

Table S2. Genetic diversity parameters among hop populations of the three Calabrian areas (CS, CZ and VV) based on SSR markers.

		N	Ne	H	I	F
Pop-CS	<i>Mean</i>	11.875	5.033	0.823	1.743	0.080
Pop-CZ	<i>Mean</i>	4.750	2.993	0.702	1.156	-0.015
Pop-VV	<i>Mean</i>	4.750	3.005	0.703	1.153	0.048

N, number of alleles; Ne, number of effective alleles; H, gene diversity; I, Shannon's information index; F, fixation index