Additional File 5: Figure S5: Genomic clines estimated in *introgress* for *A. caudacutus* and *A. nelsoni*. Each plot represents a marker and includes the name of each locus and a P value from the analysis. Solid color lines represent genomic clines for *caudacutus* homozygote genotypes, dashed lines represent genomic clines for heterozygote genotypes. Shaded areas represent 95% confidence intervals.
Additional File 5: Figure S5: Continued

Hybrid index

Hybrid index

Hybrid index

Hybrid index

Hybrid index

Hybrid index

Hybrid index

Hybrid index

Hybrid index
Additional File 5: Figure S5: Continued
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