

Table 5 Estimates of different diversity parameters involving genotypes of *C. capsularis* (152) and *C. olitorius* (140) using polymorphic SSR loci

Collection of genotypes	<i>N</i>	<i>N_a</i>	<i>N_e</i>	<i>I</i>	<i>H_e</i>
<i>C. capsularis</i>					
Indigenous	101	2.619	1.567	0.492	0.289
Exotic	51	2.08	1.584	0.462	0.282
Total	152				
Average		2.35	1.575	0.477	0.286
<i>C. olitorius</i>					
Indigenous	106	2.393	1.445	0.415	0.245
Exotic	34	2.107	1.530	0.454	0.278
Total	140				
Average		2.25	1.487	0.435	0.262

Note: *N*, sample size; *N_a*, mean no. of observed alleles per locus; *N_e*, mean no. of effective alleles per locus; *I*, mean values of Shanon's information index per locus; *H_e*, mean expected heterozygosity per locus