

Table S9 Replicate 1 internode length: mean, standard deviation, sample size, and significance groupings.

Chromosome		Internode Length			
Arm	E	F	H	O	
1S	9.6±0.69 ^a (7)	10.74±1.16 ^a (12)	8.91±1.06 ^a (17)	7.37±2.22 ^a (6)	
1L	9.86±0.92 ^a (9)	10.23±1.56 ^a (14)	9.76±1.62 ^a (10)	4.2±0.88 ^b (4)	
3S	11.29±1.19 ^a (8)	10.96±1.04 ^a (19)	11.41±0.73 ^a (16)	11.1±NA (1)	
3L	11.86±1.31 ^a (9)	11.46±1.3 ^{ab} (11)	9.29±2.23 ^b (12)	5.81±1.51 ^c (7)	
4S	11.51±0.88 ^a (12)	10.78±1.27 ^a (13)	11.09±1.29 ^a (13)	10.2±1 ^a (7)	
4L	10.39±1.04 ^a (11)	10.82±1.15 ^a (14)	11.05±0.46 ^a (11)	8.62±1.64 ^a (9)	
5S	12.21±0.84 ^a (14)	9.93±1.67 ^b (13)	8.54±1.54 ^b (16)	10.05±0.73 ^{ab} (6)	
5L	10.5±NA (1)	10.57±1.04 ^a (14)	10.56±1.18 ^a (21)	9.83±1.18 ^a (3)	
6L	9.99±0.7 ^{ab} (9)	11.04±0.8 ^{ab} (9)	11.77±2.49 ^a (11)	8.58±1.96 ^b (15)	
7L	10.91±0.52 ^a (8)	10.81±1.37 ^a (8)	10.81±1.04 ^a (11)	8.39±1.11 ^a (7)	
8L	11.49±1.45 ^{ab} (7)	11.17±1.59 ^a (11)	10.58±1.33 ^{ab} (15)	8.76±2.55 ^b (15)	
9S	12.03±1.37 ^a (12)	11.59±1.22 ^a (10)	12.68±1.02 ^a (16)	7.96±2.5 ^b (7)	
9L	9.56±1.6 ^a (9)	8.6±1.06 ^a (10)	8.57±0.97 ^a (17)	9.85±0.79 ^a (11)	
10L	12.32±0.99 ^a (13)	10.8±0.95 ^{ab} (12)	11.51±0.56 ^a (14)	8±0.72 ^b (5)	

This table contains each group's mean ± group standard deviation and the sample size, n, of each group in parenthesis (n). In superscript, linked to the standard deviation are letters "a", "b" and "c" which indicate groups not significantly different from each other at the p≤0.05 level. Dosage designations are as follows: Euploid plants (no B-A present)-E; Euploid plants (B-A present and failure of nondisjunction)-F; Hyperploid plants-H; Hypoploid plants-O