Multi-trait ridge regression BLUP with de novo GWAS improves genomic prediction for haploid induction ability and agronomic traits of haploid inducers in maize

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Supplementary Table1. The number of genotypes derived from different F_1 combinations or materials in phenotypic measurement and used in the training population of genomic prediction

No.	Plant materials	Phenotypic measurement				Training population in genomic prediction				
		DTF	HIR	PHT	PBL	DTF	HIR	PHT	PBL	Multi-trait
1	A637-derived/BHI306	10	10	10	26	10	10	10	26	10
2	B73-derived/A637-derived	6	6	6	5	6	6	6	5	5
3	B73-derived/B84-derived	7	6	7	7	7	6	7	7	6
4	B73-derived/BHI306	11	10	11	15	10	10	10	14	10
5	B73-derived/MHI	29	31	31	31	29	31	31	31	27
6	B73-derived/Mo17-derived	16	15	16	21	16	15	16	21	15
7	B73-derived/PHG83-derived	14	13	14	33	14	13	14	33	13
8	B84-derived/BHI306	8	8	8	11	8	8	8	11	8
9	FR19-derived/A637-derived	12	12	12	16	12	12	12	16	12

10	FR19-derived/B73-derived	16	16	16	23	16	16	16	23	16
11	FR19-derived/BHI306	10	10	10	23	10	10	10	23	10
12	LH82-derived/B73-derived	12	12	12	15	12	12	12	15	12
13	LH82-derived/BHI306	13	13	13	18	13	13	13	18	13
14	LH82-derived/Mo17-derived	10	9	10	16	10	9	10	16	9
15	Mo17-derived/A637-derived	9	9	9	15	9	9	9	15	9
16	Mo17-derived/BHI306	8	8	9	14	8	8	9	14	7
17	PHG83-derived/A637-derived	20	20	20	49	20	20	20	48	20
18	PHG83-derived/B84-derived	10	10	10	15	10	10	10	15	10
19	PHG83-derived/BHI306	11	10	11	28	11	10	11	28	10
20	PHG83-derived/FR19-derived	14	14	14	27	14	14	14	27	14
21	PHG83-derived/LH82-derived	5	5	5	9	5	5	5	9	5
22	PHG83-derived/MHI	35	34	34	34	35	34	34	34	34
23	PHG83-derived/Mo17-derived	8	8	8	8	8	8	8	8	8
24	Parents	9	9	9	9	0	0	0	0	0
25	MAGIC DHIs	11	11	11	11	0	0	0	0	0
26	Elite lines	8	8	8	8	0	0	0	0	0
27	Others	12	14	14	50	0	0	0	0	0