

**Table S1.** Molecular variance analysis among and within 10 *C. yunnanensis* populations using 9 SSR loci.

Source of variation	d.f.	Sum of squares	Mean square	Variance component	Percentage of variation (%)	P-value
Among populations	9	171.444	19.049	0.931	13%	0.001
Within populations	140	883.960	6.314	6.314	87%	0.001
Total	149	1055.404		7.245	100%	