

Table 1 Combined analysis in 36 safflower accessions evaluated at Holetta and Ginchi

	df	Mean square				
		DF	DM	PH (cm)	PB	SY (kg/ha)
Genotype	35	8.936**	3.386*	219.795**	0.776*	186 439.004**
Environment	1	10 746.778**	26 786.778**	4 880.248**	39.213**	4 408 055.770**
Replication(environment)	2	25.847	12.361	815.600	1.528	193 622.276
Genotype X Environment	35	3.692 ^{ns}	3.106*	108.333**	0.427*	65 990.580 ^{ns}
Error	70	2.704	1.875	79.875	0.426	56 968.517
CV (%)		1.110	0.650	7.470	11.820	60.370

Note: *, **: significant at 0.05 and 0.01 probability level respectively; CV: coefficient of variation; df: degree of freedom