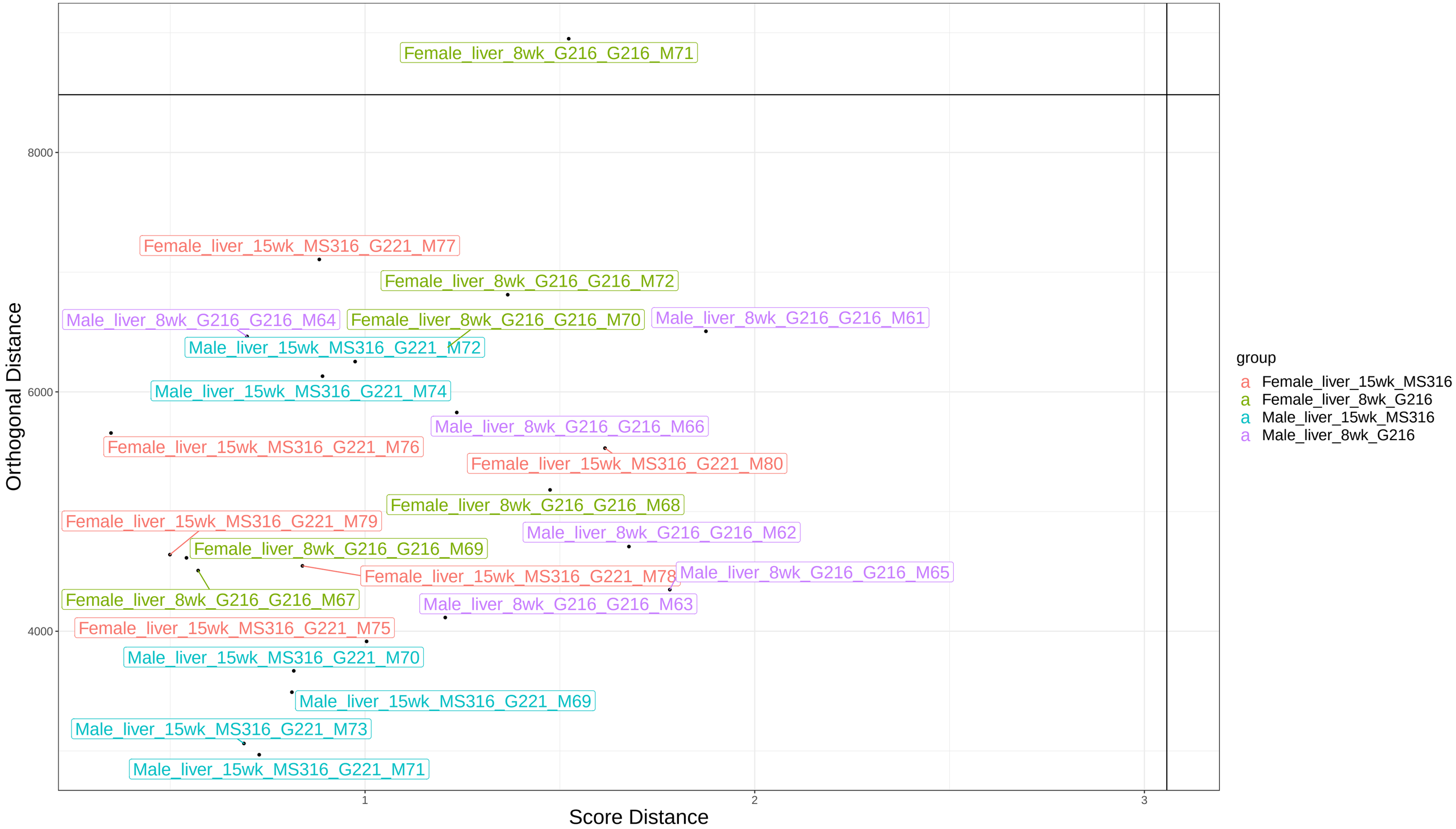
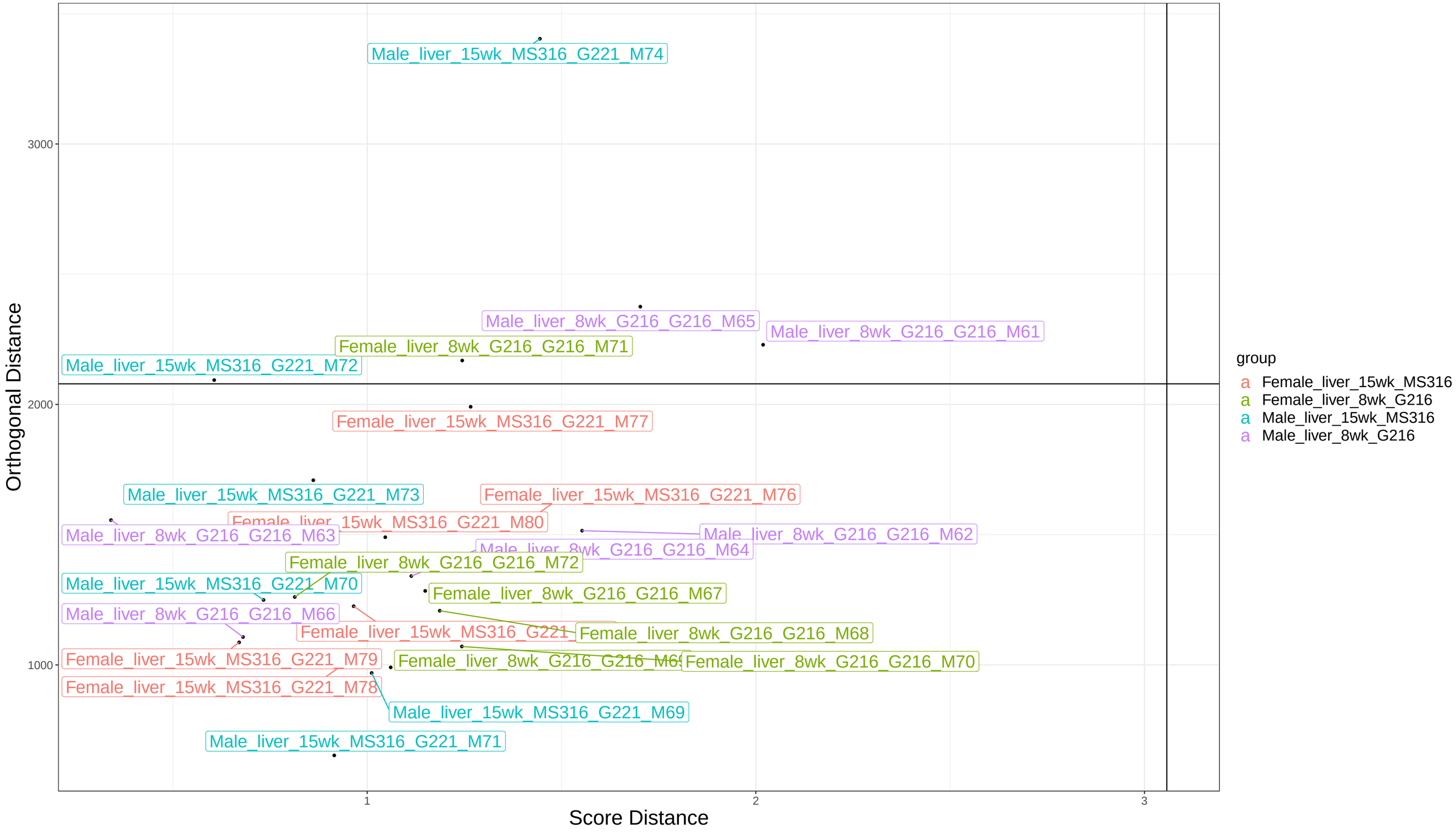
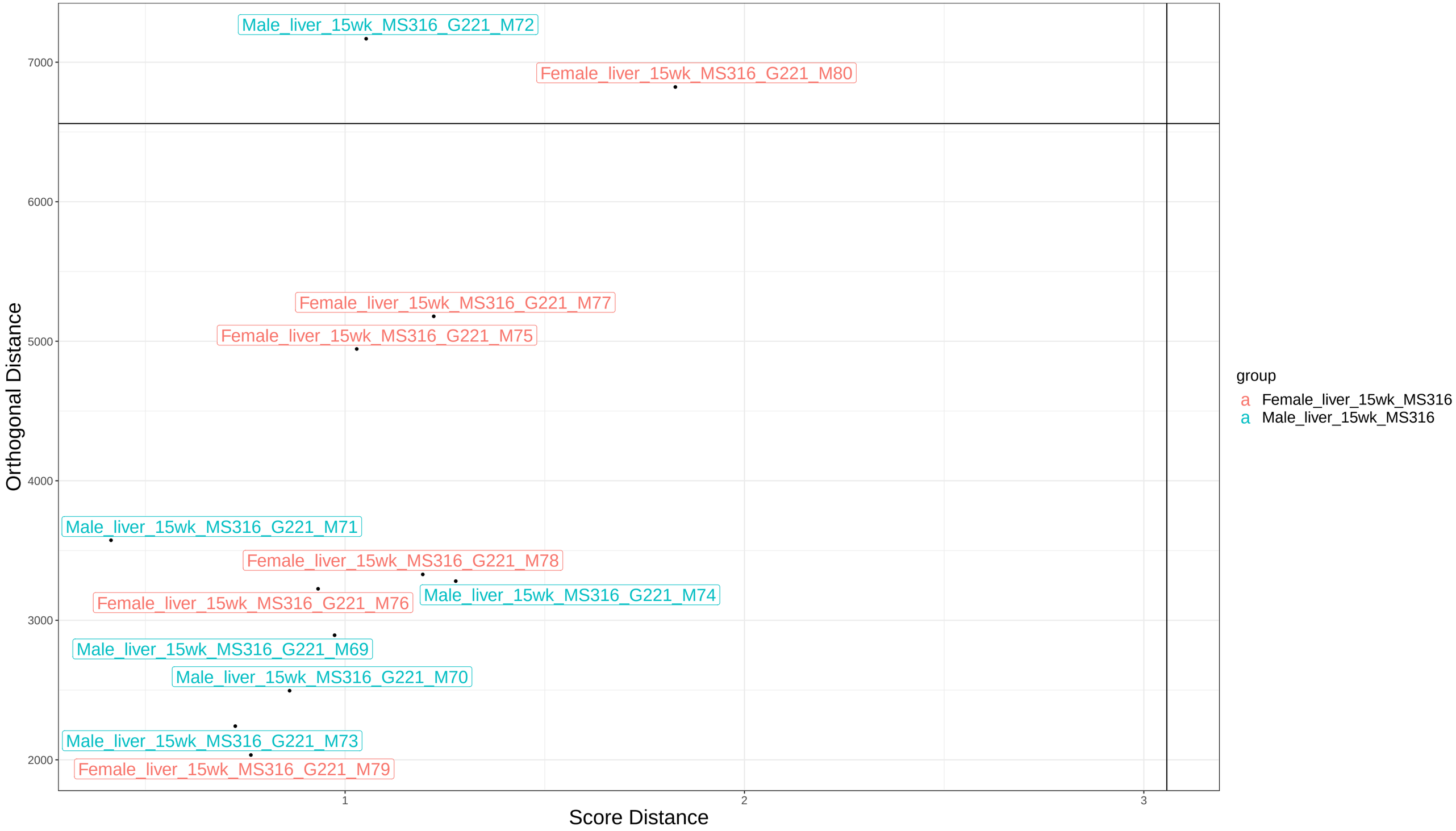
G221_RNASEQ_MS316, 0_RPCA_All_Merge,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798



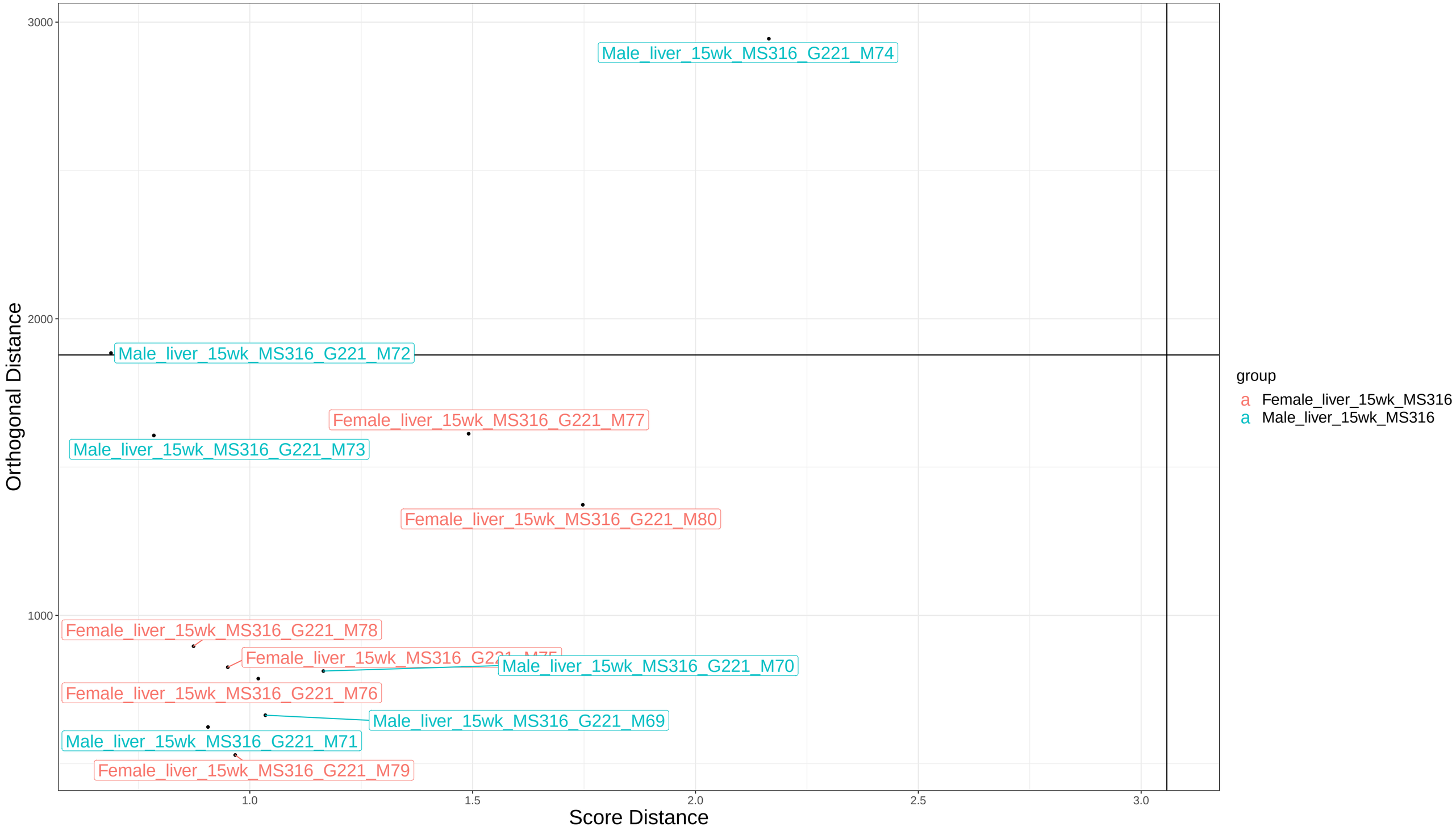
G221_RNASEQ_MS316, 0_RPCA_Significant_Union_Merge,
Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 739



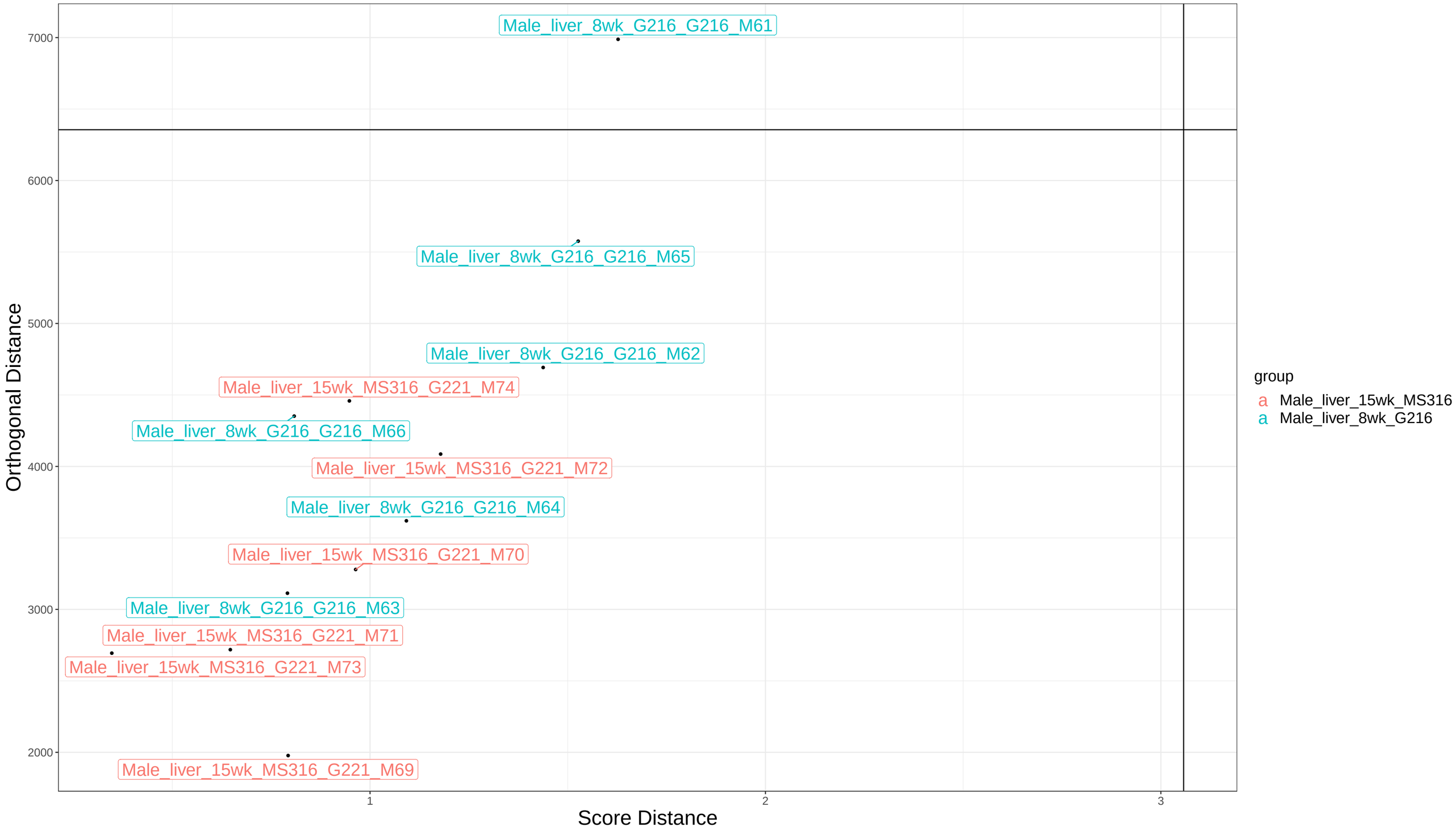
G221_RNASEQ_MS316, 1_RPCA_All_Female_liver_15wk_MS316_Male_liver_15wk_MS316,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798



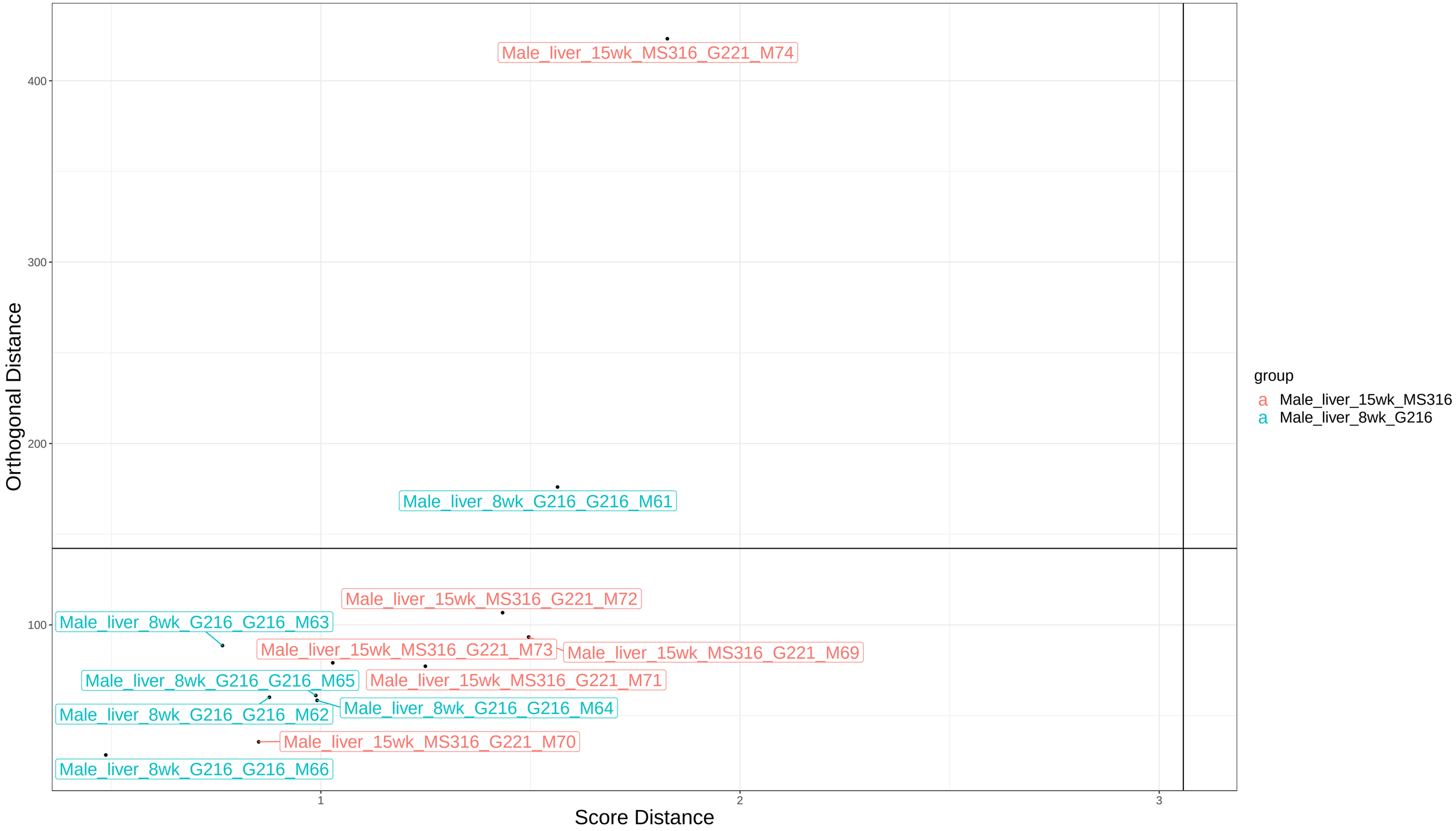
G221_RNASEQ_MS316, 1_RPCA_Significant_Female_liver_15wk_MS316_Male_liver_15wk_MS316,
Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 587



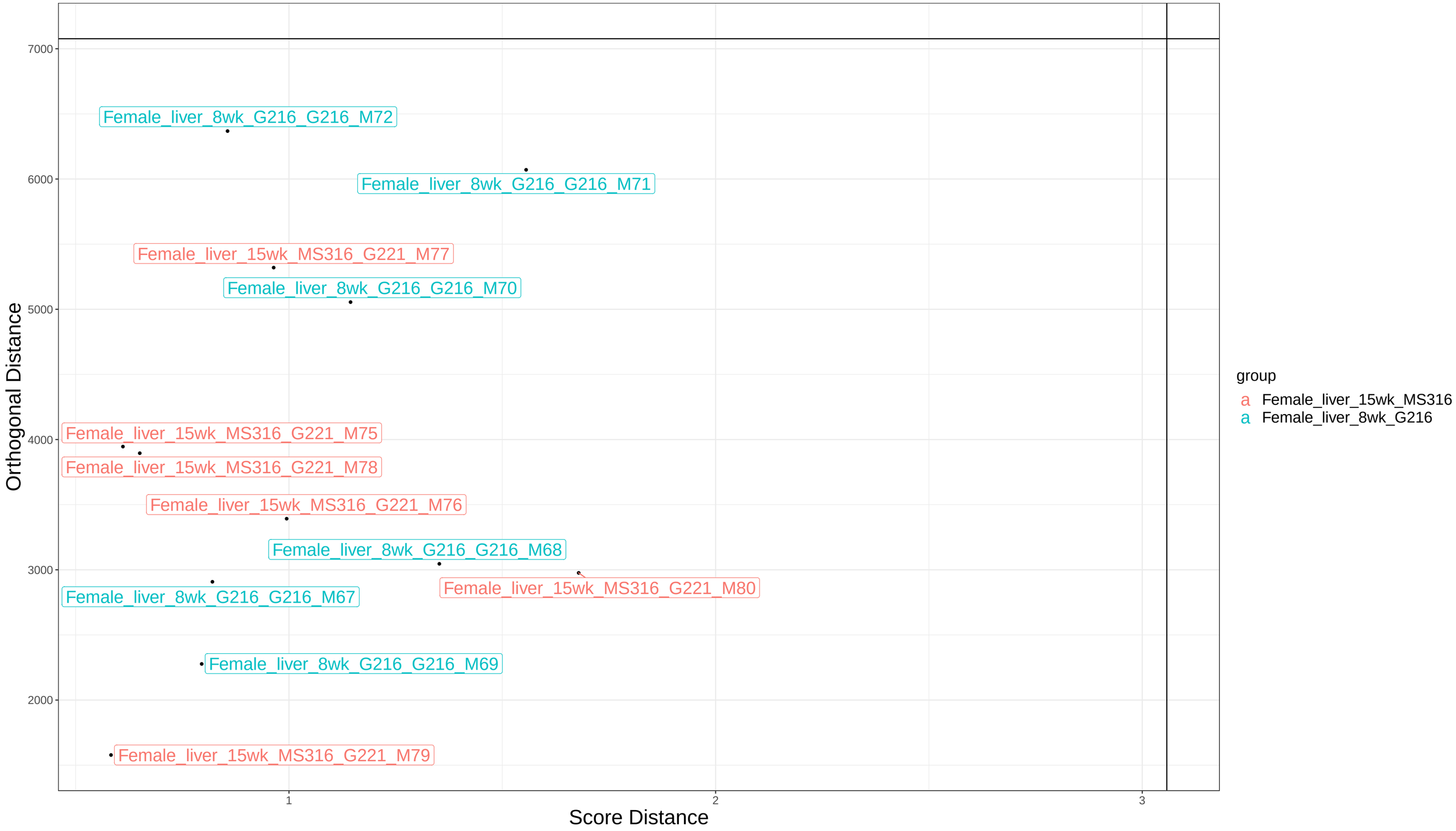
G221_RNASEQ_MS316, 2_RPCA_All_Male_liver_8wk_G216_Male_liver_15wk_MS316,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798



G221_RNASEQ_MS316, 2_RPCA_Significant_Male_liver_8wk_G216_Male_liver_15wk_MS316,
Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 110



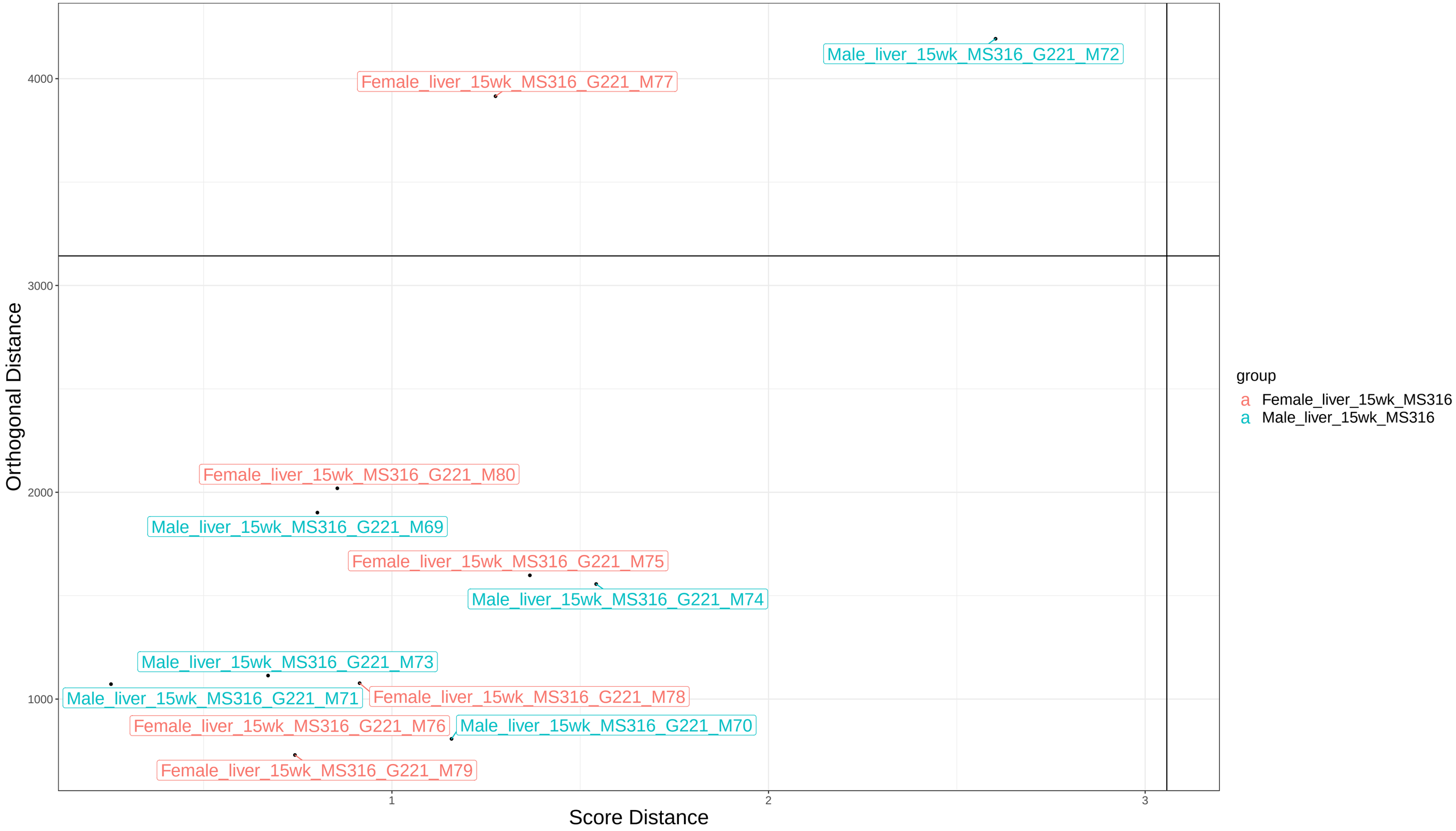
G221_RNASEQ_MS316, 3_RPCA_All_Female_liver_8wk_G216_Female_liver_15wk_MS316,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798



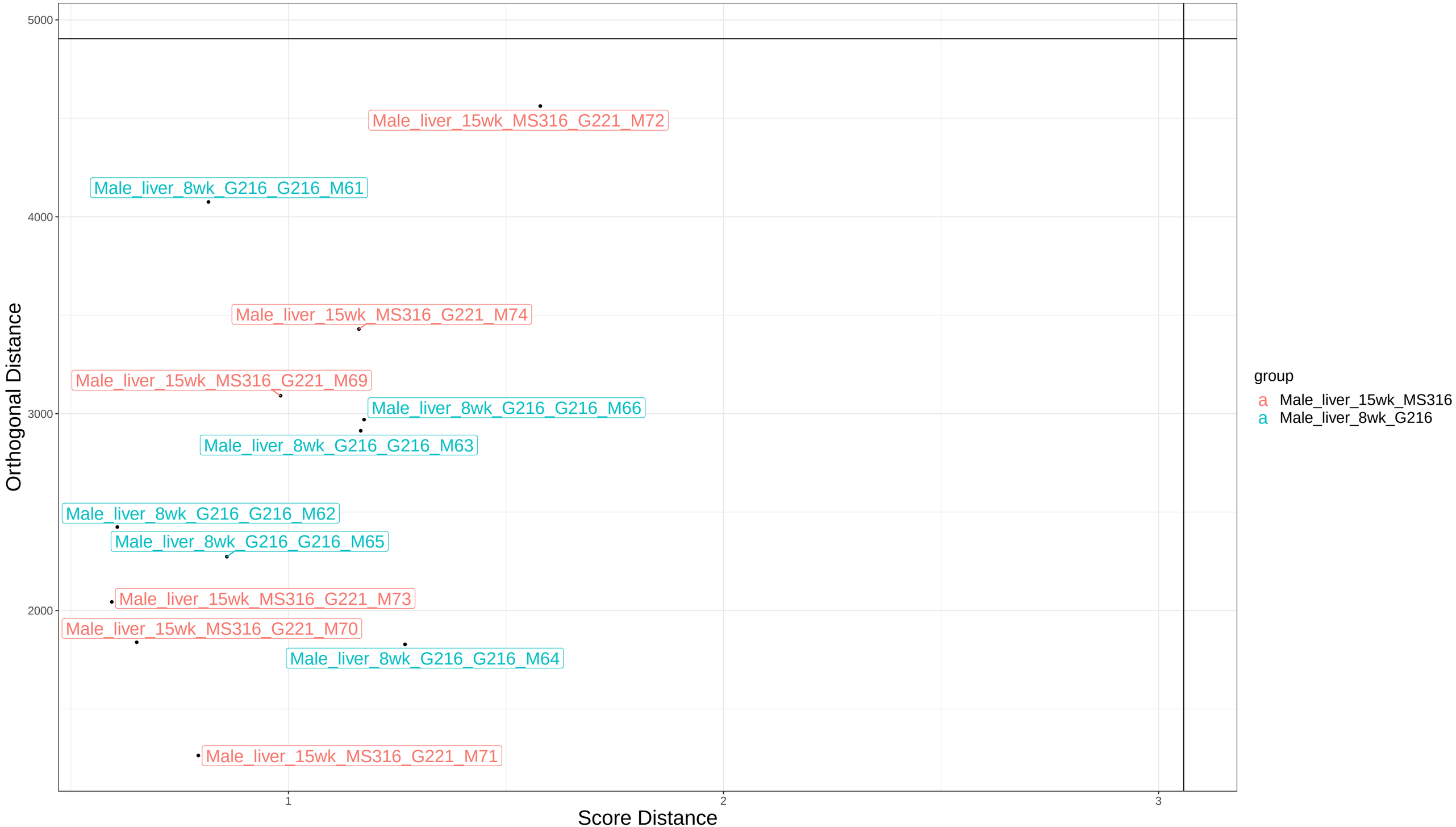
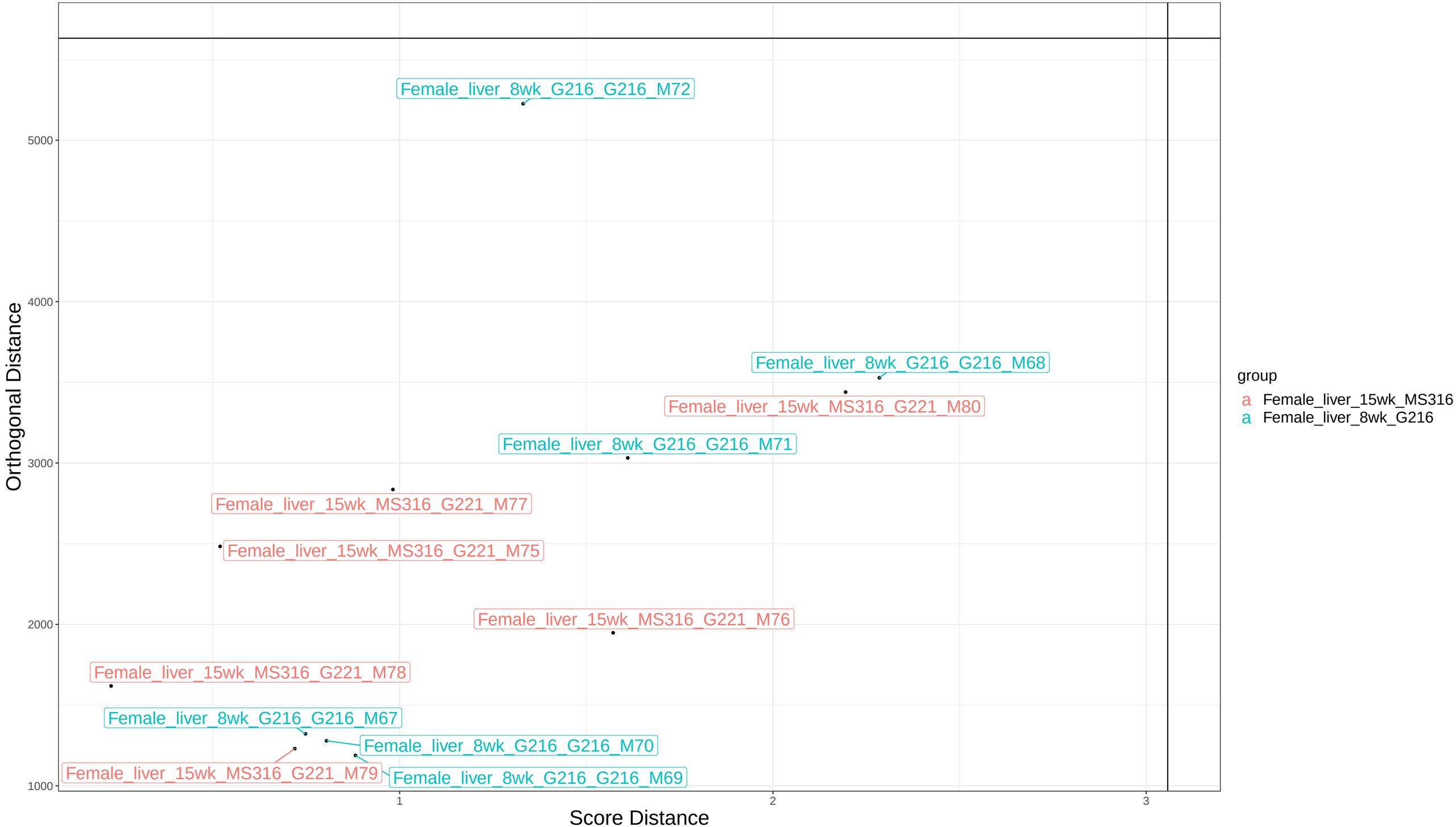
G221_RNASEQ_MS316, 3_RPCA_Significant_Female_liver_8wk_G216_Female_liver_15wk_MS316,
Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 102

G221_RNASEQ_MS316, 0_RPCA_Non−significant_Intersection_Merge,
Non−significant genes (1.2 < |FC| < 1/|1.2| and FDR >0.1, RPKM >1), RefSeqLncRNA76k_Genes: 4677

G221_RNASEQ_MS316, 1_RPCA_Non−significant_Female_liver_15wk_MS316_Male_liver_15wk_MS316,
Non−significant genes (1.2 < |FC| < 1/|1.2| and FDR >0.1, RPKM >1), RefSeqLncRNA76k_Genes: 7208



G221_RNASEQ_MS316, 2_RPCA_Non−significant_Male_liver_8wk_G216_Male_liver_15wk_MS316,
Non−significant genes (1.2 < |FC| < 1/|1.2| and FDR >0.1, RPKM >1), RefSeqLncRNA76k_Genes: 8272



G221_RNASEQ_MS316, 3_RPCA_Non−significant_Female_liver_8wk_G216_Female_liver_15wk_MS316,
Non−significant genes (1.2 < |FC| < 1/|1.2| and FDR >0.1, RPKM >1), RefSeqLncRNA76k_Genes: 8723