

Table 1 Seed fatty acid compositions (%) of the transgenic lines

Line	C16:0	C18:0	C18:1	C18:2	C18:3	C20:1	C22:1	VLCFAs
<i>BnFAE1-Ri</i>	3.91±0.25*	1.77±0.26**	61.50±0.76**	17.30±0.79**	9.31±0.33*	4.38±0.23**	1.27±0.07**	6.21±0.28**
<i>BnFAD2-Ri</i>	2.22±0.08**	1.25±0.18	32.07±0.46**	1.44±0.10**	2.67±0.10**	13.53±0.51*	45.43±0.77	60.34±0.36**
<i>BnFAE1-Ri/BnFAD2-Ri</i>	2.94±0.01**	1.70±0.10**	82.76±0.25**	2.37±0.12**	3.46±0.07**	4.98±0.20**	1.61±0.25**	6.83±0.19**
<i>AtFAE1-SE</i>	1.13±0.06**	0.62±0.04**	12.62±0.45**	10.08±0.46	6.70±0.16**	5.43±0.49**	60.16±0.55**	68.85±0.06**
<i>BnFAD2-Ri/AtFAE1-SE</i>	1.45±0.09**	0.97±0.05	22.52±0.47**	1.86±0.18**	3.03±0.10**	6.26±0.33**	60.52±0.61**	70.18±0.76**
CY2	2.57±0.04	1.04±0.08	20.53±0.53	10.73±0.08	8.20±0.29	11.73±0.62	43.90±0.66	56.92±0.45

Note: C16:0: Palmitic acid; C18:0: Stearic acid; C18:1: Oleic acid; C18:2: Linoleic acid; C18:3: Linolenic acid; C20:1: Eicosenoic acid; C22:1: Erucic acid; VLCFAs: Very long chain fatty acids including C20:0 (Arachidic acid), C20:1, C22:0 (Behenic acid), C22:1 and C24:0 (Lignoceric acid); Each value is the mean ± standard deviation of three replications; * and ** indicate statistically significant differences compared with CY2 at 0.05 and 0.01 levels, respectively