

表 3 耐盐性位点的方差分析

Table 3 Distribution of salt tolerance by ANOVA analysis

性状 Trait	连锁群 Group	标记 Marker	位置 Location	F 值 F-value	加性效应* Additive effect	贡献率 Contribution (%)
发芽率	C1	Satt180	127.77	4.77	-0.06	19.26
GR	C1	SOYGPATR	10.34	7.44	-0.03	27.11
	D1b	Satt271	137.06	4.49	0.05	18.32
	E	Satt231	70.23	6.99	0.10	25.89
	E	Satt411	12.92	10.75	0.05	34.96
	I	Sat_170	75.00	4.59	-0.01	20.32
	N	Satt255	76.49	5.32	0.03	21.88
	O	Sat_108*	129.30	7.24	0.10	27.59
发芽势	E	Satt411	12.92	12.35	-0.09	38.18
GE	E	Satt231	70.23	8.60	0.11	30.07
	N	Satt549	70.60	4.57	0.09	18.59
	N	Satt255	76.49	6.04	0.04	24.12
	O	Sat_108*	129.3	12.36	0.12	39.42
	发芽指数	C1	Satt338	123.79	4.88	-1.30
GI	E	Satt231	70.23	6.72	1.30	25.14
	E	Satt411	12.92	12.66	-1.43	38.77
	N	Satt255	76.49	4.46	0.18	19.01
	O	Sat_108*	129.3	14.83	1.48	43.83
隶属函数	C1	SOYGPATR	10.34	4.95	-0.16	19.85
MF	E	Satt231	70.23	7.97	0.37	28.49
	E	Satt411	12.92	12.75	-0.33	38.94
	N	Satt255	76.49	5.62	0.10	22.82
	O	Sat_108*	129.3	11.91	0.41	38.54

注: *: 加性效应为红丰 11 等位基因被供体 Harosoy 等位基因替代后的效应

Note: *: Additive effect was associated with Hong-feng11 alleles replaced by the alleles of Harosoy