214	0.2714
xiii.	<i>XDH-1*</i>	091	0.4071	0.5571	0.3714
		100	0.5929	0.4429	0.5143
		114	----	----	0.1143

2. Private Alleles (Population Specific Alleles)

				<i>GLDH*080</i> <i>LDH-2*134</i>	<i>EST-1*083</i> <i>AAT-2*126</i> <i>GLDH*117</i> <i>XDH-1*114</i>
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3. Parameters of Genetic Variation

i.	H obs		0.1724	0.1908	0.1704
ii.	H exp		0.3465	0.3969	0.3475
iii.	P _(0.95)		0.48	0.52	0.52
iv.	P _(0.99)		0.48	0.52	0.52
v.	A _n		1.857	2.071	2.071

2.2. Allele Frequencies of Eight Polymorphic Microsatellite Loci, Private Alleles and Parameter of Genetic Variation for *Horabagrus brachysoma* from Three Rivers.

1. Allele Frequencies

	Locus	Allele size (bp)	Meenachil	Chalakkudy	Nethravathi
i.	<i>Phy01</i>	162	0.0643	0.1071	0.0143
		170	0.0286	0.1214	0.0357
		176	0.1500	0.1500	0.6500
		180	0.5357	0.2143	0.0643
		184	0.1500	0.1714	0.1500
		190	0.0357	0.1571	0.0857
		196	0.0357	0.0786	0.0000
ii.	<i>Phy05</i>	146	0.0286	0.1643	0.4214
		150	0.0429	0.1071	0.1286
		155	0.1857	0.3071	0.3571
		162	0.5643	0.3143	0.0786
		166	0.1143	0.1000	0.0143
		170	0.0643	0.0071	0.0000
iii.	<i>Phy07-1</i>	270	0.2071	0.1786	0.3143
		275	0.6071	0.6000	0.4214
		280	0.0786	0.1143	0.1500
		285	0.1071	0.1071	0.1143
iv.	<i>Cma3</i>	147	0.0857	0.0500	0.0786
		151	0.1071	0.0286	0.3857
		155	0.1929	0.0357	0.2643
		159	0.2143	0.3071	0.1643
		163	0.1571	0.2643	0.0214
		166	0.2286	0.2643	0.0857
		170	0.0143	0.0500	0.0000
v.	<i>Cma4-2</i>	172	0.2286	0.0286	0.0357
		175	0.4643	0.4357	0.2143
		178	0.2929	0.5214	0.6500
		182	0.0143	0.0143	0.1000
vi.	<i>Cga06-1</i>	218	0.0000	0.0071	0.2571

		226	0.0143	0.0000	0.0643
		234	0.6429	0.4000	0.5429
		240	0.2929	0.3357	0.0857
		244	0.0500	0.2571	0.0500
vii	D33-2	192	0.0000	0.0214	0.0000
		200	0.0929	0.1357	0.4071
		212	0.9071	0.8429	0.5929
viii.	D38-1	252	0.2929	0.1929	0.0786
		272	0.1071	0.2214	0.3857
		295	0.6000	0.5786	0.5357
		310	0.0000	0.0071	0.0000

2. Private Alleles (Population Specific Alleles)

				D33-2-192 D38-1-310	
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3. Parameters of Genetic Variation

i.	H obs		0.4179	0.5018	0.4964
ii.	H exp		0.5710	0.6287	0.6079
iii.	Fis		-	-	-
iv.	P _(0.95)		1.000	1.000	1.000
v	P _(0.99)		1.000	1.000	1.000
vi	A _n		4.6250	4.8750	4.3750

Abbreviations used in the table:

H obs = Observed heterozygosity

H exp = Expected heterozygosity

Fis = Inbreeding coefficient

P_{HW} = Probability value of significant deviation from HWE

Pscore = Probability value of significant heterozygosity deficiency

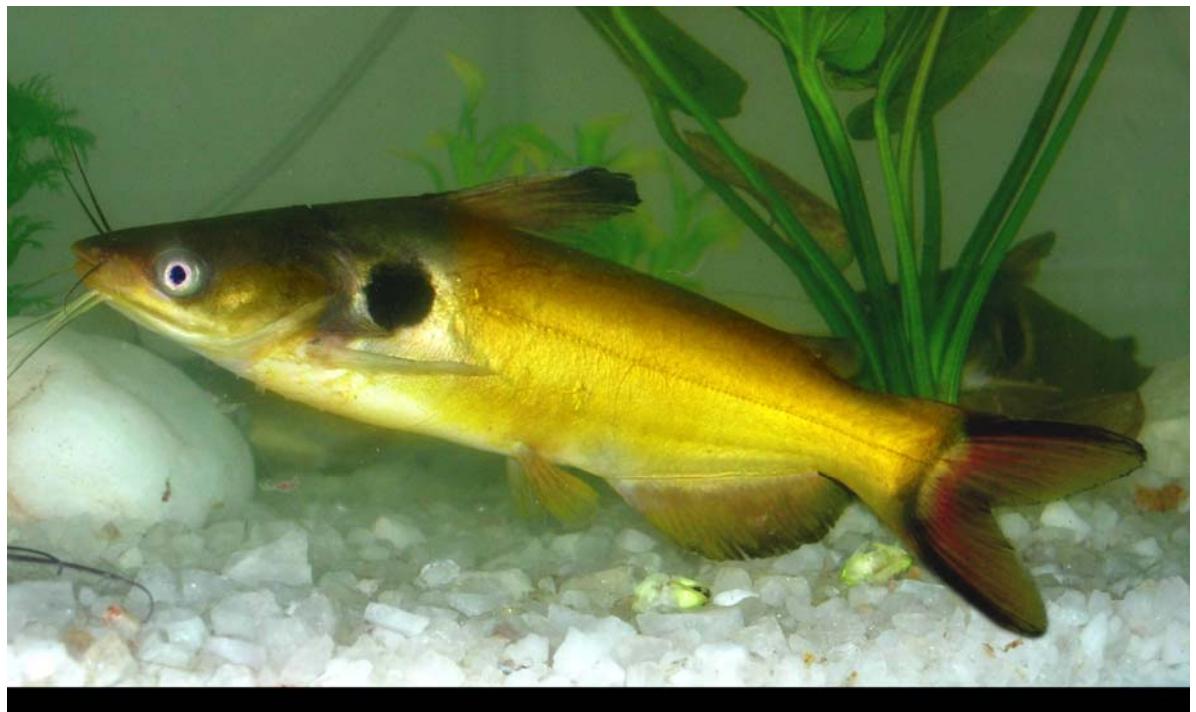
P_(0.95) = Polymorphism at 0.95 criteria

P_(0.99) = Polymorphism at 0.99 criteria

A_n = Mean number of alleles per locus

2.3. Parameters of Genetic Divergence; Allelic Heterogeneity at Allozyme and Microsatellite Loci and Coefficient of Genetic Differentiation (Fst) between Three Population Pairs

S. No.	Population Pair	Loci Exhibiting Significant Allelic Heterogeneity (P<0.0001)	Fst (P<0.0001)
Microsatellite loci			
1	Meenachil & Chalakkudy	<i>Phy01, Phy05, Cma4-2, Cga06-1</i>	0.0454
2	Meenachil & Nethravathi	<i>Phy01, Phy05, Cma3, Cma4-2, Cga06-1, D33-2</i>	0.2189
3	Chalakkudy & Nethravathi	<i>Phy01, Phy05, Cma3, Cga06-1, D33-2, D38-1</i>	0.1865
Allozyme Loci			
1	Meenachil & Chalakkudy	<i>AAT-2*, EST-3*, G₆PDH*, GLDH*, SOD*</i>	0.0952
2	Meenachil & Nethravathi	<i>EST-1*, EST-2*, GLDH*, ODH-2*, SOD*</i>	0.2640
3	Chalakkudy & Nethravathi	<i>AAT-2*, EST-1*, EST-3*, G₃PDH*, GLDH*, LDH-2*, ODH-2*</i>	0.2418



Lateral view of Sun Catfish, *Horabagrus brachysoma*