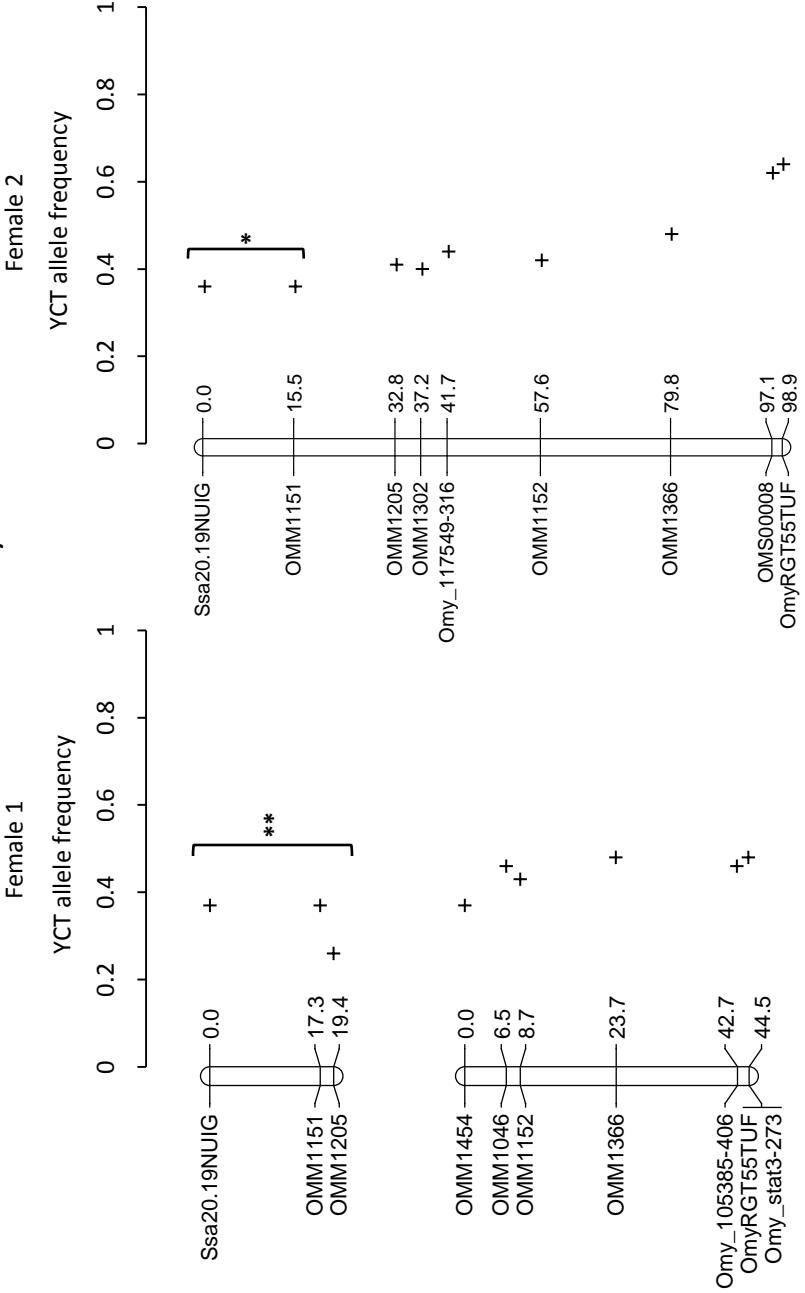
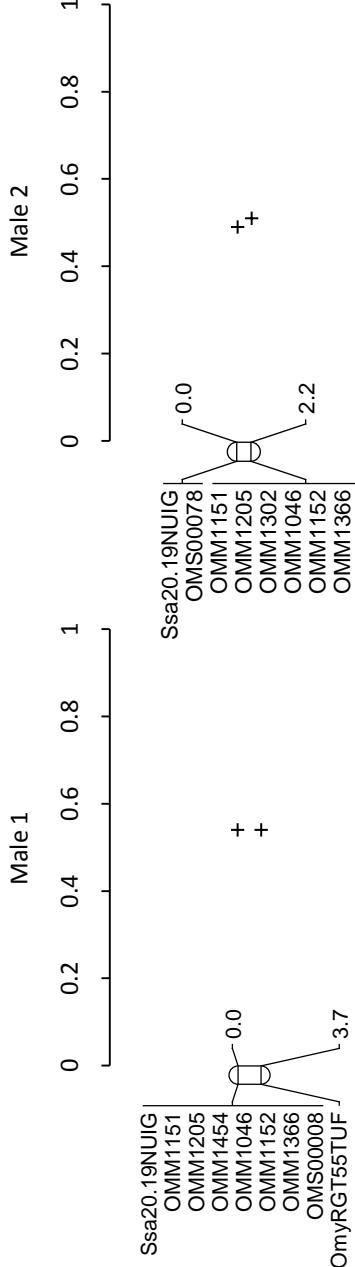


Additional file 2. Parent-specific F_1 hybrid linkage maps in p to q orientation and Yellowstone cutthroat trout (YCT) allele frequencies. Map distances (cM) represent the recombination fraction (theta) between loci. A 25 cM sliding window was used to identify blocks of loci containing YCT allele frequencies that deviated significantly from expected frequencies. Yellowstone cutthroat trout allele frequencies are plotted for each locus mapped within linkage groups. * indicates $P < 0.05$ and ** indicates $P < 0.01$.

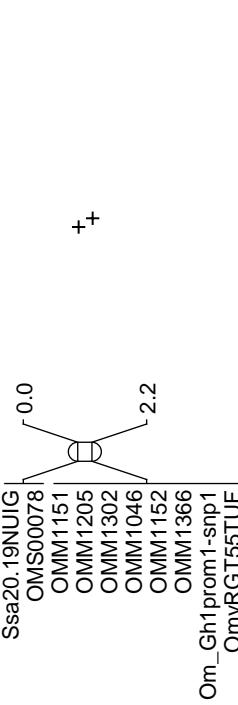
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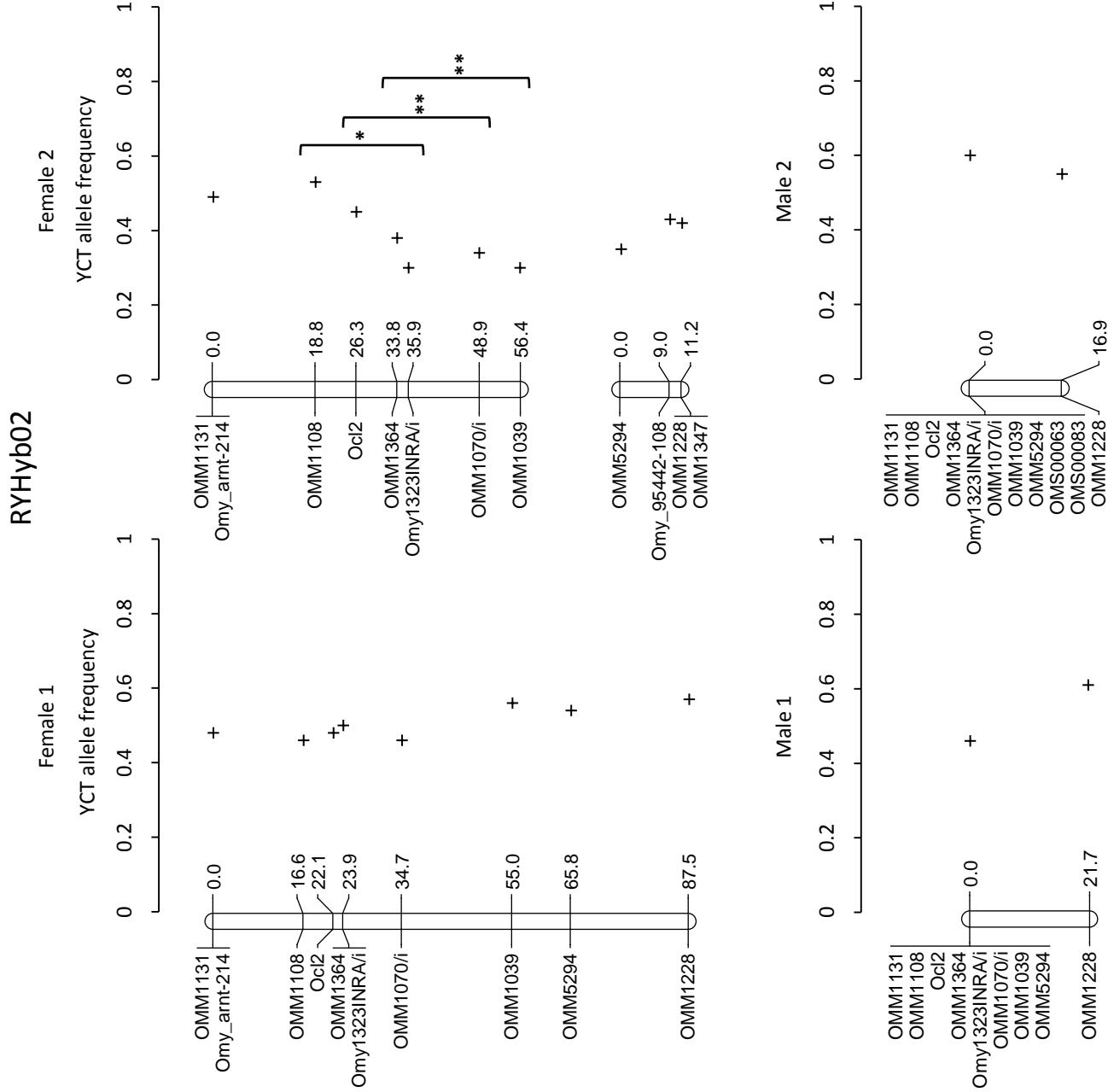


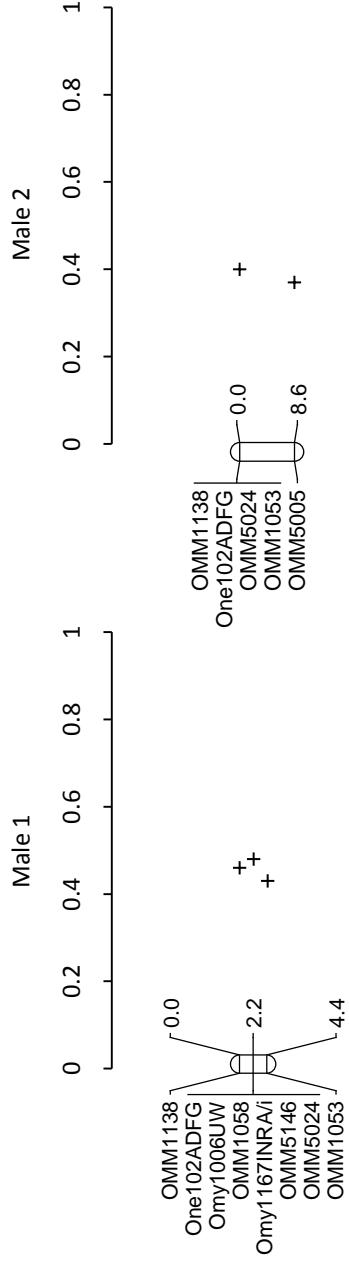
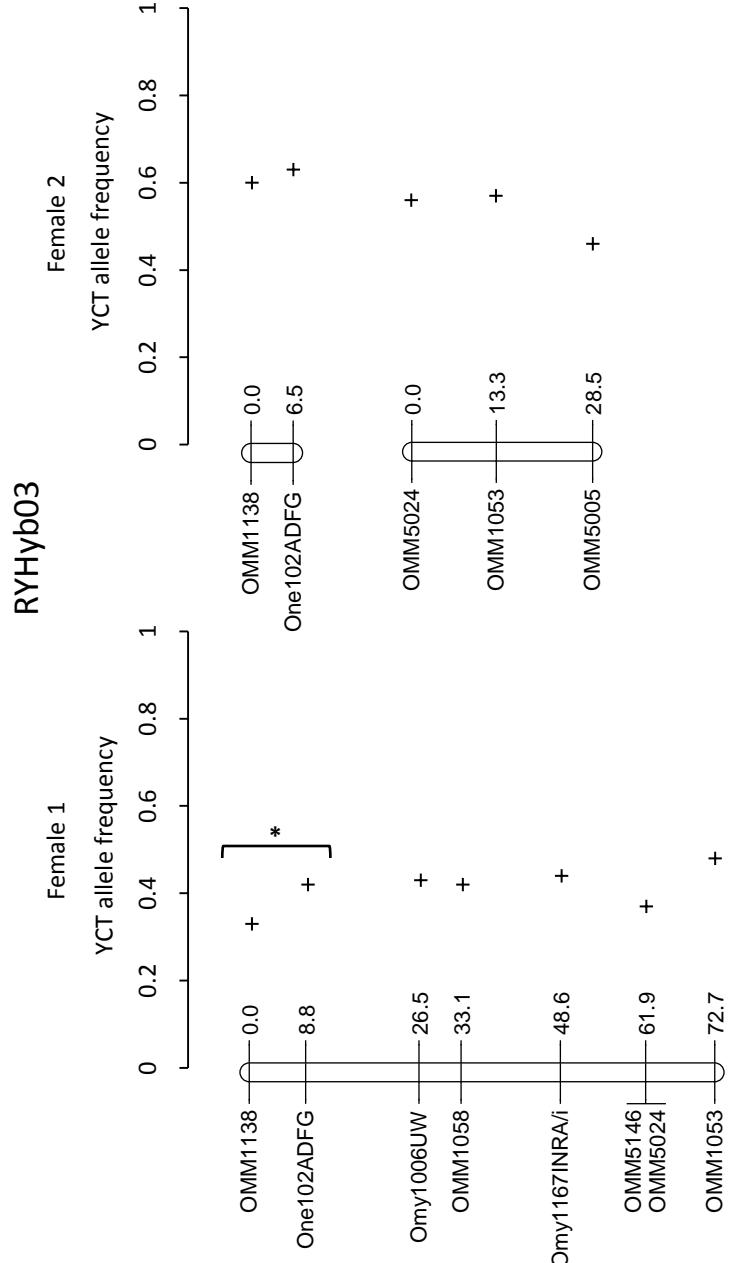
Male 1

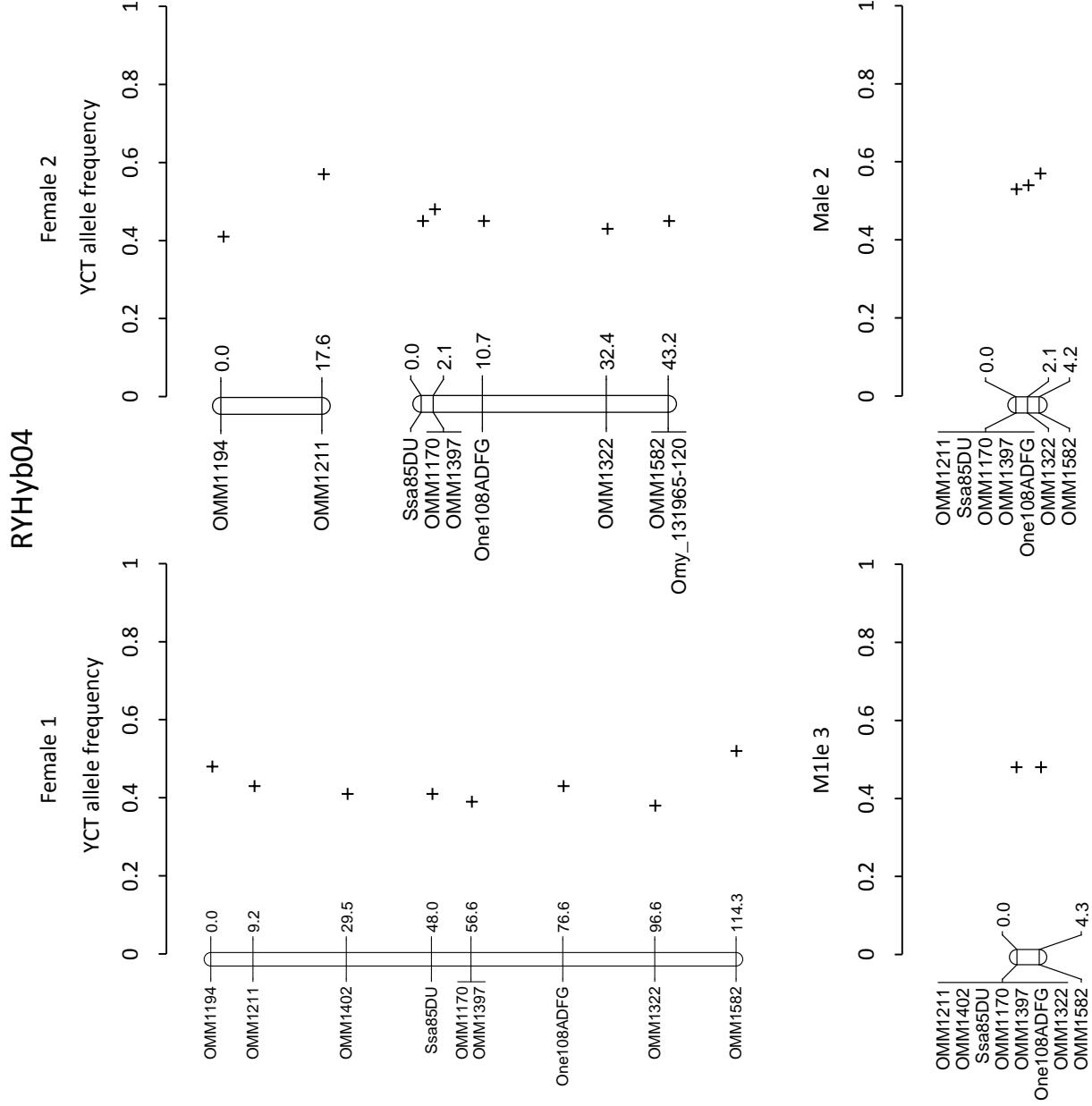


Male 2

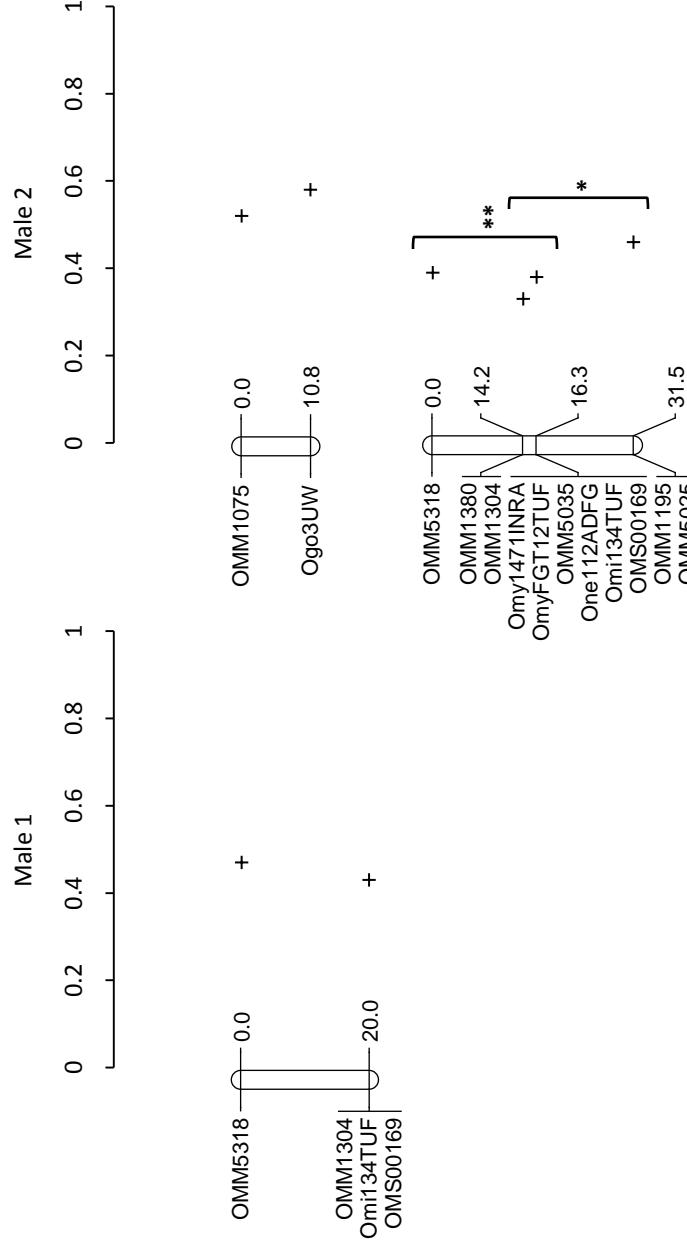
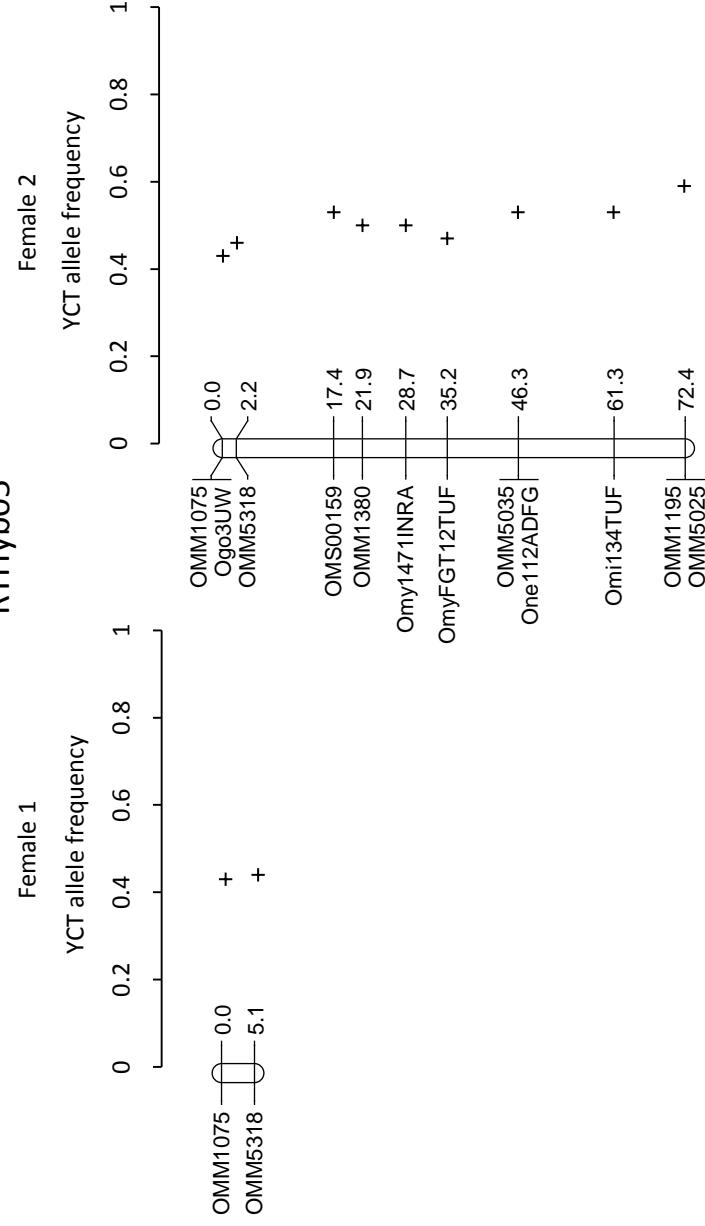




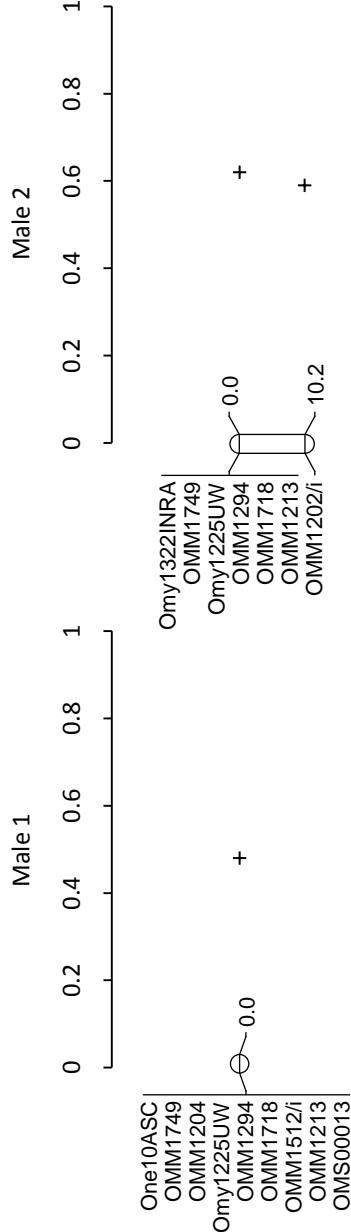
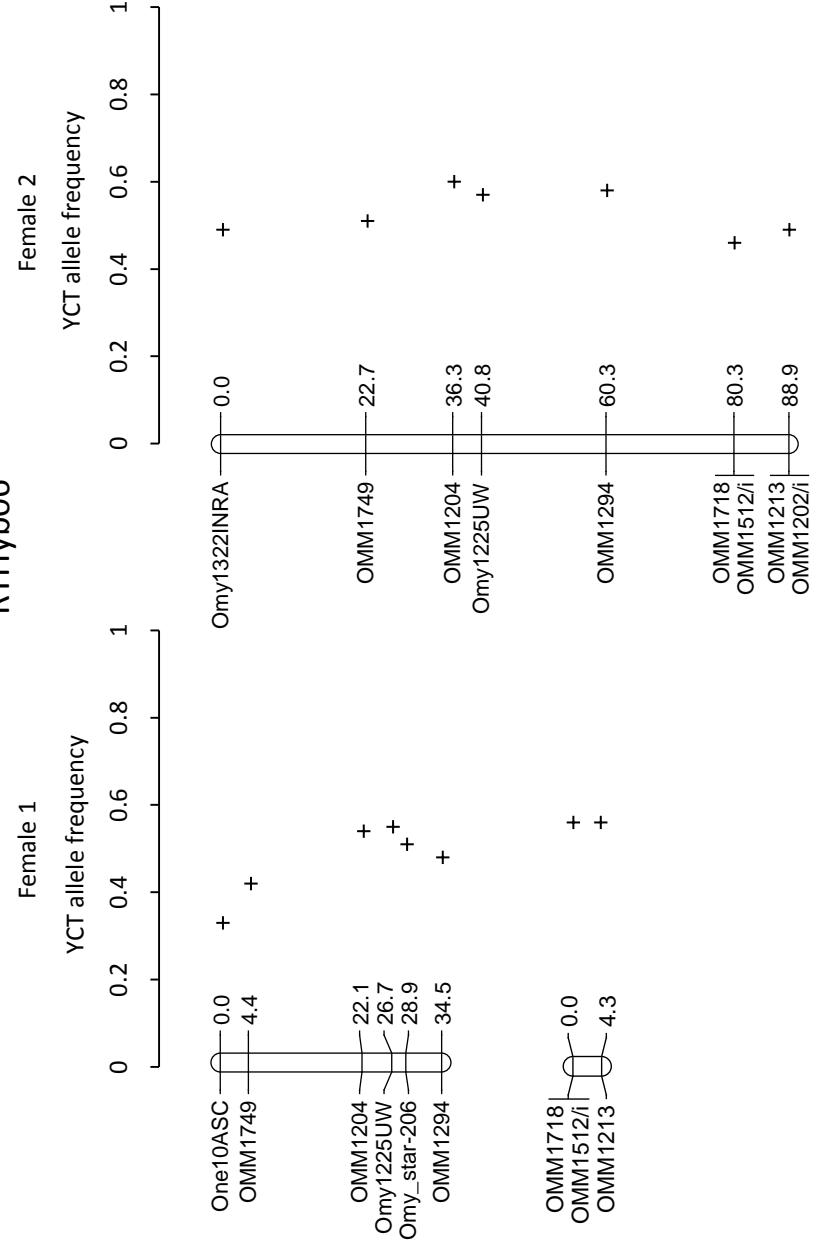


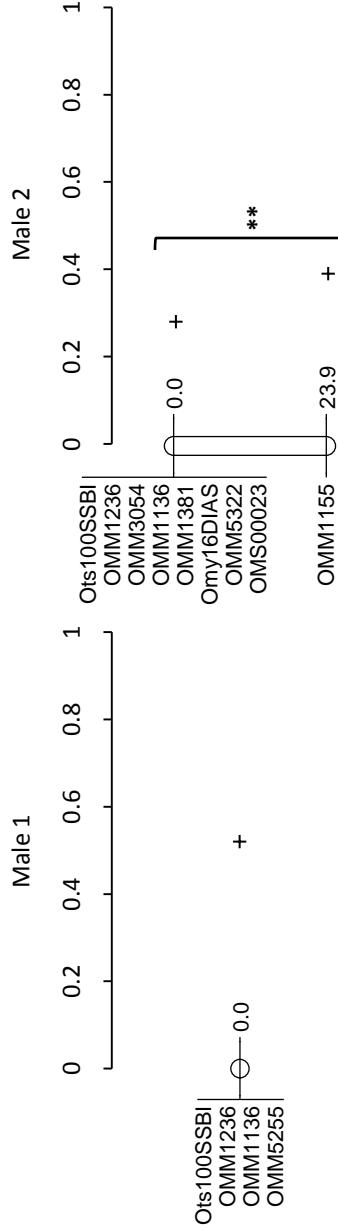
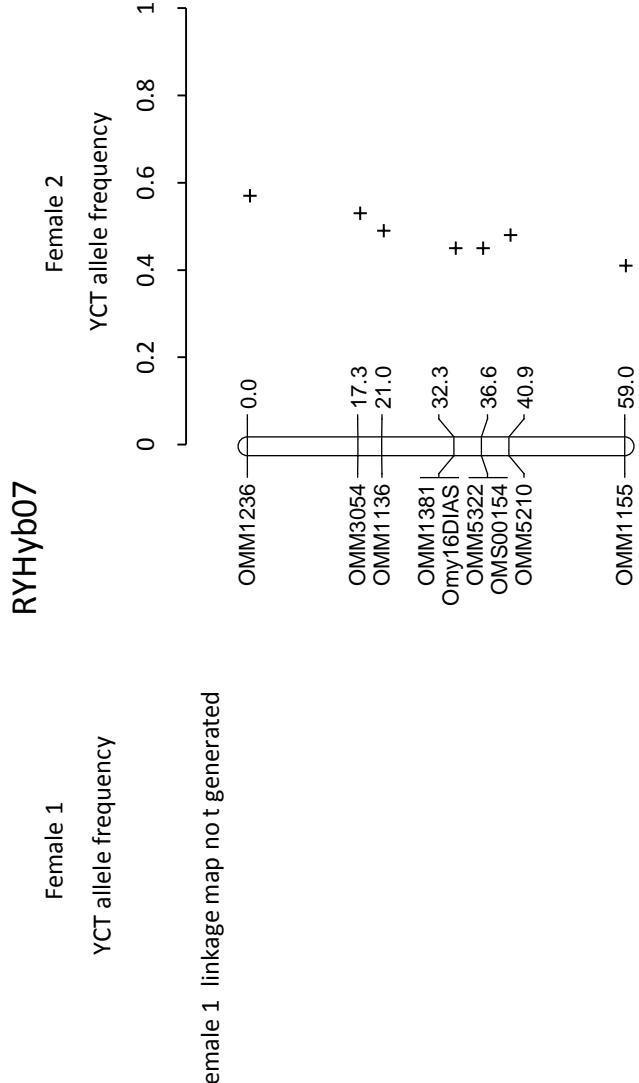


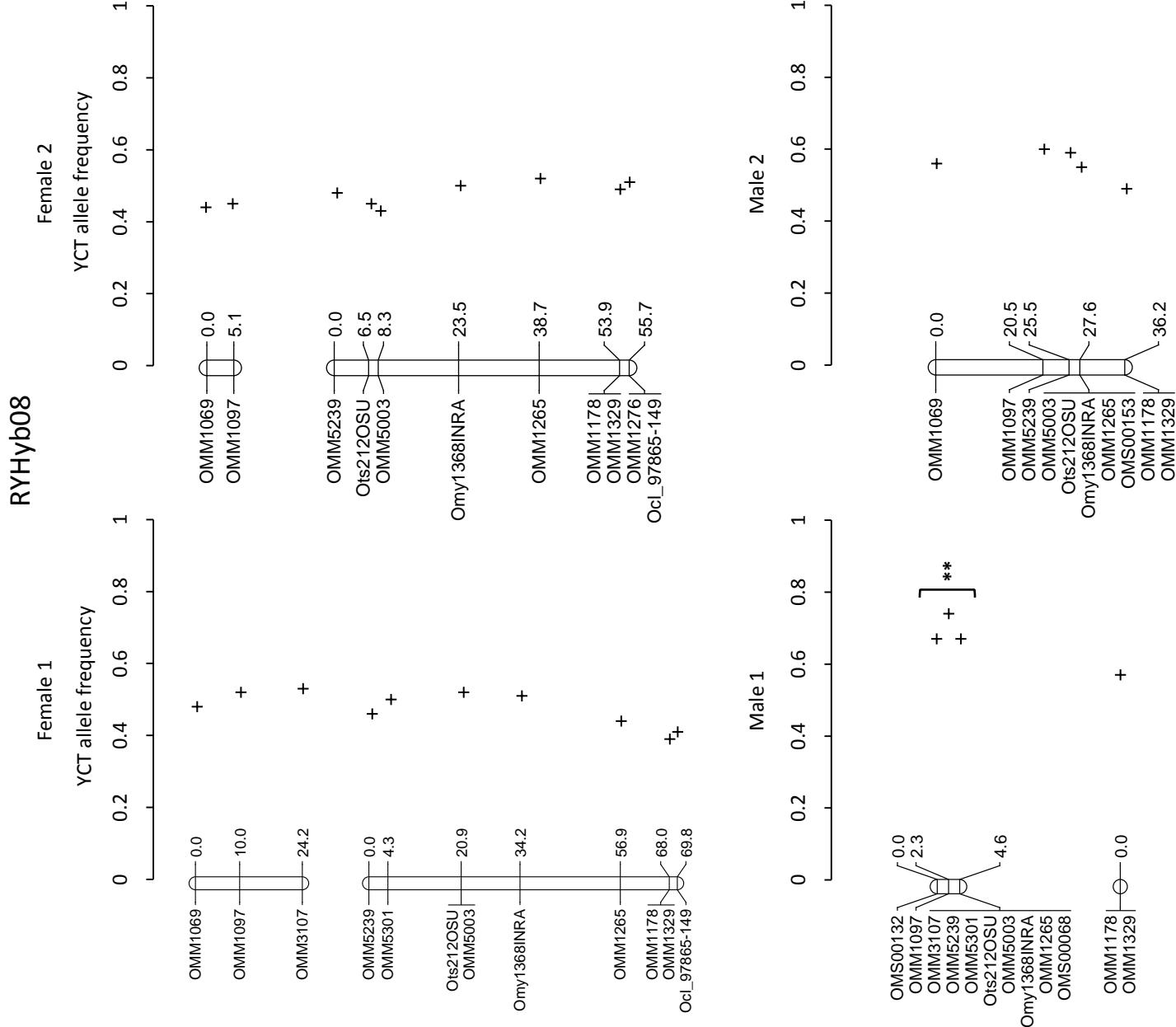
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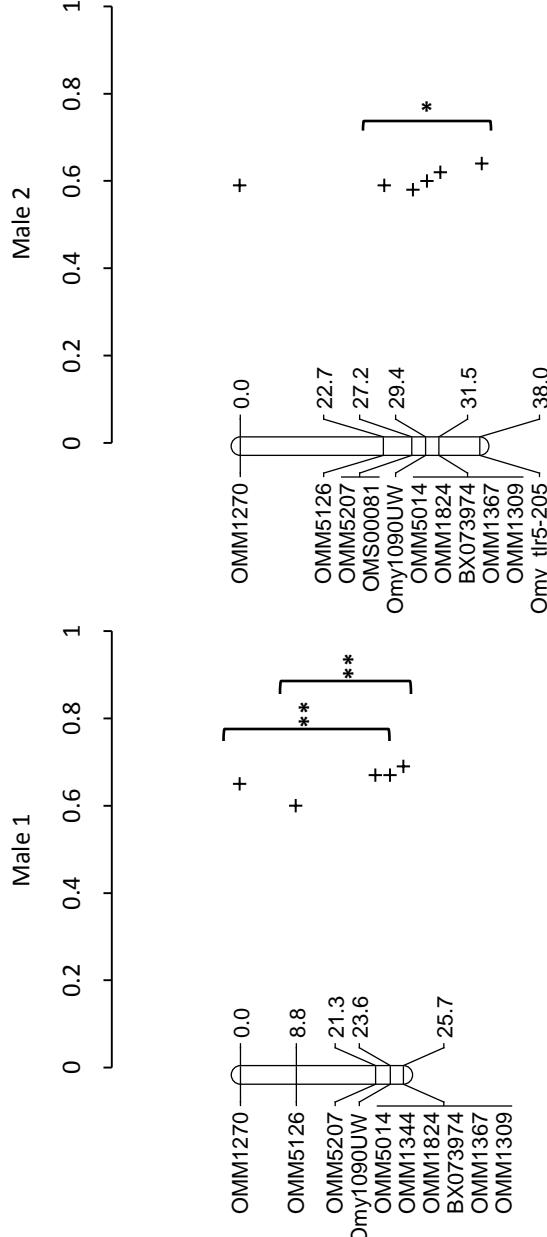
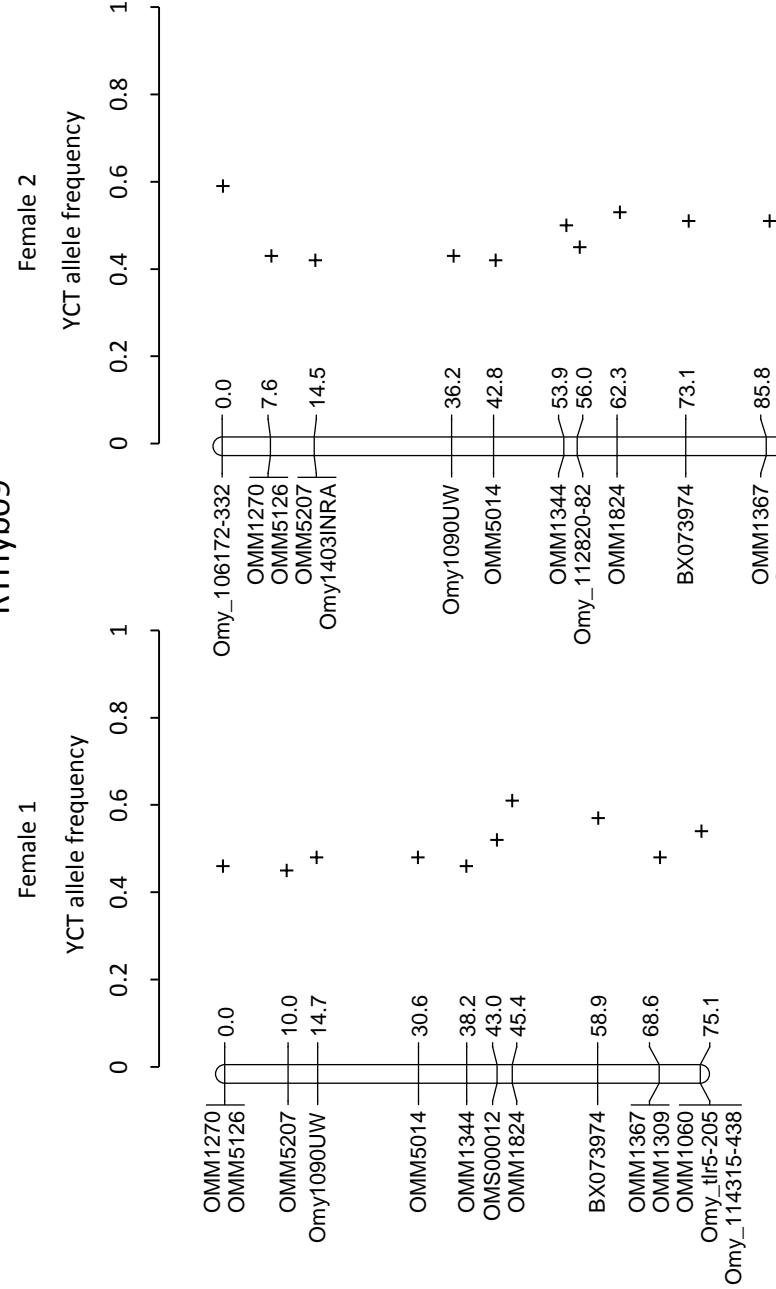
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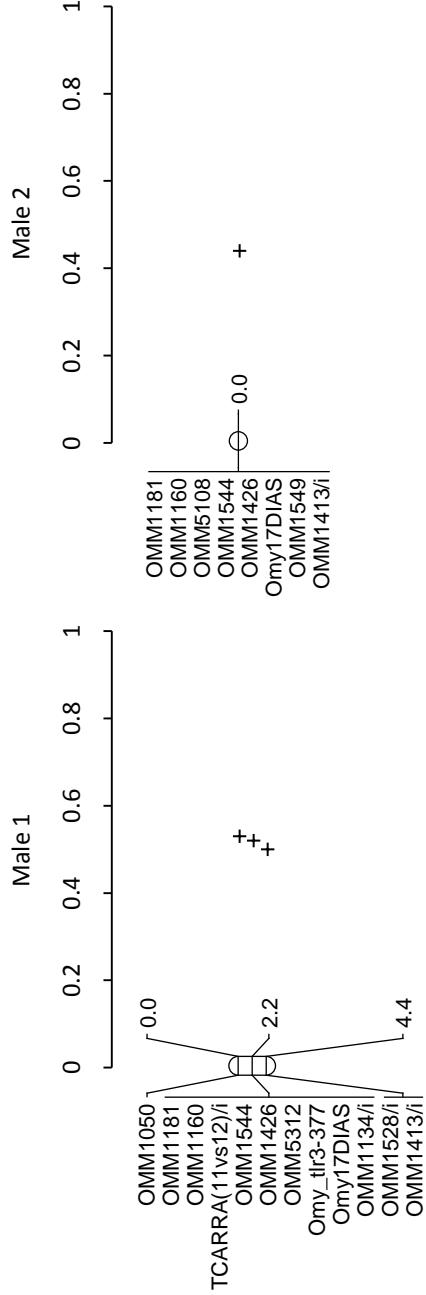
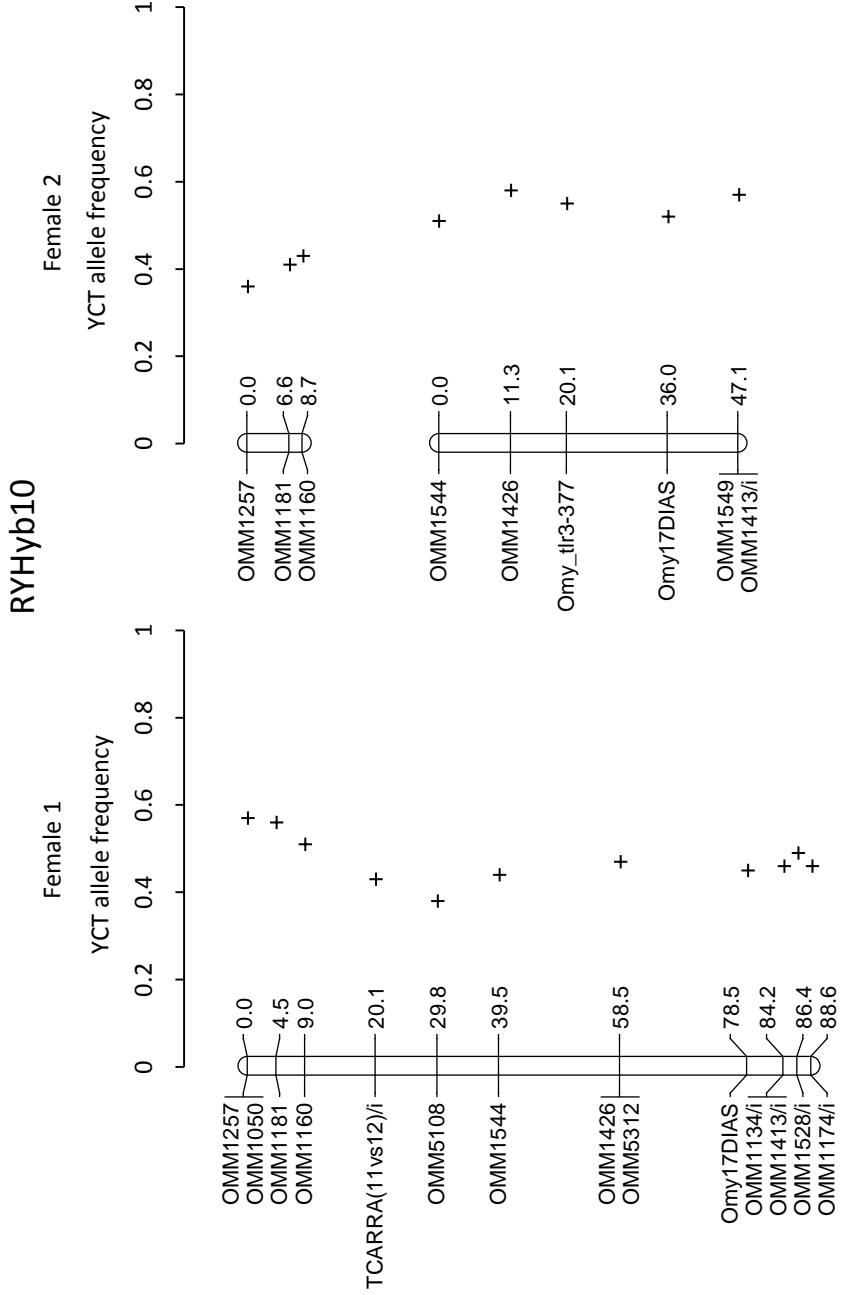


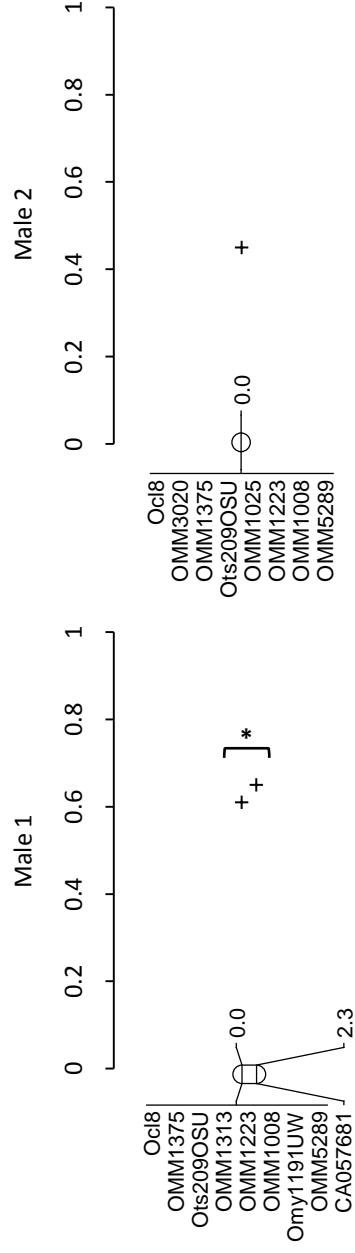
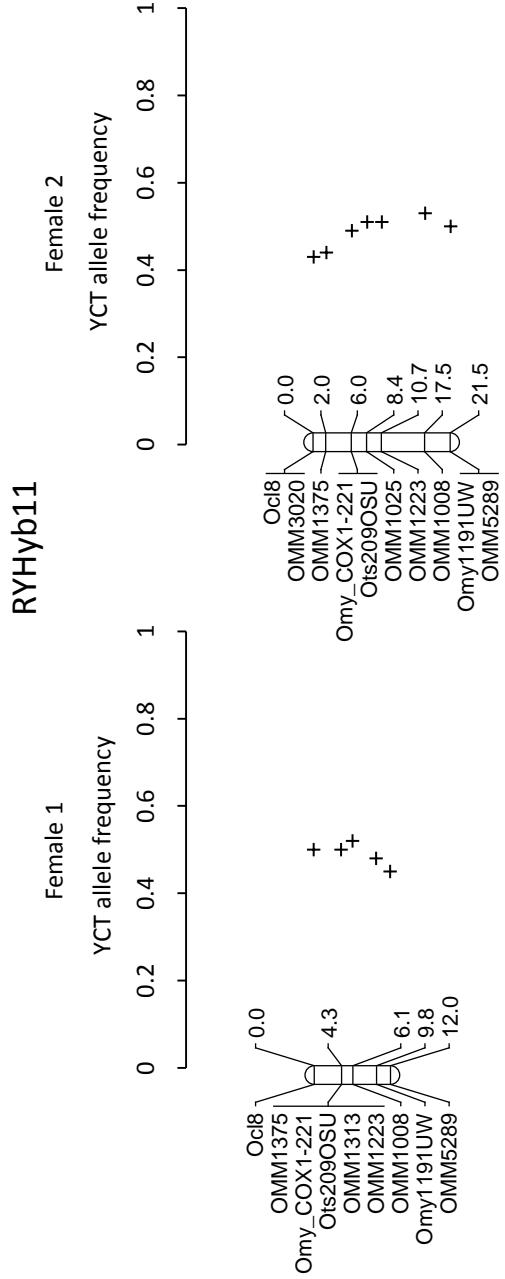


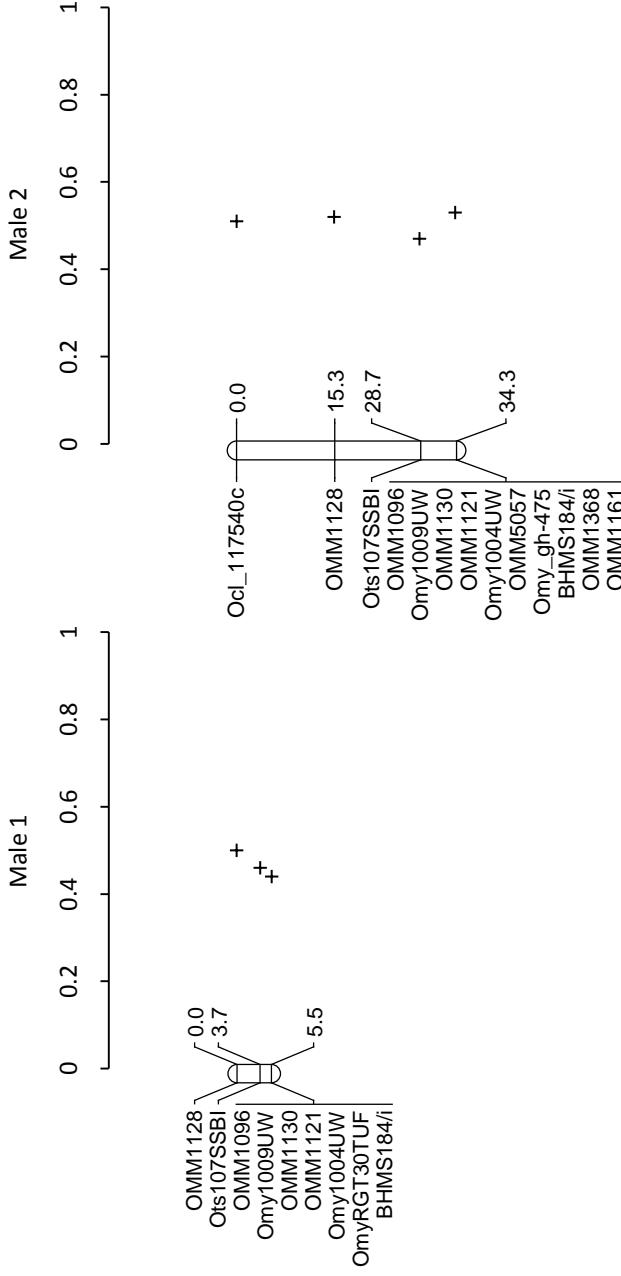
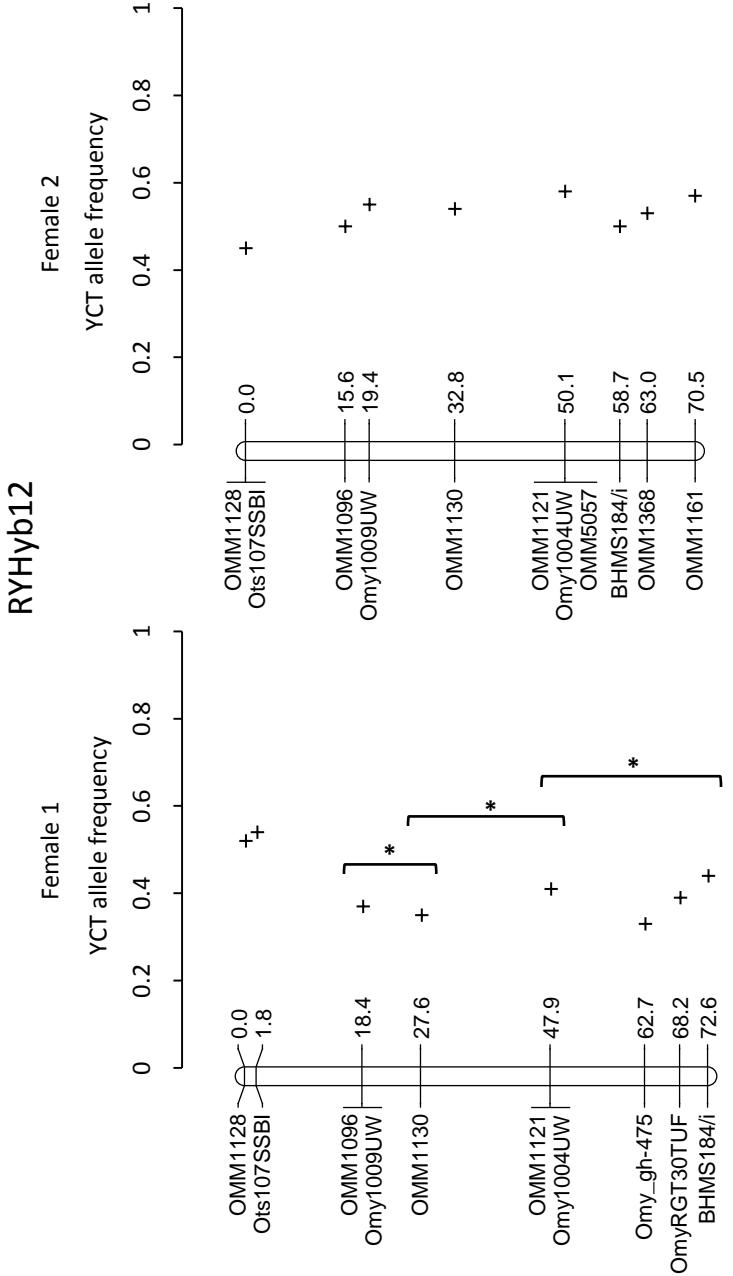


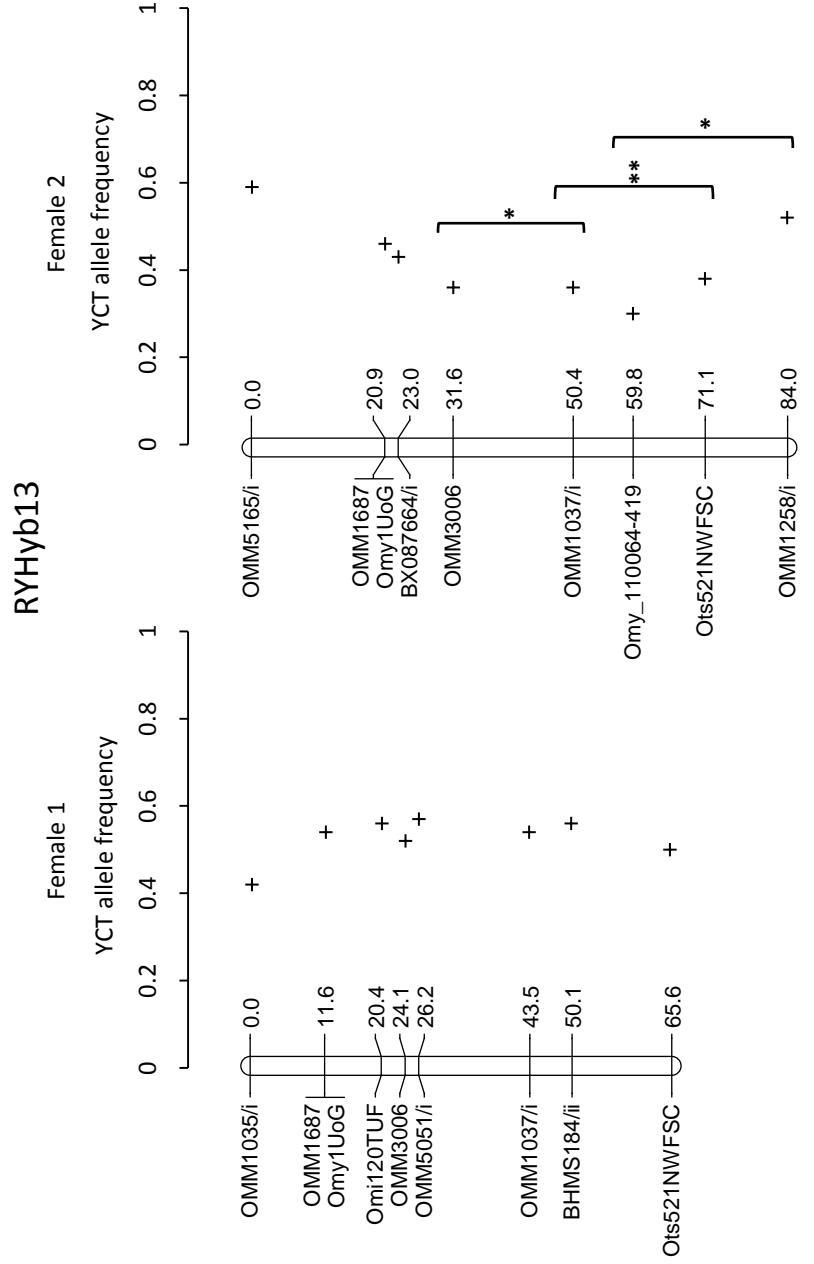
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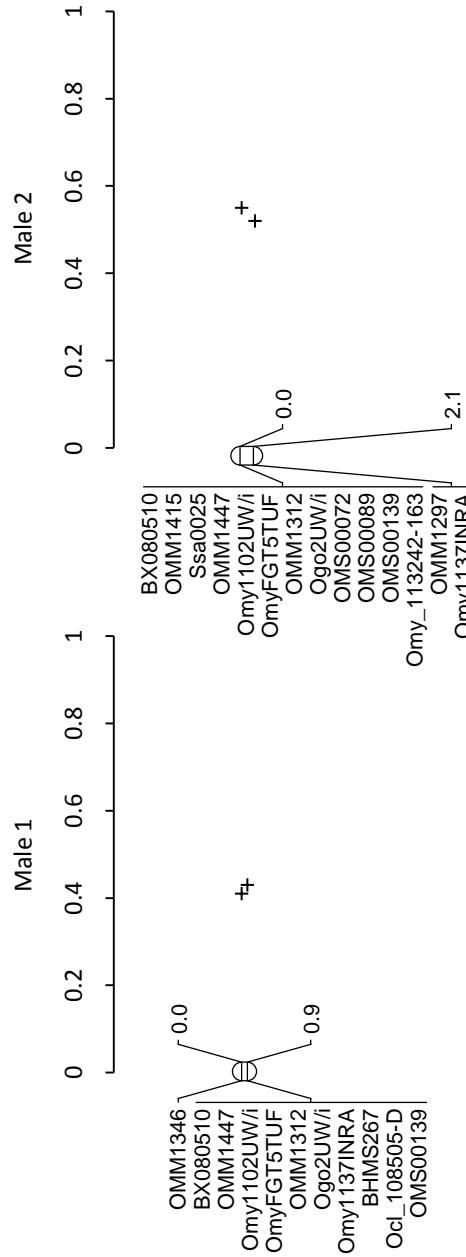
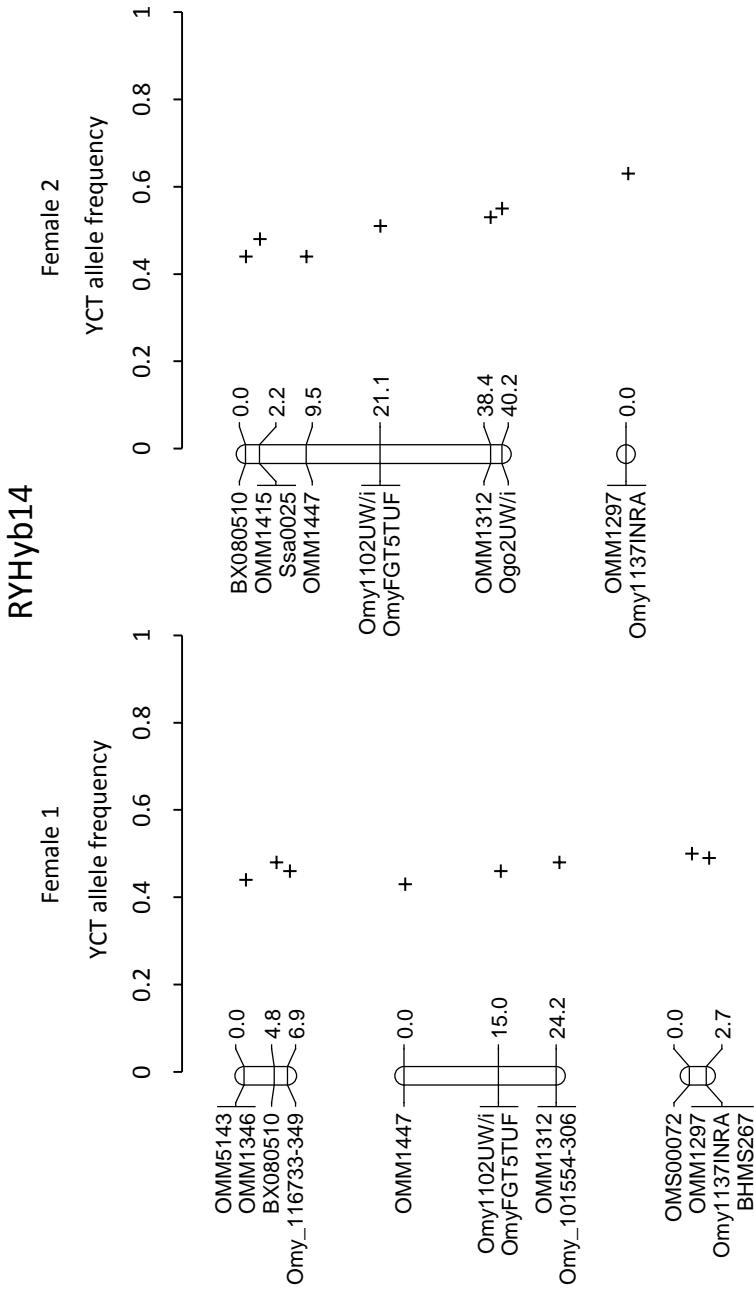


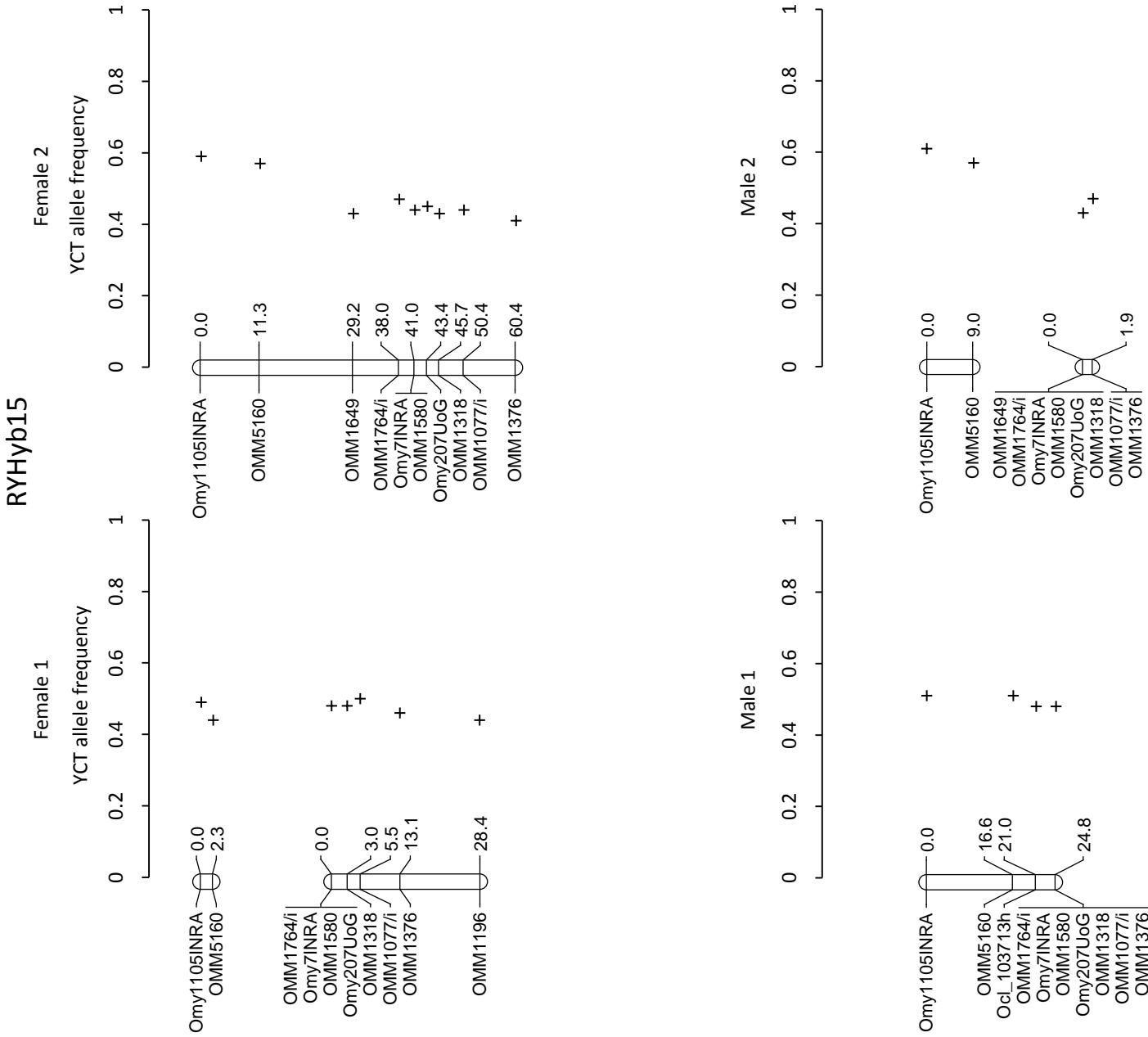




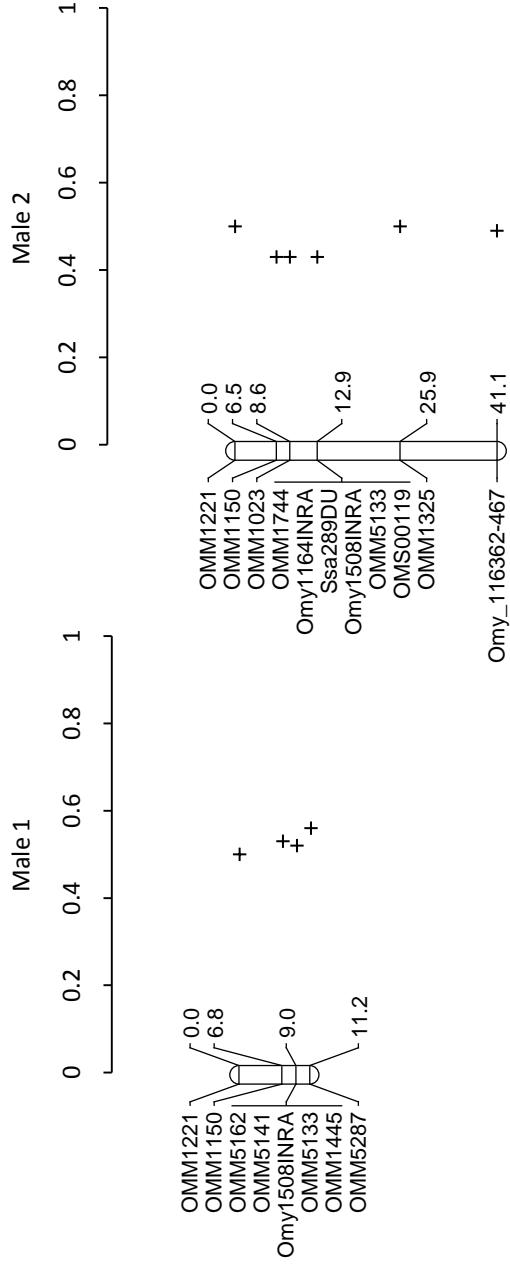
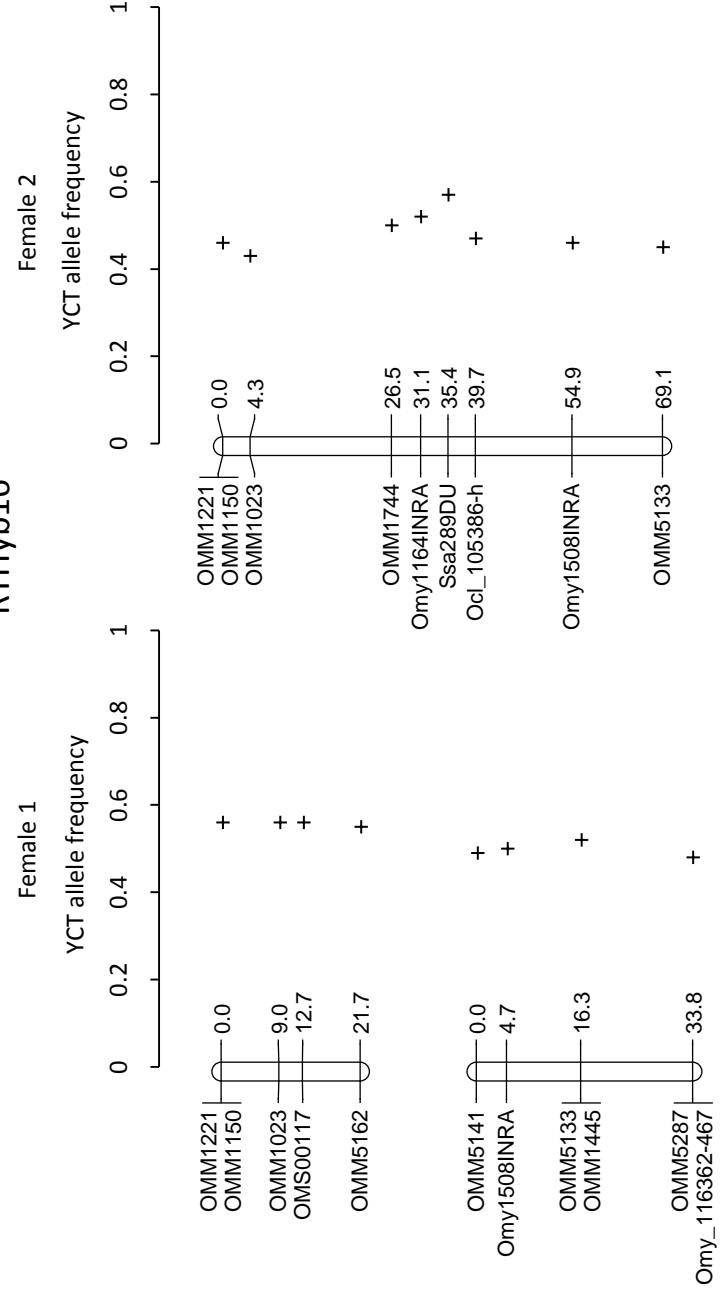




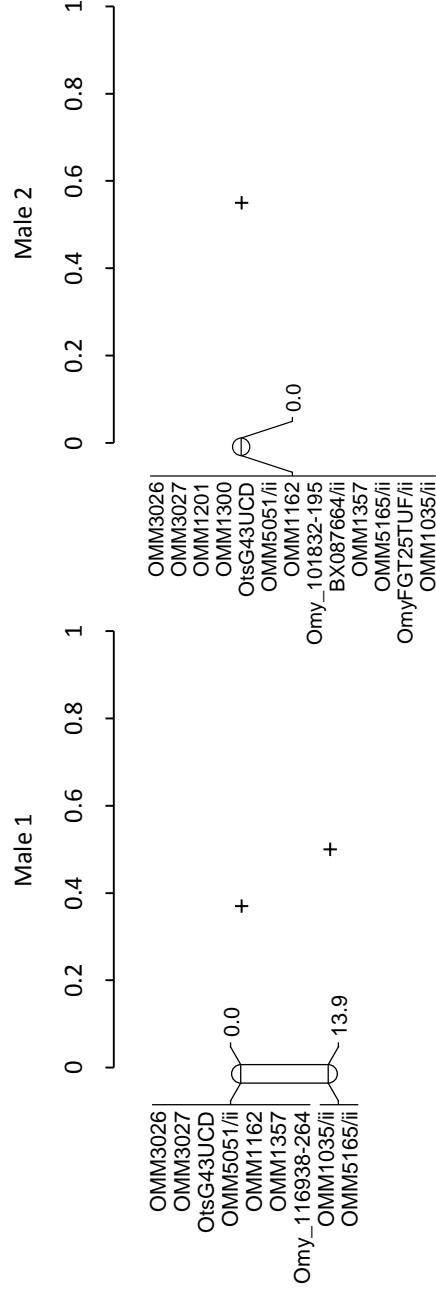
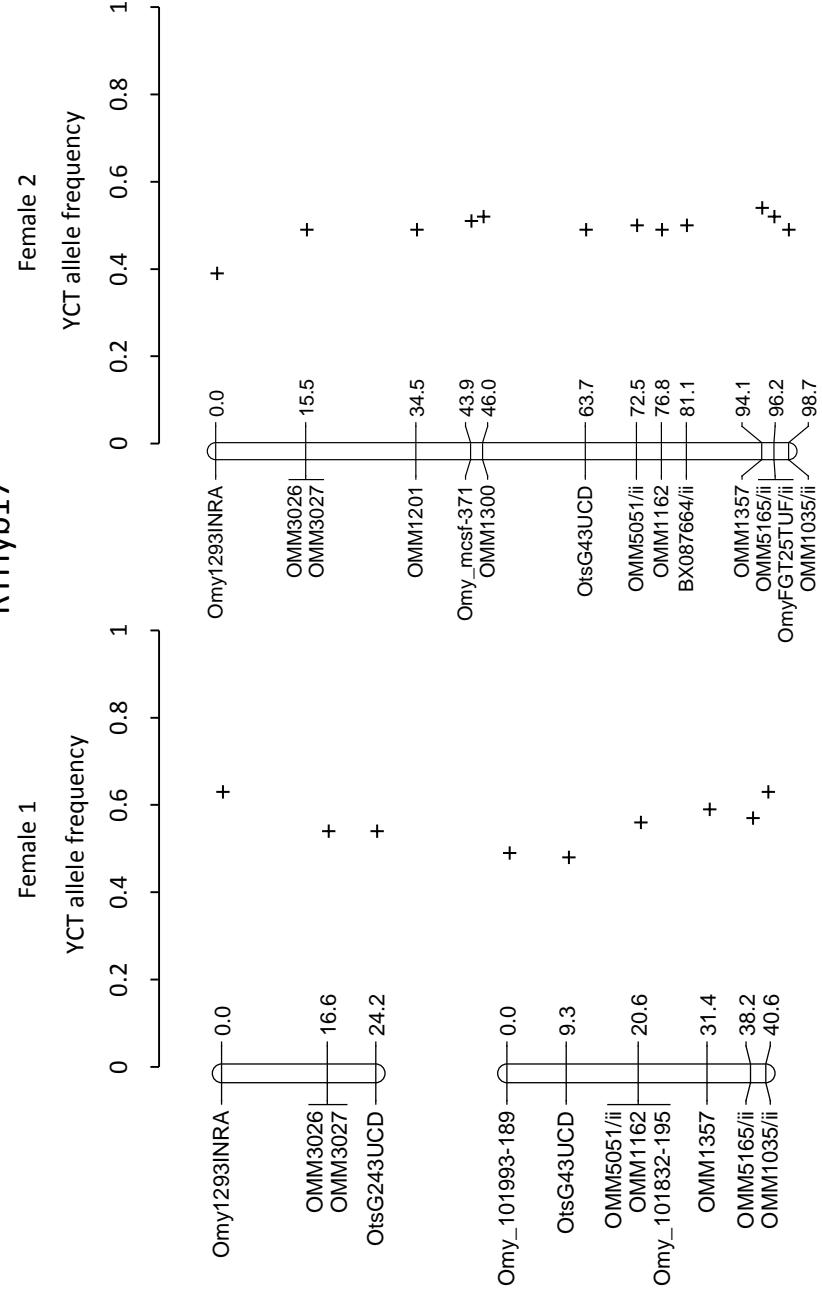


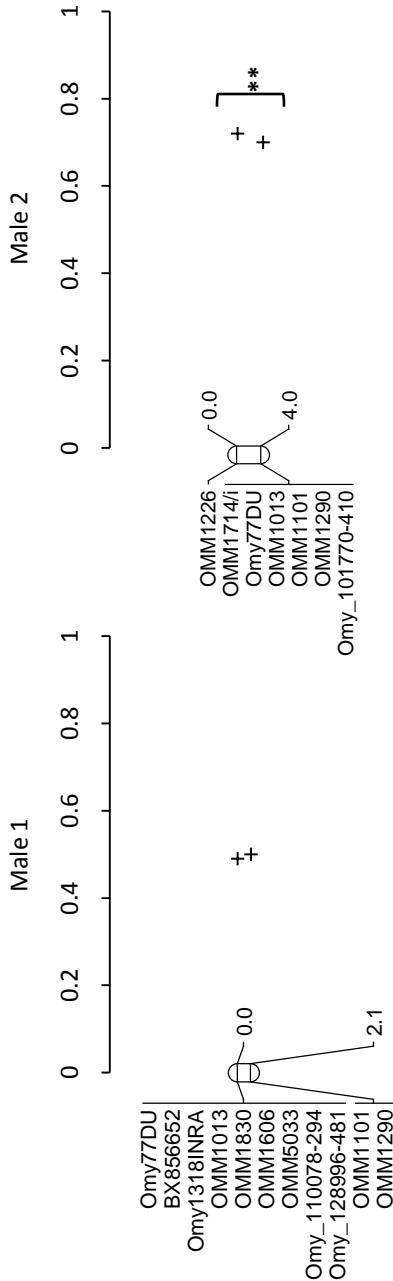
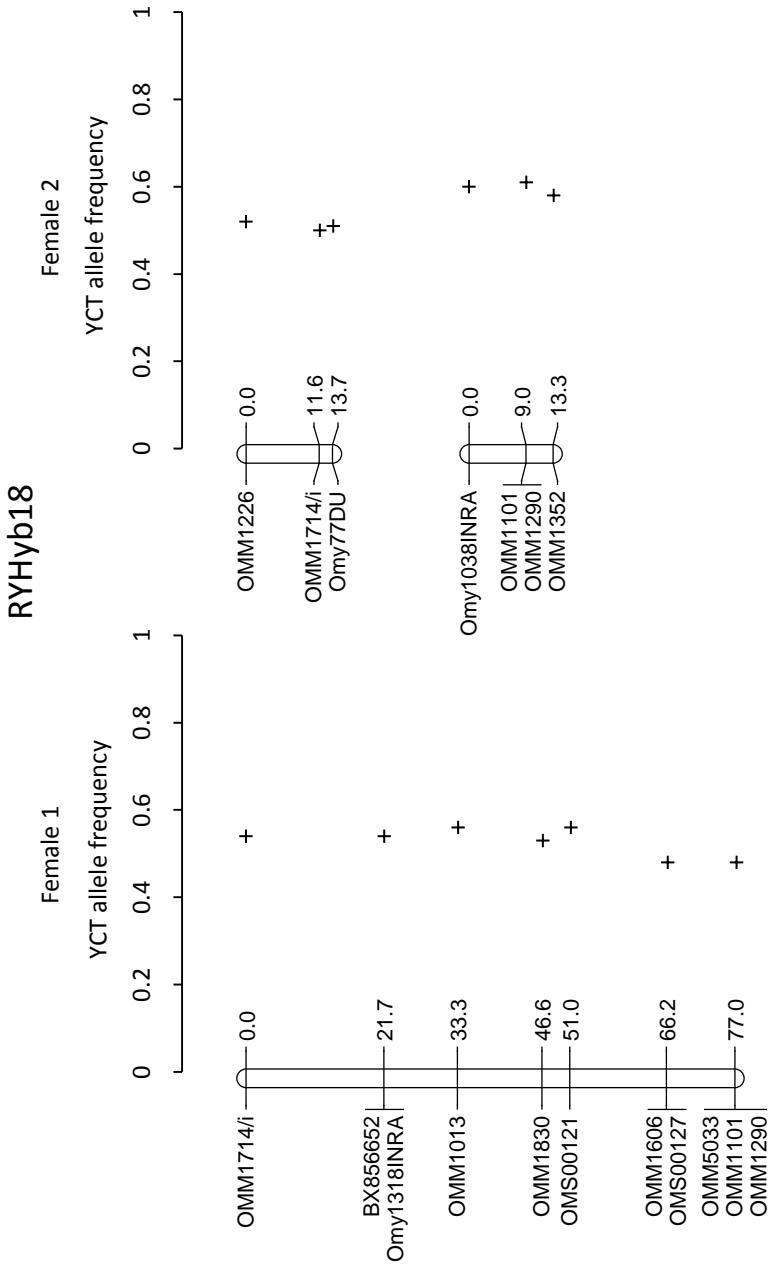


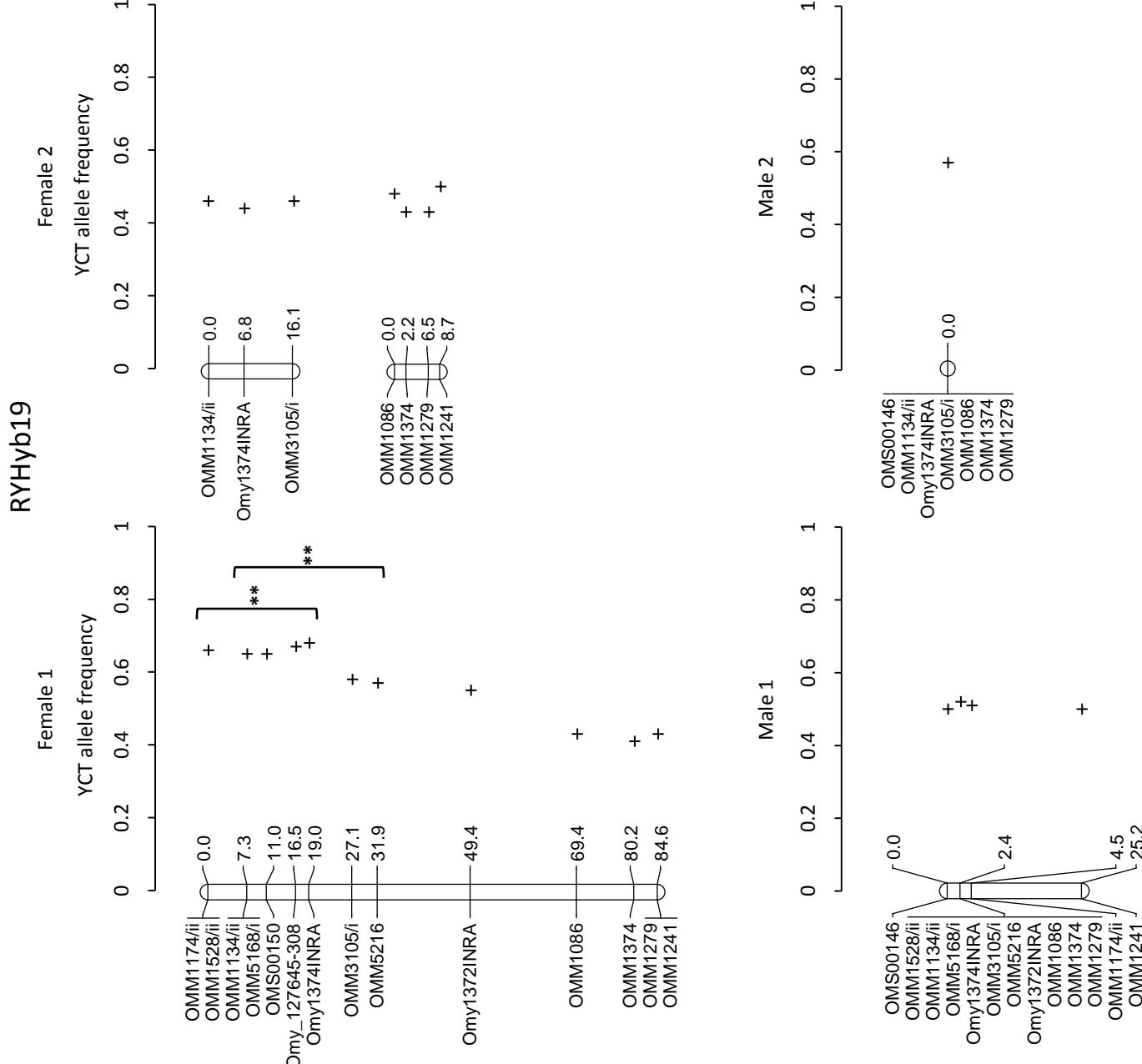
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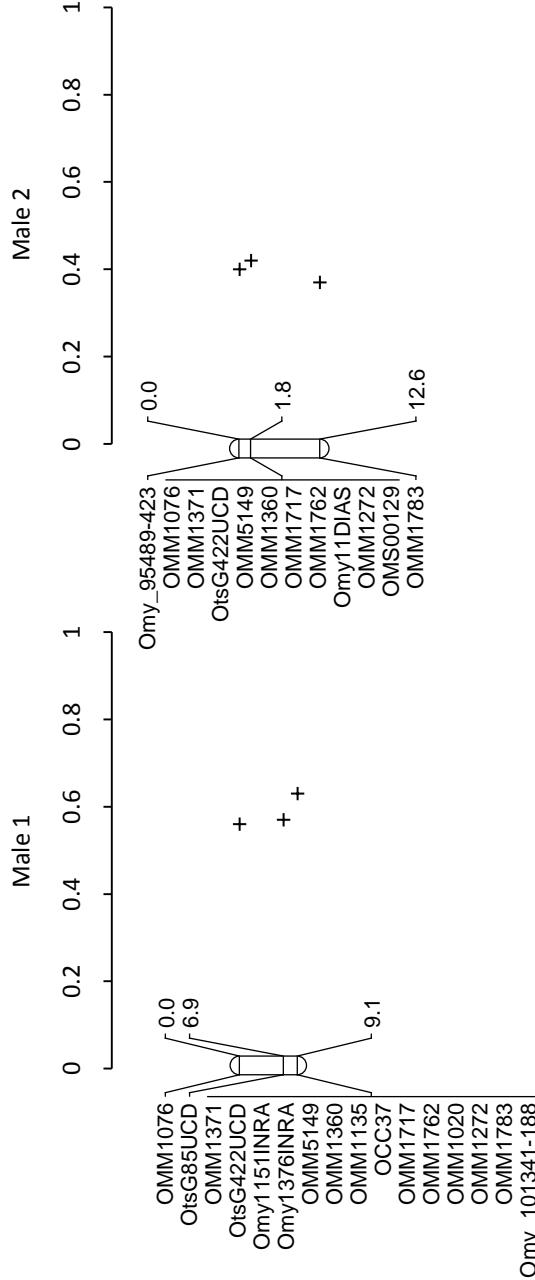
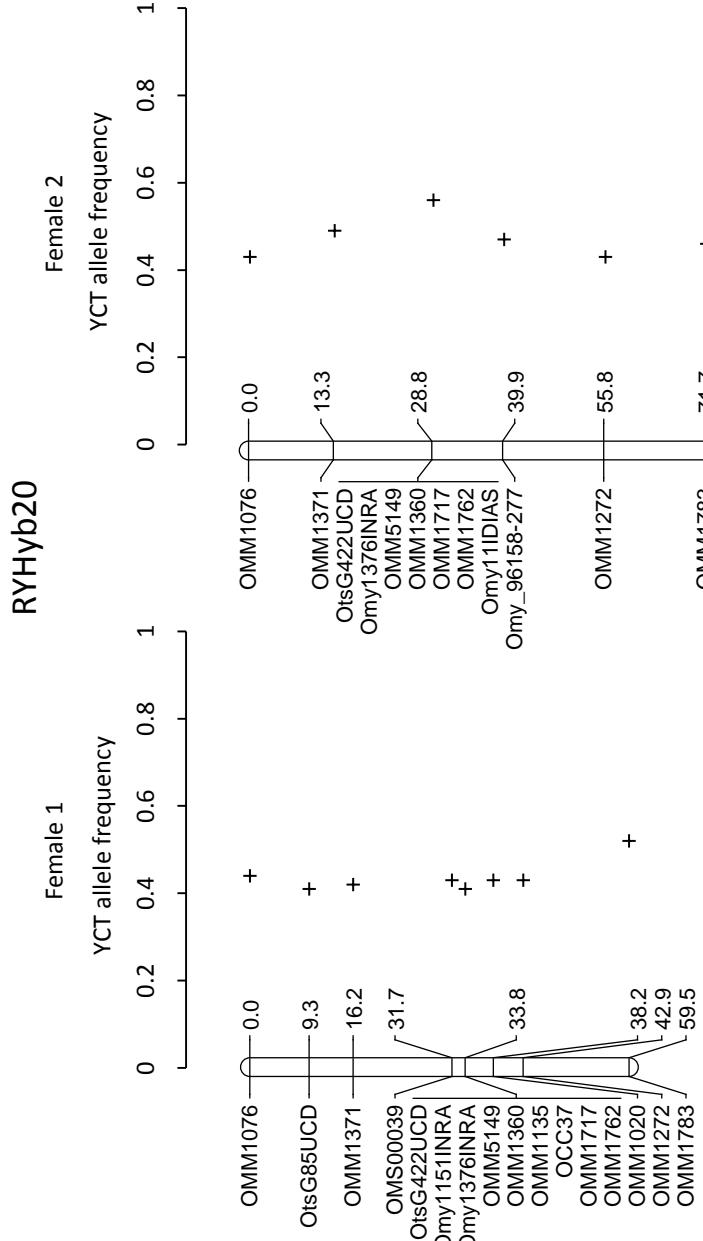


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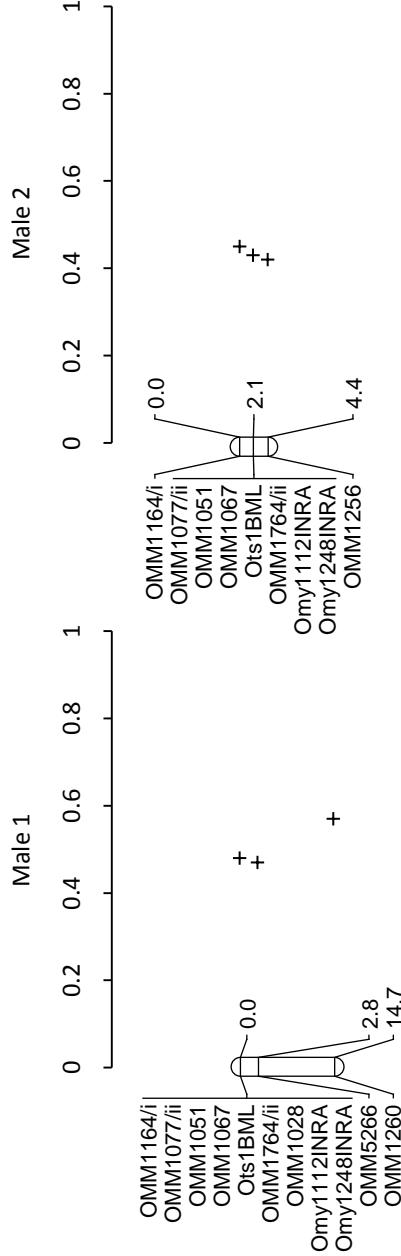
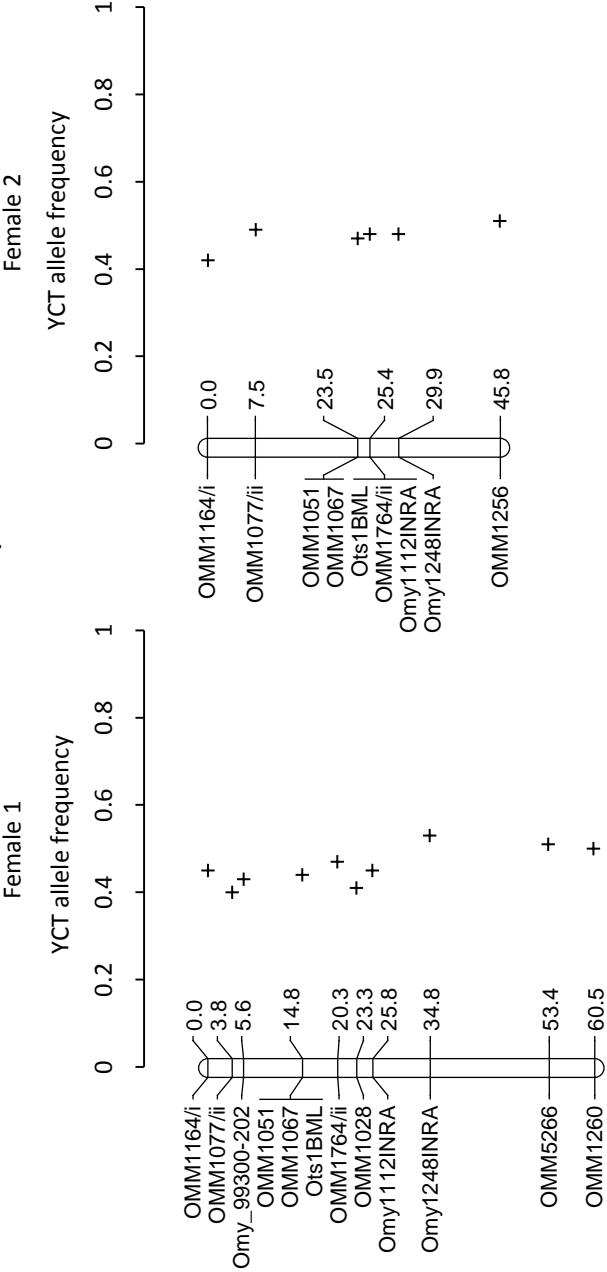




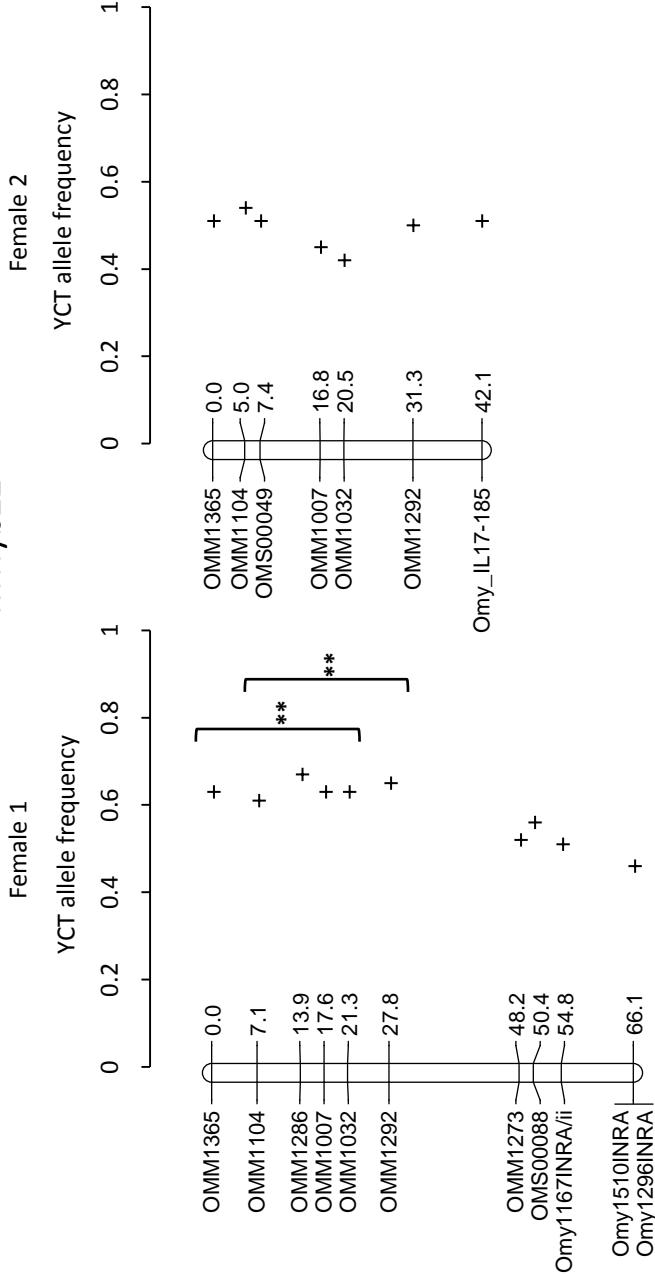


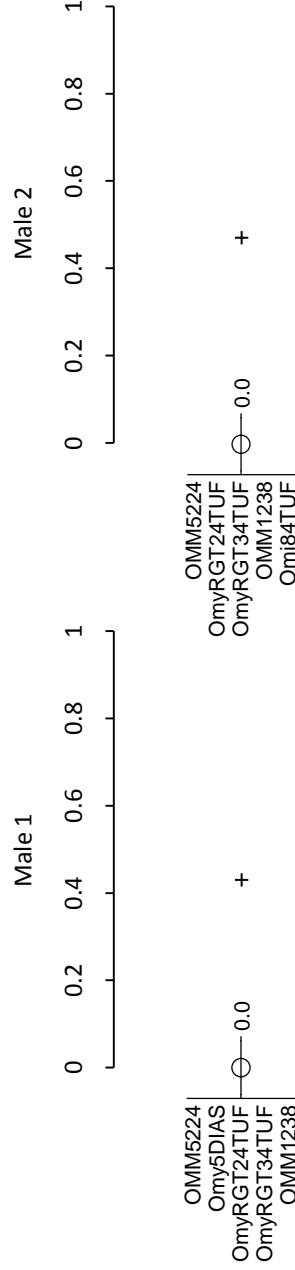
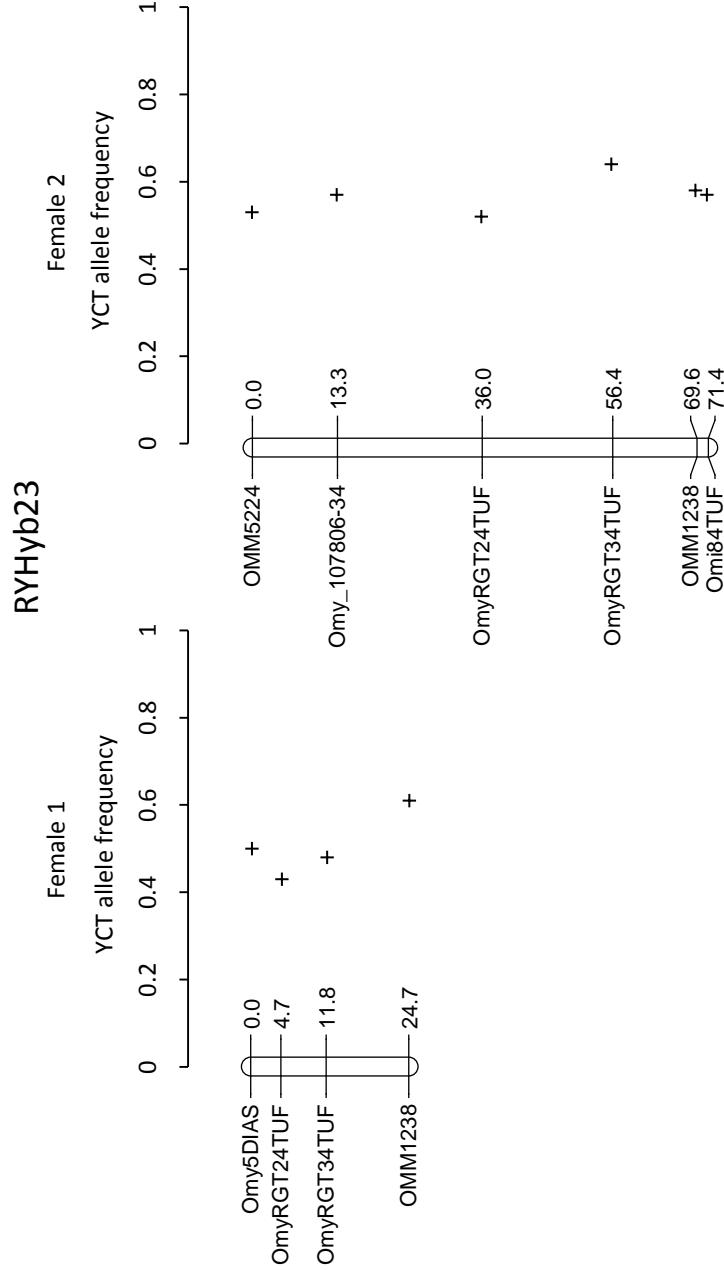


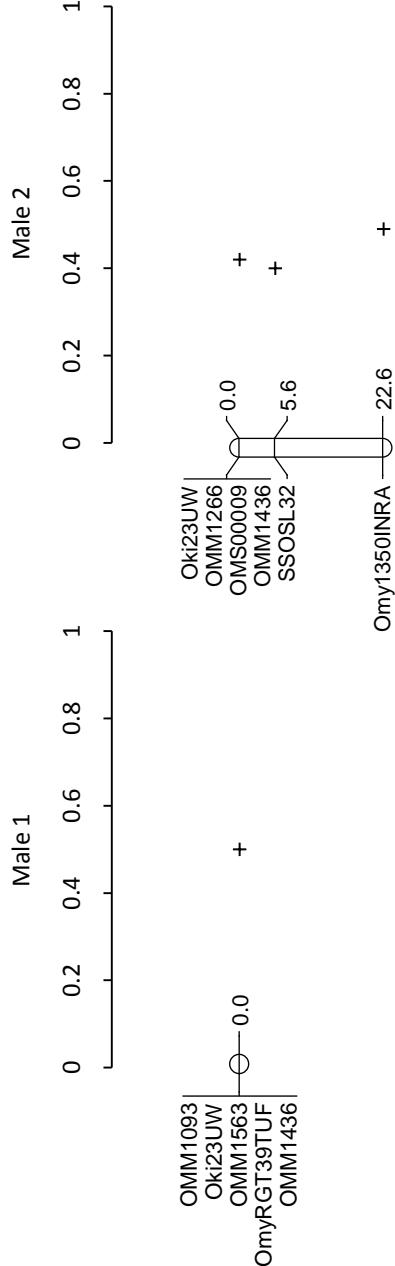
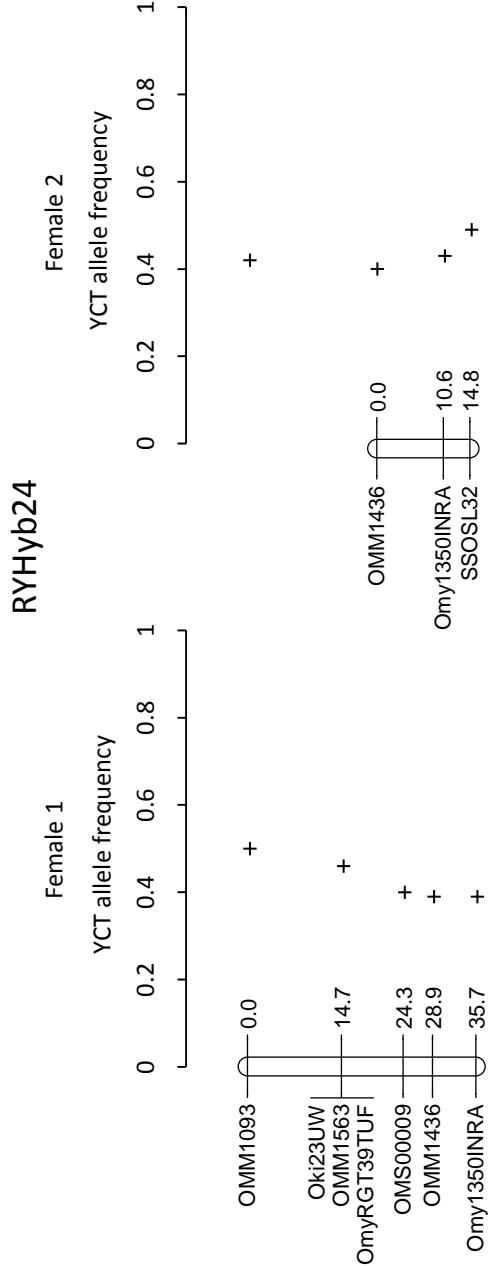
RYHyb21



RYHyb22







RYHyb25 and RYHyb29 polymorphism

