

Table 3. Percent of sampled colonies assigned to genotypes based on ten SSR markers.

Populations	Number colonies	Percent of colonies assigned to genotypes	
		<i>R. chinensis</i>	<i>R. flaviceps</i>
<i>R. chinensis</i> (possessing mtDNA haplotypes of <i>R. chinensis</i>)			
Chengdu	10	0.851	0.149
Nanjing	5	0.974	0.026
Wuhan	14	0.978	0.022
<i>R. flaviceps</i> (possessing mtDNA haplotypes of <i>R. flaviceps</i>)			
Chengdu	15	0.237	0.763
Nanjing	17	0.048	0.952
Wuhan	6	0.015	0.985