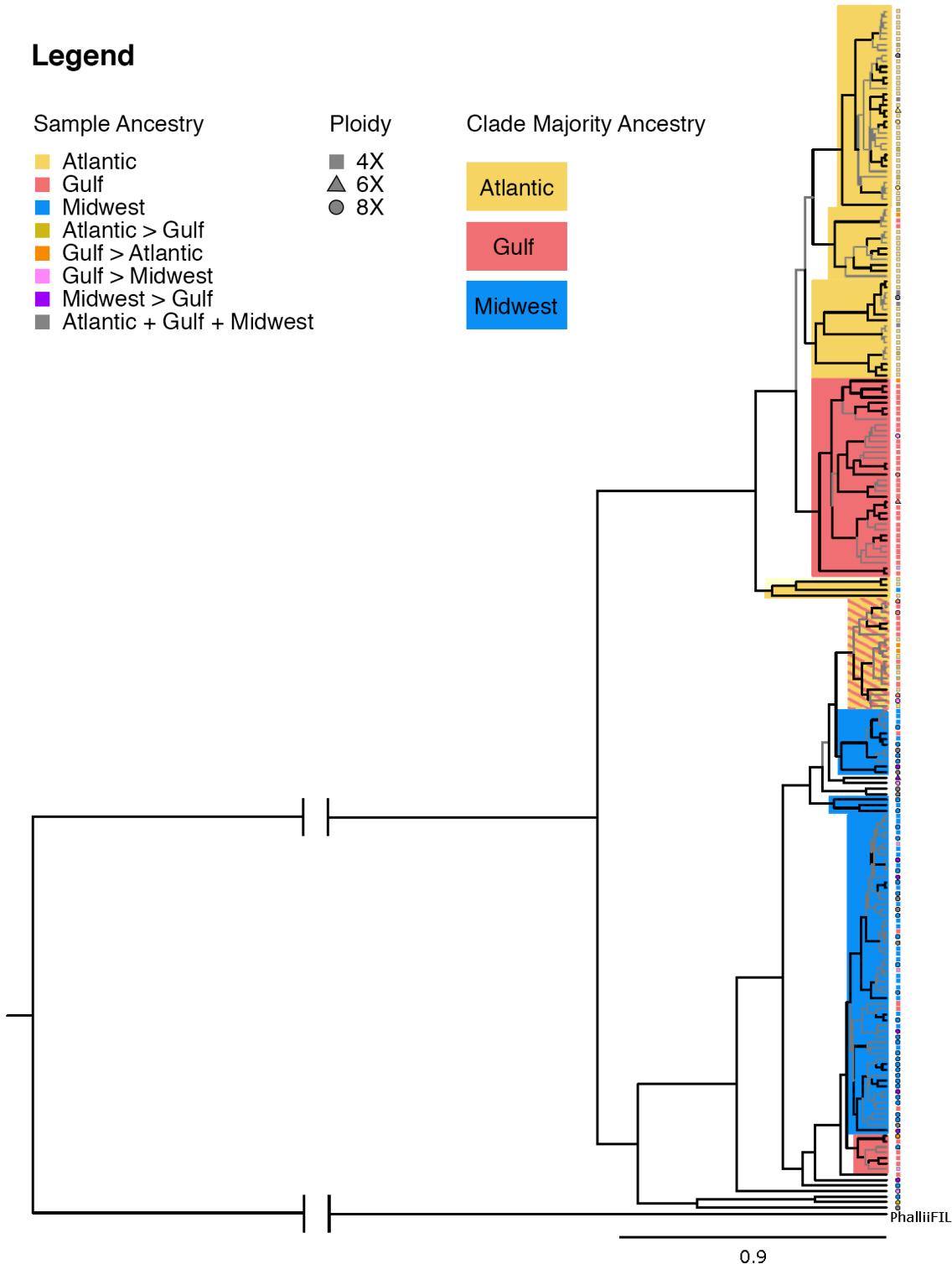


a

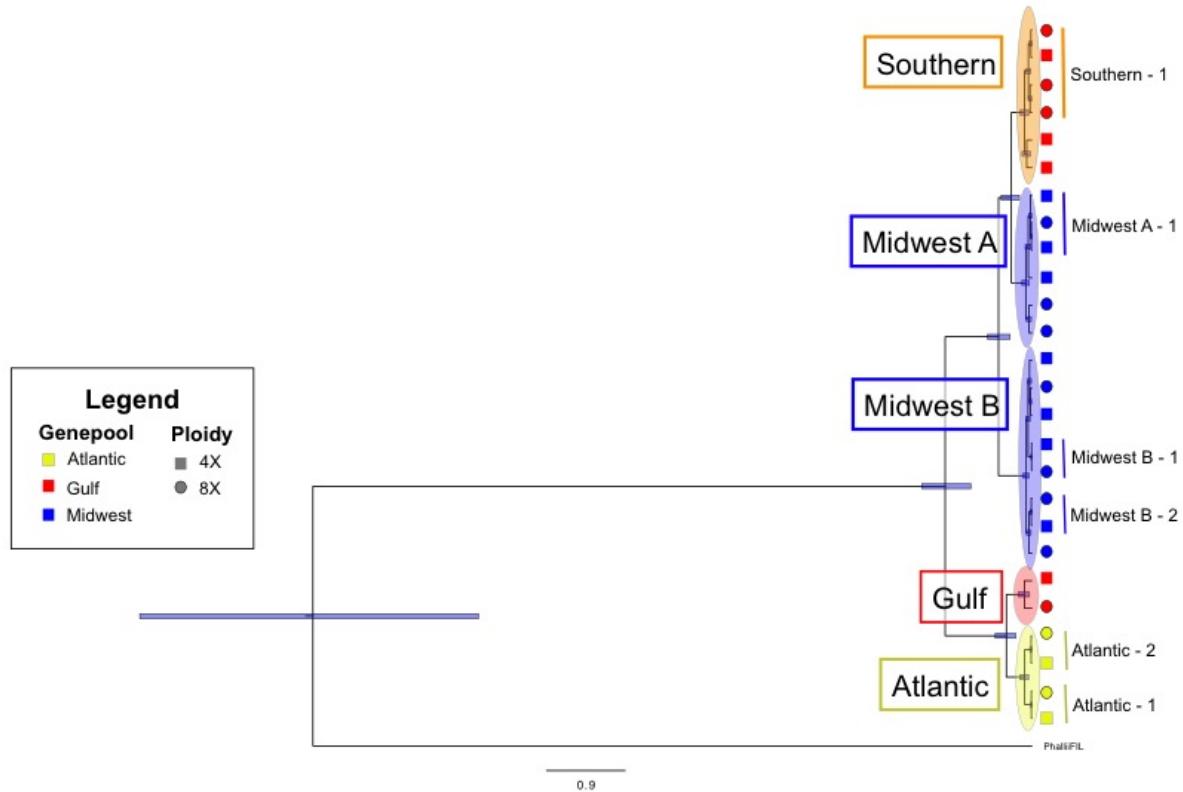
Legend

Sample Ancestry	Ploidy	Clade Majority Ancestry
Atlantic	■ 4X	Atlantic
Gulf	△ 6X	Gulf
Midwest	● 8X	Midwest
Atlantic > Gulf		
Gulf > Atlantic		
Gulf > Midwest		
Midwest > Gulf		
Atlantic + Gulf + Midwest		

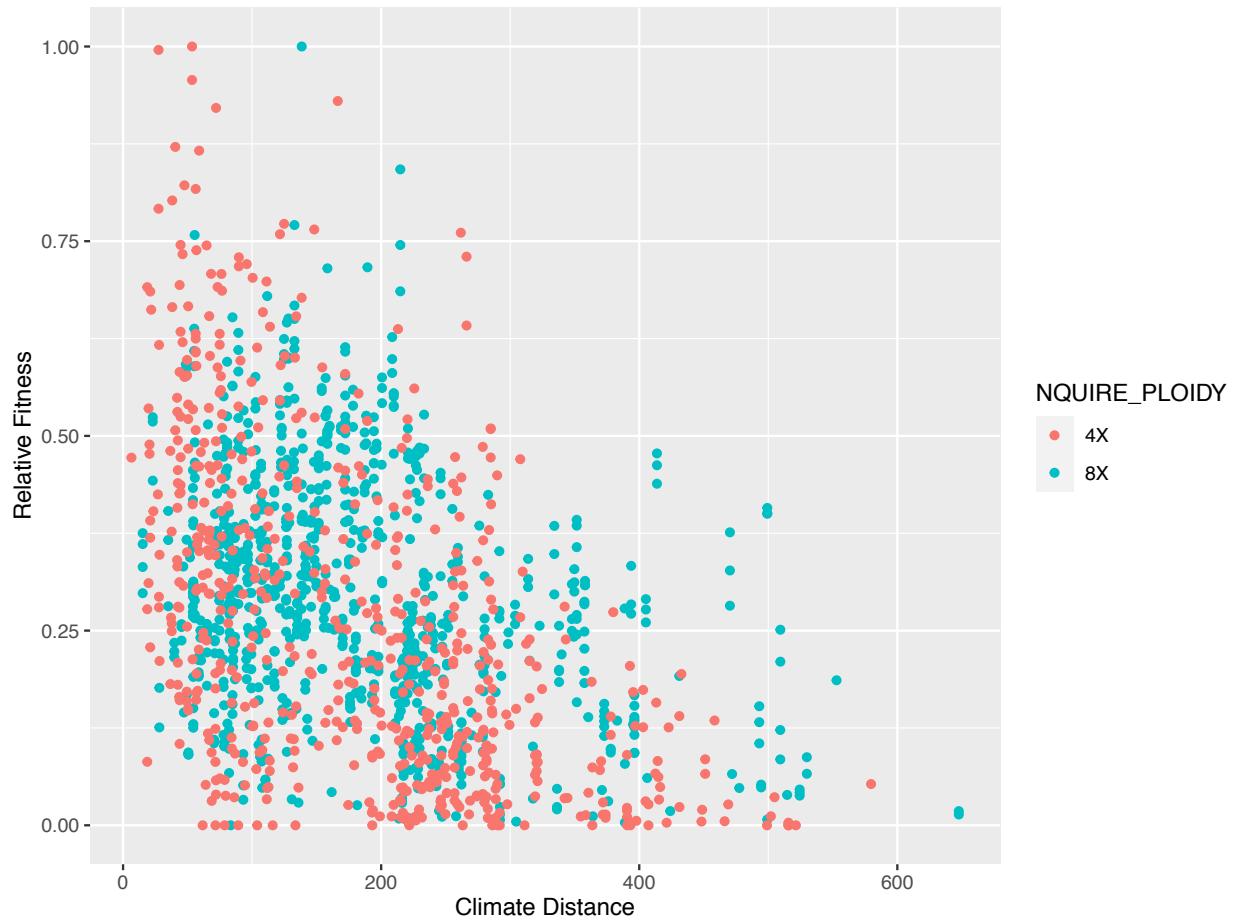


Supplemental Figure 1. Enlarged version of figure 2a with individual points that indicate the nuclear ancestry and ploidy of each sample. Gray branches indicate less than .95 posterior

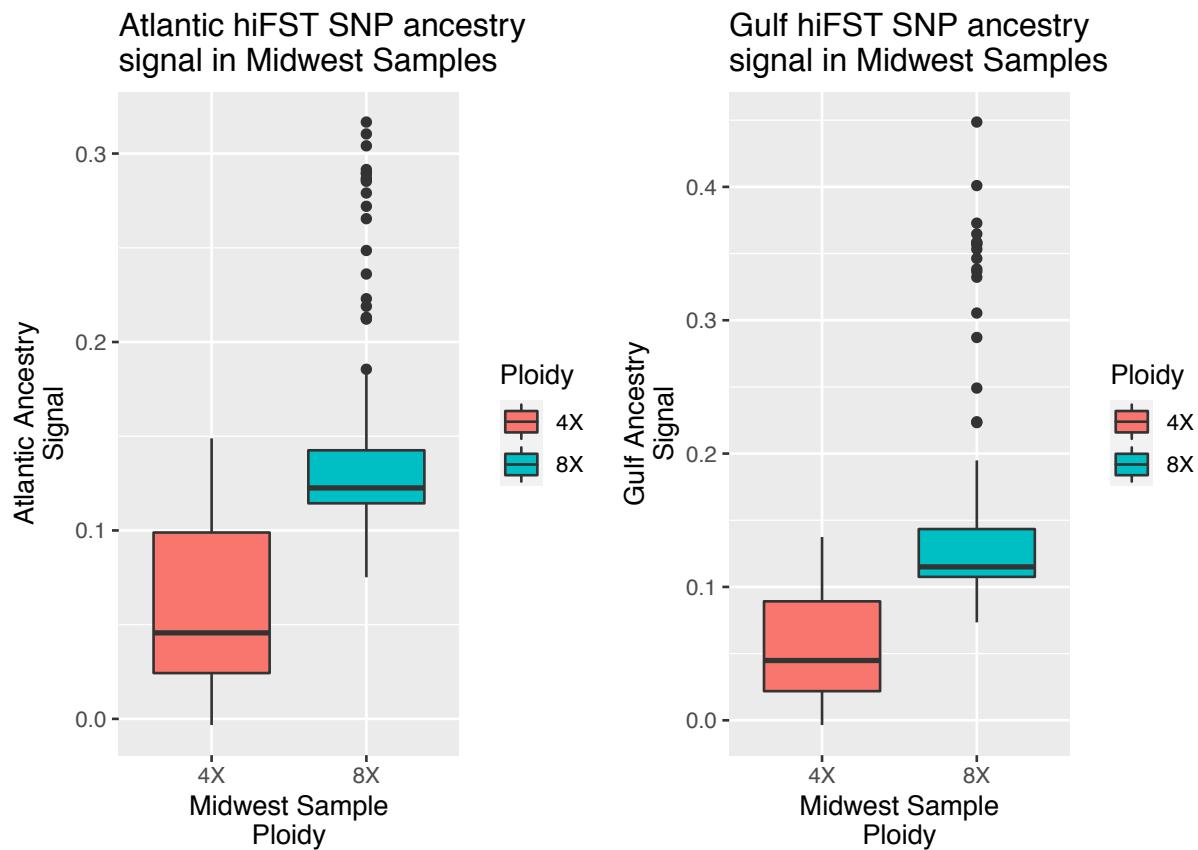
probability support, and black branches indicate greater than .95 posterior probability support. The outgroup is *P. halli* with an estimated divergence date of 8.35 mya.



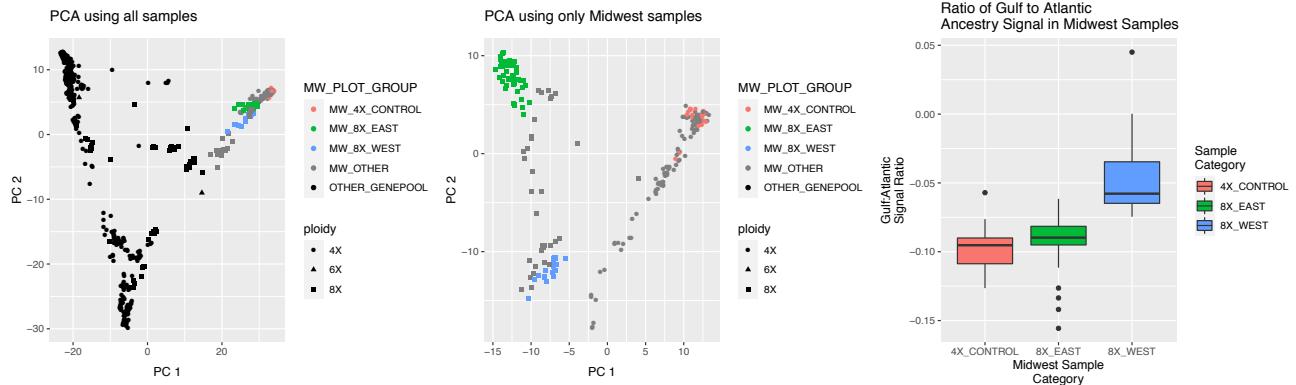
Supplemental Figure 2. Chloroplast tree using samples (see Supplemental Table 4) chosen to date the divergence of identified switchgrass groups using BEAST v2.6.6 (see Supplemental Table 2 for divergence date estimates).



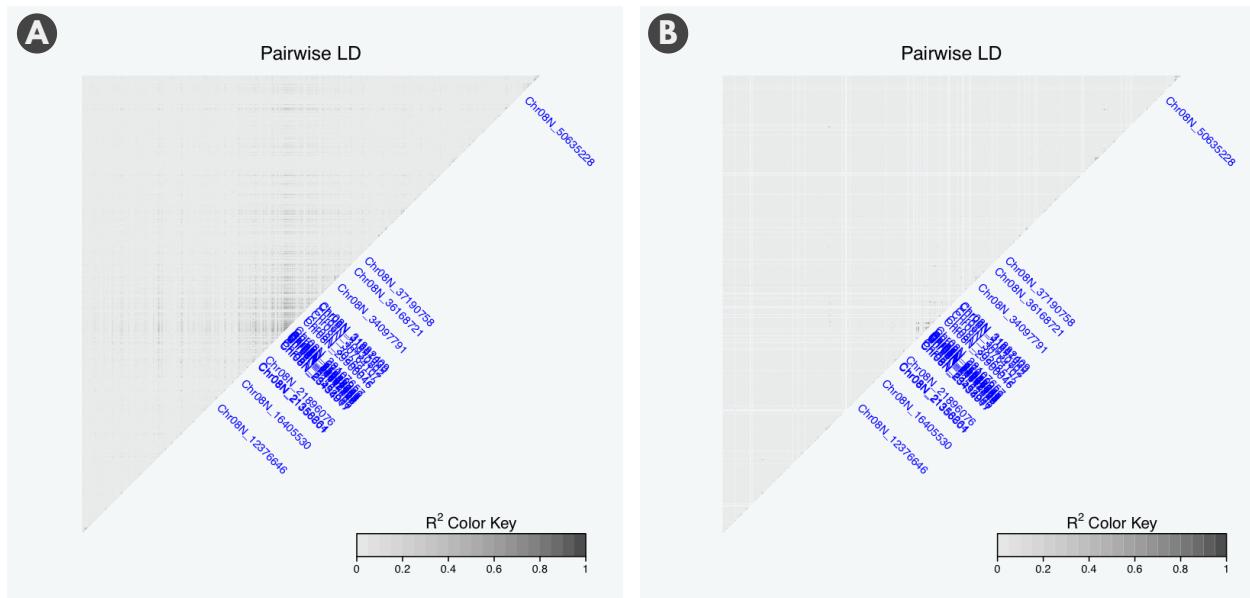
Supplemental Figure 3. Individual points used for plotting the linear relationship of relative fitness versus climate distance in Figure 3b.



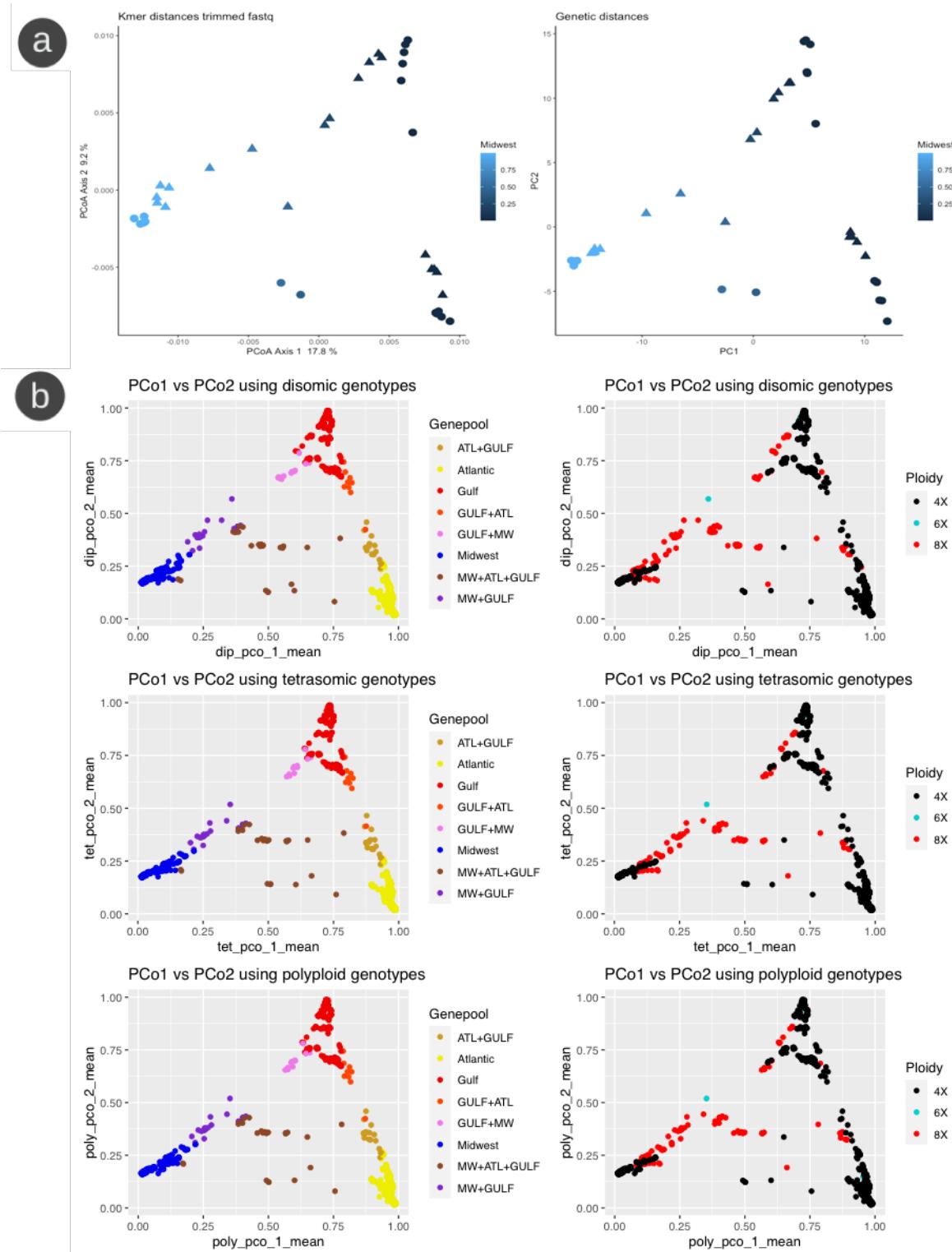
Supplemental Figure 4. Southern genepool ancestry signal in Midwest genepool. Signal of Atlantic and Gulf ancestry in 63 Midwest 4X and 105 Midwest 8X switchgrass, based on ancestry-informative alleles from SNPs with $Fst > 0.5$ between Atlantic or Gulf and Midwest 4X training sets.



Supplemental Figure 5. Comparison of Southern genepool ancestry signal in Midwest 8X. A subset of Midwest 8X samples with similar levels of Southern genepool ancestry were assigned to "East" and "West" groups based on PCA differentiation and geographic distribution. Red = Midwest 4X "control" samples with low Southern genepool ancestry signal, N = 20; green = Midwest 8X from eastern US, N = 52; blue = Midwest 8X mainly from western US, N = 20. (first panel) PCA plot of 706 samples in overall analysis; grey = Midwest samples not included in analysis in (third panel), black = Atlantic, Gulf, or Admixed samples. (second panel) PCA plot of Midwest samples; grey = Midwest samples not included in analysis in (third panel). (third panel) Ratio of Gulf to Atlantic ancestry signal. Midwest 4X control samples used to show background pattern; higher value on y-axis indicates higher ratio of Gulf to Atlantic ancestry signals.



Supplemental Figure 6. Linkage disequilibrium (LD) on Chromosome 08N for SNPs used in RDA analysis. Overlapping outlier SNPs are identified with blue text. **A)** Patterns of LD in the Midwest subpopulation. **B)** Patterns of LD in the Atlantic subpopulation.



Supplemental Figure 7. Population structure results are robust to genotyping methods. **A)** Strong similarity between PCoA generated from reference-free k-mer approach (left) and PCA generated using genotypes used in full analysis (right) shows that population structure patterns are not substantially biased by ploidy differences. Shape indicates ploidy (circle = 4X, triangle =

8X) and color indicates proportion of Midwest ancestry. The k-mer approach is computationally intensive, so the comparison uses a subset of 36 samples. B) Comparisons of PCoA results using different genotyping models for the same 50k SNPs. Top row uses disomic genotypes where all samples are assigned diploid (RR) genotypes. Second row uses tetrasomic genotypes where all samples are assigned tetraploid (RRRR) genotypes. Third row uses polyploid genotypes where 4X samples are assigned diploid genotypes and 8X individuals are assigned tetraploid genotypes. Left column shows samples colored by ancestry, right column shows samples colored by ploidy. Patterns are similar when using all three genotyping models.

Library ID	Ploidy	Genepool	Clade	Subclade
ASHOA	8X	Gulf	Southern	Southern - 1
IGQU	4X	Gulf	Southern	Southern - 1
JBEP	8X	Gulf	Southern	Southern - 1
IEMK	4X	Gulf	Southern	NA
IENZ	4X	Gulf	Southern	NA
JBEH	8X	Gulf	Southern	Southern - 1
IIJX	4X	Midwest	Midwest A	Midwest A - 1
IGSE	8X	Midwest	Midwest A	Midwest A - 1
IENF	4X	Midwest	Midwest A	Midwest A - 1
IIJM	4X	Midwest	Midwest A	NA
JBKD	8X	Midwest	Midwest A	NA
IIAQ	8X	Midwest	Midwest A	NA
IIJT	4X	Midwest	Midwest B	Midwest B - 2
IGRS	8X	Midwest	Midwest B	Midwest B - 2
IIBZ	8X	Midwest	Midwest B	NA
IIIN	4X	Midwest	Midwest B	NA
IIBS	8X	Midwest	Midwest B	NA
IGPJ	4X	Midwest	Midwest B	NA
IIAY	8X	Midwest	Midwest B	Midwest B - 1
IGRH	4X	Midwest	Midwest B	Midwest B - 1
IEYR	4X	Atlantic	Atlantic	Atlantic - 1
IEYM	8X	Atlantic	Atlantic	Atlantic - 1
IEYE	4X	Atlantic	Atlantic	Atlantic - 2
IEYD	8X	Atlantic	Atlantic	Atlantic - 2
IIBT	8X	Gulf	Gulf	NA
IEME	4X	Gulf	Gulf	NA

Supplemental Table 4. List of libraries used to estimate divergence dates in BEAST v2.6.6.