Table 2 Analysis of variance for treatments (parents and crosses) of log_{10} transformed values in 9×9 diallel crosses (excluding reciprocals) for AUDPC† and FDS†† in F_1 and F_2 generations

Source of variation	Degree of freedom	Mean square			
		F_1		F_2	
		AUDPC	FDS	AUDPC	FDS
Replication	2	0.02	0.01	0.16**	0.20**
Treatment	44	0.74**	0.28**	0.44**	0.18**
Error	88	0.01	0.01	0.02	0.02

[†] Area Under Disease Progress Curve and †† Final Disease Severity

^{**} Significant at P<0.01