

Analysis of Epistasis among QTLs on Heading Date based on Single Segment Substitution Lines in Rice

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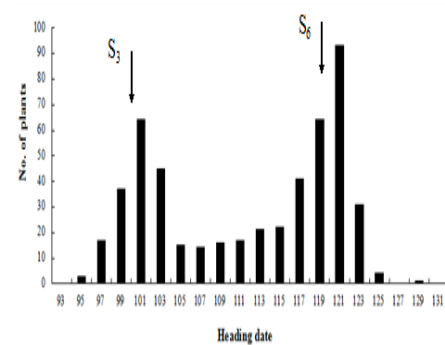
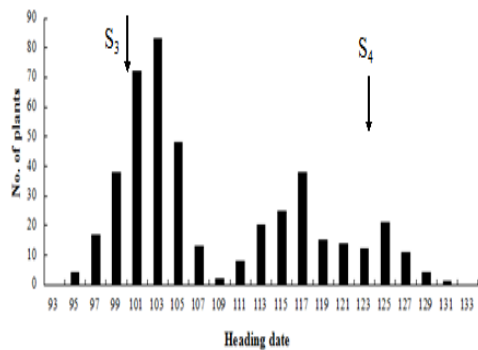
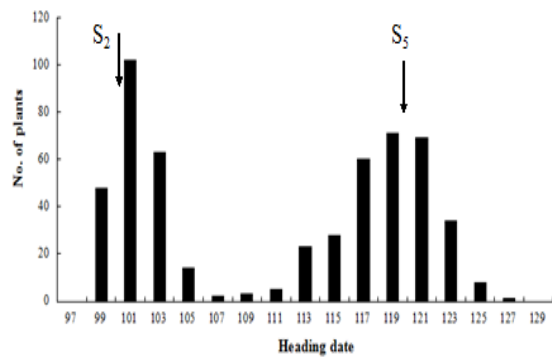
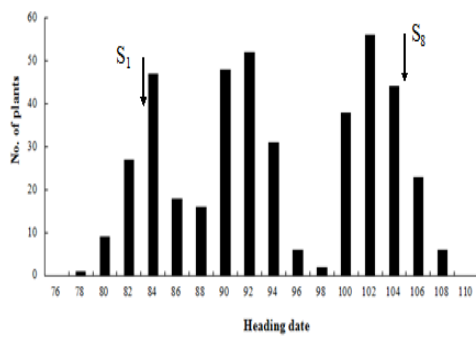
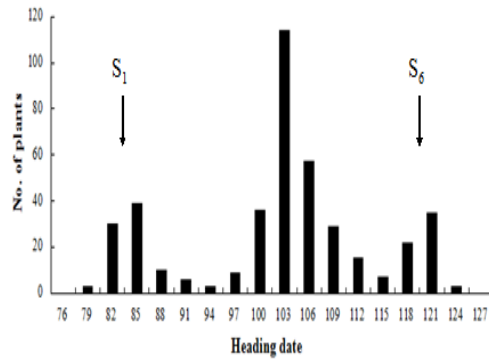
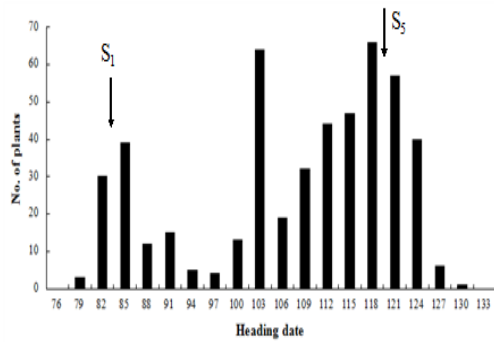
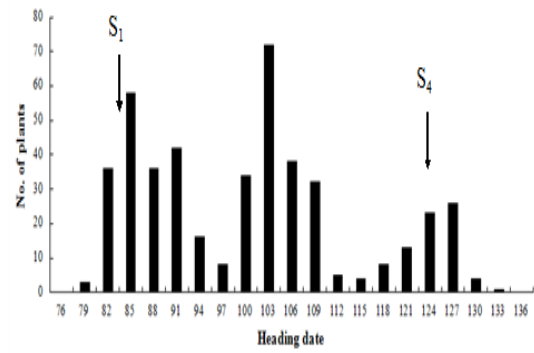
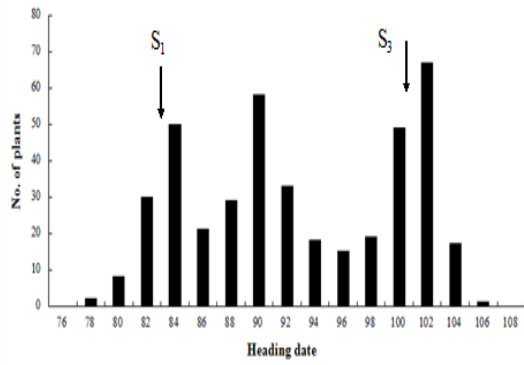
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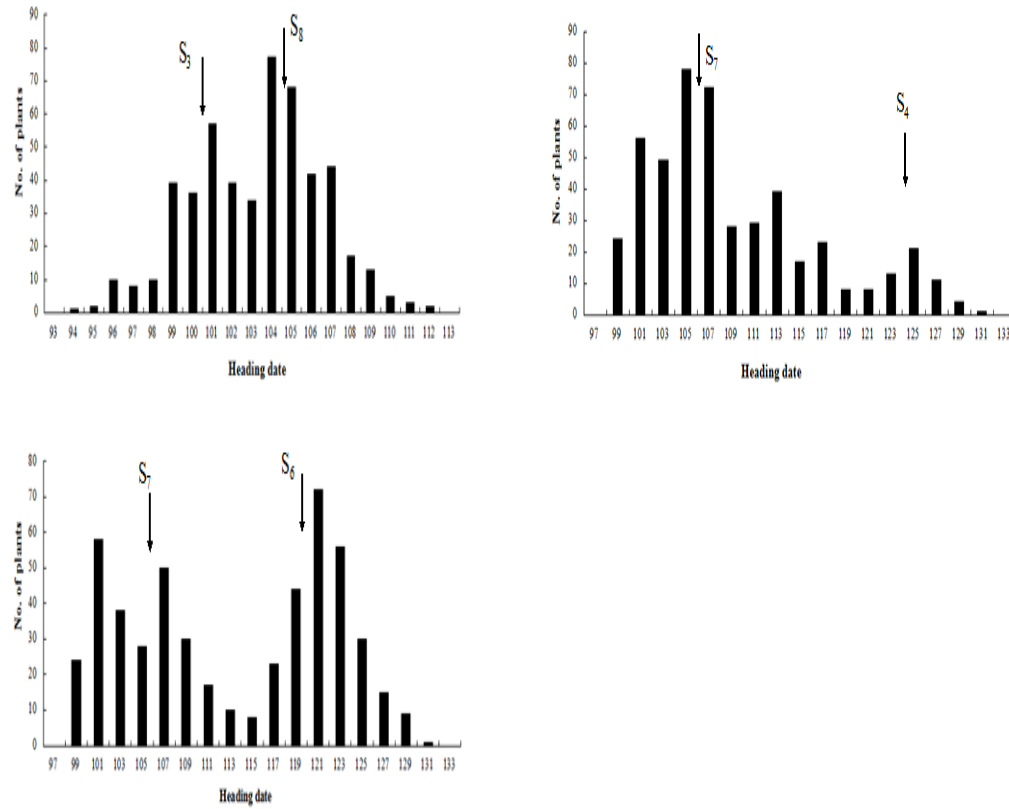
Supplementary Table 1 Analysis of variance on heading date in the both cropping seasons of 2014. Superscripts “*” and “**” indicated the significances at 5% and 1% level, respectively.

Source of variation	Degree of freedom	Sum of squares	Mean square	F-value	P-value
Environment	1	80449.20	80449.20	62066.24**	1.028E-293
Genotype	60	25732.63	428.88	330.88**	3.310E-202
Genotype×Environment	60	5159.32	85.99	66.34**	8.207E-121
Block	2	4.31	2.15	1.66	0.19
Residual error	242	313.68	1.30	-	-

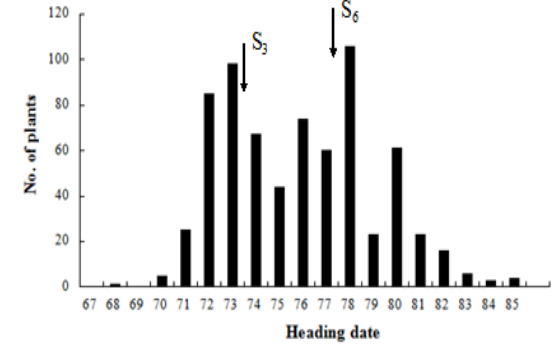
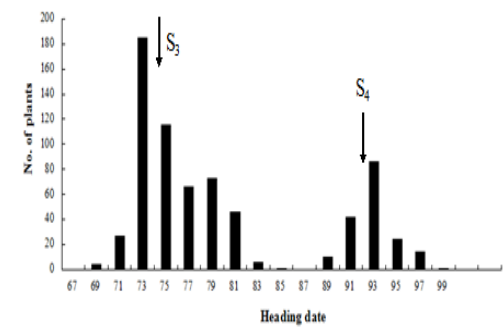
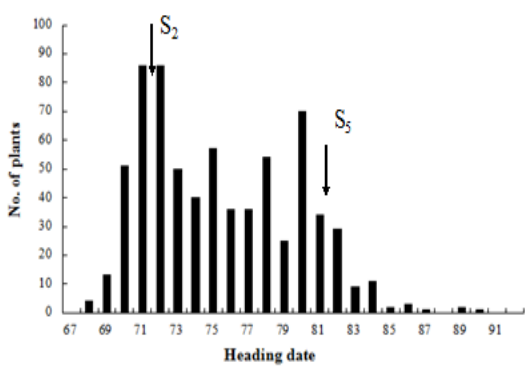
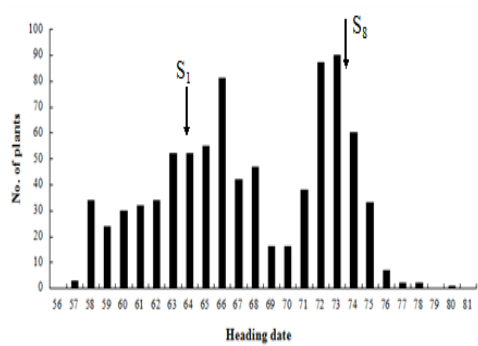
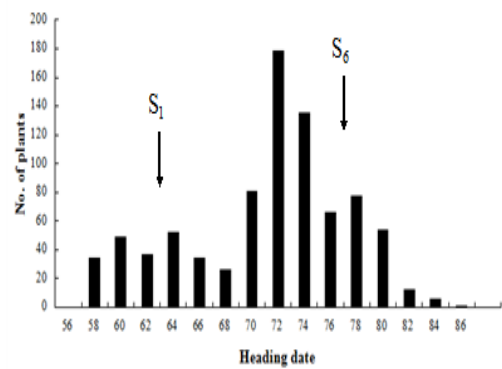
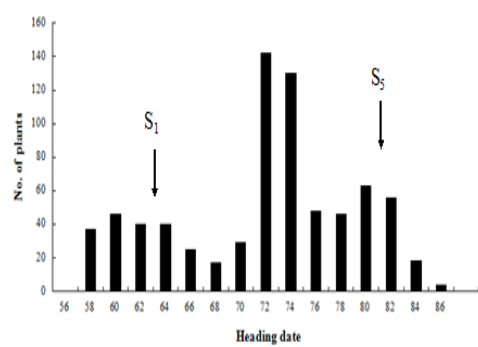
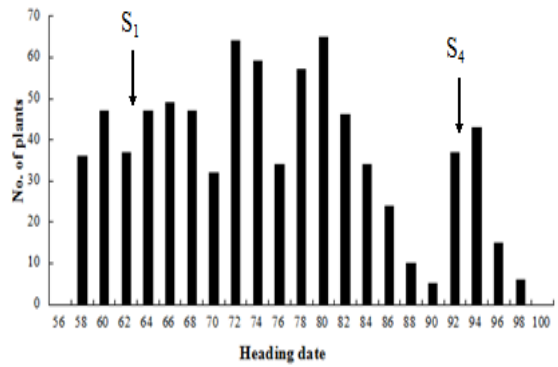
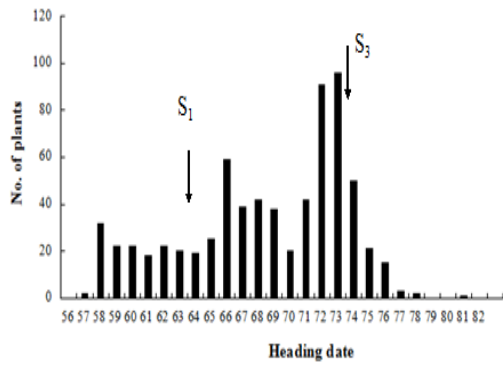
Supplementary Table 2 The genetic effect values of pyramiding materials based on genotypic values of HJX74 in the both seasons of 2014 (day). SSSL was the abbreviation of single segment substitution line. S_i represented the code of SSSL _{i} . *AABB*, *AABb*, *AaBB* and *AaBb* were putative genotypes of SSSL combination. Sign “-” indicated to improve flowering. Superscripts “* and **” indicated the significances at 5% and 1% level, respectively.

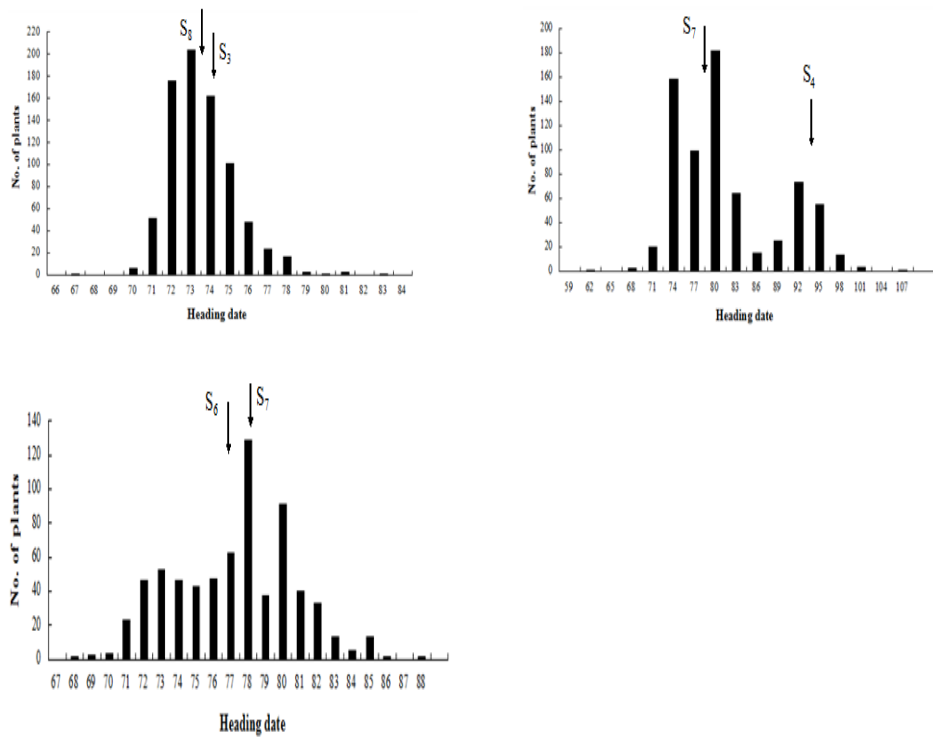
SSSL combination	The early season				The late season			
	<i>AABB</i>	<i>AABb</i>	<i>AaBB</i>	<i>AaBb</i>	<i>AABB</i>	<i>AABb</i>	<i>AaBB</i>	<i>AaBb</i>
S_1/S_3	-9.44**	-15.8**	-10.49**	-10.84**	-4.81**	-7.37**	-3.44**	-3.63**
S_1/S_4	-15.09**	5.17**	-11.02**	-1.14	-3.87**	10.32**	0.27	11.85**
S_1/S_5	4.48**	13.42**	6.50**	15.10**	-2.61**	3.07**	0.56	2.76**
S_1/S_6	-4.44**	1.07	2.13*	2.96**	-4.22**	-1.09	-0.41	1.97*
S_1/S_8	-16.06**	-8.80**	-13.91	-11.25**	-7.87**	-7.43**	-5.09**	-5.03**
S_2/S_5	9.12**	16.85**	12.18**	17.10**	2.02*	2.97**	4.02**	5.78**
S_3/S_4	1.83*	13.87**	1.57*	13.27**	5.39**	18.60**	5.52**	22.37**
S_3/S_6	12.59**	18.58**	19.78**	12.95**	5.06**	6.15**	3.22**	4.15**
S_3/S_8	3.22**	3.97**	3.97**	6.18**	2.39*	1.33	0.24	1.37
S_4/S_7	11.28**	12.70**	4.60**	4.21**	11.29**	18.02**	8.42**	7.38**
S_6/S_7	20.82**	22.47**	14.1**	16.61**	5.53**	8.05**	5.04**	8.03**





Supplementary Fig. 1 Heading date frequency distributions of F₂ populations from two single segment substitution lines in the early season of 2014. S_i represented single segment substitution line SSSL_i





Supplementary Fig. 2 Heading date frequency distributions of F₂ populations from two single segment substitution lines in the late season of 2014. S_i represented single segment substitution line SSSL_i.