

Evaluation of Genomic Prediction for Pasmov Resistance in Flax

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Supplementary figures and tables

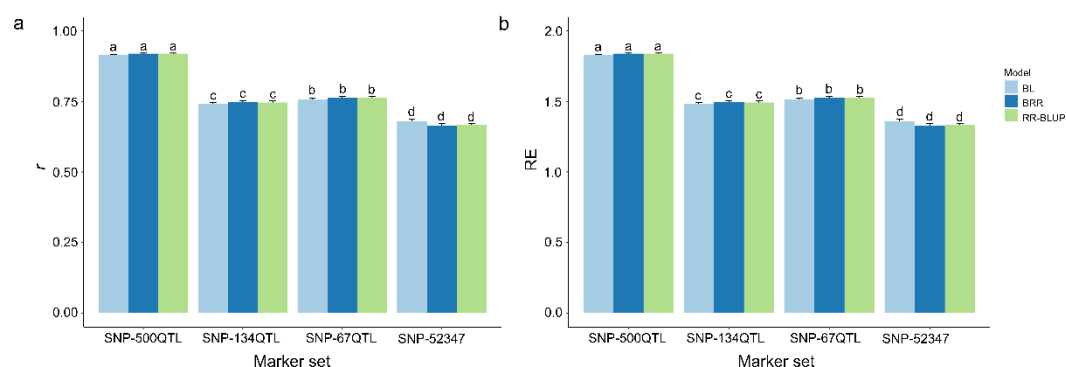


Figure S1. The accuracies (r) (a) and relative efficiency (RE) (b) of prediction models built with combinations of four marker sets and three statistical models—ridge regression best linear unbiased prediction (RR-BLUP), Bayesian LASSO (BL), and Bayesian ridge regression (BRR)—with average pasmo severity (PS) across five years (PS-mean dataset) using random five-fold cross-validation. The statistical significances of r among different statistical models are labeled with letters. Different letters show significant difference at a 5% probability level.

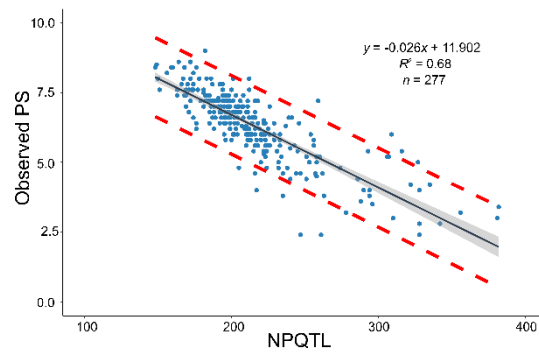


Figure S2. Linear regression of observed PS to the number of positive-effect QTL (NPQTL) in the 277 accessions that were used as a training population for genomic prediction. The grey band represents the 95% confidence interval. The red dashed lines represent upper and lower bounds of the 95% prediction interval. The average width of the prediction interval for all predicted values was 2.70.

Table S1. Analysis of variance of the accuracy (r) and relative efficiency (RE) for different marker sets and pasmo severity (PS) datasets using random five-fold cross-validation.

Source of variance	Degree of freedom	r			RE		
		MS	F	P value	MS	F	P value
Marker set (M)	3	237.8	51,955.8	<2.2e-16	951.1	51,955.8	<2.2e-16
PS dataset (P)	5	73.5	16,063.1	<2.2e-16	294.05	16,063.2	<2.2e-16
M $\times$ P	15	2.4	526.9	<2.2e-16	9.65	526.9	<2.2e-16
Residuals	60,186	0.005			0.02		

MS: mean square; F : F value.

Table S2. Genomic prediction (GP) of pasmo severity (PS) for 93 flax cultivars and breeding lines using the GP model built with SNP-500QTL and DS-mean of 277 accessions as training data.

Cultivars /Lines	Improvement status ¹	Observed PS ²	Predicted PS by GP ³	NPQTL ⁴	Predicted PS by NPQTL ⁵
CN100952	U	8.2 ± 1.1	8.6	152	7.9 (6.5 - 9.4)
CN101052	B	3.4 ± 0.5	3.6	329	3.3 (1.9 - 4.8)
CN101053	B	3.0 ± 0.7	3.3	359	2.6 (1.1 - 4.0)
CN101392	C	6.0 ± 1.9	5.9	230	5.9 (4.5 - 7.3)
CN18988	C	3.2 ± 1.1	3.8	346	2.9 (1.5 - 4.3)
CN98072	C	5.4 ± 2.8	6.1	245	5.5 (4.1 - 6.9)
CN100929	C	2.6 ± 0.5	3.1	356	2.6 (1.2 - 4.1)
CN18983	C	2.8 ± 0.8	3.4	350	2.8 (1.4 - 4.2)
CN19001	C	2.0 ± 0.0	3.0	354	2.7 (1.3 - 4.1)
CN97610	C	6.2 ± 1.1	6.2	211	6.4 (5.0 - 7.8)
CN97616	C	8.4 ± 0.9	7.7	179	7.2 (5.8 - 8.7)
CN101405	U	3.6 ± 0.5	3.7	305	4.0 (2.5 - 5.4)
CN101055	B	5.8 ± 1.6	4.8	261	5.1 (3.7 - 6.5)
CN101116	B	7.2 ± 1.5	6.8	216	6.3 (4.9 - 7.7)
CN101094	C	3.6 ± 1.5	4.3	311	3.8 (2.4 - 5.2)
CN35791	C	5.2 ± 2.2	5.1	267	5.0 (3.5 - 6.4)
CN97530	L	6.8 ± 1.5	6.7	192	6.9 (5.5 - 8.3)
CN97533	L	7.2 ± 1.6	7.0	171	7.5 (6.0 - 8.9)
CN101394	U	6.4 ± 2.7	5.2	265	5.0 (3.6 - 6.4)
CN101395	U	4.4 ± 2.3	4.6	311	3.8 (2.4 - 5.2)
CN101402	U	6.8 ± 1.9	7.1	199	6.7 (5.3 - 8.1)
CN101406	U	3.8 ± 1.3	4.1	288	4.4 (3.0 - 5.8)
CN101382	U	7.0 ± 1.2	7.0	203	6.6 (5.2 - 8.0)
CN32546	C	7.0 ± 2.1	6.7	217	6.3 (4.8 - 7.7)
CN33393	C	3.8 ± 1.6	3.7	367	2.4 (0.9 - 3.8)
CN97665	B	7.8 ± 1.3	8.0	145	8.1 (6.7 - 9.5)
CN98903	B	4.5 ± 1.7	4.9	263	5.1 (3.6 - 6.5)
CN98946	C	4.8 ± 1.1	5.6	252	5.3 (3.9 - 6.8)
CN101338	U	5.8 ± 0.8	6.4	212	6.4 (5.0 - 7.8)
CN97953	C	6.0 ± 1.6	6.3	201	6.7 (5.3 - 8.1)
CN101373	U	5.6 ± 1.5	5.9	222	6.1 (4.7 - 7.5)
CN101469	B	6.4 ± 0.5	6.6	205	6.6 (5.2 - 8.0)
CN101536	B	4.4 ± 1.5	5.3	217	6.3 (4.8 - 7.7)
CN101542	B	8.4 ± 0.5	7.6	183	7.1 (5.7 - 8.6)
CN101560	B	4.6 ± 2.6	5.1	260	5.1 (3.7 - 6.6)
CN101580	B	5.8 ± 1.6	5.7	216	6.3 (4.9 - 7.7)
CN101594	B	6.2 ± 1.3	6.1	195	6.8 (5.4 - 8.2)

CN18980	C	6.4 ± 2.6	5.5	224	6.1 (4.7 - 7.5)
CN19004	C	6.6 ± 2.3	6.9	213	6.4 (5.0 - 7.8)
CN19157	C	6.0 ± 1.0	6.3	203	6.6 (5.2 - 8.0)
CN33388	C	6.0 ± 0.7	5.5	245	5.5 (4.1 - 6.9)
CN97671	C	7.6 ± 1.1	7.9	177	7.3 (5.9 - 8.7)
Linola989	C	4.8 ± 1.7	5.4	255	5.3 (3.9 - 6.7)
PrairieBlue	C	5.8 ± 1.3	5.3	247	5.5 (4.1 - 6.9)
CN98683	C	6.0 ± 2.2	6.3	246	5.5 (4.1 - 6.9)
CN100881	C	7.2 ± 0.8	7.0	188	7.0 (5.6 - 8.4)
CN98475	C	6.4 ± 1.1	7.0	208	6.5 (5.1 - 7.9)
CN98712	C	6.8 ± 1.8	6.8	198	6.8 (5.3 - 8.2)
CN98767	C	7.2 ± 1.5	6.8	198	6.8 (5.3 - 8.2)
CN98773	C	5.4 ± 1.5	5.8	215	6.3 (4.9 - 7.7)
CN101367	U	1.8 ± 0.4	2.6	351	2.8 (1.3 - 4.2)
CN101325	U	7.0 ± 1.2	6.4	199	6.7 (5.3 - 8.1)
CN97300	C	7.2 ± 0.8	7.3	198	6.8 (5.3 - 8.2)
CN98275	C	7.0 ± 1.4	7.1	173	7.4 (6.0 - 8.8)
CN98278	C	6.4 ± 1.5	6.5	211	6.4 (5.0 - 7.8)
CN98854	C	7.4 ± 1.1	6.9	189	7.0 (5.6 - 8.4)
CN98250	C	7.0 ± 0.7	6.6	204	6.6 (5.2 - 8.0)
CN98254	C	7.0 ± 0.7	6.4	204	6.6 (5.2 - 8.0)
CN98440	C	6.0 ± 0.7	5.6	220	6.2 (4.8 - 7.6)
CN98468	C	7.0 ± 1.2	6.8	191	6.9 (5.5 - 8.3)
CN98974	C	6.6 ± 1.5	6.6	209	6.5 (5.1 - 7.9)
CN98240	L	5.3 ± 1.7	6.2	215	6.3 (4.9 - 7.7)
CN101310	U	7.4 ± 0.9	6.9	179	7.2 (5.8 - 8.7)
CN98569	U	6.4 ± 2.4	6.5	210	6.4 (5.0 - 7.8)
CN101240	B	6.4 ± 1.8	6.2	217	6.3 (4.8 - 7.7)
CN101026	B	5.4 ± 2.2	5.5	242	5.6 (4.2 - 7.0)
CN97064	C	5.4 ± 2.5	5.9	210	6.4 (5.0 - 7.8)
CN97092	C	5.2 ± 0.8	5.9	214	6.3 (4.9 - 7.7)
CN97103	C	5.0 ± 1.0	4.7	229	5.9 (4.5 - 7.4)
CN101289	B	4.4 ± 1.5	4.7	286	4.5 (3.0 - 5.9)
CN101299	B	3.0 ± 0.7	3.5	297	4.2 (2.8 - 5.6)
CN101307	B	7.0 ± 1.0	7.4	186	7.1 (5.7 - 8.5)
CN97487	C	7.2 ± 1.5	7.4	168	7.5 (6.1 - 8.9)
CN97520	C	5.6 ± 2.2	5.9	214	6.3 (4.9 - 7.7)
CN101375	U	7.0 ± 1.2	6.6	212	6.4 (5.0 - 7.8)
CN97147	C	8.2 ± 0.8	8.0	157	7.8 (6.4 - 9.2)
CN100837	U	6.0 ± 0.7	5.9	238	5.7 (4.3 - 7.1)
CN101332	U	7.2 ± 1.3	7.3	196	6.8 (5.4 - 8.2)
CN30861	C	6.6 ± 1.8	6.8	190	7.0 (5.6 - 8.4)

CN97404B	B	6.6 ± 1.3	7.0	191	6.9 (5.5 - 8.3)
CN33399	C	5.8 ± 1.6	6.3	237	5.7 (4.3 - 7.1)
CN97403	C	7.4 ± 1.3	7.2	172	7.4 (6.0 - 8.8)
CN97407	C	6.8 ± 2.3	6.1	208	6.5 (5.1 - 7.9)
CN97921	C	5.4 ± 1.3	6.1	227	6.0 (4.6 - 7.4)
CN98541	C	6.4 ± 1.5	5.9	214	6.3 (4.9 - 7.7)
CN98812	C	5.6 ± 1.7	6.4	251	5.4 (4.0 - 6.8)
CN97341	C	7.0 ± 1.0	6.5	196	6.8 (5.4 - 8.2)
CN101572	B	5.0 ± 1.4	4.9	243	5.6 (4.2 - 7.0)
CN101016	C	4.6 ± 0.9	5.2	228	6.0 (4.6 - 7.4)
CN100885	U	5.8 ± 1.5	5.9	233	5.8 (4.4 - 7.2)
CN100790	C	8.0 ± 0.7	7.3	187	7.0 (5.6 - 8.4)
CN97402	B	6.8 ± 1.6	6.5	194	6.9 (5.4 - 8.3)
CN98644	B	7.3 ± 1.0	7.1	194	6.9 (5.4 - 8.3)

¹ B: Breeding lines; C: Cultivars; L: Landrace; U: Unknown. ² Average PS across five years (2012-2016) and their standard deviations. ³ Predicted PS based on the GP model built with the marker set SNP-500QTL, the PS-mean dataset of 277 accessions, and the statistical model RR-BLUP. ⁴ NPQTL: number of positive-effect QTL. ⁵ Predicted by NPQTL (the equation is $y = -0.026x + 11.902$ based on 277 accessions as training data). The values in parentheses are prediction intervals.