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In the article "Genetic Structure of Korean Wild Populations of the Medaka *Oryzias latipes* Inferred from Allozymic Variation" by Yusuke Takehana, Sang-Rin Jeon and Mitsuru Sakaizumi, which appeared on pages 977–988 of volume 21 No. 9 (2004), errors were found in Figure 5 on page 982. The correct version of Figure 5 is printed here.

	Ch	China-West Korean Population				hybrid populations		East Korean Population					
	10 (10)	24 (6)	36 (5)	38 (5)	7 (5)	35 (5)	6 (5)	34 (6)	52 (5)	62 (5)	86 (7)	92 (7)	
ACP*													
AMY*													
CK-A*													
GPI-A*			•		•	•							
LDH-A*													
LDH-C*	· ()												
PGM*									•				
SOD*							\bigcirc						
TF*													
mtDNA	(4)	(7)	(5)	(1)	(2)	(2)	(3)	(6)	(3)	(5)	(10)	(3)	