TABLE S2 Heritability estimates in the epiRILs and natural accessions

trait name	Population	n	H <sup>2</sup>	(s.e)	X <sup>2</sup>	df	P value
flowering time	epiRILs	5567	0.2997	0.0109	1225.14	1	< 0.0001
	accessions	356	0.8602	0.0142	610.91	1	< 0.0001
fruit size (main stem)	epiRILs	1317	< 0.0001	0.0054	0	1	0.99804
	accessions	86	0.6017	0.0790	48.54	1	< 0.0001
height from soil to 1st fruit	epiRILs	4535	0.1571	0.0112	408.38	1	< 0.0001
	accessions	292	0.4164	0.0430	118.42	1	< 0.0001
maximal height	epiRILs	4543	0.0628	0.0102	138.96	1	< 0.0001
	accessions	293	0.4032	0.0428	113.56	1	< 0.0001
height of main stem	epiRILs	4561	0.0393	0.0091	87	1	< 0.0001
	accessions	292	0.3593	0.0447	95.89	1	< 0.0001
fruit size (primary branch)	epiRILs	1326	0.0392	0.0060	16.49	1	< 0.0001
	accessions	86 0.	1858	0.0861	9.15	1	0.00249
# of fruits (main stem)	epiRILs	1330	< 0.0001	0.0050	0	1	0.99633
	accessions	88	0.1394	0.0807	6.52	1	0.01067
dry biomass	epiRILs	4586	0.0184	0.0083	39.18	1	< 0.0001
	accessions	294	0.0075	0.0229	1.09	1	0.29624
rosette diameter	epiRILs	2789	0.0160	0.0116	*	*	*
	accessions	177	0.0069	0.0378	*	*	*
# of fruits (primary branch)	epiRILs	1333	< 0.0001	0.0030	0	1	0.99817
	accessions	87	< 0.0001	0.0262	0	1	0.99991

<sup>\*</sup> No values available because  $H^2$  were based on average estimates in this case (see Supporting Methods). However, 95% confidence ( $\pm 1.96$  x s.e.) intervals include zero suggesting non-significance.

Provided are the broad-sense heritability estimates ( $H^2$ ) for the epiRILs and natural accessions. Estimates were obtained from a random effects model as described in the Supporting Methods. \* No values available because  $H^2$  were based on average estimates in this case (see Supporting methods). However, 95% confidence ( $\pm 1.96 \times s.e.$ ) intervals include zero suggesting non-significance.

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