Supplementary Material

Constitutively overexpressing a tomato fructokinase gene (*LeFRK1*) in cotton (*Gossypium hirsutum* L. cv. Coker 312) positively affects plant vegetative growth, boll number and seed cotton yield

Thiya Mukherjee^A, Mariana Ivanova^A, Marisela Dagda^A, Yoshinori Kanayama^B, David Granot^C and A. Scott Holaday^{A,D}

Table S1. Number of matured bolls/plant, seed-cotton mass/boll (SCM/boll), and calculated values of seed-cotton mass/plant (SCM/plant) for cotton plants over-expressing *LeFRK1* (lines 2-2a, 19-3a and 35-1a) and segregating nulls (N, *LeFRK1* non-expressing) from experiment 1

The values are the mean \pm standard deviation (n = 4-5 plants/line). Asterisks indicate P-values comparing means for the *LeFRK1* lines with the means for the nulls. **P < 0.01, and *** P < 0.001 based on a Student's t-test

Genotype	Bolls	SCM/boll (g)	SCM/plant (g)
		(8)	(5)
N	8±2	2.72 ± 0.35	22±6
2-2a	11 ± 4	$4.23 \pm 0.43 ***$	46±11**
19-3a	$14 \pm 2**$	$3.72 \pm 0.26**$	51±10***
35-1a	14±3**	$3.49 \pm 0.18**$	49±14**

^ADepartment of Biological Sciences, Texas Tech University, Lubbock, Texas 79409, USA.

^BGraduate School of Agricultural Science, Tohoku University Aoba-ku, Sendai 981-8555, Japan.

^CDepartment of Vegetables Research, Institute of Plant Sciences, Agricultural Research Organisation, The Volcani Center, PO Box 6, Bet Dagan 50250, Israel.

^DCorresponding author. Email: scott.holaday@ttu.edu