Table (3): Estimates of heritability, expected and observed response to selection for stomata frequency and coefficient of variation for different families selected for high and low directions in populations 1 and 2 in the first (F_3) and second (F_4) cycles of selection.

Selection direction	Families	Population(1)							Population (2)							
		F ₃														
		bop♣	h²ф	h ² !!	Ex.resp.%	O.resp. %	Variance♠	C.V.	bop♣	h²ф	h ² !!	Ex.resp.%	O.resp. %	Variance♠	C.V.	
Low	1	0.45	0.1867	0.3176	7.5060	3.6276	23.06*	5.806	0.266	0.2210	0.2885	5.3439	5.4560	28.75 ns	6.969	
	2		0.1843		7.5060	3.5808	39.35 ^{ns}	7.580		0.4349		5.0221	10.0895	32.42 ns	7.782	
	3		0.5820		5.8815	8.8622	17.22**	5.306		0.2097		6.1419	5.9489	30.32 ns	6.316	
High	1		0.7124		8.4735	15.6282	35.45 ns	6.000		0.3387	0.4580	6.3414	9.9235	58.38 ns	8.541	
	2		0.6378	0.7642	9.5490	15.7684	29.01*	5.421		0.4502		5.2216	10.8600	54.70 ns	8.198	
	3		0.9426		6.8445	16.7032	24.10*	4.901		0.5851		5.2216	14.1134	29.00 ns	5.799	
Bulk						•	68.25	9.625					•	52.22	8.880	
Selection direction	Families		F_4													
		bop♣	h²ф	h ² !!	Ex.resp.%	O.resp. %	Variance ♠	C.V.	bop♣	h²ф	h ² !!	Ex.resp.%	O.resp. %	Variance♠	C.V.	
Low	1	0.497	0.576	0.484	4.328	6.548	27.60 ns	6.648	0.363	0.699	0.620	3.082	9.300	39.41 ns	7.341	
	2		0.324		5.678	4.839	38.87 ns	7.748		0.411		4.309	7.643	39.25 ns	8.014	
	3		0.552		5.948	8.635	33.71 ns	7.515		0.748		3.508	11.326	40.96 ns	8.187	
High	1		0.573		9.752	14.697	26.18*	5.275		0.564		4.656	11.326	42.62 ns	6.984	
	2		0.524		9.302	12.805	42.20 ns	6.810		0.442		4.975	9.485	66.28 ns	8.732	
	3		0.430		9.392	10.617	34.13 ns	6.245		0.316		5.403	7.367	99.30 ns	11.067	
Bulk					·	1	63.22	9.403			L	ı	ı	75.31	9.960	

 $_{op}$ *= mid-parent offspring regression (F₃ /F₄), φ= Family realized heritability, !!= realized heritability in the high or low directions, Ex.resp.% = expected response to selection, O.resp.% = observed response to selection. (ns= variances non-significantly different from the controls' variances, $_{family}$ variances were tested against bulk variances using F test: *= significantly different from the control (P<0.05), **= high-significantly different from the control, based on F test).