Conservation and divergence of autonomous pathway genes in the flowering regulatory network of *Beta vulgaris*

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Supplementary Material

Supplementary Table 1. Primer sequences.

Primer name	Sequence (5'->3')
A479	ATTTTGCCGATTTCGGAAC
A477	TGCAGAGCTTCATGGAACTG
B476	GGAGCTGCAAATTCCCGTCC
A478	ATACCAGTTGCAGGACCAGG
B478	GCGTGGACCGCTTGCTGCAACT
B477	GACTTGTGCCTCCAAGCCTTAA
A039	AACCCTTGAAGAGCAGTTTTGGC
A064	AGAGGTATTATACTATCGATCTACC
A066	GGTAGACGAGAAGTACAGCCAG
A042	CAGCAGCAACTCTAGGTTTAACAACC
A043	ACTATCCTTCAGAAATCGACATCGG
A067	GCTTCGGGATTCAACAGATCTC
A033	GTTTGGGCTGTTAAATAAGGTTGC
A065	CCAAACTCAACCCTCAACAGTG
A396	ATGGCAGAGGCAGACAAGG
A397	ATATCCATAATTTCCACCGTAAACTG
A836	ATGACGGAGAAAGGTAGAGGA
A837	TTAAGAGGGGCATGTCGTTAC
A823	GTTACTTCTTACCATCTGATCC
A825	TTGAACATGAACCCCACCACC
A067	GCTTCGGGATTCAACAGATCTC
A108	CGTGTACATTCAACCTGTTGAAG
B391	CGAAGCTGATTAATCATGGAGG
A046	GCAAGGAGCTCGAGATCAGGC
A043	ACTATCCTTCAGAAATCGACATCGG
B392	GAGCAGTTGACGTCCCTAAG
A068	TACCAGAAGTTGGCGCTGGAG
A069	TAGGGGTAGGTGAAGGAAGAG
A045	CCATGGGATATTGAGATAGACTACG
B396	TTACCGCCTCCGATCACCAC
B042	GTCTGTGAACCCTCAAAAGTC
B043	ATGTTGGTACCTGCCCAAAAG
A066	GGTAGACGAGAAGTACAGCCAG
A042	CAGCAGCAACTCTAGGTTTAACAACC
A830	TATGCTCCGCAGCAGCAACTCTAG
A831	TTCAGAGAGGTAAAGGCGCTGTCGAT
B336	CGATAACCTGGTCAAGTCC
B337	CAGTCTCAAGGTGTTCCTCC
B281	GAGAGACGGTATTCCGTATG
B282	GCATGTGGACTCTAAGAAGG
B349	CTAGCTGCACCACTAACTGC
B350	CGTTAAGAGCTGGAAGCACC
A680	GCCGCCTCGAGCAATATATGGCAGAGGCAGAG
A652	CACCTTTAATACCTATAATTCAAACTAGTTAGTAC
A896	CAACGGAATTCCTACCGTACTACCCCTC
A821	TCTAAATGCGAGTTAACTGGAGC
A901	CGACTCGAGCTAGTCATGGCTGAAGC
A938	GACCCGGGTCATCAGTAACCGTAGCCTGAGC
A875	CAAGATCTCTCTGCCGACAGTGG
A876	AATATCATGCGATCATAGGCGTC
B474	GCTTTGACCGACCACTTCGC
B475	ACGCCGAGAGCAACTTGAA

Supplementary Table 2. Days to bolting and total number of leaves at bolting in 53 T2 families derived from transformation with *BvFLK* or *FLK* and non-transformed controls (Col-0 and *flk-1*). For T2 families in the Col-0 background, plants which bolted five or more days earlier than the mean of the Col-0 control plants (37.88±1.50 days to bolting) were classified as early bolting, all other plants as late bolting. For T2 families in the *flk-1* background, plants which bolted as late as the *flk-1* mutant controls (63-77 days to bolting) were considered as late bolting, whereas plants which bolted much earlier (25-44 days to bolting) and thus within a similar time range as Col-0 control plants (35-40 days to bolting) were classified as early bolting. The observed segregation ratios were tested by X^2 analysis for deviation from the null hypotheses for dominant-recessive inheritance of one or two transgene loci (3:1 or 15:1, respectively). Six putative single transgene-locus families (indicated in bold letters) were tested by PCR for segregation of the transgene.

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
Col-0; 35S::*BvFLK;* 3/1	28	16
Col-0; 35S::*BvFLK;* 3/2	31	16
Col-0; 35S::*BvFLK;* 3/3	28	13
Col-0; 35S::*BvFLK;* 3/4	28	13
Col-0; 35S::*BvFLK;* 3/5	28	15
Col-0; 35S::*BvFLK;* 3/6	31	18
Col-0; 35S::*BvFLK;* 3/7	31	15
Col-0; 35S::*BvFLK;* 3/8	31	21
Col-0; 35S::*BvFLK;* 3/9	28	18
Col-0; 35S::*BvFLK;* 3/10	31	16
Col-0; 35S::*BvFLK;* 3/11	35	15
Col-0; 35S::*BvFLK;* 3/12	28	13
Col-0; 35S::*BvFLK;* 3/13	28	13
Col-0; 35S::*BvFLK;* 3/14	28	11
Col-0; 35S::*BvFLK;* 3/15	31	16
Col-0; 35S::*BvFLK;* 3/16	28	14
Col-0; 35S::*BvFLK;* 3/17	28	14
Segregation ratio (early bolting: late bolting)	16:1	
X^2 for 3:1	3.31 (ns) ***a***	
X^2 for 15:1	0.01 (ns)	
Col-0; 35S::*BvFLK* ; 5/1	31	20
Col-0; 35S::*BvFLK* ; 5/2	-*C*	-
Col-0; 35S::*BvFLK* ; 5/3	31	21
Col-0; 35S::*BvFLK* ; 5/4	31	19
Col-0; 35S::*BvFLK* ; 5/5	31	18
Col-0; 35S::*BvFLK* ; 5/6	31	22
Col-0; 35S::*BvFLK* ; 5/7	28	17
Col-0; 35S::*BvFLK* ; 5/8	28	14
Col-0; 35S::*BvFLK* ; 5/9	31	17
Col-0; 35S::*BvFLK* ; 5/10	31	18
Col-0; 35S::*BvFLK* ; 5/11	31	21
Col-0; 35S::*BvFLK* ; 5/12	31	17
Col-0; 35S::*BvFLK* ; 5/13	31	12
Col-0; 35S::*BvFLK* ; 5/14	31	18
Col-0; 35S::*BvFLK* ; 5/15	39	15

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
Col-0; 35S::*BvFLK;* 4/1	25	14
Col-0; 35S::*BvFLK;* 4/2	28	11
Col-0; 35S::*BvFLK;* 4/3	28	15
Col-0; 35S::*BvFLK;* 4/4	28	15
Col-0; 35S::*BvFLK;* 4/5	28	16
Col-0; 35S::*BvFLK;* 4/6	31	15
Col-0; 35S::*BvFLK;* 4/7	40	22
Col-0; 35S::*BvFLK;* 4/8	28	17
Col-0; 35S::*BvFLK;* 4/9	31	15
Col-0; 35S::*BvFLK;* 4/10	31	19
Col-0; 35S::*BvFLK;* 4/11	31	17
Col-0; 35S::*BvFLK;* 4/12	31	12
Col-0; 35S::*BvFLK;* 4/13	31	17
Col-0; 35S::*BvFLK;* 4/14	38	16
Col-0; 35S::*BvFLK;* 4/15	28	14
Col-0; 35S::*BvFLK;* 4/16	31	14
Col-0; 35S::*BvFLK;* 4/17	24	14
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	
Col-0; 35S::*BvFLK* ; 6/1	31	13
Col-0; 35S::*BvFLK* ; 6/2	28	16
Col-0; 35S::*BvFLK* ; 6/3	35	13
Col-0; 35S::*BvFLK* ; 6/4	-	-
Col-0; 35S::*BvFLK* ; 6/5	24	9
Col-0; 35S::*BvFLK* ; 6/6	31	14
Col-0; 35S::*BvFLK* ; 6/7	28	15
Col-0; 35S::*BvFLK* ; 6/8	28	13
Col-0; 35S::*BvFLK* ; 6/9	28	15
Col-0; 35S::*BvFLK* ; 6/10	28	13
Col-0; 35S::*BvFLK* ; 6/11	39	17
Col-0; 35S::*BvFLK* ; 6/12	28	13
Col-0; 35S::*BvFLK* ; 6/13	31	12
Col-0; 35S::*BvFLK* ; 6/14	31	14
Col-0; 35S::*BvFLK* ; 6/15	31	15

Col-0; 35S::*BvFLK* ; 5/16	31	15
Col-0; 35S::*BvFLK* ; 5/17	31	18
Segregation ratio (early bolting: late bolting)	15:1	
X^2 for 3:1	3.00 (ns)	
X^2 for 15:1	0.00 (ns)	
Col-0; 35S::*BvFLK* ; 7/1	**28**	**16**
Col-0; 35S::*BvFLK* ; 7/2 [b]	**40**	**18**
Col-0; 35S::*BvFLK* ; 7/3	**28**	**11**
Col-0; 35S::*BvFLK* ; 7/4	**28**	**13**
Col-0; 35S::*BvFLK* ; 7/5	**28**	**16**
Col-0; 35S::*BvFLK* ; 7/6	**29**	**15**
Col-0; 35S::*BvFLK* ; 7/7	**33**	**15**
Col-0; 35S::*BvFLK* ; 7/8	**24**	**15**
Col-0; 35S::*BvFLK* ; 7/9	**28**	**14**
Col-0; 35S::*BvFLK* ; 7/10 [b]	**39**	**18**
Col-0; 35S::*BvFLK* ; 7/11	**24**	**15**
Col-0; 35S::*BvFLK* ; 7/12 [b]	**38**	**19**
Col-0; 35S::*BvFLK* ; 7/13	**25**	**17**
Col-0; 35S::*BvFLK* ; 7/14	**24**	**17**
Col-0; 35S::*BvFLK* ; 7/15	**31**	**17**
Col-0; 35S::*BvFLK* ; 7/16	**31**	**17**
Col-0; 35S::*BvFLK* ; 7/17 [b]	**35**	**19**
Segregation ratio (early bolting: late bolting)	13:4	
X^2 for 3:1	0.02 (ns)	
X^2 for 15:1	8.56 **	
Col-0; 35S::*BvFLK* ; 12/1	25	12
Col-0; 35S::*BvFLK* ; 12/2	31	18
Col-0; 35S::*BvFLK* ; 12/3	25	13
Col-0; 35S::*BvFLK* ; 12/4	25	14
Col-0; 35S::*BvFLK* ; 12/5	31	18
Col-0; 35S::*BvFLK* ; 12/6	25	16
Col-0; 35S::*BvFLK* ; 12/7	28	21
Col-0; 35S::*BvFLK* ; 12/8	28	12
Col-0; 35S::*BvFLK* ; 12/9	28	14
Col-0; 35S::*BvFLK* ; 12/10	24	13
Col-0; 35S::*BvFLK* ; 12/11	28	12
Col-0; 35S::*BvFLK* ; 12/12	25	17
Col-0; 35S::*BvFLK* ; 12/13	28	15
Col-0; 35S::*BvFLK* ; 12/14	28	14
Col-0; 35S::*BvFLK* ; 12/15	28	14
Col-0; 35S::*BvFLK* ; 12/16	-	-
Col-0; 35S::*BvFLK* ;12/17	24	17
Segregation ratio (early bolting: late bolting)	16: 0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
Col-0; 35S::*BvFLK* ; 14/1	28	15
Col-0; 35S::*BvFLK* ; 14/2	28	14

Col-0; 35S::*BvFLK* ; 6/16	28	13
Col-0; 35S::*BvFLK* ; 6/17	28	14
Segregation ratio (early bolting: late bolting)	14:2	
X^2 for 3:1	1.33 (ns)	
X^2 for 15:1	1.07 (ns)	
Col-0; 35S::*BvFLK* ; 10/1	28	11
Col-0; 35S::*BvFLK* ; 10/2	31	12
Col-0; 35S::*BvFLK* ; 10/3	28	9
Col-0; 35S::*BvFLK* ; 10/4	38	15
Col-0; 35S::*BvFLK* ; 10/5	28	12
Col-0; 35S::*BvFLK* ; 10/6	28	16
Col-0; 35S::*BvFLK* ; 10/7	28	16
Col-0; 35S::*BvFLK* ; 10/8	28	14
Col-0; 35S::*BvFLK* ; 10/9	28	11
Col-0; 35S::*BvFLK* ; 10/10	31	14
Col-0; 35S::*BvFLK* ; 10/11	28	13
Col-0; 35S::*BvFLK* ; 10/12	31	18
Col-0; 35S::*BvFLK* ; 10/13	28	21
Col-0; 35S::*BvFLK* ; 10/14	-	-
Col-0; 35S::*BvFLK* ; 10/15	24	13
Col-0; 35S::*BvFLK* ; 10/16	28	17
Col-0; 35S::*BvFLK* ; 10/17	28	17
Segregation ratio (early bolting: late bolting)	15: 1	
X^2 for 3:1	3.00 (ns)	
X^2 for 15:1	0.00 (ns)	
Col-0; 35S::*BvFLK* ; 13/1	28	14
Col-0; 35S::*BvFLK* ; 13/2	31	16
Col-0; 35S::*BvFLK* ; 13/3	39	18
Col-0; 35S::*BvFLK* ; 13/4	28	13
Col-0; 35S::*BvFLK* ; 13/5	31	14
Col-0; 35S::*BvFLK* ; 13/6	31	16
Col-0; 35S::*BvFLK* ; 13/7	28	14
Col-0; 35S::*BvFLK* ; 13/8	28	13
Col-0; 35S::*BvFLK* ; 13/9	31	16
Col-0; 35S::*BvFLK* ; 13/10	38	18
Col-0; 35S::*BvFLK* ; 13/11	31	18
Col-0; 35S::*BvFLK* ; 13/12	31	15
Col-0; 35S::*BvFLK* ; 13/13	31	16
Col-0; 35S::*BvFLK* ; 13/14	31	17
Col-0; 35S::*BvFLK* ; 13/15	25	15
Col-0; 35S::*BvFLK* ; 13/16	31	18
Col-0; 35S::*BvFLK* ; 13/17	28	14
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	
Col-0; 35S::*BvFLK* ; 21/1	38	18
Col-0; 35S::*BvFLK* ; 21/2	28	19

Col-0; 35S::*BvFLK* ; 14/3	38	16
Col-0; 35S::*BvFLK* ; 14/4	31	15
Col-0; 35S::*BvFLK* ; 14/5	31	17
Col-0; 35S::*BvFLK* ; 14/6	28	14
Col-0; 35S::*BvFLK* ; 14/7	39	12
Col-0; 35S::*BvFLK* ; 14/8	31	18
Col-0; 35S::*BvFLK* ; 14/9	31	17
Col-0; 35S::*BvFLK* ; 14/10	31	15
Col-0; 35S::*BvFLK* ; 14/11	31	15
Col-0; 35S::*BvFLK* ; 14/12	28	14
Col-0; 35S::*BvFLK* ; 14/13	31	16
Col-0; 35S::*BvFLK* ; 14/14	31	18
Col-0; 35S::*BvFLK* ; 14/15	28	12
Col-0; 35S::*BvFLK* ; 14/16	31	14
Col-0; 35S::*BvFLK* ; 14/17	31	18
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	

Col-0; 35S::*BvFLK* ; 21/3	28	17
Col-0; 35S::*BvFLK* ; 21/4	24	15
Col-0; 35S::*BvFLK* ; 21/5	25	14
Col-0; 35S::*BvFLK* ; 21/6	28	13
Col-0; 35S::*BvFLK* ; 21/7	28	14
Col-0; 35S::*BvFLK* ; 21/8	24	12
Col-0; 35S::*BvFLK* ; 21/9	28	12
Col-0; 35S::*BvFLK* ; 21/10	24	14
Col-0; 35S::*BvFLK* ; 21/11	28	16
Col-0; 35S::*BvFLK* ; 21/12	25	16
Col-0; 35S::*BvFLK* ; 21/13	25	15
Col-0; 35S::*BvFLK* ; 21/14	24	13
Col-0; 35S::*BvFLK* ; 21/15	25	16
Col-0; 35S::*BvFLK* ; 21/16	39	21
Col-0; 35S::*BvFLK* ; 21/17	28	17
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
Col-0; endo::*BvFLK* ; 2/1	25	14
Col-0; endo::*BvFLK* ; 2/2	38	17
Col-0; endo::*BvFLK* ; 2/3	28	16
Col-0; endo::*BvFLK* ; 2/4	28	16
Col-0; endo::*BvFLK* ; 2/5	25	14
Col-0; endo::*BvFLK* ; 2/6	35	6
Col-0; endo::*BvFLK* ; 2/7	24	14
Col-0; endo::*BvFLK* ; 2/8	-	-
Col-0; endo::*BvFLK* ; 2/9	25	13
Col-0; endo::*BvFLK* ; 2/10	24	14
Col-0; endo::*BvFLK* ; 2/11	28	15
Col-0; endo::*BvFLK* ; 2/12	25	12
Col-0; endo::*BvFLK* ; 2/13	31	18
Col-0; endo::*BvFLK* ; 2/14	25	14
Col-0; endo::*BvFLK* ; 2/15	28	15
Col-0; endo::*BvFLK* ; 2/16	24	12
Col-0; endo::*BvFLK* ; 2/17	25	14
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
Col-0; endo::*BvFLK* ; 4/1	28	13
Col-0; endo::*BvFLK* ; 4/2	31	18
Col-0; endo::*BvFLK* ; 4/3	31	17
Col-0; endo::*BvFLK* ; 4/4	28	16
Col-0; endo::*BvFLK* ; 4/5	28	15
Col-0; endo::*BvFLK* ; 4/6	28	18
Col-0; endo::*BvFLK* ; 4/7	25	9
Col-0; endo::*BvFLK* ; 4/8	25	12
Col-0; endo::*BvFLK* ; 4/9	28	16
Col-0; endo::*BvFLK* ; 4/10	31	15
Col-0; endo::*BvFLK* ; 4/11	28	17
Col-0; endo::*BvFLK* ; 4/12	31	19
Col-0; endo::*BvFLK* ; 4/13	28	14
Col-0; endo::*BvFLK* ; 4/14	24	13
Col-0; endo::*BvFLK* ; 4/15	31	18
Col-0; endo::*BvFLK* ; 4/16	28	15
Col-0; endo::*BvFLK* ; 4/17	28	14
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
Col-0; endo::*BvFLK* ; 6/1 [b]	**38**	**18**
Col-0; endo::*BvFLK* ; 6/2	**28**	**15**
Col-0; endo::*BvFLK* ; 6/3	**28**	**15**
Col-0; endo::*BvFLK* ; 6/4	**25**	**12**
Col-0; endo::*BvFLK* ; 6/5	**28**	**14**

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
Col-0; endo::*BvFLK* ; 3/1	31	18
Col-0; endo::*BvFLK* ; 3/2	31	14
Col-0; endo::*BvFLK* ; 3/3	28	15
Col-0; endo::*BvFLK* ; 3/4	31	16
Col-0; endo::*BvFLK* ; 3/5	31	18
Col-0; endo::*BvFLK* ; 3/6	28	16
Col-0; endo::*BvFLK* ; 3/7	28	15
Col-0; endo::*BvFLK* ; 3/8	28	17
Col-0; endo::*BvFLK* ; 3/9	31	18
Col-0; endo::*BvFLK* ; 3/10	28	13
Col-0; endo::*BvFLK* ; 3/11	28	15
Col-0; endo::*BvFLK* ; 3/12	31	18
Col-0; endo::*BvFLK* ; 3/13	31	19
Col-0; endo::*BvFLK* ; 3/14	31	15
Col-0; endo::*BvFLK* ; 3/15	31	17
Col-0; endo::*BvFLK* ; 3/16	-	-
Col-0; endo::*BvFLK* ; 3/17	31	15
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
Col-0; endo::*BvFLK* ; 5/1	28	16
Col-0; endo::*BvFLK* ; 5/2	35	15
Col-0; endo::*BvFLK* ; 5/3	28	17
Col-0; endo::*BvFLK* ; 5/4	28	13
Col-0; endo::*BvFLK* ; 5/5	31	17
Col-0; endo::*BvFLK* ; 5/6	38	15
Col-0; endo::*BvFLK* ; 5/7	31	19
Col-0; endo::*BvFLK* ; 5/8	28	18
Col-0; endo::*BvFLK* ; 5/9	31	19
Col-0; endo::*BvFLK* ; 5/10	28	16
Col-0; endo::*BvFLK* ; 5/11	28	14
Col-0; endo::*BvFLK* ; 5/12	28	18
Col-0; endo::*BvFLK* ; 5/13	31	16
Col-0; endo::*BvFLK* ; 5/14	28	14
Col-0; endo::*BvFLK* ; 5/15	28	16
Col-0; endo::*BvFLK* ; 5/16	28	13
Col-0; endo::*BvFLK* ; 5/17	28	19
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	
Col-0; endo::*BvFLK* ; 11/1	31	23
Col-0; endo::*BvFLK* ; 11/2	31	24
Col-0; endo::*BvFLK* ; 11/3	31	18
Col-0; endo::*BvFLK* ; 11/4	28	15
Col-0; endo::*BvFLK* ; 11/5	28	13

Col-0; endo::*BvFLK* ; 6/6 [b]	**28**	**16**
Col-0; endo::*BvFLK* ; 6/7	**28**	**16**
Col-0; endo::*BvFLK* ; 6/8	**38**	**16**
Col-0; endo::*BvFLK* ; 6/9	**28**	**17**
Col-0; endo::*BvFLK* ; 6/10	**28**	**18**
Col-0; endo::*BvFLK* ; 6/11	**28**	**13**
Col-0; endo::*BvFLK* ; 6/12	**31**	**14**
Col-0; endo::*BvFLK* ; 6/13 [b]	**37**	**16**
Col-0; endo::*BvFLK* ; 6/14 [b]	**38**	**18**
Col-0; endo::*BvFLK* ; 6/15	**28**	**16**
Col-0; endo::*BvFLK* ; 6/16	**31**	**17**
Col-0; endo::*BvFLK* ; 6/17	**31**	**16**
Segregation ratio (early bolting: late bolting)	13:4	
X^2 for 3:1	0.02 (ns)	
X^2 for 15:1	8.56 **	
Col-0; endo::*BvFLK* ; 12/1	31	15
Col-0; endo::*BvFLK* ; 12/2	38	15
Col-0; endo::*BvFLK* ; 12/3	31	18
Col-0; endo::*BvFLK* ; 12/4	35	15
Col-0; endo::*BvFLK* ; 12/5	31	17
Col-0; endo::*BvFLK* ; 12/6	28	15
Col-0; endo::*BvFLK* ; 12/7	31	17
Col-0; endo::*BvFLK* ; 12/8	31	16
Col-0; endo::*BvFLK* ; 12/9	31	18
Col-0; endo::*BvFLK* ; 12/10	28	11
Col-0; endo::*BvFLK* ; 12/11	31	18
Col-0; endo::*BvFLK* ; 12/12	31	11
Col-0; endo::*BvFLK* ; 12/13	31	17
Col-0; endo::*BvFLK* ; 12/14	31	19
Col-0; endo::*BvFLK* ; 12/15	31	17
Col-0; endo::*BvFLK* ; 12/16	31	18
Col-0; endo::*BvFLK* ; 12/17	31	19
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	
Col-0; endo::*BvFLK* ; 15/1	31	18
Col-0; endo::*BvFLK* ; 15/2	28	13
Col-0; endo::*BvFLK* ; 15/3	28	16
Col-0; endo::*BvFLK* ; 15/4	-	-
Col-0; endo::*BvFLK* ; 15/5	25	14
Col-0; endo::*BvFLK* ; 15/6	28	16
Col-0; endo::*BvFLK* ; 15/7	25	16
Col-0; endo::*BvFLK* ; 15/8	28	12
Col-0; endo::*BvFLK* ; 15/9	24	18
Col-0; endo::*BvFLK* ; 15/10	38	18
Col-0; endo::*BvFLK* ; 15/11	31	12
Col-0; endo::*BvFLK* ; 15/12	-	-
Col-0; endo::*BvFLK* ; 15/13	31	14

Col-0; endo::*BvFLK* ; 11/6	-	-
Col-0; endo::*BvFLK* ; 11/7	31	19
Col-0; endo::*BvFLK* ; 11/8	31	19
Col-0; endo::*BvFLK* ; 11/9	28	18
Col-0; endo::*BvFLK* ; 11/10	31	20
Col-0; endo::*BvFLK* ; 11/11	31	18
Col-0; endo::*BvFLK* ; 11/12	31	17
Col-0; endo::*BvFLK* ; 11/13	31	21
Col-0; endo::*BvFLK* ; 11/14	28	15
Col-0; endo::*BvFLK* ; 11/15	28	16
Col-0; endo::*BvFLK* ; 11/16	31	20
Col-0; endo::*BvFLK* ; 11/17	31	19
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
Col-0; endo::*BvFLK* ; 13/1	28	16
Col-0; endo::*BvFLK* ; 13/2	28	18
Col-0; endo::*BvFLK* ; 13/3	28	14
Col-0; endo::*BvFLK* ; 13/4	24	14
Col-0; endo::*BvFLK* ; 13/5	24	12
Col-0; endo::*BvFLK* ; 13/6	28	14
Col-0; endo::*BvFLK* ; 13/7	28	12
Col-0; endo::*BvFLK* ; 13/8	28	16
Col-0; endo::*BvFLK* ; 13/9	28	16
Col-0; endo::*BvFLK* ; 13/10	31	23
Col-0; endo::*BvFLK* ; 13/11	28	16
Col-0; endo::*BvFLK* ; 13/12	25	14
Col-0; endo::*BvFLK* ; 13/13	28	18
Col-0; endo::*BvFLK* ; 13/14	31	20
Col-0; endo::*BvFLK* ; 13/15	31	17
Col-0; endo::*BvFLK* ; 13/16	25	13
Col-0; endo::*BvFLK* ; 13/17	-	-
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
Col-0; endo::*BvFLK* ; 16/1	31	15
Col-0; endo::*BvFLK* ; 16/2	38	15
Col-0; endo::*BvFLK* ; 16/3	31	18
Col-0; endo::*BvFLK* ; 16/4	35	15
Col-0; endo::*BvFLK* ; 16/5	31	17
Col-0; endo::*BvFLK* ; 16/6	28	15
Col-0; endo::*BvFLK* ; 16/7	31	17
Col-0; endo::*BvFLK* ; 16/8	31	16
Col-0; endo::*BvFLK* ; 16/9	31	18
Col-0; endo::*BvFLK* ; 16/10	28	11
Col-0; endo::*BvFLK* ; 16/11	35	18
Col-0; endo::*BvFLK* ; 16/12	31	11
Col-0; endo::*BvFLK* ; 16/13	31	17

Col-0; endo::*BvFLK* ; 15/14	31	14
Col-0; endo::*BvFLK* ; 15/15	28	15
Col-0; endo::*BvFLK* ; 15/16	25	15
Col-0; endo::*BvFLK* ; 15/17	31	13
Segregation ratio (early bolting: late bolting)	15:0	
X^2 for 3:1	5.00 *	
X^2 for 15:1	1.00 (ns)	

Col-0; endo::*BvFLK* ; 16/14	31	19
Col-0; endo::*BvFLK* ; 16/15	31	17
Col-0; endo::*BvFLK* ; 16/16	31	18
Col-0; endo::*BvFLK* ; 16/17	31	19
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
Col-0; 35S::*AtFLK;* 1/1	31	17
Col-0; 35S::*AtFLK;* 1/2	31	18
Col-0; 35S::*AtFLK;* 1/3	31	15
Col-0; 35S::*AtFLK;* 1/4	28	14
Col-0; 35S::*AtFLK;* 1/5	31	17
Col-0; 35S::*AtFLK;* 1/6	31	15
Col-0; 35S::*AtFLK;* 1/7	38	15
Col-0; 35S::*AtFLK;* 1/8	31	15
Col-0; 35S::*AtFLK;* 1/9	35	14
Col-0; 35S::*AtFLK;* 1/10	31	15
Col-0; 35S::*AtFLK;* 1/11	31	15
Col-0; 35S::*AtFLK;* 1/12	28	13
Col-0; 35S::*AtFLK;* 1/13	40	18
Col-0; 35S::*AtFLK;* 1/14	28	14
Col-0; 35S::*AtFLK;* 1/15	31	17
Col-0; 35S::*AtFLK;* 1/16	31	16
Col-0; 35S::*AtFLK;* 1/17	28	14
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	
Col-0; 35S::*AtFLK* ; 3/1	25	11
Col-0; 35S::*AtFLK* ; 3/2	28	13
Col-0; 35S::*AtFLK* ; 3/3	31	18
Col-0; 35S::*AtFLK* ; 3/4	28	15
Col-0; 35S::*AtFLK* ; 3/5	25	13
Col-0; 35S::*AtFLK* ; 3/6	28	12
Col-0; 35S::*AtFLK* ; 3/7	25	13
Col-0; 35S::*AtFLK* ; 3/8	25	15
Col-0; 35S::*AtFLK* ; 3/9	28	14
Col-0; 35S::*AtFLK* ; 3/10	25	12
Col-0; 35S::*AtFLK* ; 3/11	24	12
Col-0; 35S::*AtFLK* ; 3/12	28	13
Col-0; 35S::*AtFLK* ; 3/13	28	12
Col-0; 35S::*AtFLK* ; 3/14	28	11
Col-0; 35S::*AtFLK* ; 3/15	25	13
Col-0; 35S::*AtFLK* ; 3/16	24	13
Col-0; 35S::*AtFLK* ; 3/17	25	13
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
Col-0; 35S::*AtFLK* ; 5/1	**25**	**15**
Col-0; 35S::*AtFLK* ; 5/2	**27**	**14**
Col-0; 35S::*AtFLK* ; 5/3	**27**	**15**
Col-0; 35S::*AtFLK* ; 5/4	**27**	**18**
Col-0; 35S::*AtFLK* ; 5/5	**27**	**13**

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
Col-0; 35S::*AtFLK* ; 2/1	25	12
Col-0; 35S::*AtFLK* ; 2/2	25	13
Col-0; 35S::*AtFLK* ; 2/3	25	15
Col-0; 35S::*AtFLK* ; 2/4	25	16
Col-0; 35S::*AtFLK* ; 2/5	25	13
Col-0; 35S::*AtFLK* ; 2/6	25	13
Col-0; 35S::*AtFLK* ; 2/7	25	15
Col-0; 35S::*AtFLK* ; 2/8	25	17
Col-0; 35S::*AtFLK* ; 2/9	25	13
Col-0; 35S::*AtFLK* ; 2/10	28	15
Col-0; 35S::*AtFLK* ; 2/11	28	12
Col-0; 35S::*AtFLK* ; 2/12	28	12
Col-0; 35S::*AtFLK* ; 2/13	28	15
Col-0; 35S::*AtFLK* ; 2/14	24	12
Col-0; 35S::*AtFLK* ; 2/15	28	15
Col-0; 35S::*AtFLK* ; 2/16	28	16
Col-0; 35S::*AtFLK* ; 2/17	28	14
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
Col-0; 35S::*AtFLK* ; 4/1	25	11
Col-0; 35S::*AtFLK* ; 4/2	28	15
Col-0; 35S::*AtFLK* ; 4/3	31	14
Col-0; 35S::*AtFLK* ; 4/4	28	16
Col-0; 35S::*AtFLK* ; 4/5	31	15
Col-0; 35S::*AtFLK* ; 4/6	28	14
Col-0; 35S::*AtFLK* ; 4/7	28	12
Col-0; 35S::*AtFLK* ; 4/8	35	15
Col-0; 35S::*AtFLK* ; 4/9	28	13
Col-0; 35S::*AtFLK* ; 4/10	31	14
Col-0; 35S::*AtFLK* ; 4/11	28	15
Col-0; 35S::*AtFLK* ; 4/12	31	16
Col-0; 35S::*AtFLK* ; 4/13	31	18
Col-0; 35S::*AtFLK* ; 4/14	38	17
Col-0; 35S::*AtFLK* ; 4/15	28	14
Col-0; 35S::*AtFLK* ; 4/16	28	12
Col-0; 35S::*AtFLK* ; 4/17	28	14
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	
Col-0; 35S::*AtFLK* ; 6/1	31	13
Col-0; 35S::*AtFLK* ; 6/2	31	15
Col-0; 35S::*AtFLK* ; 6/3	31	15
Col-0; 35S::*AtFLK* ; 6/4	28	15
Col-0; 35S::*AtFLK* ; 6/5	31	15

Col-0; 35S::*AtFLK* ; 5/6	**25**	**13**
Col-0; 35S::*AtFLK* ; 5/7	**25**	**15**
Col-0; 35S::*AtFLK* ; 5/8	**28**	**14**
Col-0; 35S::*AtFLK* ; 5/9	**28**	**14**
Col-0; 35S::*AtFLK* ; 5/10 [b]	**38**	**18**
Col-0; 35S::*AtFLK* ; 5/11	**31**	**14**
Col-0; 35S::*AtFLK* ; 5/12 [b]	**39**	**18**
Col-0; 35S::*AtFLK* ; 5/13 [b]	**38**	**19**
Col-0; 35S::*AtFLK* ; 5/14 [b]	**40**	**21**
Col-0; 35S::*AtFLK* ; 5/15	**28**	**16**
Col-0; 35S::*AtFLK* ; 5/16	**25**	**14**
Col-0; 35S::*AtFLK* ; 5/17	**31**	**20**
Segregation ratio (early bolting: late bolting)	13:4	
X^2 for 3:1	0.02 (ns)	
X^2 for 15:1	8.56 **	
Col-0; 35S::*AtFLK* ; 7/1	31	18
Col-0; 35S::*AtFLK* ; 7/2	28	14
Col-0; 35S::*AtFLK* ; 7/3	28	16
Col-0; 35S::*AtFLK* ; 7/4	28	14
Col-0; 35S::*AtFLK* ; 7/5	35	13
Col-0; 35S::*AtFLK* ; 7/6	31	14
Col-0; 35S::*AtFLK* ; 7/7	28	13
Col-0; 35S::*AtFLK* ; 7/8	28	11
Col-0; 35S::*AtFLK* ; 7/9	25	11
Col-0; 35S::*AtFLK* ; 7/10	28	13
Col-0; 35S::*AtFLK* ; 7/11	31	10
Col-0; 35S::*AtFLK* ; 7/12	-	-
Col-0; 35S::*AtFLK* ; 7/13	28	15
Col-0; 35S::*AtFLK* ; 7/14	31	18
Col-0; 35S::*AtFLK* ; 7/15	28	15
Col-0; 35S::*AtFLK* ; 7/16	28	13
Col-0; 35S::*AtFLK* ; 7/17	31	17
Segregation ratio (early bolting: late bolting)	15:1	
X^2 for 3:1	3.00 (ns)	
X^2 for 15:1	0.00 (ns)	
Col-0; 35S::*AtFLK* ; 9/1	28	17
Col-0; 35S::*AtFLK* ; 9/2	28	19
Col-0; 35S::*AtFLK* ; 9/3	31	17
Col-0; 35S::*AtFLK* ; 9/4	25	15
Col-0; 35S::*AtFLK* ; 9/5	28	17
Col-0; 35S::*AtFLK* ; 9/6	25	15
Col-0; 35S::*AtFLK* ; 9/7	28	15
Col-0; 35S::*AtFLK* ; 9/8	25	17
Col-0; 35S::*AtFLK* ; 9/9	31	17
Col-0; 35S::*AtFLK* ; 9/10	28	15
Col-0; 35S::*AtFLK* ; 9/11	35	16
Col-0; 35S::*AtFLK* ; 9/12	35	15
Col-0; 35S::*AtFLK* ; 9/13	28	20

Col-0; 35S::*AtFLK* ; 6/6	38	15
Col-0; 35S::*AtFLK* ; 6/7	28	16
Col-0; 35S::*AtFLK* ; 6/8	25	12
Col-0; 35S::*AtFLK* ; 6/9	28	14
Col-0; 35S::*AtFLK* ; 6/10	39	16
Col-0; 35S::*AtFLK* ; 6/11	31	15
Col-0; 35S::*AtFLK* ; 6/12	28	13
Col-0; 35S::*AtFLK* ; 6/13	28	16
Col-0; 35S::*AtFLK* ; 6/14	31	16
Col-0; 35S::*AtFLK* ; 6/15	28	14
Col-0; 35S::*AtFLK* ; 6/16	31	17
Col-0; 35S::*AtFLK* ; 6/17	28	18
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	
Col-0; 35S::*AtFLK;* 8/1	31	18
Col-0; 35S::*AtFLK;* 8/2	31	17
Col-0; 35S::*AtFLK;* 8/3	28	16
Col-0; 35S::*AtFLK;* 8/4	28	14
Col-0; 35S::*AtFLK;* 8/5	25	12
Col-0; 35S::*AtFLK;* 8/6	27	16
Col-0; 35S::*AtFLK;* 8/7	27	17
Col-0; 35S::*AtFLK;* 8/8	25	12
Col-0; 35S::*AtFLK;* 8/9	28	18
Col-0; 35S::*AtFLK;* 8/10	28	16
Col-0; 35S::*AtFLK;* 8/11	28	15
Col-0; 35S::*AtFLK;* 8/12	29	13
Col-0; 35S::*AtFLK;* 8/13	28	13
Col-0; 35S::*AtFLK;* 8/14	25	14
Col-0; 35S::*AtFLK;* 8/15	31	17
Col-0; 35S::*AtFLK;* 8/16	31	16
Col-0; 35S::*AtFLK;* 8/17	28	15
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	

Col-0; 35S::*AtFLK* ; 9/14	25	15
Col-0; 35S::*AtFLK* ; 9/15	25	15
Col-0; 35S::*AtFLK* ; 9/16	28	16
Col-0; 35S::*AtFLK* ; 9/17	28	21
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
flk-1; 35S::BvFLK; 1/1	28	13
flk-1; 35S::BvFLK; 1/2	28	12
flk-1; 35S::BvFLK; 1/3	28	16
flk-1; 35S::BvFLK; 1/4	28	17
flk-1; 35S::BvFLK; 1/5	28	14
flk-1; 35S::BvFLK; 1/6	28	15
flk-1; 35S::BvFLK; 1/7	28	20
flk-1; 35S::BvFLK; 1/8	25	15
flk-1; 35S::BvFLK; 1/9	28	15
flk-1; 35S::BvFLK; 1/10	28	15
flk-1; 35S::BvFLK; 1/11	38	15
flk-1; 35S::BvFLK; 1/12	25	14
flk-1; 35S::BvFLK; 1/13	28	15
flk-1; 35S::BvFLK; 1/14	25	14
flk-1; 35S::BvFLK; 1/15	28	17
flk-1; 35S::BvFLK; 1/16	28	14
flk-1; 35S::BvFLK; 1/17	28	18
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; 35S::BvFLK; 5/1	28	19
flk-1; 35S::BvFLK; 5/2	28	13
flk-1; 35S::BvFLK; 5/3	28	14
flk-1; 35S::BvFLK; 5/4	28	17
flk-1; 35S::BvFLK; 5/5	28	17
flk-1; 35S::BvFLK; 5/6	28	15
flk-1; 35S::BvFLK; 5/7	28	18
flk-1; 35S::BvFLK; 5/8	28	17
flk-1; 35S::BvFLK; 5/9	31	16
flk-1; 35S::BvFLK; 5/10	28	17
flk-1; 35S::BvFLK; 5/11	35	13
flk-1; 35S::BvFLK; 5/12	25	18
flk-1; 35S::BvFLK; 5/13	28	16
flk-1; 35S::BvFLK; 5/14	31	15
flk-1; 35S::BvFLK; 5/15	25	20
flk-1; 35S::BvFLK; 5/16	25	18
flk-1; 35S::BvFLK; 5/17	28	19
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; 35S::BvFLK; 9/1	31	15
flk-1; 35S::BvFLK; 9/2	28	16
flk-1; 35S::BvFLK; 9/3	28	16
flk-1; 35S::BvFLK; 9/4	28	12
flk-1; 35S::BvFLK; 9/5	24	13

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
flk-1; 35S::BvFLK; 2/1	31	18
flk-1; 35S::BvFLK; 2/2	31	17
flk-1; 35S::BvFLK; 2/3	28	18
flk-1; 35S::BvFLK; 2/4	-	-
flk-1; 35S::BvFLK; 2/5	28	20
flk-1; 35S::BvFLK; 2/6	25	16
flk-1; 35S::BvFLK; 2/7	28	17
flk-1; 35S::BvFLK; 2/8	25	21
flk-1; 35S::BvFLK; 2/9	25	18
flk-1; 35S::BvFLK; 2/10	28	15
flk-1; 35S::BvFLK; 2/11	31	14
flk-1; 35S::BvFLK; 2/12	31	16
flk-1; 35S::BvFLK; 2/13	28	14
flk-1; 35S::BvFLK; 2/14	25	13
flk-1; 35S::BvFLK; 2/15	31	17
flk-1; 35S::BvFLK; 2/16	28	15
flk-1; 35S::BvFLK; 2/17	28	16
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
flk-1; 35S::BvFLK; 7/1	35	11
flk-1; 35S::BvFLK; 7/2	35	16
flk-1; 35S::BvFLK; 7/3	35	14
flk-1; 35S::BvFLK; 7/4	35	14
flk-1; 35S::BvFLK; 7/5	31	11
flk-1; 35S::BvFLK; 7/6	28	17
flk-1; 35S::BvFLK; 7/7	31	17
flk-1; 35S::BvFLK; 7/8	31	18
flk-1; 35S::BvFLK; 7/9	28	15
flk-1; 35S::BvFLK; 7/10	28	16
flk-1; 35S::BvFLK; 7/11	38	16
flk-1; 35S::BvFLK; 7/12	31	19
flk-1; 35S::BvFLK; 7/13	31	17
flk-1; 35S::BvFLK; 7/14	31	20
flk-1; 35S::BvFLK; 7/15	28	14
flk-1; 35S::BvFLK; 7/16	28	17
flk-1; 35S::BvFLK; 7/17	28	18
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; 35S::BvFLK; 10/1	28	16
flk-1; 35S::BvFLK; 10/2	28	17
flk-1; 35S::BvFLK; 10/3	28	15
flk-1; 35S::BvFLK; 10/4	28	16
flk-1; 35S::BvFLK; 10/5	28	17

flk-1; 35S::BvFLK; 9/6	28	18
flk-1; 35S::BvFLK; 9/7	28	18
flk-1; 35S::BvFLK; 9/8	24	116
flk-1; 35S::BvFLK; 9/9	24	16
flk-1; 35S::BvFLK; 9/10	28	17
flk-1; 35S::BvFLK; 9/11	28	17
flk-1; 35S::BvFLK; 9/12	25	16
flk-1; 35S::BvFLK; 9/13	28	19
flk-1; 35S::BvFLK; 9/14	24	17
flk-1; 35S::BvFLK; 9/15	28	15
flk-1; 35S::BvFLK; 9/16	28	18
flk-1; 35S::BvFLK; 9/17	28	16
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; 35S::BvFLK; 11/1	25	17
flk-1; 35S::BvFLK; 11/2	25	17
flk-1; 35S::BvFLK; 11/3	28	17
flk-1; 35S::BvFLK; 11/4	28	15
flk-1; 35S::BvFLK; 11/5	28	18
flk-1; 35S::BvFLK; 11/6	25	13
flk-1; 35S::BvFLK; 11/7	28	17
flk-1; 35S::BvFLK; 11/8	28	13
flk-1; 35S::BvFLK; 11/9	28	17
flk-1; 35S::BvFLK; 11/10	35	20
flk-1; 35S::BvFLK; 11/11	28	17
flk-1; 35S::BvFLK; 11/12	25	17
flk-1; 35S::BvFLK; 11/13	28	18
flk-1; 35S::BvFLK; 11/14	28	19
flk-1; 35S::BvFLK; 11/15	225	17
flk-1; 35S::BvFLK; 11/16	35	18
flk-1; 35S::BvFLK; 11/17	28	17
Segregation ratio (early bolting: late bolting)	17: 0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; 35S::BvFLK; 14/1	31	19
flk-1; 35S::BvFLK; 14/2	28	17
flk-1; 35S::BvFLK; 14/3	31	17
flk-1; 35S::BvFLK; 14/4	31	19
flk-1; 35S::BvFLK; 14/5	28	16
flk-1; 35S::BvFLK; 14/6	28	17
flk-1; 35S::BvFLK; 14/7	35	18
flk-1; 35S::BvFLK; 14/8	31	17
flk-1; 35S::BvFLK; 14/9	35	16
flk-1; 35S::BvFLK; 14/10	28	18
flk-1; 35S::BvFLK; 14/11	28	17
flk-1; 35S::BvFLK; 14/12	35	17
flk-1; 35S::BvFLK; 14/13	35	17

flk-1; 35S::BvFLK; 10/6	28	15
flk-1; 35S::BvFLK; 10/7	28	20
flk-1; 35S::BvFLK; 10/8	28	16
flk-1; 35S::BvFLK; 10/9	28	15
flk-1; 35S::BvFLK; 10/10	25	16
flk-1; 35S::BvFLK; 10/11	25	17
flk-1; 35S::BvFLK; 10/12	28	14
flk-1; 35S::BvFLK; 10/13	28	18
flk-1; 35S::BvFLK; 10/14	25	16
flk-1; 35S::BvFLK; 10/15	28	21
flk-1; 35S::BvFLK; 10/16	28	18
flk-1; 35S::BvFLK; 10/17	25	17
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; 35S::BvFLK; 12/1	28	16
flk-1; 35S::BvFLK; 12/2	28	18
flk-1; 35S::BvFLK; 12/3	31	15
flk-1; 35S::BvFLK; 12/4	28	14
flk-1; 35S::BvFLK; 12/5	28	18
flk-1; 35S::BvFLK; 12/6	28	13
flk-1; 35S::BvFLK; 12/7	35	18
flk-1; 35S::BvFLK; 12/8	28	16
flk-1; 35S::BvFLK; 12/9	28	19
flk-1; 35S::BvFLK; 12/10	28	17
flk-1; 35S::BvFLK; 12/11	28	17
flk-1; 35S::BvFLK; 12/12	28	15
flk-1; 35S::BvFLK; 12/13	28	16
flk-1; 35S::BvFLK; 12/14	31	12
flk-1; 35S::BvFLK; 12/15	28	16
flk-1; 35S::BvFLK; 12/16	38	15
flk-1; 35S::BvFLK; 12/17	28	16
Segregation ratio (early bolting: late bolting)	17: 0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; 35S::BvFLK; 18/1	**28**	**18**
flk-1; 35S::BvFLK; 18/2	**35**	**17**
flk-1; 35S::BvFLK; 18/3	**35**	**17**
flk-1; 35S::BvFLK; 18/4	**35**	**16**
flk-1; 35S::BvFLK; 18/5	**35**	**15**
flk-1; 35S::BvFLK; 18/6	**35**	**16**
flk-1; 35S::BvFLK; 18/7	**38**	**15**
flk-1; 35S::BvFLK; 18/8	**38**	**16**
flk-1; 35S::BvFLK; 18/9	**28**	**17**
flk-1; 35S::BvFLK; 18/10	**35**	**18**
flk-1; 35S::BvFLK; 18/11 [b]	**69**	**71**
flk-1; 35S::BvFLK; 18/12	**31**	**12**
flk-1; 35S::BvFLK; 18/13	**35**	**18**

flk-1; 35S::BvFLK; 14/14	35	15
flk-1; 35S::BvFLK; 14/15	28	17
flk-1; 35S::BvFLK; 14/16	31	15
flk-1; 35S::BvFLK; 14/17	31	14
Segregation ratio (early bolting: late bolting)	17: 0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	

flk-1; 35S::BvFLK; 18/14	**35**	**15**
flk-1; 35S::BvFLK; 18/15	**35**	**18**
flk-1; 35S::BvFLK; 18/16 [b]	**66**	**65**
flk-1; 35S::BvFLK; 18/17	**-**	**-**
Segregation ratio (early bolting: late bolting)	14:2	
X^2 for 3:1	1.33 (ns)	
X^2 for 15:1	1.07 (ns)	

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
flk-1; endo::BvFLK; 1/1	28	23
flk-1; endo::BvFLK; 1/2	28	13
flk-1; endo::BvFLK; 1/3	35	18
flk-1; endo::BvFLK; 1/4	28	18
flk-1; endo::BvFLK; 1/5	31	13
flk-1; endo::BvFLK; 1/6	31	15
flk-1; endo::BvFLK; 1/7	31	13
flk-1; endo::BvFLK; 1/8	28	19
flk-1; endo::BvFLK; 1/9	28	15
flk-1; endo::BvFLK; 1/10	31	14
flk-1; endo::BvFLK; 1/11	31	15
flk-1; endo::BvFLK; 1/12	31	15
flk-1; endo::BvFLK; 1/13	35	17
flk-1; endo::BvFLK; 1/14	28	16
flk-1; endo::BvFLK; 1/15	28	15
flk-1; endo::BvFLK; 1/16	28	17
flk-1; endo::BvFLK; 1/17	35	17
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; endo::BvFLK; 3/1	28	19
flk-1; endo::BvFLK; 3/2	35	16
flk-1; endo::BvFLK; 3/3	31	14
flk-1; endo::BvFLK; 3/4	35	14
flk-1; endo::BvFLK; 3/5	35	15
flk-1; endo::BvFLK; 3/6	35	16
flk-1; endo::BvFLK; 3/7	31	14
flk-1; endo::BvFLK; 3/8	38	16
flk-1; endo::BvFLK; 3/9	31	16
flk-1; endo::BvFLK; 3/10	28	14
flk-1; endo::BvFLK; 3/11	28	12
flk-1; endo::BvFLK; 3/12	31	17
flk-1; endo::BvFLK; 3/13	-	-
flk-1; endo::BvFLK; 3/14	28	18
flk-1; endo::BvFLK; 3/15	28	16
flk-1; endo::BvFLK; 3/16	28	13
flk-1; endo::BvFLK; 3/17	31	13
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
flk-1; endo::BvFLK; 5/1	**35**	**13**
flk-1; endo::BvFLK; 5/2	**35**	**15**
flk-1; endo::BvFLK; 5/3	**35**	**15**
flk-1; endo::BvFLK; 5/4	**31**	**17**
flk-1; endo::BvFLK; 5/5	**28**	**15**

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
flk-1; endo::BvFLK; 2/1	31	17
flk-1; endo::BvFLK; 2/2	31	15
flk-1; endo::BvFLK; 2/3	-	-
flk-1; endo::BvFLK; 2/4	31	14
flk-1; endo::BvFLK; 2/5	35	13
flk-1; endo::BvFLK; 2/6	31	17
flk-1; endo::BvFLK; 2/7	35	16
flk-1; endo::BvFLK; 2/8	31	15
flk-1; endo::BvFLK; 2/9	35	28
flk-1; endo::BvFLK; 2/10	31	14
flk-1; endo::BvFLK; 2/11	38	12
flk-1; endo::BvFLK; 2/12	35	16
flk-1; endo::BvFLK; 2/13	31	15
flk-1; endo::BvFLK; 2/14	31	16
flk-1; endo::BvFLK; 2/15	28	15
flk-1; endo::BvFLK; 2/16	31	15
flk-1; endo::BvFLK; 2/17	31	14
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
flk-1; endo::BvFLK; 4/1	28	18
flk-1; endo::BvFLK; 4/2	28	15
flk-1; endo::BvFLK; 4/3	28	13
flk-1; endo::BvFLK; 4/4	28	16
flk-1; endo::BvFLK; 4/5	31	17
flk-1; endo::BvFLK; 4/6	31	19
flk-1; endo::BvFLK; 4/7	28	15
flk-1; endo::BvFLK; 4/8	28	15
flk-1; endo::BvFLK; 4/9	25	14
flk-1; endo::BvFLK; 4/10	28	16
flk-1; endo::BvFLK; 4/11	25	18
flk-1; endo::BvFLK; 4/12	28	17
flk-1; endo::BvFLK; 4/13	38	15
flk-1; endo::BvFLK; 4/14	28	15
flk-1; endo::BvFLK; 4/15	28	16
flk-1; endo::BvFLK; 4/16	28	14
flk-1; endo::BvFLK; 4/17	28	17
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; endo::BvFLK; 6/1	28	14
flk-1; endo::BvFLK; 6/2	39	15
flk-1; endo::BvFLK; 6/3	38	15
flk-1; endo::BvFLK; 6/4	38	16
flk-1; endo::BvFLK; 6/5	28	16

flk-1; endo::BvFLK; 5/6	**38**	**14**
flk-1; endo::BvFLK; 5/7	**35**	**15**
flk-1; endo::BvFLK; 5/8 [b]	**71**	**64**
flk-1; endo::BvFLK; 5/9	**35**	**21**
flk-1; endo::BvFLK; 5/10	**28**	**13**
flk-1; endo::BvFLK; 5/11 [b]	**67**	**69**
flk-1; endo::BvFLK; 5/12	**38**	**18**
flk-1; endo::BvFLK; 5/13 [b]	**69**	**68**
flk-1; endo::BvFLK; 5/14	**28**	**14**
flk-1; endo::BvFLK; 5/15 [b]	**67**	**66**
flk-1; endo::BvFLK; 5/16	**31**	**19**
flk-1; endo::BvFLK; 5/17	**31**	**17**
Segregation ratio (early bolting: late bolting)	13:4	
X^2 for 3:1	0.02 (ns)	
X^2 for 15:1	8.56 **	
flk-1; endo::BvFLK; 7/1	31	14
flk-1; endo::BvFLK; 7/2	28	17
flk-1; endo::BvFLK; 7/3	28	20
flk-1; endo::BvFLK; 7/4	28	20
flk-1; endo::BvFLK; 7/5	28	18
flk-1; endo::BvFLK; 7/6	28	17
flk-1; endo::BvFLK; 7/7	28	16
flk-1; endo::BvFLK; 7/8	28	21
flk-1; endo::BvFLK; 7/9	28	16
flk-1; endo::BvFLK; 7/10	28	16
flk-1; endo::BvFLK; 7/11	28	17
flk-1; endo::BvFLK; 7/12	28	15
flk-1; endo::BvFLK; 7/13	28	15
flk-1; endo::BvFLK; 7/14	28	18
flk-1; endo::BvFLK; 7/15	28	16
flk-1; endo::BvFLK; 7/16	31	19
flk-1; endo::BvFLK; 7/17	35	20
Segregation ratio (early bolting: late bolting)	17: 0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; endo::BvFLK; 9/1	35	17
flk-1; endo::BvFLK; 9/2	28	15
flk-1; endo::BvFLK; 9/3	38	16
flk-1; endo::BvFLK; 9/4	31	16
flk-1; endo::BvFLK; 9/5	38	17
flk-1; endo::BvFLK; 9/6	28	15
flk-1; endo::BvFLK; 9/7	38	13
flk-1; endo::BvFLK; 9/8	31	19
flk-1; endo::BvFLK; 9/9	28	11
flk-1; endo::BvFLK; 9/10	28	10
flk-1; endo::BvFLK; 9/11	28	12
flk-1; endo::BvFLK; 9/12	28	12
flk-1; endo::BvFLK; 9/13	28	16

flk-1; endo::BvFLK; 6/6	38	11
flk-1; endo::BvFLK; 6/7	31	15
flk-1; endo::BvFLK; 6/8	38	15
flk-1; endo::BvFLK; 6/9	39	15
flk-1; endo::BvFLK; 6/10	38	14
flk-1; endo::BvFLK; 6/11	35	16
flk-1; endo::BvFLK; 6/12	31	11
flk-1; endo::BvFLK; 6/13	31	15
flk-1; endo::BvFLK; 6/14	35	15
flk-1; endo::BvFLK; 6/15	38	17
flk-1; endo::BvFLK; 6/16	31	15
flk-1; endo::BvFLK; 6/17	31	16
Segregation ratio (early bolting: late bolting)	17:0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	
flk-1; endo::BvFLK; 8/1	38	18
flk-1; endo::BvFLK; 8/2	28	14
flk-1; endo::BvFLK; 8/3	31	15
flk-1; endo::BvFLK; 8/4	31	17
flk-1; endo::BvFLK; 8/5	31	15
flk-1; endo::BvFLK; 8/6	31	16
flk-1; endo::BvFLK; 8/7	28	15
flk-1; endo::BvFLK; 8/8	31	19
flk-1; endo::BvFLK; 8/9	31	15
flk-1; endo::BvFLK; 8/10	38	18
flk-1; endo::BvFLK; 8/11	31	17
flk-1; endo::BvFLK; 8/12	28	11
flk-1; endo::BvFLK; 8/13	38	14
flk-1; endo::BvFLK; 8/14	31	16
flk-1; endo::BvFLK; 8/15	28	13
flk-1; endo::BvFLK; 8/16	28	14
flk-1; endo::BvFLK; 8/17	28	16
Segregation ratio (early bolting: late bolting)	17: 0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	

flk-1; endo::BvFLK; 9/14	28	19
flk-1; endo::BvFLK; 9/15	28	13
flk-1; endo::BvFLK; 9/16	31	14
flk-1; endo::BvFLK; 9/17	28	20
Segregation ratio (early bolting: late bolting)	17: 0	
X^2 for 3:1	6.08 **	
X^2 for 15:1	1.14 (ns)	

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
flk-1; 35S::AtFLK; 1/1	28	18
flk-1; 35S::AtFLK; 1/2	28	15
flk-1; 35S::AtFLK; 1/3	28	16
flk-1; 35S::AtFLK; 1/4	28	16
flk-1; 35S::AtFLK; 1/5	28	12
flk-1; 35S::AtFLK; 1/6	28	12
flk-1; 35S::AtFLK; 1/7	28	13
flk-1; 35S::AtFLK; 1/8	35	15
flk-1; 35S::AtFLK; 1/9	-	-
flk-1; 35S::AtFLK; 1/10	28	10
flk-1; 35S::AtFLK; 1/11	31	17
flk-1; 35S::AtFLK; 1/12	28	15
flk-1; 35S::AtFLK; 1/13	28	14
flk-1; 35S::AtFLK; 1/14	31	17
flk-1; 35S::AtFLK; 1/15	31	19
flk-1; 35S::AtFLK; 1/16	35	18
flk-1; 35S::AtFLK; 1/17	31	16
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
flk-1; 35S::AtFLK; 3/1	**28**	**16**
flk-1; 35S::AtFLK; 3/2	**38**	**18**
flk-1; 35S::AtFLK; 3/3	**35**	**18**
flk-1; 35S::AtFLK; 3/4	**38**	**17**
flk-1; 35S::AtFLK; 3/5	**35**	**21**
flk-1; 35S::AtFLK; 3/6	**25**	**15**
flk-1; 35S::AtFLK; 3/7	**44**	**22**
flk-1; 35S::AtFLK; 3/8	**44**	**18**
flk-1; 35S::AtFLK; 3/9	**31**	**15**
flk-1; 35S::AtFLK; 3/10	**31**	**19**
flk-1; 35S::AtFLK; 3/11	**28**	**17**
flk-1; 35S::AtFLK; 3/12	**35**	**16**
flk-1; 35S::AtFLK; 3/13	**31**	**16**
flk-1; 35S::AtFLK; 3/14	**31**	**15**
flk-1; 35S::AtFLK; 3/15	**38**	**14**
flk-1; 35S::AtFLK; 3/16 [b]	**71**	**73**
flk-1; 35S::AtFLK; 3/17 [b]	**68**	**64**
Segregation ratio (early bolting: late bolting)	15:2	
X^2 for 3:1	1.59 (ns)	
X^2 for 15:1	0.86 (ns)	

T2 plants (genetic background; transgene cassette; family number / plant number)	Number of days to bolting	Total number of leaves at bolting
flk-1; 35S::AtFLK; 2/1	35	15
flk-1; 35S::AtFLK; 2/2	38	13
flk-1; 35S::AtFLK; 2/3	28	14
flk-1; 35S::AtFLK; 2/4	35	18
flk-1; 35S::AtFLK; 2/5	38	17
flk-1; 35S::AtFLK; 2/6	44	18
flk-1; 35S::AtFLK; 2/7	31	15
flk-1; 35S::AtFLK; 2/8	31	13
flk-1; 35S::AtFLK; 2/9	31	19
flk-1; 35S::AtFLK; 2/10	28	13
flk-1; 35S::AtFLK; 2/11	44	19
flk-1; 35S::AtFLK; 2/12	31	14
flk-1; 35S::AtFLK; 2/13	28	14
flk-1; 35S::AtFLK; 2/14	35	18
flk-1; 35S::AtFLK; 2/15	38	17
flk-1; 35S::AtFLK; 2/16	38	16
flk-1; 35S::AtFLK; 2/17		
Segregation ratio (early bolting: late bolting)	16:0	
X^2 for 3:1	5.33 **	
X^2 for 15:1	1.07 (ns)	
flk-1; 35S::AtFLK; 4/1	31	16
flk-1; 35S::AtFLK; 4/2	-	-
flk-1; 35S::AtFLK; 4/3	28	12
flk-1; 35S::AtFLK; 4/4	25	17
flk-1; 35S::AtFLK; 4/5	35	19
flk-1; 35S::AtFLK; 4/6	31	15
flk-1; 35S::AtFLK; 4/7	28	13
flk-1; 35S::AtFLK; 4/8	31	17
flk-1; 35S::AtFLK; 4/9	28	10
flk-1; 35S::AtFLK; 4/10	28	15
flk-1; 35S::AtFLK; 4/11	35	15
flk-1; 35S::AtFLK; 4/12	35	15
flk-1; 35S::AtFLK; 4/13	35	16
flk-1; 35S::AtFLK; 4/14	35	14
flk-1; 35S::AtFLK; 4/15	35	12
flk-1; 35S::AtFLK; 4/16	-	-
flk-1; 35S::AtFLK; 4/17	35	13
Segregation ratio (early bolting: late bolting)	15:0	
X^2 for 3:1	5.00 *	
X^2 for 15:1	1.00 (ns)	

Genotype / plant number	Number of days to bolting	Total number of leaves at bolting
Col-0/1	37	18
Col-0/2	40	17
Col-0/3	39	17
Col-0/4	38	16
Col-0/5	38	19
Col-0/6	39	18
Col-0/7	39	17
Col-0/8	40	16
Col-0/9	38	16
Col-0/10	35	17
Col-0/11	36	19
Col-0/12	40	19
Col-0/13	36	18
Col-0/14	37	18
Col-0/15	38	18
Col-0/16	37	18
Col-0/17	37	17

Genotype / plant number	Number of days to bolting	Total number of leaves at bolting
flk-1/1	74	67
flk-1/2	74	67
flk-1/3	74	68
flk-1/4	77	67
flk-1/5	67	72
flk-1/6	69	69
flk-1/7	66	71
flk-1/8	77	67
flk-1/9	71	69
flk-1/10	69	71
flk-1/11	72	67
flk-1/12	65	65
flk-1/13	74	65
flk-1/14	65	67
flk-1/15	65	69
flk-1/16	65	67
flk-1/17	63	65

[a] Non-significant (α>0.05)

[b] Non-transgenic T2 plants as tested by PCR

[c] '-' indicates plants which died before bolting

* α=0.05; ** α=0.01

Supplementary Table 3. Unpaired t-test for number of days to bolting (DTB) and total number of leaves at bolting (TNL) between *BvFLK* or *FLK* transformants and non-transgenic *A. thaliana* plants within six T2 populations.

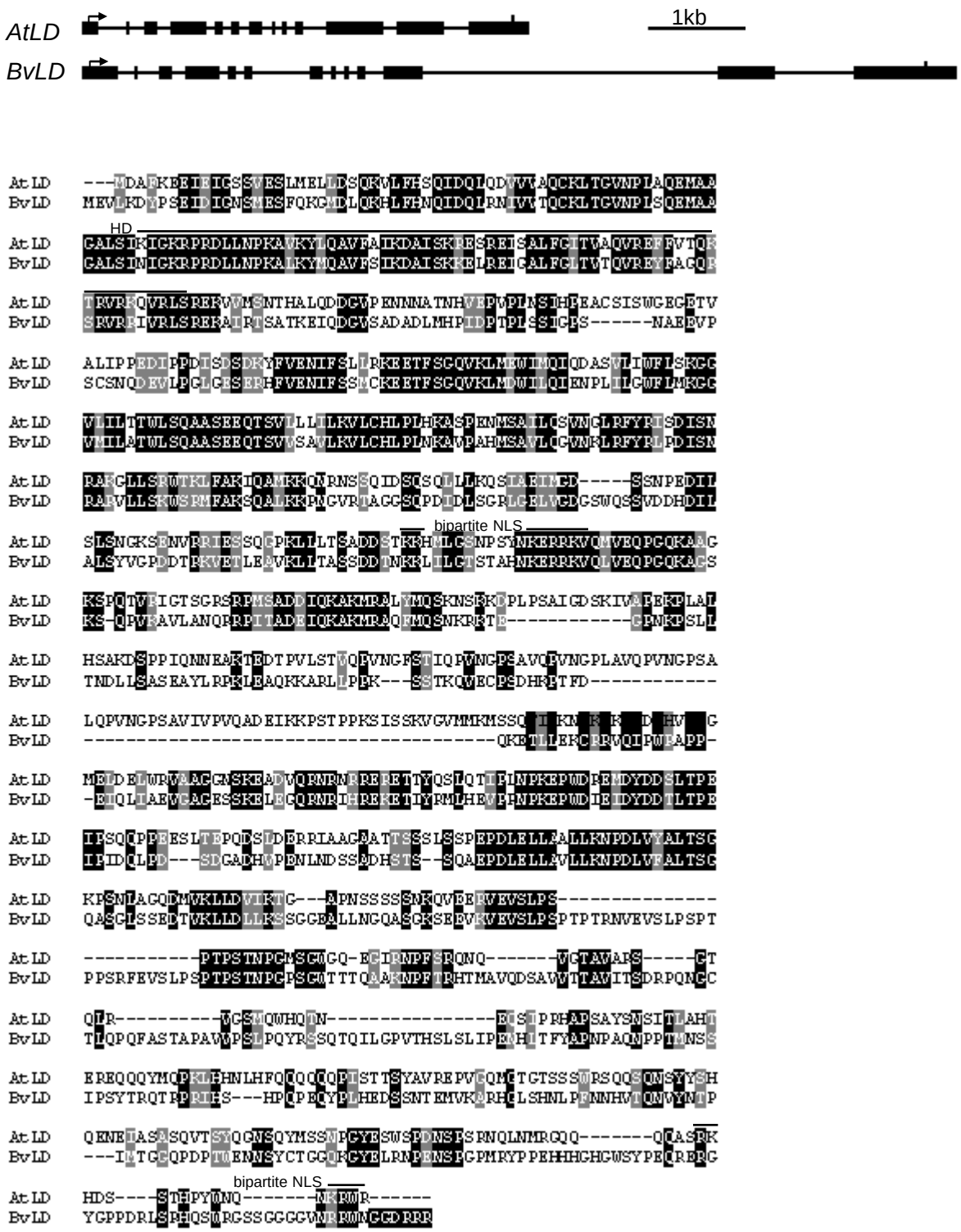
T2 family (genetic background; transgene cassette; family number)	DTB (mean ± standard deviation)		T-test value for DTB (probability)	TNL (mean ± standard deviation)		T-test value for TNL (probability)
	Transgenic plants [a]	Non-transgenic plants		Transgenic plants	Non-transgenic plants	
Col-0; 35S::*BvFLK*; #7	27.77 ±2.89 (n=13[b])	38.00 ±2.16 (n=4[b])	6.48 (0.00)	15.23 ±1.79	18.50 ±0.58	3.53 (0.00)
Col-0; endo::*BvFLK*; #6	28.46 ±1.66 (n=13[b])	38.00 ±0.82 (n=4[b])	10.89 (0.00)	15.31 ±1.70	17.00 ±1.15	1.84 (0.09)
Col-0; 35S::*AtFLK*; #5	27.23 ±2.05 (n=13[b])	38.75 ±0.96 (n=4[b])	10.71 (0.00)	15.00 ±2.00	19.00 ±1.41	3.69 (0.00)
flk-1; 35S::*BvFLK*; #18	34.14 ±3.06 (n=14[b])	67.50 ±2.12 (n=2[b])	14.70 (0.00)	16.29 ±1.68	68.00 ±4.24	34.56 (0.00)
flk-1; endo::*BvFLK*; #5	32.92 ±3.62 (n=13[b])	68.50 ±1.91 (n=4[b])	18.60 (0.00)	15.85 ±2.41	66.75 ±2.22	37.52 (0.00)
flk-1; 35S::*AtFLK*; #3	34.13 ±5.60 (n=15[b])	69.50 ±2.12 (n=2[b])	8.63 (0.00)	17.13 ±2.26	68.50 ±6.36	24.94 (0.00)

[a] Transgenic plants were identified by PCR analysis as described in Materials and Methods.
[b] Number of, respectively, transgenic or non-transgenic plants within a given T2 population.

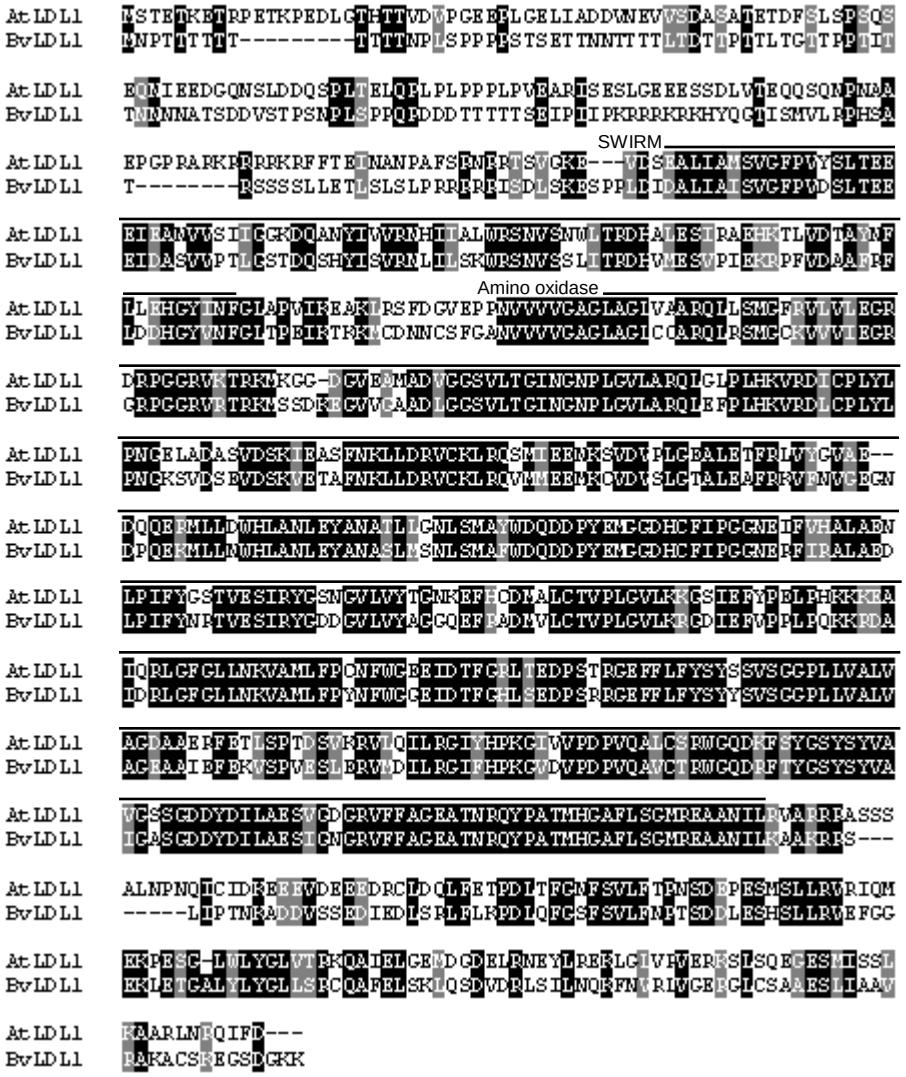
Supplementary Table 4 Unpaired t-test for number of days to bolting (DTB) and total number of leaves at bolting (TNL) between *BvFVE1* transformants and non-transgenic *A. thaliana* plants within four T2 populations.

T2 family (genetic background; transgene cassette; family number)	DTB (mean ± standard deviation)		T-test value for DTB (probability)	TNL (mean ± standard deviation)		T-test value for TNL (probability)
	Transgenic plants [a]	Non-transgenic plants		Transgenic plants	Non-transgenic plants	
Col-0; 35S::*BvFVE1*; #9	35.22 ±3.95 (n=23[b])	36.50 ±9.19 (n=2[b])	0.40 (0.69)	15.65 ±3.39	19.00 ±8.49	1.57 (0.13)
Col-0; 35S::*BvFVE1*; #16	31.43 ±4.24 (n=14[b])	31.73 ±1.68 (n=11[b])	0.22 (0.82)	12.86 ±2.57	12.36 ±1.12	0.59 (0.56)
fve-7; 35S::*BvFVE1*; #1	66.62 ±7.08 (n=13[b])	65.75 ±4.33 (n=8[b])	0.31 (0.76)	33.31 ±9.27	36.88 ±6.24	0.38 (0.70)
fve-7; 35S::*BvFVE1*; #32	59.95 ±7.26 (n=21[b])	-[c]	2.11 (0.04)[d]	32.76 ±7.91	-[c]	1.32 (0.20)[d]

[a] Transgenic plants were identified by PCR analysis as described in Materials and Methods.
[b] Number of, respectively, transgenic or non-transgenic plants within a given T2 population.
[c] T2 family #32 did not include non-transgenic plants.
[d] The means of DTB and TNL in transgenic plants was compared against the respective means in the non-transgenic plant group of family #1 which was grown in parallel.

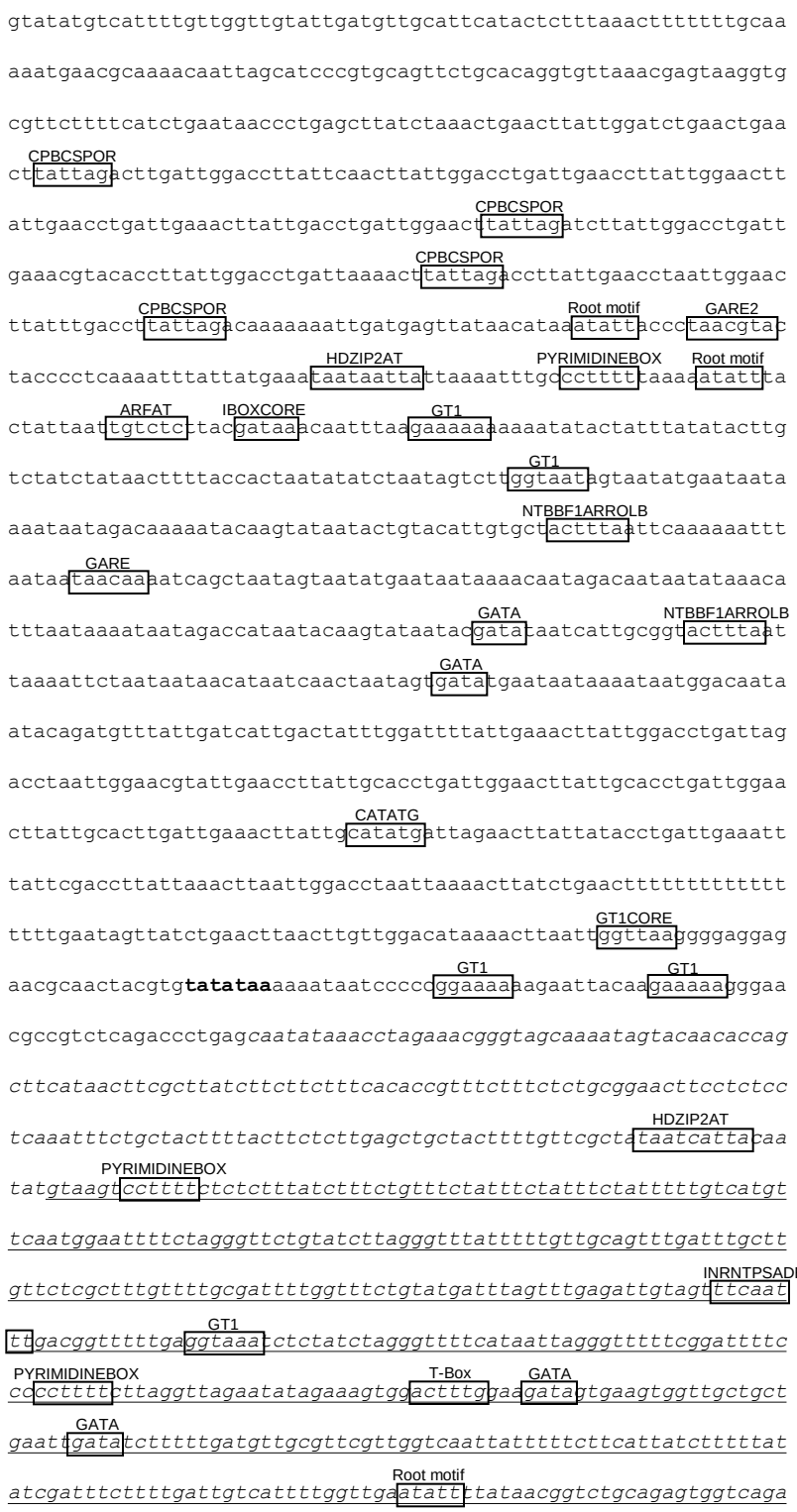


Suppl. Fig. 1



Suppl. Fig. 1. Sequence and structure of the autonomous pathway gene homologs *BvLD* and *BvLDL1*. (A) Exon-intron structure of *BvLD*, *BvLDL1* and the respective *A. thaliana* genes (*LD*, accession number CAJ53849; *LDL1*, accession number NP_176471). Exons are indicated as black rectangles, the position of start and stop codons is indicated by arrows and vertical bars, respectively. (B) Pairwise sequence alignments and domain organization. The alignments were generated using ClustalW2 (http://www.ebi.ac.uk/Tools/clustalw2/index.html). Identical and similar residues are highlighted by black or grey boxes, respectively. The position of protein domains according to Pfam 22.0 (http://pfam.sanger.ac.uk/) is marked by horizontal lines above the alignment. Two putative nuclear localization signals (NLS) in *LD* according to Lee *et al*. (1994) are indicated. HD, homeo domain; SWIRM, SWI3, RSC8 and MOIRA domain.

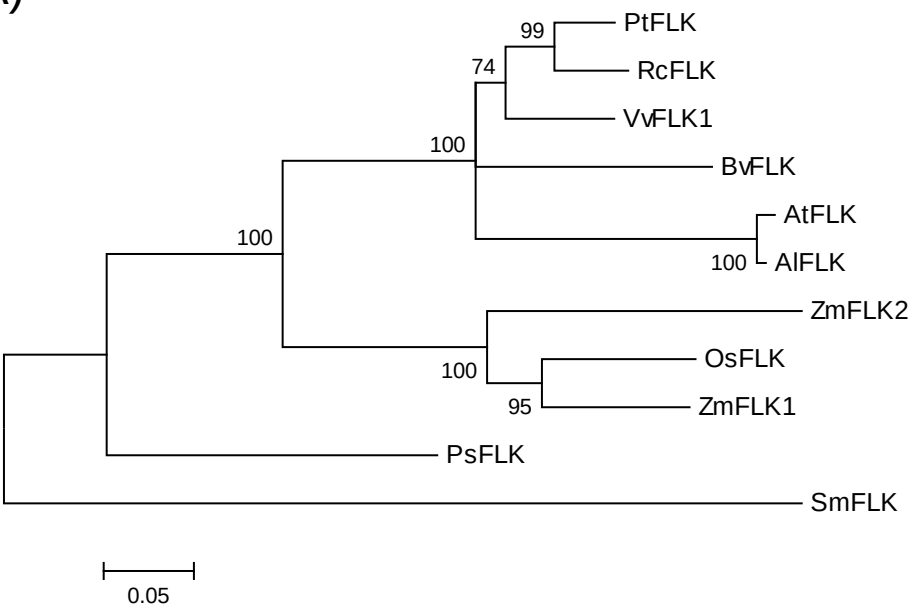
Suppl. Fig. 1



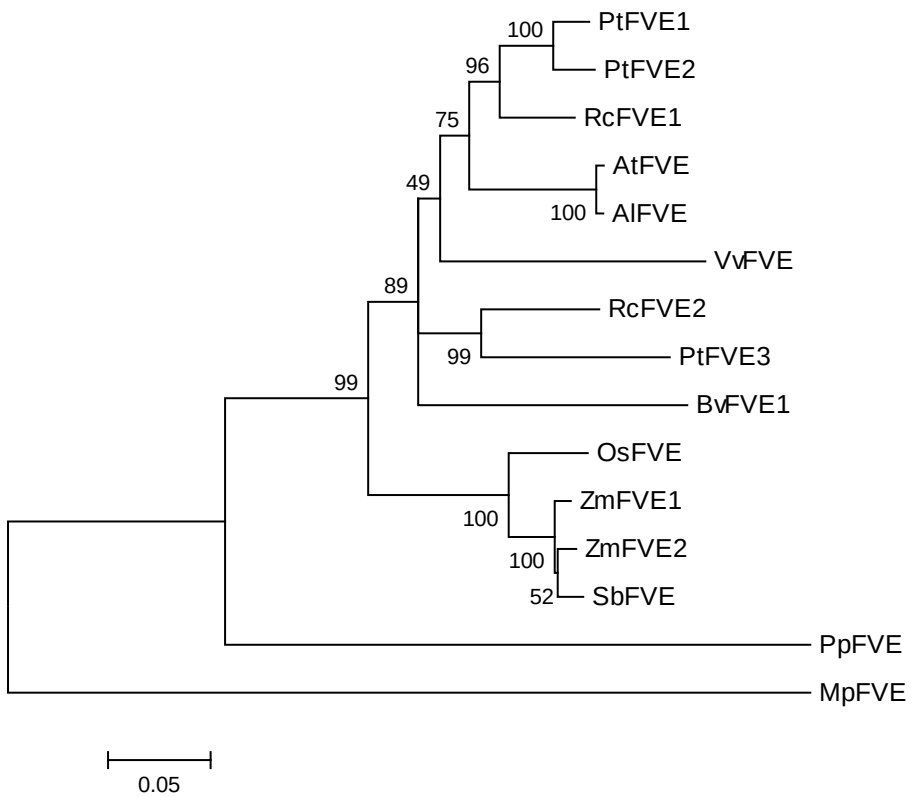
Suppl. Fig. 2. *BvFLK* promoter and 5' UTR. 1860 bp of the genomic sequence upstream of the start codon are shown. Bold letters indicate a putative TATA-box according to the transcription start site prediction program TSSP (http://www.softberry.ru/berry.phtml). The 5' UTR is indicated in italics. An intron within the 5' UTR is underlined. Transcription and splicing of the 5' UTR was verified by RT-PCR and sequencing. Root motifs (Elmayan and Tepfer 1995) and putative phytohormone- (ARFAT (Nag *et al*., 2005), CATATG motif (Xu *et al*., 1997), CPBCSPOR (Fusada *et al*., 2005), GARE (Ogawa *et al*., 2003), GARE2 (Sutoh and Yamauchi 2003), PYRIMIDINEBOX (Mena *et al*., 2002), NTBBF1ARROLB (Baumann *et al*., 1999)) and light-regulated promoter elements (GT1 and GT1CORE consensus sequences (Zhou 1999), IBOX core motif (Terzaghi and Cashmore, 1995), GATA box (Reyes *et al*., 2004), INRNTPSADB (Nakamura *et al*., 2002), T-box (Chan *et al*., 2001), HDZIP2AT (Ohgishi *et al*., 2001)) are boxed. We note that all four CPBCSPOR cytokinin-response elements (TATTAG) are clustered ~0.8-1.0 kb upstream of the transcription start site and the flanking nucleotides are conserved for these four elements, thus giving rise to a 9 nt motif, CTTATTAGA.

Suppl. Fig. 2

(A)


PtFLK
RcFLK
VvFLK1
BvFLK
AtFLK
AlFLK
ZmFLK2
OsFLK
ZmFLK1
PsFLK
SmFLK
99
74
100
100
100
100
95
0.05

(B)


PtFVE1
PtFVE2
RcFVE1
AtFVE
AlFVE
VvFVE
RcFVE2
PtFVE3
BvFVE1
OsFVE
ZmFVE1
ZmFVE2
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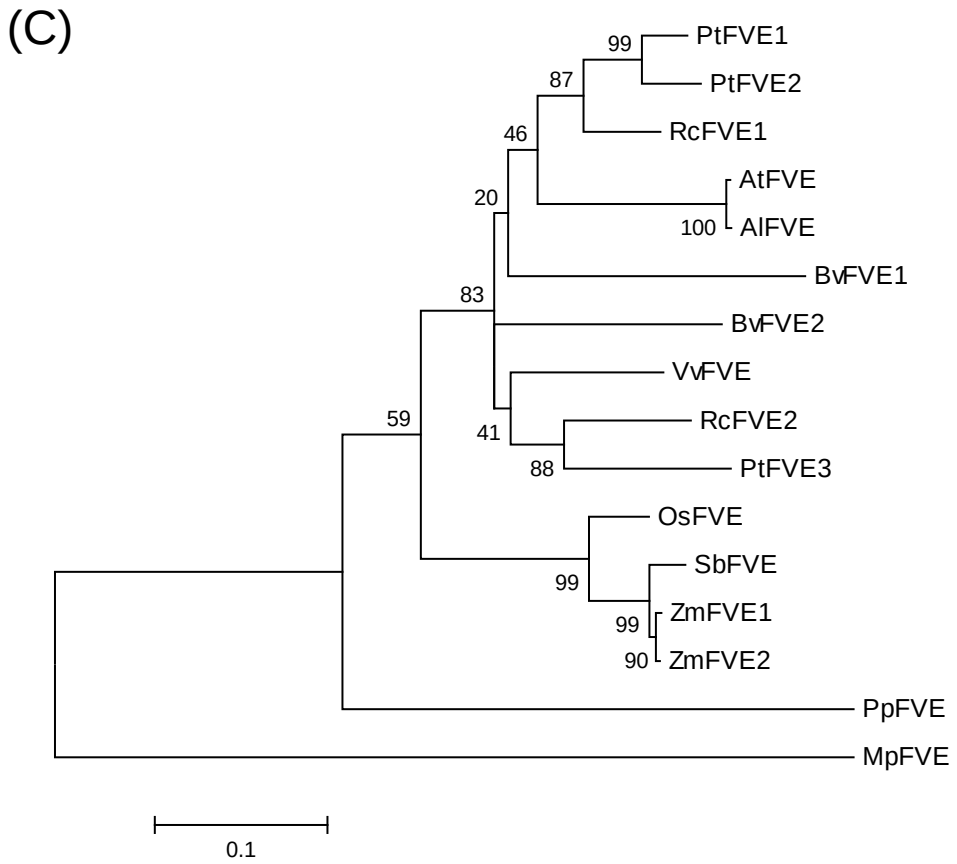
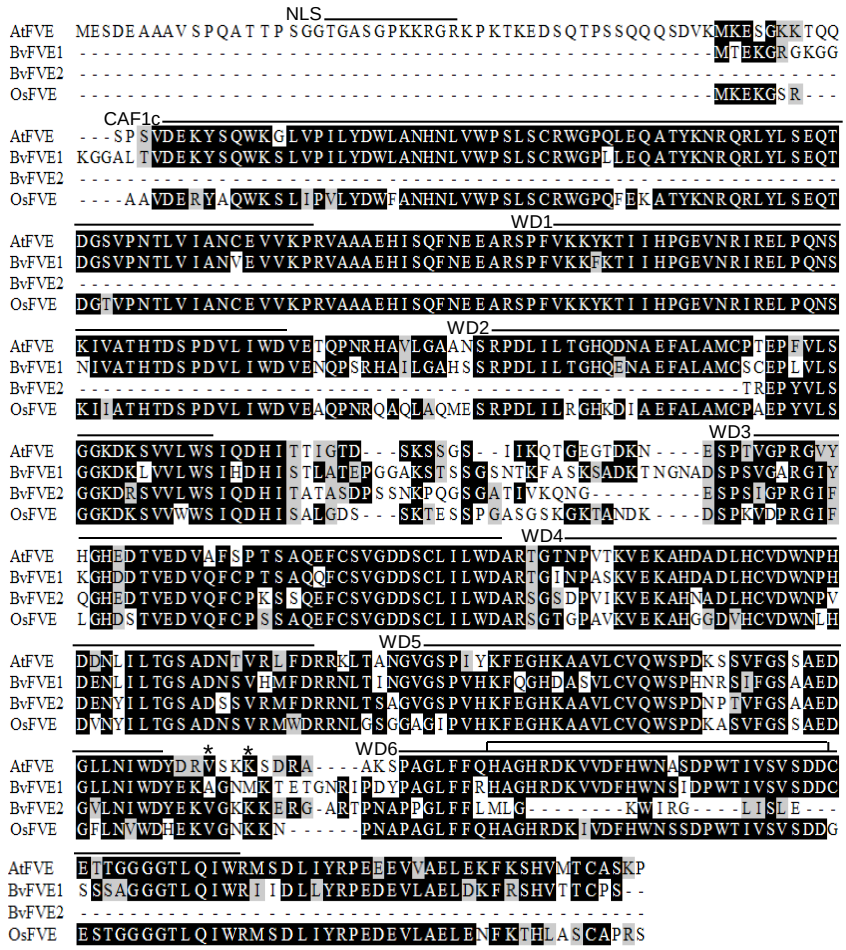
99
100
100
52
0.05



Suppl. Fig. 3. Phylogenetic analysis of *FLK* (A) and *FVE* (B,C). Included are all putative plant orthologs retrieved from the NCBI RefSeq protein database (http://www.ncbi.nlm.nih.gov/refseq) using blastp and bidirectional best hit analysis as described in Materials and Methods. (A) Unrooted neighbor-joining tree including *A. thaliana* FLK (AtFLK), BvFLK, *Arabidopsis lyrata* FLK (AlFLK, XP_002884454.1), *Oryza sativa* FLK (OsFLK, AAL31692.1), *Picea sitchensis* FLK (PsFLK, ABK24418.1), *Populus trichocarpa* FLK (PtFLK, XP_002319087.1), *Ricinus communis* FLK (RcFLK, XP_002521945.1), *Selaginella moellendorffii* FLK (SmFLK, XP_002968912.1), *Vitis vinifera* FLK (VvFLK, XP_002269249.1), *Zea mays* FLK1 (ZmFLK1, NP_001151605.1), and *Z. mays* FLK2 (ZmFLK2, NP_001148654.1). (B) Unrooted neighbor-joining tree including *A. thaliana* FVE (AtFVE), BvFVE1, *A. lyrata* FVE (AlFVE, XP_002886035.1), *Micromonas pusilla* FVE (MpFVE, XP_003058120.1), *Physcomitrella patens* FVE (PpFVE, XP_001776478.1), *P. trichocarpa* FVE1 (PtFVE1, XP_002332825.1), *P. trichocarpa* FVE2 (PtFVE2, XP_002330494.1), *P. trichocarpa* FVE3 (PtFVE3, XP_002327543.1), *R. communis* FVE1 (RcFVE1, XP_002514113.1), *R. communis* FVE2 (RcFVE2, XP_002527406.1), *Sorghum bicolor* FVE (SbFVE, XP_002456237.1), *V. vinifera* FVE (VvFVE, XP_002282044.1), *Z. mays* FVE1 (ZmFVE1, NP_001105067.1), and *Z. mays* FVE2 (ZmFVE2, NP_001105191.1). (C) Unrooted neighbor-joining tree including a partial predicted protein sequence of BvFVE2 (corresponding to nucleotide positions 3 – 629 of EST EG550040) and all proteins in (B), except that for tree construction only the partial amino acid sequences were used that align to the partial BvFVE2 sequence. Bootstrap values as percentages of 1000 replicates are given at the branching points of the trees.



Suppl. Fig. 4. Multiple sequence alignment including *A. thaliana* FVE (AtFVE), BvFVE1, the putative translation product of the largest open reading frame in the *B. vulgaris* EST EG550040 (BvFVE2), and OsFVE. Asterisks indicate a valine and a lysine residue which are conserved between FVE, BvFVE2 and OsFVE. NLS, putative nuclear localization signal (Ausin *et al.*, 2004); CAF1c, CAF1 subunit C / histone binding protein RBBP4 domain; WD, WD40 repeat domain. A potential zinc binding site (unfilled box) in WD6 (Kenzior and Folk, 1998) is also indicated.