

Table S7. Allele specific expression results of *ZmYAB2.1* with five or more reads from Lemmon *et al.* (2014)

F1 Hybrid	Percentage of Reads Belonging to Maize (%)	Total Number of Reads
B73 x TIL14	0	65
W22 x TIL14	0	15
Ki3 x TIL11	28	39
Ki3 x TIL03	10	31

References

- Cook, J. P., M. D. McMullen, J. B. Holland, F. Tian, P. Bradbury *et al.*, 2012 Genetic architecture of maize kernel composition in the nested association mapping and inbred association panels. *Plant Physiol.* 158: 824–834.
- Darling, A. C. E., B. Mau, F. R. Blattner, and N. T. Perna, 2004 Mauve: multiple alignment of conserved genomic sequence with rearrangements. *Genome Res.* 14: 1394–1403.
- Lemmon, Z. H., R. Bukowski, Q. Sun, and J. F. Doebley, 2014 The role of *cis* regulatory evolution in maize domestication. *PLoS Genet.* 10: e1004745.