

Evolution and phylogeography analysis of diploid and polyploid *Misgurnus anguillicaudatus* populations across China

Jia Zhong, Shaokui Yi, Laiyan Ma, Weimin Wang*

College of Fisheries, Key Lab of Agricultural Animal Genetics, Breeding and Reproduction of Ministry of Education/Key Lab of Freshwater Animal Breeding, Ministry of Agriculture, Huazhong Agricultural University, Wuhan 430070, China

* Corresponding author. E-mail address: wangwm@mail.hzau.edu.cn (W. Wang).

Table S1. Material examined for *Misgurnus anguillicaudatus* arranged by geographic basin. The sampled number and its different ploidy percentage (%) are shown.

Code	Locality	Sampled number	% of triploids	% of tetraploids
Pearl River basin				
BS	Baise city, West River, Guangxi	45	-	-
CH	Conghua district, Pearl River Delta, Guangdong	45	-	-
CAZ	Chaozhou city, Hanjiang River, Guangdong	45	-	-
COZ	Chongzuo city, West River, Guangxi	45	-	-
DA	Du'an Yao Autonomous county, West River, Guangxi	45	-	-
FOS	Foshan city, Pearl River Delta, Guangdong	45	-	-
GG	Guigang city, West River, Guangxi	45	-	-
GP	Guiping city, West River, Guangxi	45	-	-
HY	Heyuan city, East River, Guangdong	41	-	-
HEZ	Hezhou city, West River, Guangxi	45	-	-
HUZ	Huizhou city, Pearl River Delta, Guangdong	45	-	-
LD	Luodian county, West River, Guizhou	48	-	-
MZ	Meizhou city, Hanjiang River, Guangdong	45	-	-
NN	Nanning city, West River, Guangxi	45	-	-
PL	Pingle county, West River, Guangxi	45	-	8.89
RJ	Rongjiang county, West River, Guizhou	45	-	-
SG	Saoguan city, North River, Guizhou	45	-	-
SJ	Dong Autonomous County of Sanjiang, West River, Guangxi	45	-	-
SHZ	Shenzhen city, Pearl River Delta, Guangdong	45	-	-

TE	Tiane county, West River, Guangxi	22	-	-
TX	Tengxian county, West River, Guangxi	45	-	-
XYI	Xingyi city, West River, Guizhou	45	-	-
XZ	Xiangzhou county, West River, Guangxi	55	-	7.27
YGD	Yingde city, North River, Guangdong	45	-	-
YZ	Yizhou district, West River, Guangxi	45	-	-
ZQ	Zhaoqing city, West River, Guangdong	45	-	-
ZS	Zhongshan city, Pearl River Delta, Guangdong	45	-	-
GL	Guilin city, West River, Guangxi	45	4.44	68.89
LP	Lipu city, West River, Guangxi	45	-	-
LZ	Liuzhou city, West River, Guangxi	45	-	-
Yangtze River basin				
SZ	Suizhou city, Han River, Hubei	41	4.88	9.76
ZX	Zhongxiang city, Han River, Hubei	43	4.65	9.30
ES	Enshi Tujia and Miao Autonomous Prefecture, Han River, Hubei	27	3.70	25.93
YD	Yidu city, Middle Yangtze River, Hubei	45	-	-
YUX	Yunyang district, Han River, Hubei	45	-	-
LHK	Laohekou city, Han River, Hubei	46	-	30.43
YM	Yunmeng county, Han River, Hubei	42	2.38	-
LIY	Liuyang city, Dongting Lake, Hunan	45	-	-
HEY	Hengyang city, Dongting Lake, Hunan	45	-	-
SY	Shaoyang city, Dongting Lake, Hunan	56	1.79	14.29
HH	Huaihua city, Dongting Lake, Hunan	60	5	8.33
ZJJ	Zhangjiajie city, Dongting Lake, Hunan	70	2.86	5.71
CHD	Changde city, Dongting Lake, Hunan	65	-	1.54
YUY	Yueyang city, Dongting Lake, Hunan	45	-	-
WN	Wuning county, Poyang Lake, Jiangxi	45	8.89	62.22
JJ	Jiujiang city, Poyang Lake, Jiangxi	45	-	44.44
XUC	Xuancheng city, Lower Yangtze River, Anhui	45	-	-
WX	Wuxi city, Lower Yangtze River, Jiangsu	45	-	-
HZ	Huzhou city, Taihu Lake, Zhejiang	45	-	-
AQ	Anqing city, Lower Yangtze River, Anhui	45	-	-
JA	Jian city, Poyang Lake, Jiangxi	45	-	-
WUH	Wuhu city, Lower Yangtze River, Anhui	43	-	-
CAH	Chaohu city, Lower Yangtze River, Anhui	46	-	-
TOL	Tongling city, Lower Yangtze River, Anhui	45	-	-
XG	Xingguo county, Poyang Lake, Jiangxi	45	-	-
JDZ	Jingdezhen city, Poyang Lake, Jiangxi	45	-	11.11
GX	Guixi city, Poyang Lake, Jiangxi	45	-	-
FZ	Fuzhou city, Poyang Lake, Jiangxi	45	-	13.33
XY	Xinyu city, Poyang Lake, Jiangxi	45	-	6.67
YX	Yongxiu county, Poyang Lake, Jiangxi	45	-	8.89
NT	Nantong city, Poyang Lake, Jiangxi	45	-	-

SH	Shanghai city, Lower Yangtze River, Shanghai	45	-	-
JX	Jinxian county, Poyang Lake, Jiangxi	45	-	-
MY	Mianyang city, Upper Yangtze River, Sichuan	45	-	-
DJY	Dujiangyan city, Upper Yangtze River, Sichuan	47	-	-
LS	Leshan city, Upper Yangtze River, Sichuan	45	-	-
NEJ	Neijiang city, Upper Yangtze River, Sichuan	45	-	-
LUZ	Luzhou city, Upper Yangtze River, Sichuan	46	-	-
CS	Chishui city, Upper Yangtze River, Guizhou	44	-	-
ZY	Zunyi city, Upper Yangtze River, Guizhou	45	-	-
NP	Nanping county, Upper Yangtze River, Chongqing	45	-	-
WL	Wulong county, Upper Yangtze River, Chongqing	43	-	-
YB	Yibin city, Upper Yangtze River, Sichuan	45	-	-
WZ	Wanzhou district, Upper Yangtze River, Chongqing	45	-	-
NC	Nanchong city, Upper Yangtze River, Sichuan	45	-	-
GY	Guangyuan city, Upper Yangtze River, Sichuan	45	-	-
DZ	Dazhou city, Upper Yangtze River, Sichuan	45	-	-
LZH	Liangzihu Lake, Middle Yangtze River, Hubei	45	-	-
HS	Huangshi city, Middle Yangtze River, Hubei	36	2.78	2.78
XZX	Xingzi county, Poyang Lake, Jiangxi	28	-	7.14
WXU	Wuxue city, Middle Yangtze River, Hubei	45	-	-
HUK	Hukou county, Poyang Lake, Jiangxi	45	-	-
XT	Xiantao city, Han River, Hubei	40	-	2.50
QJ	Qianjiang city, Han River, Hubei	45	-	-
SAY	Shayang county, Han River, Hubei	45	-	-
CD	Caidian district, Han River, Hubei	45	-	28.89
HOH	Honghu city, Han River, Hubei	45	-	-
HAC	Hanchuan city, Han River, Hubei	45	-	-
JY	Jiayu county, Middle Yangtze River, Hubei	36	-	25.00
YL	Yangluo county, Middle Yangtze River, Hubei	53	-	9.43
HP	Huangpi county, Middle Yangtze River, Hubei	41	-	7.32
Huaihe River basin				
XNY	Xinyang city, Upper Huaihe River, Henan	45	-	-
XX	Xixian county, Upper Huaihe River, Henan	42	-	-
TB	Tongbai county, Upper Huaihe River, Henan	45	-	-
LA	Luan city, Upper Huaihe River, Anhui	45	-	-
GS	Gushi county, South coast of the Huaihe river, Henan	45	-	-
GZ	Guzhen county, Middle Huaihe River, Anhui	48	-	-
WH	Wuhe county, Middle Huaihe River, Anhui	45	-	-
HB	Huaibin county, Middle Huaihe River, Henan	45	-	-
HN	Huainan city, Middle Huaihe River, Anhui	45	-	-
LB	Lingbi county, Middle Huaihe River, Anhui	45	-	-
YIS	Yishui county, Lower Huaihe River, Shandong	45	-	-
ZC	Zhucheng city, Lower Huaihe River, Shandong	45	-	-

SQ	Suqian city, Lower Huaihe River, Jiangsu	45	-	-
TC	Tancheng county, Lower Huaihe River, Shandong	45	-	-
ZZ	Zaozhuang city, Lower Huaihe River, Shandong	45	-	-
LY	Linyi city, Lower Huaihe River, Shandong	45	-	-
WF	Weifang city, Lower Huaihe River, Shandong	45	-	-
GAY	Gaoyou city, Upper Huaihe River, Jiangsu	45	-	-
HA	Huaian city, Middle Huaihe River, Anhui	45	-	-
FN	Funing county, Upper Huaihe River, Jiangsu	45	-	-
XS	Xiangshui county, Upper Huaihe River, Jiangsu	45	-	-
YS	Yingshang county, Middle Huaihe River, Anhui	45	-	-
XUY	Xuyi county, Lower Huaihe River, Jiangsu	45	-	-
XC	Xiangcheng city, Middle Huaihe River, Henan	45	-	-
Yellow River basin				
DP	Dongping county, Lower Yellow River, Shandong	45	-	-
LJ	Lijin county, Lower Yellow River, Shandong	45	-	-
HC	Hancheng city, Middle Yellow River, Shanxi	41	-	-
HD	Hongdong county, Middle Yellow River, Shanxi	45	-	-
KF	Kaifeng city, Lower Yellow River, Henan	45	-	-
LN	Luoning county, Middle Yellow River, Henan	45	-	-
QY	Qinyang city, Middle Yellow River, Henan	45	-	-
SMX	Sanmenxia city, Middle Yellow River, Henan	45	-	-
XAY	Xianyang city, Middle Yellow River, Shanxi	47	-	-
YCA	Yichuan county, Middle Yellow River, Henan	45	-	-
YIC	Yinchuan city, Upper Yellow River, Ningxia	46	-	-
Haihe River basin				
CED	Chengde city, Luanhe River, Hebei	45	-	-
SJZ	Shijiazhuang city, Ziya River, Hebei	45	-	-
DIZ	Dingzhou city, Ziya River, Hebei	45	-	-
AX	Anxin county, Daqing River, Hebei	24	16.67	-
HUR	Huairou district, Chaobai River, Beijing	45	-	-
SAH	Sanhe city, Jiyun River, Hebei	45	-	-
QX	Qianxi county, Luanhe River, Hebei	45	-	-
NH	Ninghe district, Chaobai River, Tianjin	45	-	-
NL	Nanle county, Zhangwei River, Henan	48	2.08	-
GT	Guantao county, Zhangwei River, Hebei	45	4.44	-
LC	Liaocheng city, Zhangwei River, Shandong	45	4.44	8.89
YUC	Yucheng city, Tuhai River, Shandong	45	-	-
ZH	Zhanhua district, Tuhai River, Shandong	45	-	-
QYU	Qingyun county, Tuhai River, Shandong	45	-	-
DEZ	Dezhou city, Zhangwei River, Shandong	45	-	-
HES	Hengshui city, Ziya River, Hebei	45	-	-
CZ	Cangzhou city, Daqing River, Hebei	45	-	4.44
BZ	Bazhou city, Yongding River, Hebei	45	-	-
ZUZ	Zhuozhou city, Daqing River, Hebei	45	-	-

Liaohe River basin					
PJ	Panjin city, Liaohe River, Liaoning	30	-	-	-
BX	Benxi city, Taizi River, Liaoning	45	-	-	-
FS	Fushun city, Hunhe River, Liaoning	45	-	-	-
TL	Tieling city, Liaohe River, Liaoning	22	-	-	-
ZW	Zhangwu county, Liuhe River, Liaoning	45	-	-	-
SL	Shuangliao city, West Liaohe River, Jilin	45	-	-	-
HR	Huanren county, Hunhe River, Liaoning	45	-	-	-
Songhua River basin					
JL	Jilin city, Songhua River, Jilin	45	-	-	-
SOY	Songyuan city, Songhua River, Jilin	45	-	-	-
QQ	Qiqihaer city, Nen River, Heilongjiang	45	-	-	-
NJ	Nenjiang county, Nen River, Heilongjiang	45	-	-	-
HEB	Haerbin city, Mudan River, Heilongjiang	70	-	-	-
MDJ	Mudanjiang city, Mudan River, Heilongjiang	70	-	-	-
YIL	Yilan county, Mudan River, Heilongjiang	45	-	-	-
FY	Fuyuan city, Wusuli River, Heilongjiang	47	-	-	-
FJ	Fujin city, Songhua River, Heilongjiang	22	-	-	-
MS	Mishan city, Wusuli River, Heilongjiang	45	-	-	-

Table S2. Haplotype distributions in 22 populations of *M. anguillicaudatus* based on *cyt b* and CR data sets. Bold numbers indicate shared haplotypes among populations.

Seq	Code	Locality	Haplotypes(diploid numbers/tetraploid numbers)
	LC	Hai River	H2(9/2), H10(1/1), H11(1/0), H138(0/1)
	CZ	Hai River	H1(1/0), H2(6/2), H3(1/0), H4(1/0), H5(1/0)
	LHK	Han River	H7(0/1), H44(8/4), H45(1/0), H146(0/1)
	ZX	Han River	H2(1/0), H7(1/0), H40(2/0), H44(3/3), H81(1/0), H82(2/0), H113(0/1)
<i>Cyt b</i>	CD	Han River	H39(1/2), H44(2/0), H59(1/0), H60(1/0), H61(1/0), H62(1/0), H63(1/0), H64(1/0), H65(1/0), H108(0/1), H109(0/2), H110(0/1), H111(0/1), H112(0/1), H139(0/1), H140(0/1), H141(0/1), H142(0/1), H152(0/1)
	HP	Han River,	H9(4/0), H40(0/1), H83(3/0), H84(1/0), H85(1/0), H102(1/0), H112(0/2)
	JY	Han River	H39(1/0), H40(1/5), H44(1/0), H69(1/0), H70(1/0), H71(1/0), H72(1/0), H73(1/0), H74(1/0), H75(1/0), H104(0/1), H112(0/1), H145(0/1)
	YL	Han River	H39(2/0), H40(0/1), H93(1/0), H94(1/0), H95(1/0), H96(2/0), H97(1/0), H103(1/0), H104(0/1), H105(0/1), H106(0/1), H107(0/1)
	ES	Han River	H19(3/0), H20(3/3), H21(4/4)
	SZ	Han River	H7(1/3), H86(7/1), H87(2/0)

ZJJ	Dongting Lake	H9(2/0), H40(0/2), H98(3/0) , H99(2/0), H100(1/0), H101(2/0) , H134(0/1), H135(0/1)
SY	Dongting Lake	H46(3/0), H47(1/0) , H48(1/0), H49(1/0), H50(1/0), H51(1/0), H52(1/1) , H58(1/0), H68(0/1), H100(0/1) , H125(0/4), H126(0/1)
HH	Dongting Lake	H2(1/0), H14(1/0), H47(0/1), H51(2/0), H52(1/0), H66(2/1) , H67(1/0), H68(2/2) , H119(0/1)
JJ	Poyang Lake	H38(1/0), H39(3/0), H40(1/1), H41(1/0) , H42(2/0), H43(1/0), H121(0/7), H122(0/1), H123(0/2)
WN	Poyang Lake	H76(1/3), H77(4/0), H78(1/0), H79(1/1), H80(1/1), H127(0/2), H128(0/1) , H147(0/1), H148(0/2), H149(0/1), H150(0/1)
YX	Poyang Lake	H38(2/0), H39(1/0) , H53(2/0), H54(4/0), H55(1/0), H132(0/2), H133(0/1), H151(0/1)
JDZ	Poyang Lake	H29(1/1), H30(1/0), H31(1/0), H32(1/0), H33(2/0), H34(1/0), H35(1/0), H36(1/2), H37(1/0), H120(0/2)
FZ	Poyang Lake	H22(1/0), H23(1/0), H24(2/0), H25(1/0), H26(1/0), H27(1/0), H28(1/0), H56(1/0), H57(1/0), H114(0/1), H115(0/1), H116(0/1), H117(0/1), H118(0/1)
XY	Poyang Lake	H41(3/0) , H88(2/0), H89(1/0), H90(1/0), H91(1/0), H92(2/0), H128(0/1) , H129(0/1), H130(0/1)
GL	Pearl River	H2(1/2), H4(0/2) , H6(1/0), H7(5/1) , H8(1/0), H9(1/0), H10(1/0) , H66(0/3) , H131(0/1) , H137(0/2), H143(0/1), H144(0/1)
PL	Pearl River	H4(1/0), H7(1/1), H9(4/0) , H12(1/0), H13(1/0), H14(1/0) , H15(1/0), H98(0/1) , H124(0/1), H136(0/1)
XZ	Pearl River	H9(0/1) , H16(2/0), H17(1/0), H18(7/0), H98(0/2), H131(0/1)
LC	Hai River	H1(4/1), H2(6/2)
CZ	Hai River	H1(3/0), H2(3/0) , H3(2/2), H4(1/0)
LHK	Han River	H5(0/2), H45(8/5) , H46(1/0)
ZX	Han River	H5(2/1), H45(1/2), H79(2/2) , H89(2/0), H90(2/0), H91(1/0), 93(0/1)
CD	Han River	H27(3/0), H45(2/0) , H68(1/0), H69(1/0), H70(1/0), H71(1/0), H72(1/0), H94(0/1), H95(0/2), H96(0/1), H97(0/1), H98(0/1), H99(0/1), H100(0/1)
HP	Han River	H9(10/0), H79(0/2), H103(0/1)
CR	JY	H27(2/0), H42(1/0), H45(2/0) , H76(2/0), H77(1/0), H78(1/0), H79(1/3), H101(0/4) , H102(0/1)
	YL	H27(3/0), H61(4/0) , H62(1/0), H63(1/0), H64(1/0), H79(0/2), H101(0/1), H103(0/1) , H104(0/1)
	ES	H32(2/0), H33(3/3), H34(1/0), H35(4/4)
	SZ	H5(7/1), H9(1/0), H10(0/3) , H80(1/0), H81(1/0)
	ZJJ	H9(1/0), H15(5/0) , H65(2/0), H66(1/0), H67(1/0), H79(0/2) , H133(0/1), H134(0/1)
SY	Dongting Lake	H47(1/0), H48(2/0), H49(1/0) , H50(1/0), H51(1/0), H52(1/0), H53(1/0), H54(1/1) , H55(1/0), H74(0/1) , H123(0/3), H124(0/1), H125(0/1), H126(0/1)

HH	Dongting Lake	H5(1/0), H49(0/1), H53(2/0), H54(2/1), H61(1/0), H73(1/1), H74(2/1) , H75(1/0), H120(0/1)
JJ	Poyang Lake	H18(1/0), H27(5/0), H28(1/0), H42(1/0) , H43(1/0), H44(1/0), H101(0/4) , H110(0/1), H111(0/1), H112(0/1), H113(0/1), H114(0/2)
WN	Poyang Lake	H73(1/0) , H82(1/6), H83(1/0), H84(1/2), H85(1/0), H86(1/1), H87(1/0), H88(1/0), H115(0/1)
YX	Poyang Lake	H26(1/0), H27(1/0), H28(2/0) , H29(1/0), H30(4/0), H31(1/0), H130(0/2), H131(0/1), H132(0/1)
JDZ	Poyang Lake	H36(2/1) , H37(1/0), H38(4/0), H39(1/0), H40(1/2), H41(1/0), H109(0/2)
FZ	Poyang Lake	H18(2/0) , H19(1/0), H20(1/0), H21(1/0), H22(1/0), H23(1/0), H24(1/0), H25(1/0), H36(0/1) , H92(1/0), H105(0/1), H106(0/1), H107(0/1), H108(0/1)
XY	Poyang Lake	H18(4/0) , H56(2/0), H57(1/0), H58(1/0), H59(1/0), H60(1/0), H127(0/1), H128(0/1), H129(0/1)
GL	Pearl River	H5(8/1) , H6(1/0), H7(1/0), H9(0/1), H11(0/3), H73(0/1) , H117(0/1), H118(0/2), H119(0/1)
PL	Pearl River	H5(0/1) , H8(1/0), H9(3/0), H10(1/0), H11(1/0), H12(1/0) , H13(1/0), H14(1/0), H66(1/0), H116(0/1) , H121(0/1), H122(0/1)
XZ	Pearl River	H9(0/2), H15(2/2) , H16(6/0), H17(2/0), H116(0/1)

Table S3. Matrix of pairwise population geographic distances (km).

Code	CZ	LC	GL	XZ	PL	HP	ES	LHK	ZX	CD	YL	JY	SZ	WN	JJ	JDZ	FZ	XY	YX	SY	HH	ZJJ
CZ	-																					
LC	216.2	-																				
GL	1578.8	1362.9	-																			
XZ	1730.7	1512.7	148.5	-																		
PL	1628.5	1410.3	81.8	119.6	-																	
HP	860.9	637.0	750.2	895.3	785.4	-																
ES	1135.5	912.9	566.6	704.0	640.0	472.4	-															
LHK	809.0	602.5	805.4	951.9	864.7	303.8	310.6	-														
ZX	883.9	666.1	699.9	848.9	751.0	173.4	311.8	157.1	-													
CD	896.3	677.5	705.0	850.8	741.3	46.7	436.7	296.9	151.7	-												
YL	876.4	658.6	741.0	884.6	772.5	31.3	489.0	332.6	197.1	51.5	-											
JY	964.3	746.1	642.9	787.1	677.0	109.4	430.5	341.6	185.4	68.9	96.6	-										
SZ	801.9	581.0	782.2	931.2	829.7	131.0	450.8	176.4	95.4	137.8	161.4	199.3	-									
WN	1018.5	805.7	660.1	795.7	677.6	194.1	554.5	474.6	321.7	179.5	181.0	136.9	317.1	-								
JJ	958.9	750.0	759.9	892.6	775.1	203.8	631.2	505.2	365.3	212.8	173.4	200.3	334.4	100.1	-							
JDZ	1006.0	806.6	823.8	947.9	827.3	322.5	751.5	626.8	488.6	336.3	295.0	322.5	452.4	200.8	123.9	-						
FZ	1152.3	946.9	683.6	798.6	678.1	377.4	716.6	663.7	528.4	368.7	346.5	324.5	505.3	190.1	197.9	167.0	-					
XY	1178.8	966.0	552.2	673.7	553.0	343.6	595.1	592.8	434.9	319.3	316.6	257.1	455.3	160.0	234.5	273.5	142.6	-				
YX	1036.5	827.2	698.7	826.5	707.4	248.7	627.4	541.2	390.4	243.5	217.2	209.0	377.2	73.5	78.9	135.9	130.3	159.6	-			
SY	1327.9	1109.3	258.1	404.1	302.2	491.9	407.1	569.2	449.8	447.4	483.1	387.6	528.1	420.4	520.5	603.1	488.1	345.7	469.3	-		
HH	1352.6	1137.1	283.1	401.5	332.6	561.5	307.0	556.7	472.8	515.9	561.0	467.6	563.3	533.4	632.7	727.3	626.7	484.8	591.1	149.4	-	
ZJJ	1177.4	962.7	432.2	577.8	500.0	423.9	162.5	378.0	304.2	379.9	429.5	347.9	398.9	448.9	539.8	650.1	588.7	457.2	518.5	230.3	178.6	-

Table S4. Genetic diversity indices in 5 lineages based on *cyt b* and CR datasets.

Sequence	Group	n	h	<i>Hd</i>	π
<i>Cyt b</i>	Lineage I	200	75	0.961	0.02120
	Lineage II	94	56	0.986	0.03953
	Lineage III	22	8	0.697	0.00249
	Lineage IV	14	6	0.802	0.00361
	Lineage V	8	2	0.250	0.00041
CR	Lineage I	198	68	0.956	0.01399
	Lineage II	94	50	0.973	0.01471
	Lineage III	22	5	0.407	0.00105
	Lineage IV	14	5	0.670	0.00469
	Lineage V	8	2	0.429	0.00072

n, number of samples; h, number of haplotypes; *Hd*, haplotype diversity; π , nucleotide diversity; numbers in bold indicate statistically significant results ($P < 0.05$).

Table S5. Genetic diversity indices in 22 populations including diploids and tetraploids based on *cyt b* and CR datasets.

Seq	Code	Region	Diploid			Tetraploid		
			n/h	<i>Hd</i>	π	n/h	<i>Hd</i>	π
CR	JY	Han River	10/7	0.933	0.0118	8/3	0.679	0.0022
	CD	Han River	10/7	0.911	0.0142	8/7	0.964	0.0116
	YL	Han River	10/5	0.800	0.0046	5/4	0.900	0.0054
	LHK	Han River	9/2	0.222	0.0004	7/2	0.476	0.0056
	ZX	Han River	10/6	0.911	0.0077	6/4	0.867	0.0072
	ES	Han River	10/4	0.778	0.0111	7/2	0.571	0.0077
	SZ	Han River	10/4	0.533	0.0166	4/1	0.000	0.0000
	HP	Han River	10/1	0.000	0.0000	3/2	0.667	0.0011
	HH	Dongting Lake	10/7	0.933	0.0168	5/5	1.000	0.0199
	SY	Dongting Lake	10/9	0.978	0.0130	8/6	0.893	0.0128

	ZJJ	Dongting Lake	10/5	0.756	0.0211	4/3	0.833	0.0117
	JJ	Poyang Lake	10/6	0.778	0.0192	10/6	0.844	0.0127
	WN	Poyang Lake	8/8	1.000	0.0313	10/4	0.644	0.0041
	JDZ	Poyang Lake	10/6	0.844	0.0061	5/3	0.800	0.0064
	FZ	Poyang Lake	10/9	0.978	0.0033	5/5	1.000	0.0091
	XY	Poyang Lake	10/6	0.844	0.0042	3/3	1.000	0.0056
	YX	Poyang Lake	10/6	0.844	0.0101	4/3	0.833	0.0079
	GL	Pearl River	10/3	0.378	0.0160	10/7	0.911	0.0217
	PL	Pearl River	10/8	0.933	0.0280	4/4	1.000	0.0245
	XZ	Pearl River	10/3	0.622	0.0470	5/3	0.800	0.0284
	LC	Hai River	10/2	0.533	0.0009	3/2	0.667	0.0011
	CZ	Hai River	9/4	0.806	0.0019	2/1	0.000	0.0000
<i>Cyt b</i>	JY	Han River	10/10	1.000	0.0088	8/4	0.643	0.0021
	CD	Han River	10/9	0.978	0.0281	13/11	0.974	0.0147
	YL	Han River	9/7	0.944	0.0058	5/5	1.000	0.0083
	LHK	Han River	10/3	0.378	0.0059	7/4	0.714	0.0132
	ZX	Han River	10/6	0.889	0.0153	4/2	0.500	0.0140
	ES	Han River	10/3	0.733	0.0117	7/2	0.571	0.0123
	SZ	Han River	10/3	0.511	0.0402	4/2	0.500	0.0008
	HP	Han River	10/5	0.800	0.0018	3/2	0.667	0.0011
	HH	Dongting Lake	10/7	0.933	0.0226	5/4	0.900	0.0248
	SY	Dongting Lake	10/8	0.933	0.0203	8/5	0.786	0.0123
	ZJJ	Dongting Lake	10/5	0.867	0.0491	4/3	0.833	0.0085
	JJ	Poyang Lake	9/6	0.889	0.0472	11/4	0.600	0.0406
	WN	Poyang Lake	8/5	0.786	0.0562	13/9	0.936	0.0076
	JDZ	Poyang Lake	10/9	0.978	0.0112	5/3	0.800	0.0089
	FZ	Poyang Lake	10/9	0.978	0.0063	5/5	1.000	0.0149
	XY	Poyang Lake	10/6	0.889	0.0071	3/3	1.000	0.0132
	YX	Poyang Lake	10/5	0.822	0.0270	4/3	0.833	0.0129
	GL	Pearl River	10/6	0.778	0.0413	13/8	0.923	0.0300
	PL	Pearl River	10/7	0.867	0.0644	4/4	1.000	0.0550
	XZ	Pearl River	10/3	0.511	0.0555	4/3	0.833	0.0635
	LC	Hai River	10/2	0.200	0.0003	3/2	0.667	0.0044
	CZ	Hai River	10/4	0.533	0.0013	2/1	0.000	0.0000

n, number of samples; h, number of haplotypes; Hd , haplotype diversity; π , nucleotide diversity.

Table S6. Hierarchical analyses of molecular variance (AMOVA) based on *cyt b* and CR datasets. F statistics (fixation indices): F_{ST} (among specimens among populations), F_{SC} (among populations within groups), and F_{CT} (among groups).

Source of variation	d.f.	Variance	% of	Fixation	P

		component	variation	index	
<i>Cyt b</i> group 1 (ploidy-based)					
Between ploidies	1	-0.286	-1.52	$F_{CT} = -0.015$	0.659
Among populations within ploidies	42	12.575	66.91	$F_{SC} = 0.659$	<0.001
Within populations	307	6.505	34.61	$F_{ST} = 0.654$	<0.001
<i>Cyt b</i> group 2 (region-based)					
Between regions	4	7.043	34.02	$F_{CT} = 0.340$	<0.001
Among populations within regions	17	5.346	25.82	$F_{SC} = 0.391$	<0.001
Within populations	329	8.312	40.15	$F_{ST} = 0.598$	<0.001
<i>Cyt b</i> group 3 (lineage-based)					
Between lineages	5	19.371	70.83	$F_{CT} = 0.708$	<0.001
Among populations within lineages	24	2.807	10.26	$F_{SC} = 0.352$	<0.001
Within populations	322	5.170	18.91	$F_{ST} = 0.811$	<0.001
CR group 1 (ploidy-based)					
Between ploidies	1	-0.100	-1.00	$F_{CT} = -0.010$	0.728
Among populations within ploidies	42	6.183	61.56	$F_{SC} = 0.609$	<0.001
Within populations	298	3.962	39.44	$F_{ST} = 0.606$	<0.001
CR group 2 (region-based)					
Between regions	4	2.577	23.97	$F_{CT} = 0.240$	<0.001
Among populations within regions	17	3.006	27.95	$F_{SC} = 0.368$	<0.001
Within populations	320	5.170	48.08	$F_{ST} = 0.519$	<0.001
CR group 3 (lineage-based)					
Between lineages	4	9.657	68.10	$F_{CT} = 0.681$	<0.001
Among populations within lineages	25	1.521	10.73	$F_{SC} = 0.336$	<0.001
Within populations	312	3.002	21.17	$F_{ST} = 0.788$	<0.001

Table S7. Pairwise F_{ST} (below diagonal), average Kimura 2-parameter (K2P) genetic distance values within populations (underlined along diagonal) and between pairs of populations (above diagonal) based on *cyt b* and CR data set. Bold numbers indicate statistical significance after sequential Bonferroni corrections.

Seq	Code	CZ	LC	GL	XZ	PL	HP	ES	LHK	ZX	CD	YL	JY	SZ	WN	JJ	JDZ	FZ	XY	YX	SY	HH	ZJJ
Cyt b	CZ	0.0011	0.0011	0.0226	0.1518	0.0733	0.1011	0.0257	0.0217	0.0199	0.0285	0.0243	0.0235	0.0208	0.0933	0.0505	0.1083	0.1146	0.1161	0.1037	0.0277	0.0227	0.0805
	LC	0.0035	0.0013	0.0227	0.1518	0.0734	0.1015	0.0258	0.0218	0.0199	0.0286	0.0244	0.0236	0.0209	0.0937	0.0508	0.1089	0.1151	0.1166	0.1043	0.0277	0.0227	0.0806
	GL	0.1640	0.2858	0.0381	0.1491	0.0742	0.0910	0.0374	0.0348	0.0336	0.0397	0.0365	0.0360	0.0358	0.0926	0.0579	0.1052	0.1104	0.1118	0.1018	0.0382	0.0352	0.0817
	XZ	0.3354	0.4740	0.1613	0.1058	0.1383	0.1399	0.1475	0.1473	0.1473	0.1437	0.1441	0.1454	0.1525	0.1409	0.1416	0.1344	0.1401	0.1400	0.1354	0.1434	0.1453	0.1304
	PL	0.2409	0.3851	0.0299	0.1385	0.0828	0.0747	0.0722	0.0723	0.0718	0.0724	0.0717	0.0721	0.0751	0.0961	0.0803	0.0988	0.1033	0.1038	0.0987	0.0713	0.0718	0.0788
	HP	0.2737	0.4167	0.0964	0.1730	0.0159	0.0433	0.0901	0.0925	0.0916	0.0880	0.0899	0.0896	0.0917	0.0980	0.0907	0.0938	0.0959	0.0962	0.0960	0.0888	0.0915	0.0883
	ES	0.3776	0.5067	0.2090	0.2972	0.2118	0.2383	0.0117	0.0134	0.0157	0.0194	0.0139	0.0155	0.0402	0.0835	0.0410	0.0955	0.1025	0.1028	0.0919	0.0205	0.0217	0.0750
	LHK	0.4618	0.5891	0.2660	0.3784	0.2869	0.3211	0.4154	0.0091	0.0123	0.0174	0.0120	0.0141	0.0367	0.0833	0.0412	0.0962	0.1030	0.1039	0.0923	0.0218	0.0217	0.0773
	ZX	0.2669	0.4049	0.1114	0.2198	0.1229	0.1485	0.2653	0.0515	0.0148	0.0190	0.0137	0.0144	0.0351	0.0836	0.0411	0.0968	0.1035	0.1044	0.0931	0.0220	0.0217	0.0760
	CD	0.2018	0.3220	0.0553	0.1295	0.0472	0.0650	0.1726	0.1966	0.0648	0.0228	0.0167	0.0179	0.0420	0.0825	0.0423	0.0939	0.1004	0.1009	0.0905	0.0250	0.0257	0.0756
	YL	0.2175	0.3574	0.0579	0.1374	0.0495	0.0696	0.1848	0.2660	0.0952	0.0023	0.0105	0.0116	0.0391	0.0804	0.0373	0.0914	0.0986	0.0992	0.0881	0.0201	0.0210	0.0747
	JY	0.2489	0.3776	0.0944	0.1726	0.0878	0.0812	0.2170	0.2644	0.0753	0.0444	0.0260	0.0119	0.0385	0.0808	0.0374	0.0925	0.0997	0.1003	0.0894	0.0204	0.0211	0.0740
	SZ	0.4036	0.5410	0.1658	0.3187	0.1980	0.2583	0.3604	0.4319	0.2708	0.1891	0.2033	0.2350	0.0340	0.0955	0.0611	0.1095	0.1147	0.1163	0.1060	0.0416	0.0377	0.0844
	WN	0.2335	0.3565	0.0836	0.1597	0.0765	0.1008	0.2031	0.2763	0.1288	0.0478	0.0500	0.0870	0.2203	0.0458	0.0672	0.0372	0.0380	0.0378	0.0414	0.0807	0.0834	0.0940
	JJ	0.2695	0.3938	0.1161	0.1944	0.1108	0.1288	0.2375	0.3116	0.1513	0.0619	0.0570	0.0820	0.2556	0.1090	0.0488	0.0683	0.0739	0.0743	0.0685	0.0432	0.0451	0.0805
	JDZ	0.2331	0.3698	0.0747	0.1540	0.0670	0.0921	0.2005	0.2802	0.1214	0.0379	0.0397	0.0780	0.2189	0.0670	0.1009	0.0107	0.0186	0.0185	0.0204	0.0932	0.0965	0.0961
	FZ	0.2088	0.3457	0.0516	0.1300	0.0430	0.0680	0.1769	0.2567	0.0975	0.0148	0.0157	0.0545	0.1950	0.0437	0.0776	0.0333	0.0099	0.0098	0.0226	0.0993	0.1028	0.1023
	XY	0.2414	0.3846	0.0782	0.1595	0.0706	0.0962	0.2073	0.2901	0.1261	0.0409	0.0429	0.0817	0.2265	0.0670	0.0945	0.0606	0.0365	0.0093	0.0225	0.0995	0.1034	0.1024
	YX	0.2566	0.3962	0.0947	0.1758	0.0879	0.1134	0.2225	0.3038	0.1429	0.0489	0.0508	0.0949	0.2418	0.0871	0.1054	0.0779	0.0539	0.0817	0.0273	0.0898	0.0930	0.0962
	SY	0.2319	0.3608	0.0784	0.1558	0.0710	0.0956	0.2006	0.2767	0.1242	0.0422	0.0441	0.0817	0.2183	0.0709	0.1042	0.0613	0.0379	0.0648	0.0817	0.0206	0.0213	0.0713
	HH	0.2185	0.3533	0.0603	0.1732	0.0818	0.1114	0.2193	0.2990	0.1365	0.0535	0.0589	0.0967	0.2381	0.0854	0.1195	0.0762	0.0524	0.0799	0.0971	0.0483	0.0225	0.0730

	ZJJ	0.2454	0.3851	0.0785	0.1293	0.0220	0.0502	0.2118	0.2930	0.1138	0.0472	0.0397	0.0421	0.2308	0.0765	0.0979	0.0670	0.0430	0.0706	0.0879	0.0673	0.0862	0.0685
CR	CZ	<u>0.0019</u>	0.0017	0.0135	0.1070	0.0302	0.0406	0.0170	0.0132	0.0131	0.0180	0.0158	0.0167	0.0097	0.0302	0.0198	0.0305	0.0321	0.0308	0.0300	0.0188	0.0171	0.0342
	LC	0.1143	<u>0.0009</u>	0.0133	0.1068	0.0300	0.0404	0.0168	0.0130	0.0129	0.0178	0.0156	0.0165	0.0095	0.0300	0.0196	0.0303	0.0319	0.0306	0.0297	0.0186	0.0169	0.0340
	GL	0.2140	0.3358	<u>0.0198</u>	0.1039	0.0305	0.0356	0.0218	0.0183	0.0182	0.0219	0.0199	0.0213	0.0161	0.0323	0.0236	0.0314	0.0324	0.0314	0.0310	0.0237	0.0223	0.0338
	XZ	0.2186	0.3486	0.2093	<u>0.0839</u>	0.0989	0.0980	0.1058	0.1052	0.1042	0.1058	0.1048	0.1062	0.1030	0.1037	0.1030	0.0992	0.1000	0.0996	0.0999	0.1050	0.1054	0.0918
	PL	0.1224	0.2568	0.0753	0.0966	<u>0.0341</u>	0.0308	0.0331	0.0304	0.0305	0.0319	0.0309	0.0331	0.0285	0.0392	0.0333	0.0358	0.0373	0.0363	0.0361	0.0341	0.0336	0.0329
	HP	0.4125	0.5385	0.3567	0.3296	0.1716	<u>0.0177</u>	0.0417	0.0387	0.0376	0.0378	0.0380	0.0396	0.0341	0.0425	0.0378	0.0368	0.0391	0.0386	0.0381	0.0405	0.0404	0.0355
	ES	0.2755	0.3998	0.2654	0.2726	0.1825	0.4468	<u>0.0098</u>	0.0090	0.0113	0.0135	0.0101	0.0130	0.0202	0.0300	0.0181	0.0296	0.0316	0.0296	0.0291	0.0172	0.0171	0.0323
	LHK	0.4621	0.5788	0.3880	0.4426	0.3495	0.6266	0.4880	<u>0.0029</u>	0.0056	0.0099	0.0062	0.0079	0.0160	0.0286	0.0147	0.0281	0.0296	0.0278	0.0273	0.0160	0.0151	0.0308
	ZX	0.1645	0.2939	0.0877	0.1674	0.0628	0.3162	0.2211	0.2569	<u>0.0072</u>	0.0120	0.0083	0.0090	0.0158	0.0287	0.0150	0.0281	0.0300	0.0284	0.0275	0.0174	0.0166	0.0308
	CD	0.1196	0.2465	0.1227	0.1238	0.0328	0.2928	0.1769	0.2729	0.0546	<u>0.0141</u>	0.0107	0.0136	0.0200	0.0302	0.0183	0.0294	0.0317	0.0299	0.0292	0.0190	0.0191	0.0328
	YL	0.1538	0.2854	0.1551	0.1571	0.0643	0.3172	0.2117	0.3812	0.0752	0.0312	<u>0.0073</u>	0.0097	0.0181	0.0279	0.0150	0.0275	0.0300	0.0281	0.0272	0.0161	0.0164	0.0311
	JY	0.1539	0.2801	0.1550	0.1570	0.0663	0.3034	0.2098	0.3090	0.0334	0.0356	0.0321	<u>0.0105</u>	0.0192	0.0300	0.0159	0.0296	0.0323	0.0308	0.0293	0.0195	0.0190	0.0328
	SZ	0.2826	0.4125	0.0179	0.2721	0.1243	0.4314	0.3317	0.4676	0.1339	0.1809	0.2165	0.2143	<u>0.0126</u>	0.0304	0.0215	0.0293	0.0314	0.0303	0.0296	0.0232	0.0218	0.0329
	WN	0.1886	0.3141	0.1851	0.1905	0.1000	0.3603	0.2427	0.4036	0.1404	0.0980	0.1301	0.1307	0.2480	<u>0.0228</u>	0.0255	0.0182	0.0203	0.0189	0.0203	0.0323	0.0326	0.0393
	JJ	0.1485	0.2721	0.1500	0.1518	0.0621	0.3170	0.2036	0.3595	0.1028	0.0205	0.0309	0.0203	0.2080	0.1260	<u>0.0175</u>	0.0223	0.0249	0.0237	0.0228	0.0233	0.0232	0.0336
	JDZ	0.1687	0.3000	0.1689	0.1714	0.0787	0.3489	0.2258	0.3954	0.1202	0.0774	0.1095	0.1107	0.2309	0.1441	0.1061	<u>0.0066</u>	0.0109	0.0090	0.0100	0.0334	0.0334	0.0372
	FZ	0.1096	0.2420	0.1133	0.1143	0.0212	0.2910	0.1694	0.3386	0.0634	0.0213	0.0524	0.0545	0.1734	0.0879	0.0441	0.0541	<u>0.0061</u>	0.0057	0.0099	0.0344	0.0335	0.0372
	XY	0.1523	0.2885	0.1537	0.1558	0.0611	0.3397	0.2119	0.3889	0.1035	0.0603	0.0925	0.0940	0.2169	0.1281	0.0754	0.1071	0.0085	<u>0.0052</u>	0.0088	0.0328	0.0321	0.0363
	YX	0.1506	0.2845	0.1521	0.1541	0.0604	0.3345	0.2095	0.3825	0.1025	0.0484	0.0784	0.0859	0.2143	0.1268	0.0654	0.1060	0.0485	0.0888	<u>0.0110</u>	0.0323	0.0318	0.0360
	SY	0.1196	0.2465	0.1227	0.1238	0.0328	0.2928	0.1769	0.3374	0.0742	0.0327	0.0634	0.0654	0.1809	0.0980	0.0613	0.0774	0.0213	0.0603	0.0597	<u>0.0167</u>	0.0168	0.0333
	HH	0.1440	0.2757	0.1130	0.1476	0.0502	0.3247	0.2023	0.3665	0.0851	0.0541	0.0692	0.0873	0.1756	0.1142	0.0830	0.1000	0.0429	0.0829	0.0820	0.0102	<u>0.0172</u>	0.0333
	ZJJ	0.1733	0.3066	0.1702	0.0793	0.0633	0.3035	0.2310	0.4042	0.0917	0.0810	0.0962	0.0855	0.2324	0.1482	0.1098	0.1278	0.0703	0.1110	0.1099	0.0810	0.1038	<u>0.0280</u>

Table S8. Migration estimates (number of migrants per generation) among lineages of *M. anguillicaudatus* calculated in MIGRATE using *cyt b* and CR.

Ploidy	From	To	Mean	95% credibility interval	Nm
Diploids	Lineage II	Lineage I	286.25	223.33-350	2.062
	Lineage III	Lineage I	267.31	203.33-332.67	1.926
	Lineage IV	Lineage I	263.6	200.67-326	1.899
	Lineage V	Lineage I	152.56	0-35.33	1.099
	Lineage I	Lineage II	269.45	206.67-332	3.553
	Lineage III	Lineage II	269.28	208-329.33	3.550
	Lineage IV	Lineage II	272.57	211.33-335.33	3.594
	Lineage V	Lineage II	270.49	202-338.67	3.566
	Lineage I	Lineage III	296.65	232-362	0.279
	Lineage II	Lineage III	310.49	244-377.33	0.292
	Lineage IV	Lineage III	298.83	234.67-362.67	0.281
	Lineage V	Lineage III	280.96	212-354.67	0.264
	Lineage I	Lineage IV	299.45	228.67-370	1.229
	Lineage II	Lineage IV	304.51	238.67-370	1.250
	Lineage III	Lineage IV	284.31	221.33-348	1.167
	Lineage V	Lineage IV	291.09	228.67-354	1.195
Tetraploids	Lineage I	Lineage V	281.97	217.33-348	0.422
	Lineage II	Lineage V	296.91	230-364.67	0.445
	Lineage III	Lineage V	286.82	223.33-350.67	0.430
	Lineage IV	Lineage V	295.06	228.67-360.67	0.442
	Lineage II	Lineage I	273.31	211.33-335.33	1.781
	Lineage III	Lineage I	271.45	210.23-324.33	1.768
	Lineage IV	Lineage I	283.46	219.33-348	1.847
	Lineage I	Lineage II	279.3	214.67-344	1.576
	Lineage III	Lineage II	294.4	226.67-362	1.661
	Lineage IV	Lineage II	281.81	218-346	1.590

Lineage I	Lineage III	296.81	228.67-364.67	0.979
Lineage II	Lineage III	300.86	233.33-368.67	0.993
Lineage IV	Lineage III	298.97	233.33-365.33	0.987
Lineage I	Lineage IV	296.44	230.67-362	1.934
Lineage II	Lineage IV	295.65	230.67-361.33	1.928
Lineage III	Lineage IV	298.95	232.67-366	1.950

Table S9. Migration estimates (number of migrants per generation) among regions of *M. anguillicaudatus* calculated in MIGRATE using *cyt b* and CR.

Ploidy	From	To	Mean	95% credibility interval	Nm
Diploids	Han River	Haihe River	316.42	250.00-384.00	0.092
	Dongting Lake	Haihe River	127.32	0.00-28.67	0.037
	Poyang Lake	Haihe River	296.55	230.67-362.67	0.086
	Pearl River	Haihe River	298.65	233.33-365.33	0.087
	Haihe Rvier	Han River	305.36	240.00-371.33	1.111
	Dongting Lake	Han River	294.50	229.33-359.33	1.071
	Poyang Lake	Han River	274.60	213.33-336.67	0.999
	Pearl River	Han River	294.50	229.33-360.67	1.071
	Haihe Rvier	Dongting Lake	284.69	222.00-348.00	0.799
	Han River	Dongting Lake	305.17	240.00-370.67	0.856
	Poyang Lake	Dongting Lake	296.74	232.67-361.33	0.832
	Pearl River	Dongting Lake	315.71	250.67-380.67	0.886
	Haihe Rvier	Poyang Lake	165.92	13.33-70.67	1.035
	Han River	Poyang Lake	287.11	225.33-350.00	1.790
	Dongting Lake	Poyang Lake	273.86	212.67-335.33	1.708
	Pearl River	Poyang Lake	263.53	205.33-322.67	1.643
	Haihe Rvier	Pearl River	304.53	240.67-368.00	0.687
Tetraploids	Han River	Pearl River	279.04	216.67-342.00	0.629
	Dongting Lake	Pearl River	303.40	238.67-369.33	0.684
	Poyang Lake	Pearl River	281.18	218.00-346.00	0.634
	Han River	Haihe River	299.75	232.67-366.67	0.694
	Dongting Lake	Haihe River	299.10	231.33-366.00	0.692
	Poyang Lake	Haihe River	300.20	233.33-367.33	0.695
	Pearl River	Haihe River	298.23	232.00-364.67	0.690
	Haihe Rvier	Han River	296.64	230.67-362.67	0.521

Haihe Rvier	Dongting Lake	298.43	232.00-366.00	0.654
Han River	Dongting Lake	303.25	236.00-371.33	0.664
Poyang Lake	Dongting Lake	297.35	230.67-365.33	0.651
Pearl River	Dongting Lake	301.41	235.33-367.33	0.660
Haihe Rvier	Poyang Lake	134.31	0.00-42.67	0.609
Han River	Poyang Lake	159.13	0.00-46.00	0.721
Dongting Lake	Poyang Lake	283.55	220.67-348.00	1.285
Pearl River	Poyang Lake	283.45	219.33-348.00	1.285
Haihe Rvier	Pearl River	307.26	240.00-376.67	0.604
Han River	Pearl River	303.70	236.67-371.33	0.597
Dongting Lake	Pearl River	299.18	232.00-366.00	0.588
Poyang Lake	Pearl River	301.16	235.33-367.33	0.592

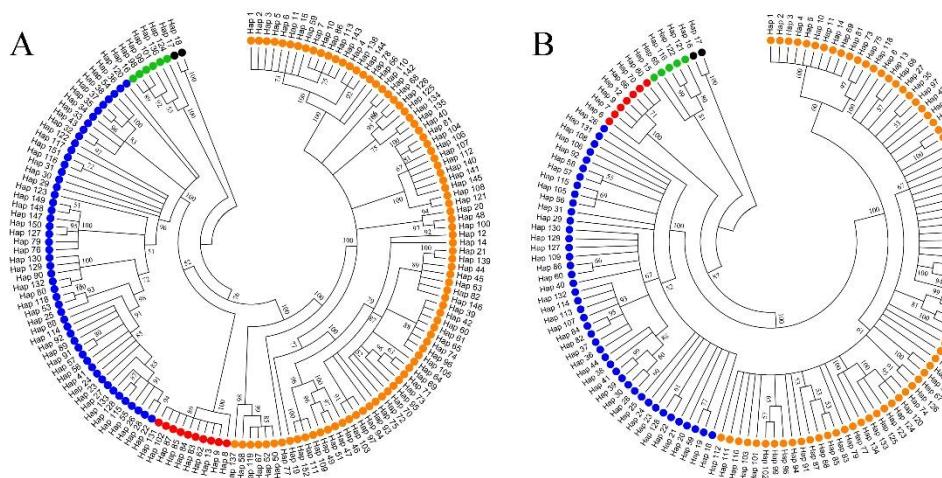


Fig. S1 Bayesian phylogenetic tree for 22 mixed-ploidy *M. anguilllicaudatus* populations based on *cyt b* (A) and CR (B) datasets. Numbers above the branches correspond to the bootstrap support values (only values > 50% are shown). Clades are highlighted with different colours: yellow - lineage I, blue - lineage II, red - lineage III, green - lineage IV and black - lineage V.

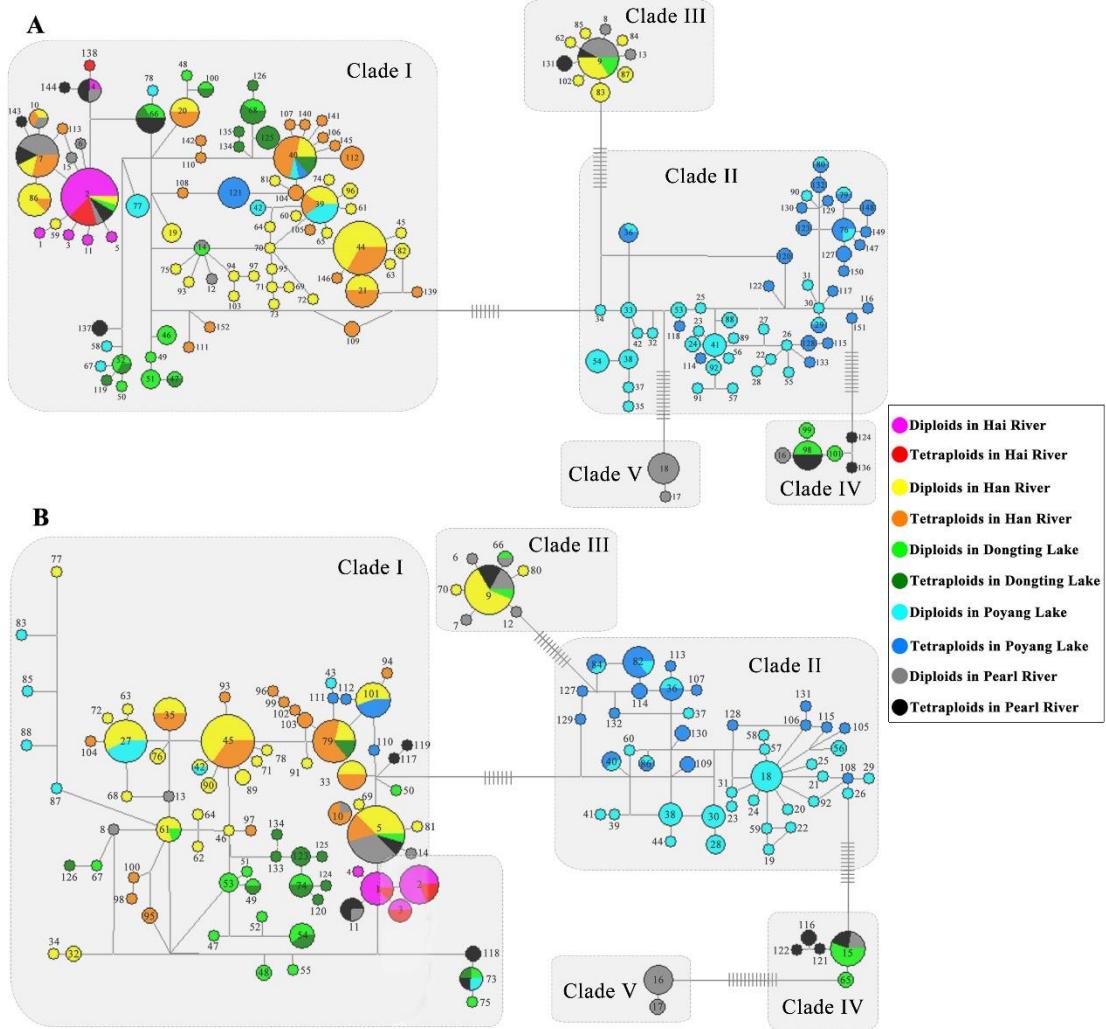


Fig. S2 Median-joining network of *cyt b* (**a**) and CR (**b**) haplotypes from 22 populations of *Misgurnus anguillicaudatus*. The circle size of a haplotype denotes the number of specimens. Colors correspond to different ploidies and regions.

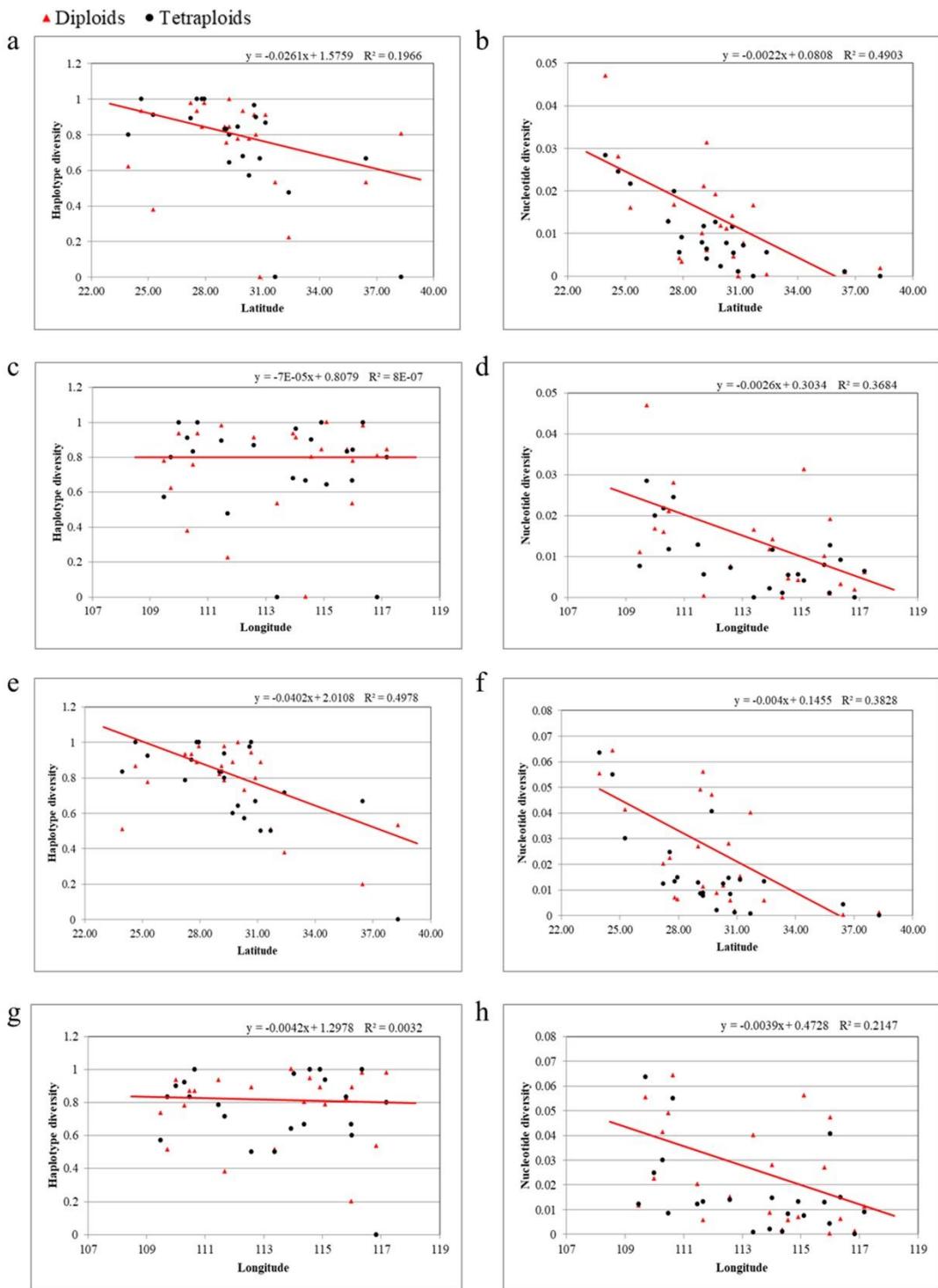


Fig. S3 Haplotype (**a, c, e, g**) and nucleotide diversity (**b, d, f, h**) of *M. anguilllicaudatus* plotted against latitude and longitude for diploid populations (red) and tetraploid populations (black) based on CR (**a, b, c, d**) and *cyt b* (**e, f, g, h**). The regression lines with equations and explained variances (R^2) are based on all population types.

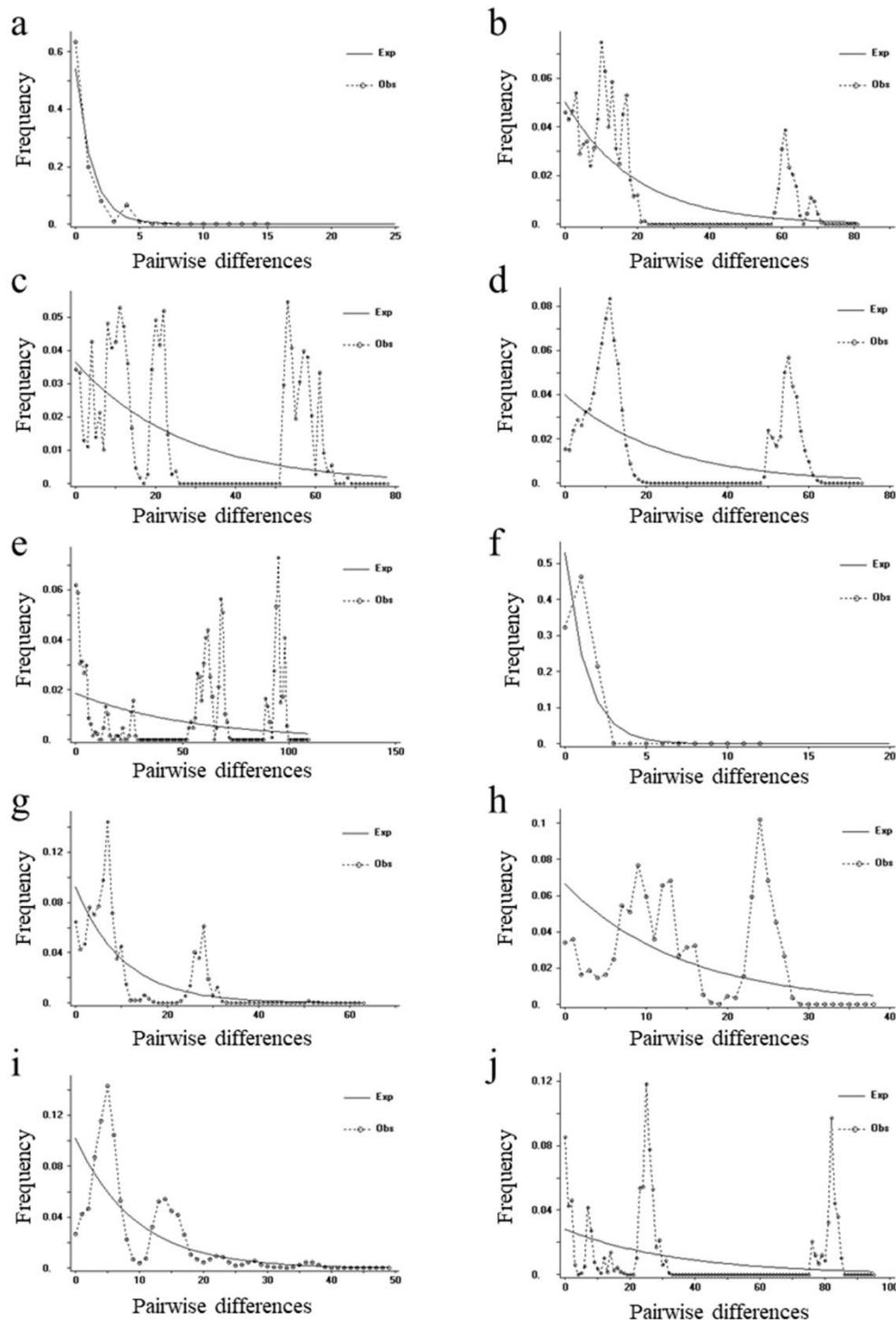


Fig. S4 Observed and expected mismatch distributions for *M. anguillicaudatus* in the (a) Haihe River, (b) Han River, (c) Dongting Lake, (d) Poyang Lake and (e) Pearl River for the cyt b dataset, and in the (f) Haihe River, (g) Han River, (h) Dongting Lake, (i) Poyang Lake and (j) Pearl River for the CR dataset. Dashed line, observed distribution; solid line, theoretical expected distribution under a population expansion model.