

表 1 中华鳖 5 个群体 11 个微卫星位点的遗传多样性参数

Table 1 The parameters of genetic diversity of five populations of *Trionyx sinensis* at 11 microsatellite loci

位点 Loci	洞庭(DT)			黄河(HH)			黄沙(HS)			日本(RB)			绿卡(LK)		
	Dongting(DT)			Huanghe(HH)			Huangsha(HS)			Riben(RB)			Lvka(LK)		
	Na	Ne	PIC	Na	Ne	PIC	Na	Ne	PIC	Na	Ne	PIC	Na	Ne	PIC
P-01	3	2.8986	0.5812	3	2.9557	0.5876	4	1.7094	0.3894	4	3.0303	0.6125	4	2.6987	0.5627
P-04	3	2.4357	0.5052	3	2.1978	0.4415	3	2.9364	0.5851	3	1.8349	0.3656	3	1.9417	0.3825
P-05	3	1.1834	0.1470	3	1.1443	0.1213	3	2.0955	0.4166	3	2.5388	0.5380	3	1.3564	0.2415
P-06	3	1.8672	0.3832	3	1.8730	0.3720	2	1.9231	0.3648	2	1.5571	0.2938	3	1.8237	0.3754
P-09	5	3.6961	0.6813	5	3.1469	0.6217	5	3.7344	0.6936	3	2.4862	0.5169	5	4.1190	0.7147
P-10	3	1.8967	0.3988	3	2.4759	0.5095	3	2.9801	0.5904	3	1.7946	0.3585	3	2.2613	0.4778
P-11	3	1.9715	0.3867	3	1.6202	0.3212	2	1.9802	0.3725	3	1.2649	0.1935	3	1.9417	0.4064
P-12	4	1.4096	0.2697	3	1.3072	0.2142	3	1.2262	0.1754	2	1.0339	0.0323	4	1.7459	0.3892
PS-04	4	1.5721	0.3362	5	1.5345	0.3333	3	1.2685	0.1990	1	1.0000	0.0000	4	1.4140	0.2750
PS-11	5	2.7607	0.5908	5	2.5388	0.5678	6	1.4252	0.2883	6	3.4749	0.6687	6	2.3018	0.5381
HLJZHB 076	6	4.2959	0.7340	4	2.6866	0.5674	5	3.2907	0.6502	4	2.1924	0.4424	6	4.7619	0.7612

注:  $N_a$ : 等位基因数;  $N_e$ : 有效等位基因数; PIC: 多态信息含量

Note:  $N_a$ : Number of alleles;  $N_e$ : Effective number of alleles, PIC: Polymorphism information content