Table 3 Estimates of genetic parameters and their proportions for AUDPC† and FDS†† in 9×9 diallel crosses (excluding reciprocals) of F_1 and F_2 generations

		Estimate			
Parameter		F ₁		F ₂	
		AUDPC	FDS	AUDPC	FDS
D	Additive genetic variance	0.17**	0.10**	0.17**	0.09**
$\mid H_1 \mid$	Dominance genetic variance	0.60**	0.19**	1.39**	0.41**
$\mid H_2 \mid$	Proportion of dominance genetic variance due to +	0.47**	0.15**	1.26**	0.38**
and – effects of genes				1.20	0.36
$\mid h^2 \mid$	Net dominance effect (as algebraic sum over all loci	0.53**	0.15**	3.70**	0.46**
in heterozygous phase in all crosses)					0.40
$\mid F \mid$	Mean covariance of additive and dominance effect	0.01	0.02	0.24	0.01**
of genes			***-		
$\mid E \mid$	Expected environmental variance	0.01	0.01	0.02	0.01**
Proportion					
$ P_1 /D ^2$	Average degree of dominance	1.87	1.40	1.43	1.06
$ H_{2} /4H_{1} $	Proportion of genes with + and – effects in parents	0.19	0.20	0.23	0.23
h^2/H_2	Number of group(s) of gene which control the trait				
and exhibit dominance		1.12	0.97	2.93	1.21
$ R_D^{\bullet}/K_R $	Proportion of dominant-recessive genes in parents	1.04	1.12	1.67	1.82
h_{ns}^2	Heritability (narrow sense)	54.45	61.31	40.09	54.55

[†] Area Under Disease Progress Curve and †† Final Disease Severity, ** Significant at P<0.01