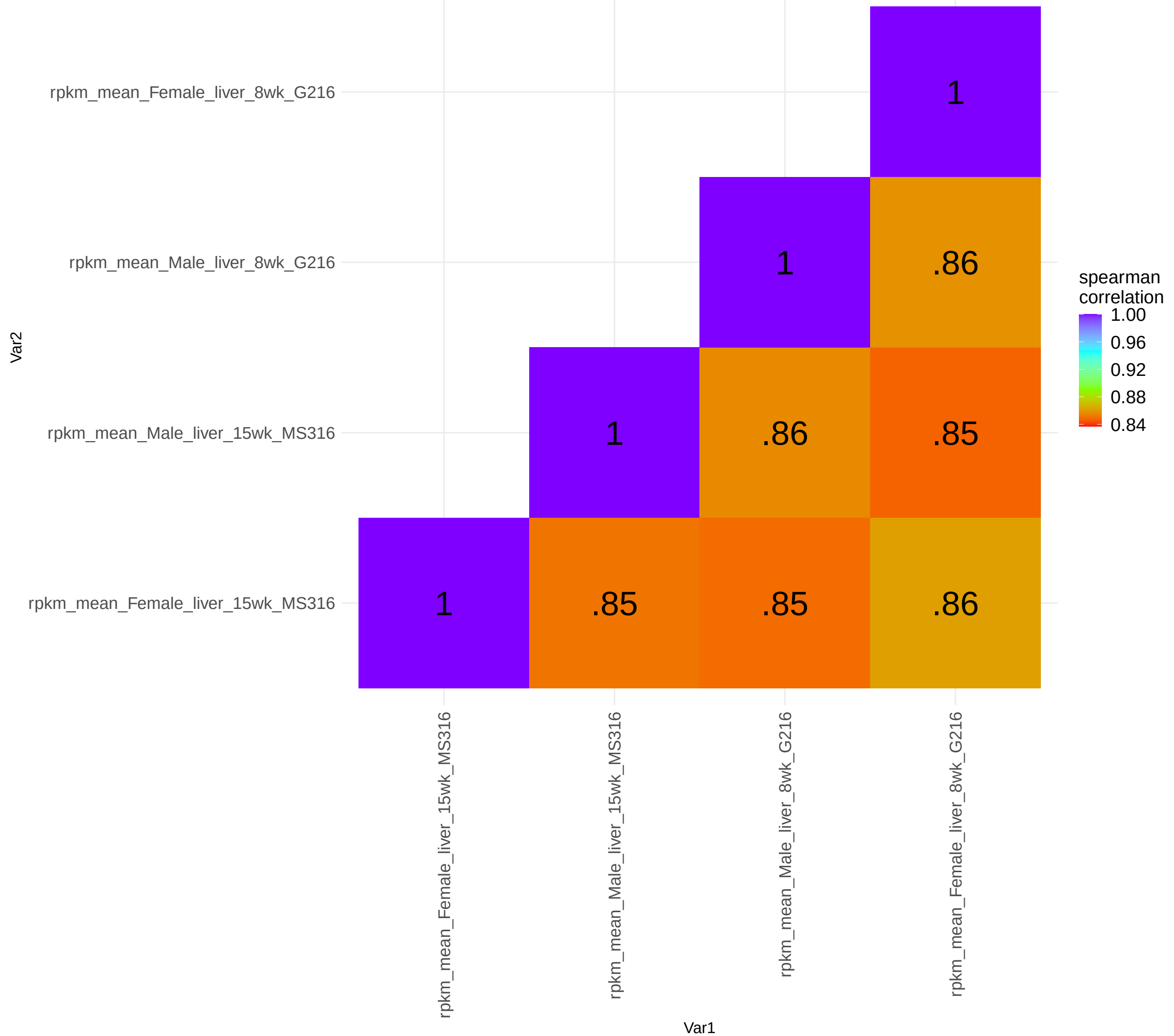
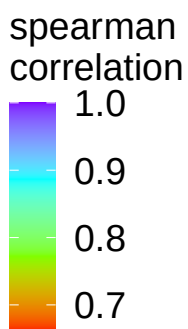
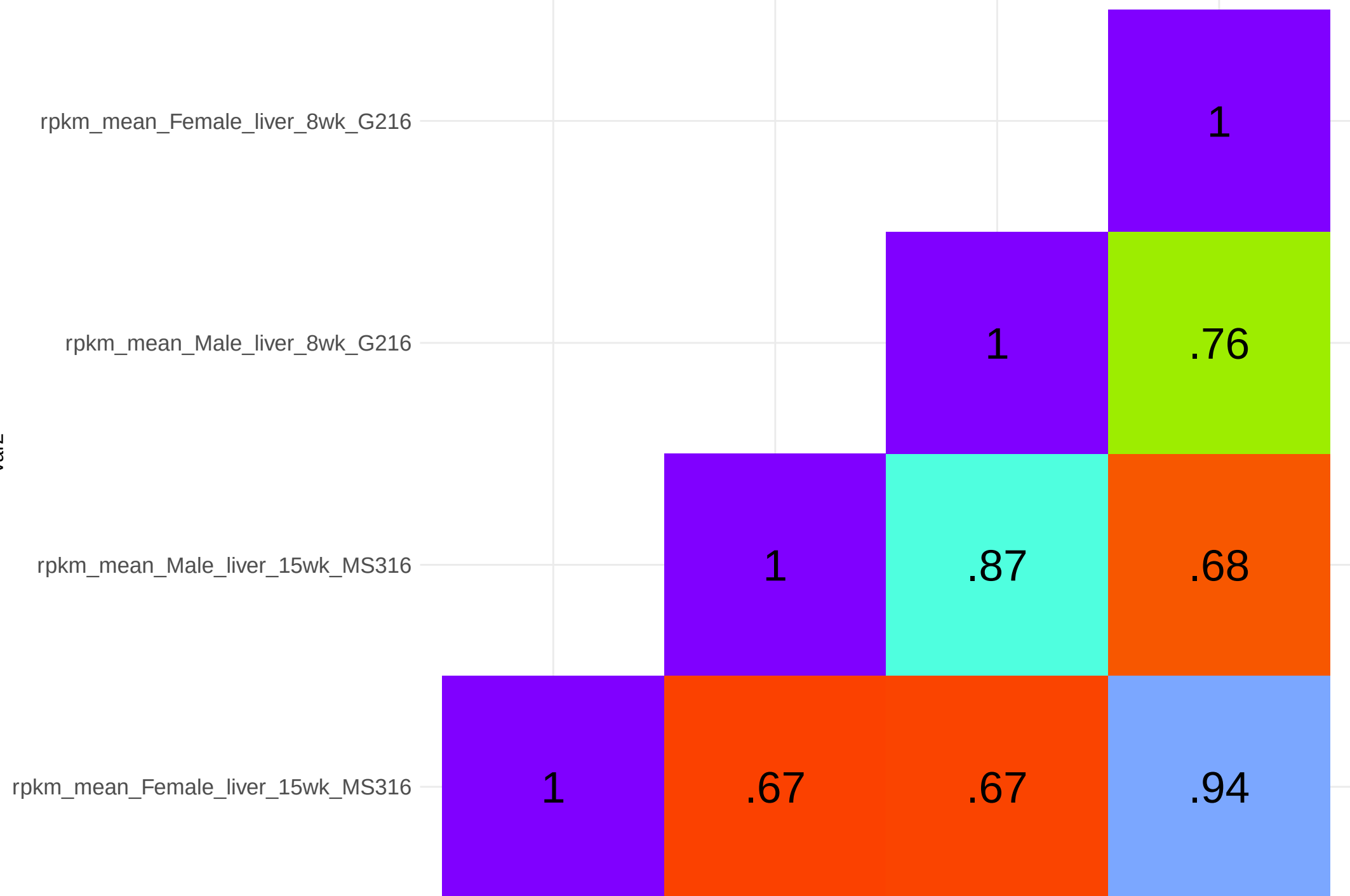


G221_RNASEQ_MS316, 00_Spearman_All_Merge,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798



G221_RNASEQ_MS316, 00_Spearman_Significant_Union_Merge,
Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 739

Var2



rpkm_mean_Female_liver_15wk_MS316

rpkm_mean_Male_liver_15wk_MS316

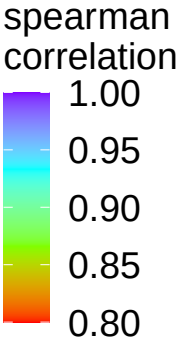
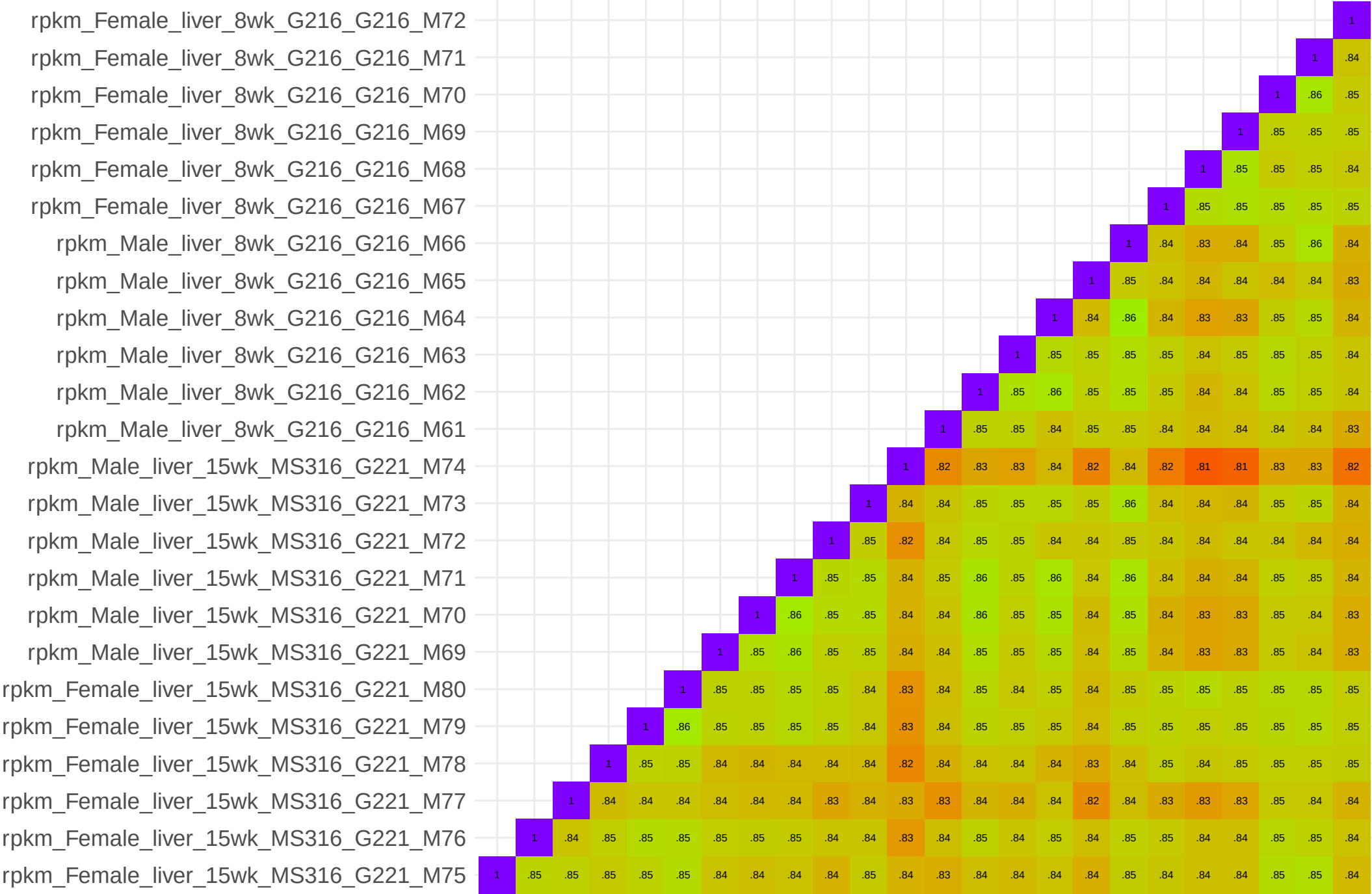
rpkm_mean_Male_liver_8wk_G216

rpkm_mean_Female_liver_8wk_G216

Var1

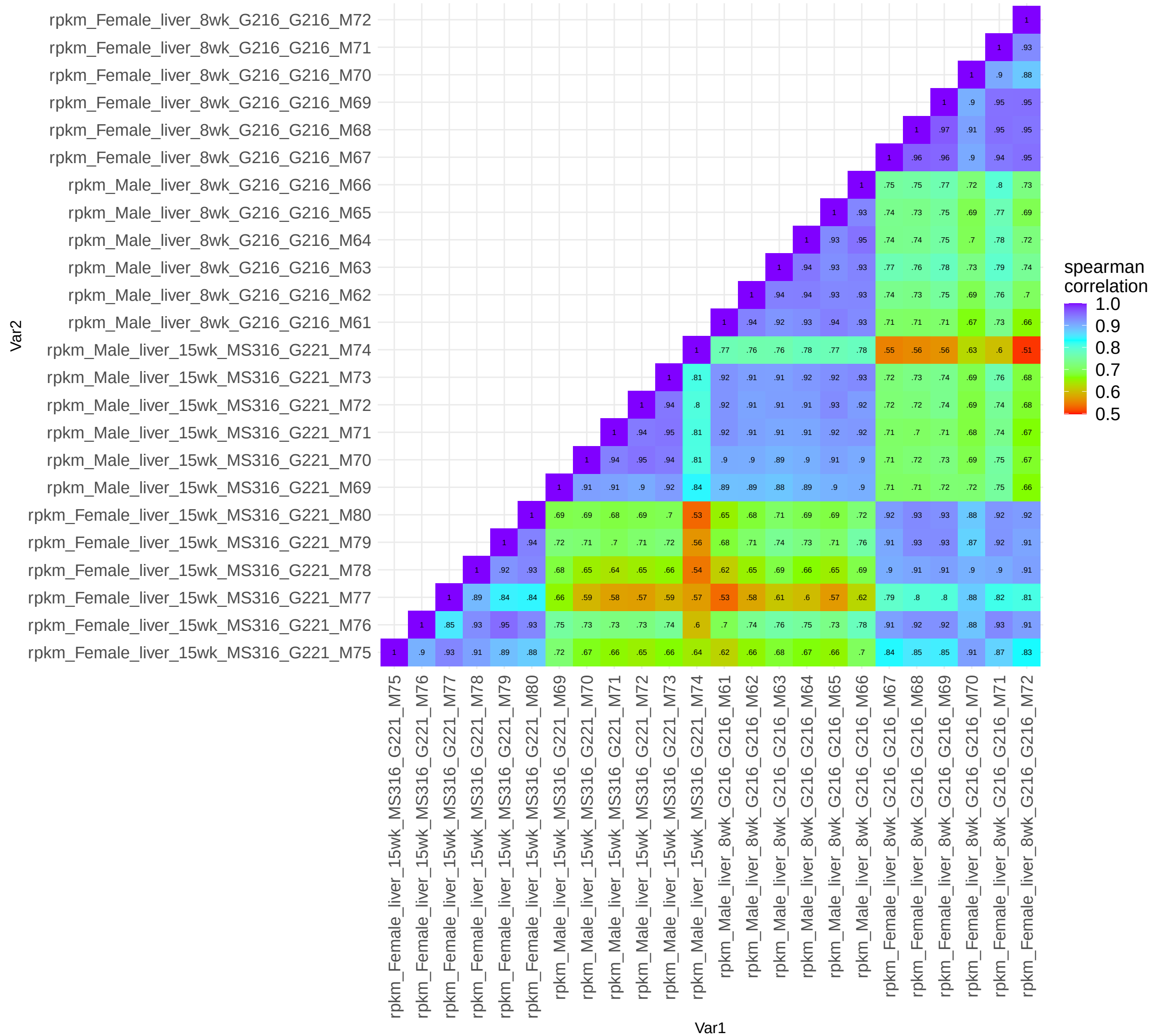
G221_RNASEQ_MS316, 0_Spearman_All_Merge,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798

Var2

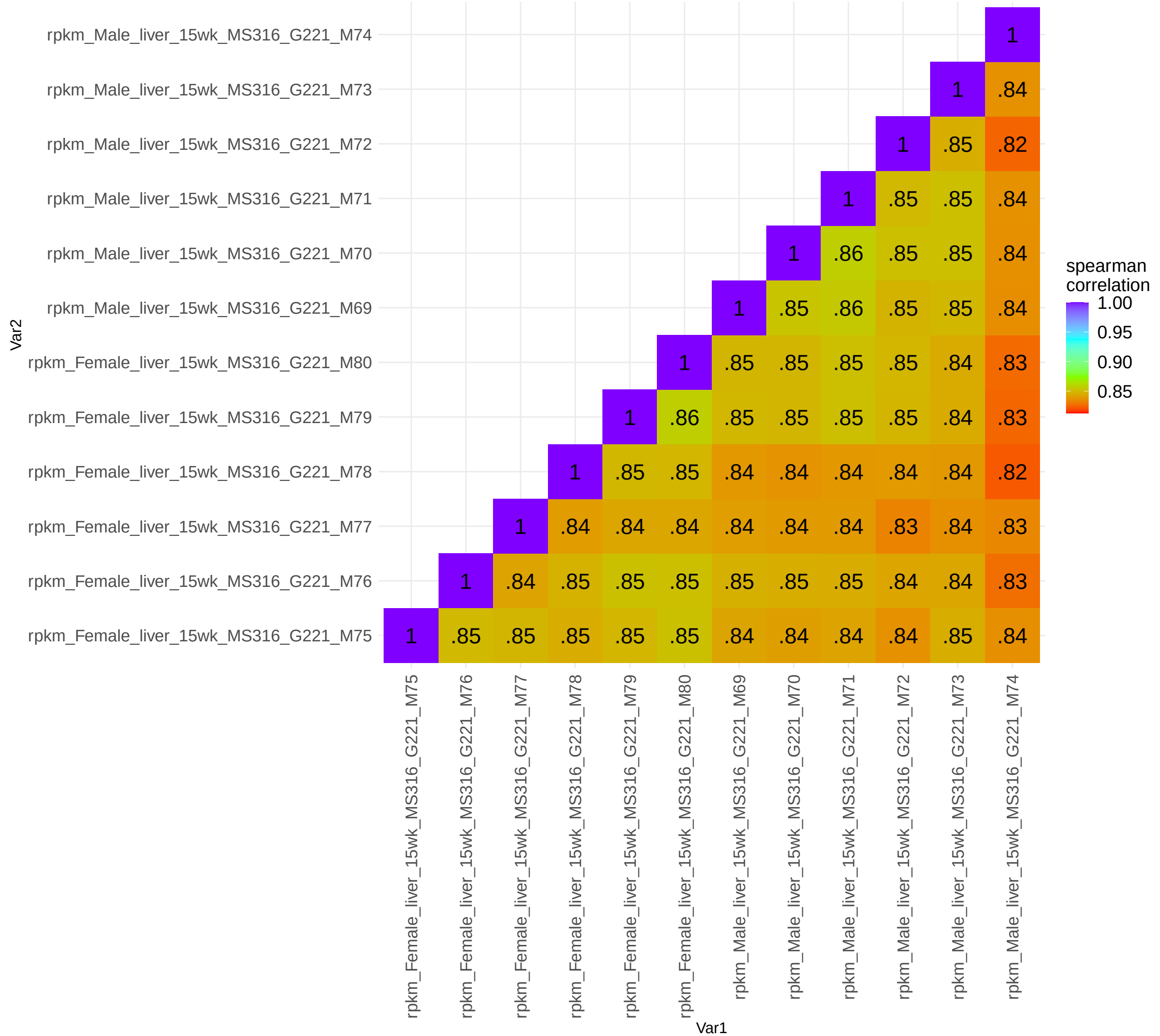


Var1

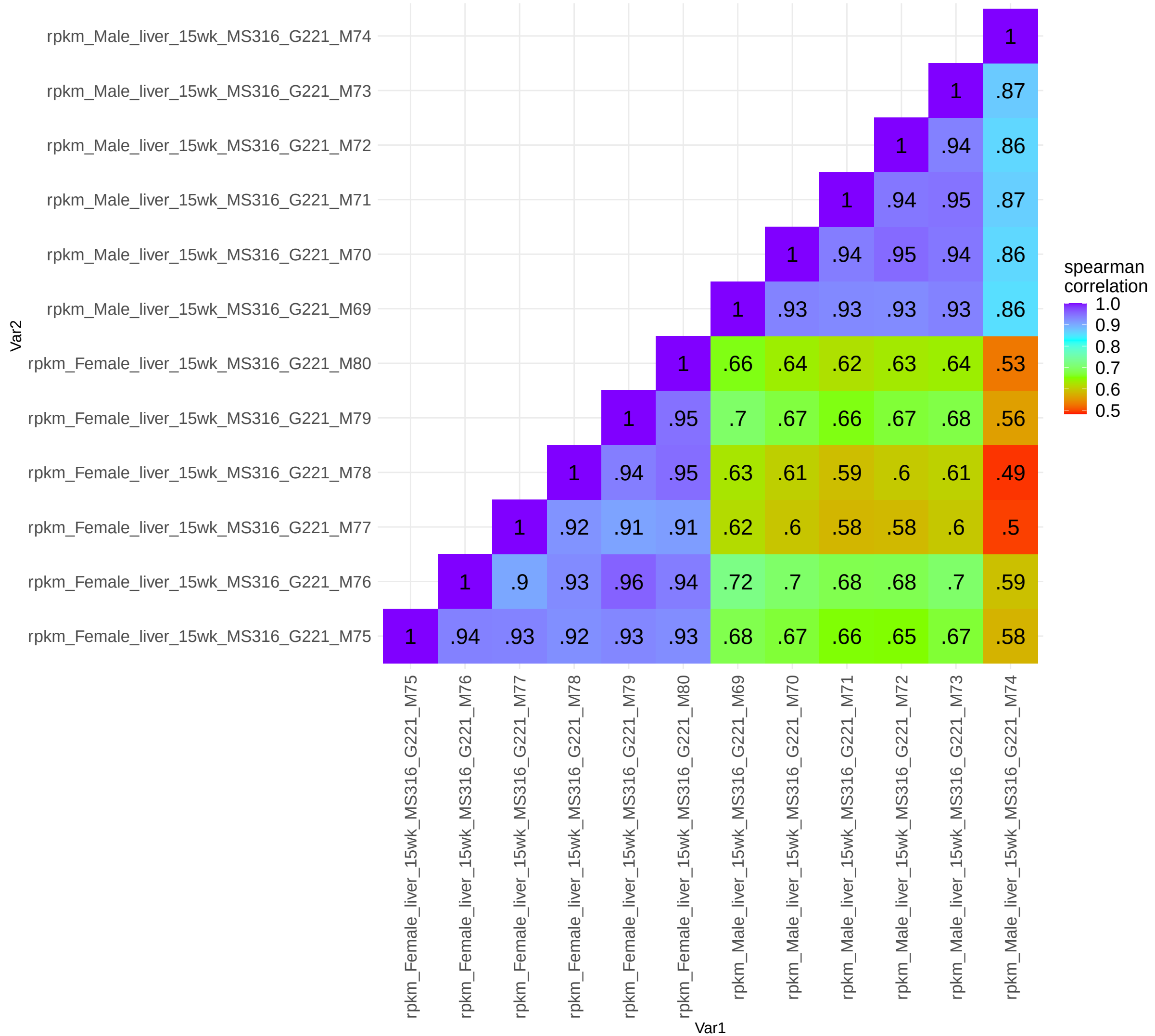
G221_RNASEQ_MS316, 0_Spearman_Significant_Union_Merge,
Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 739



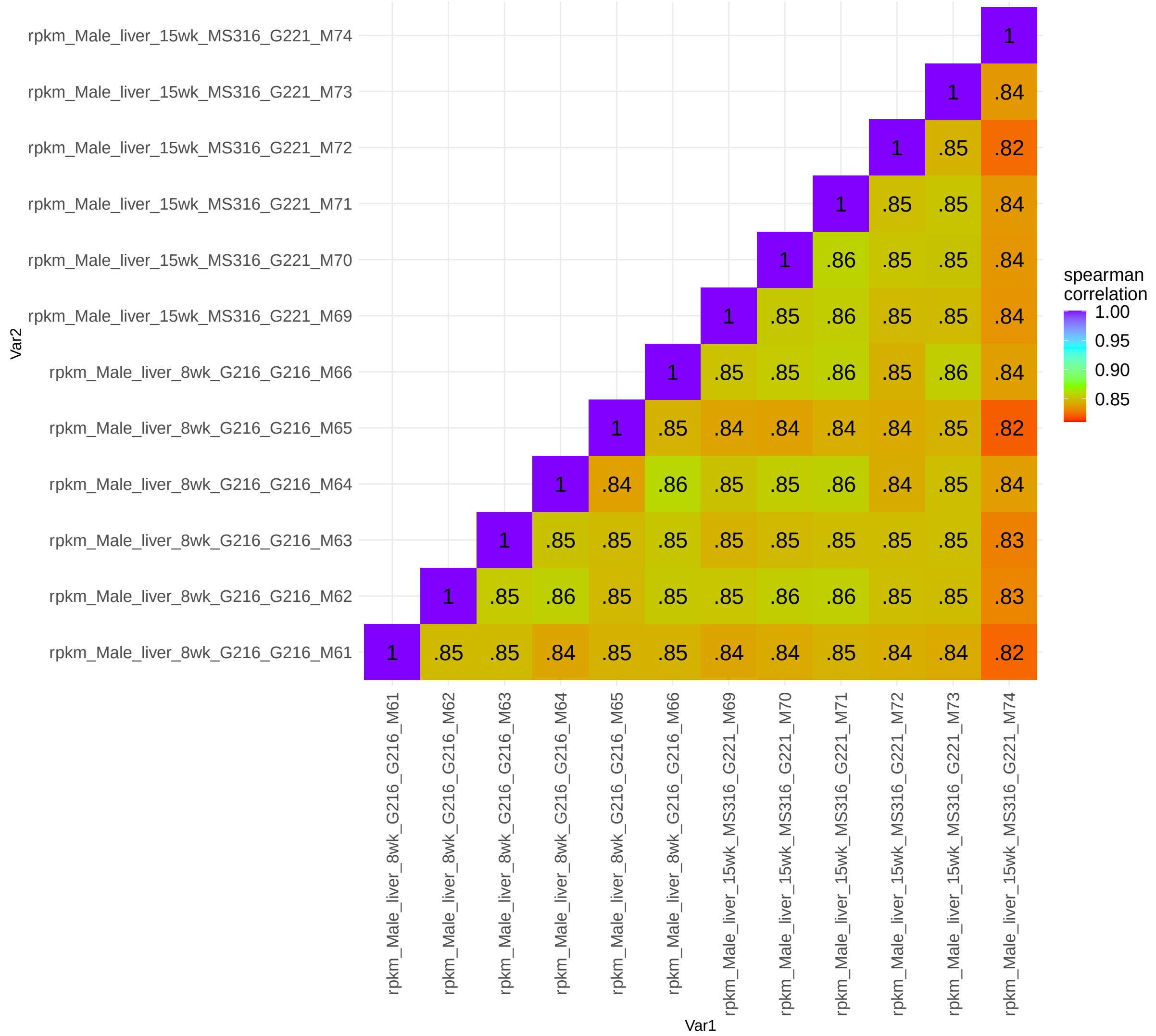
G221_RNASEQ_MS316, 1_Spearman_All_Female_liver_15wk_MS316_Male_liver_15wk_MS316,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798



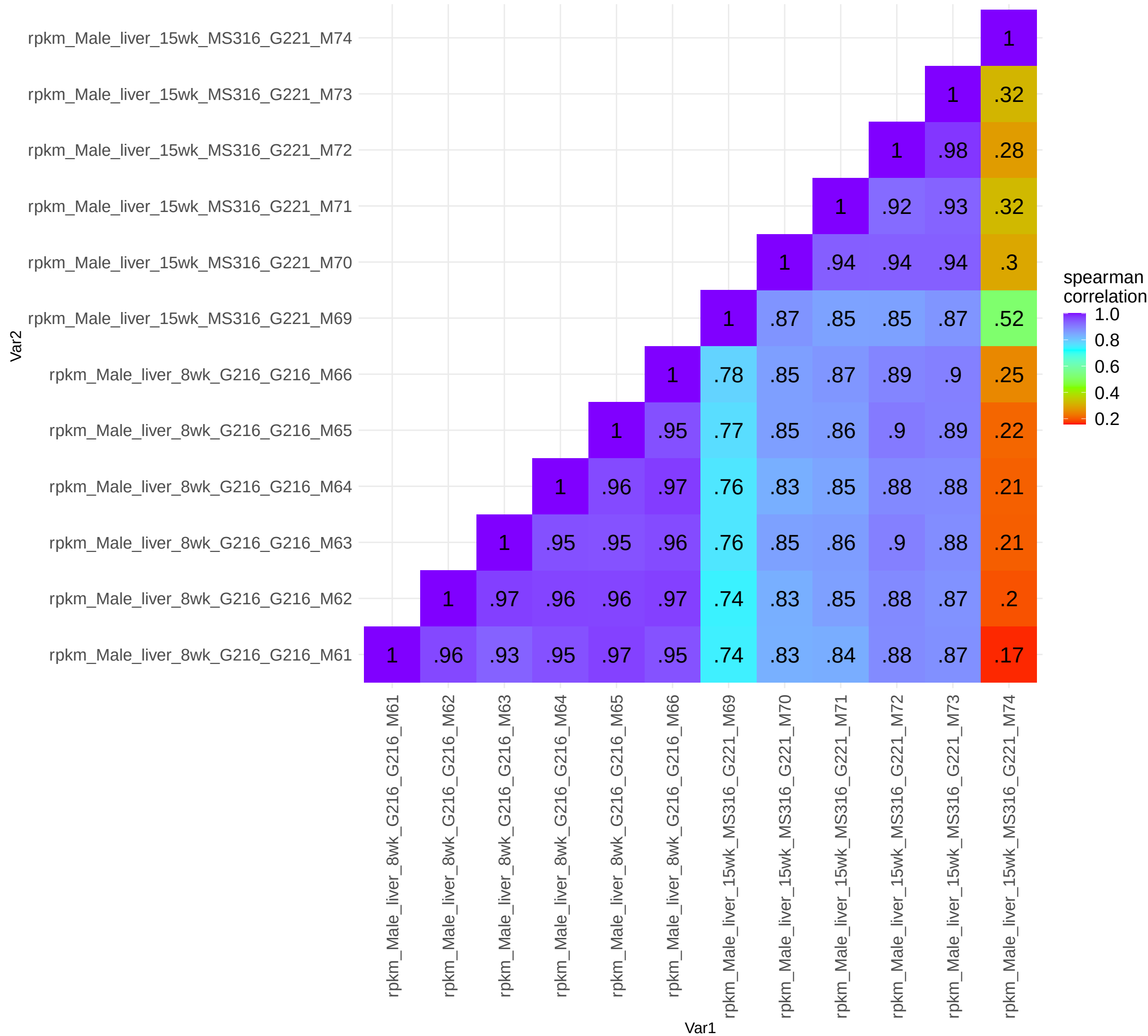
G221_RNASEQ_MS316, 1_Spearman_Significant_Female_liver_15wk_MS316_Male_liver_15wk_MS316, Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 587



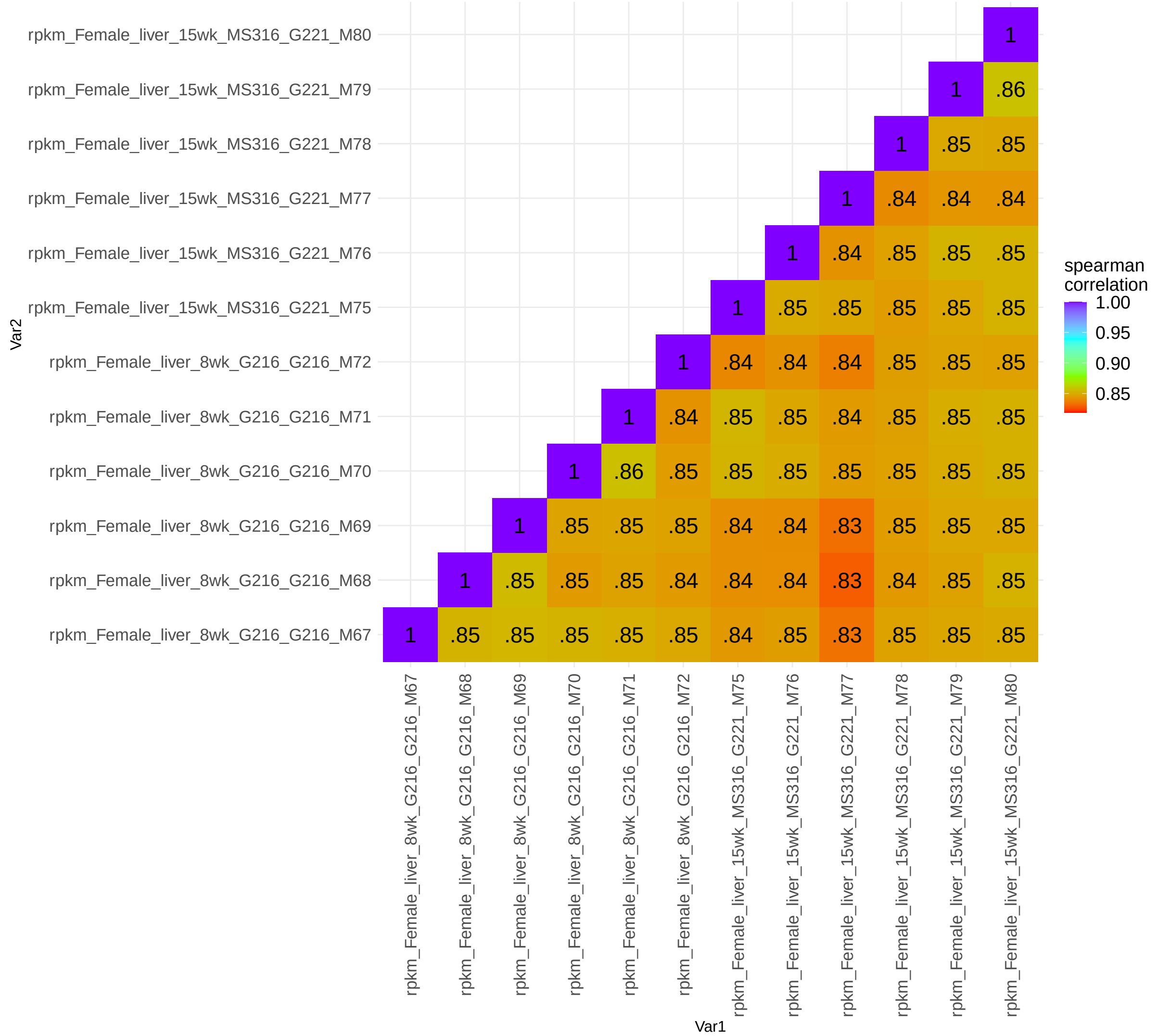
G221_RNASEQ_MS316, 2_Spearman_All_Male_liver_8wk_G216_Male_liver_15wk_MS316,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798



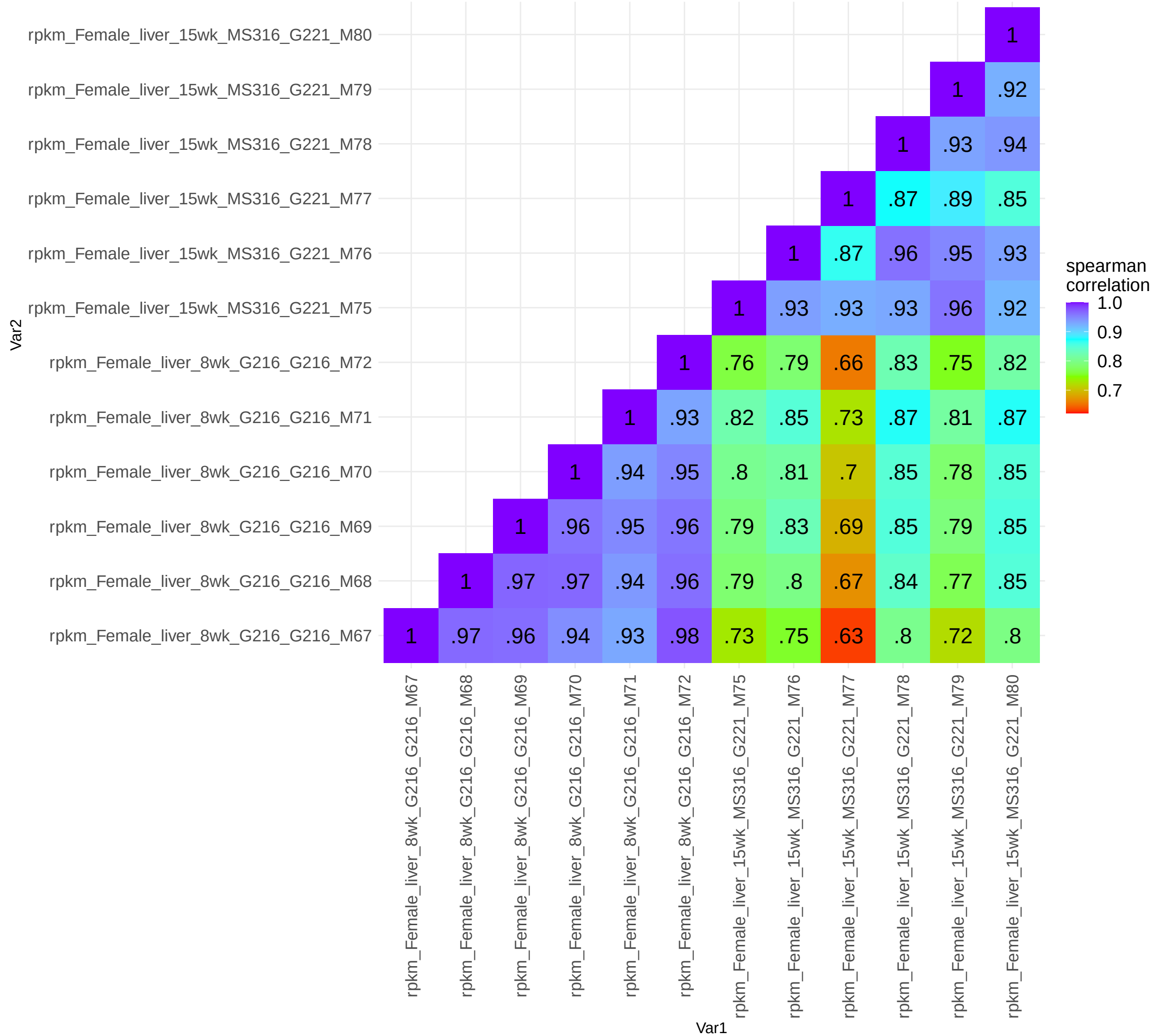
G221_RNASEQ_MS316, 2_Spearman_Significant_Male_liver_8wk_G216_Male_liver_15wk_MS316, Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 110



G221_RNASEQ_MS316, 3_Spearman_All_Female_liver_8wk_G216_Female_liver_15wk_MS316,
All genes (without filter), RefSeqLncRNA76k_Genes: 75798

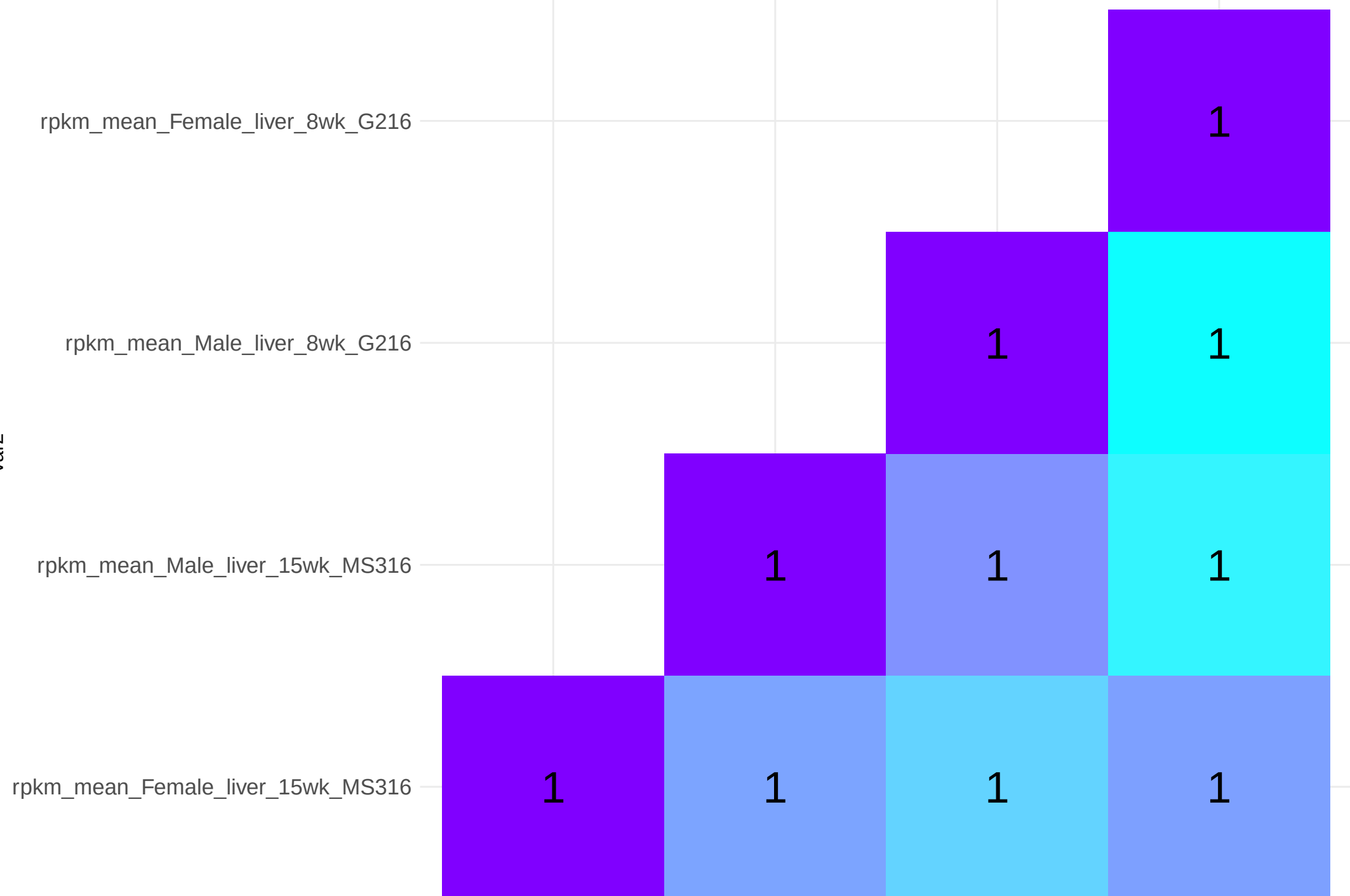


G221_RNASEQ_MS316, 3_Spearman_Significant_Female_liver_8wk_G216_Female_liver_15wk_MS316, Significant genes (|FC|>2 and FDR<0.05), RefSeqLncRNA76k_Genes: 102



G221_RNASEQ_MS316, 00_Spearman_Non-significant_Intersection_Merge,
Non-significant genes ($1.2 < |FC| < 1/1.2$ and $FDR > 0.1$, $RPKM > 1$), RefSeqLncRNA76k_Genes: 4677

Var2



spearman
correlation

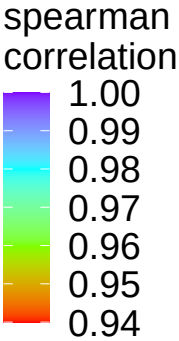
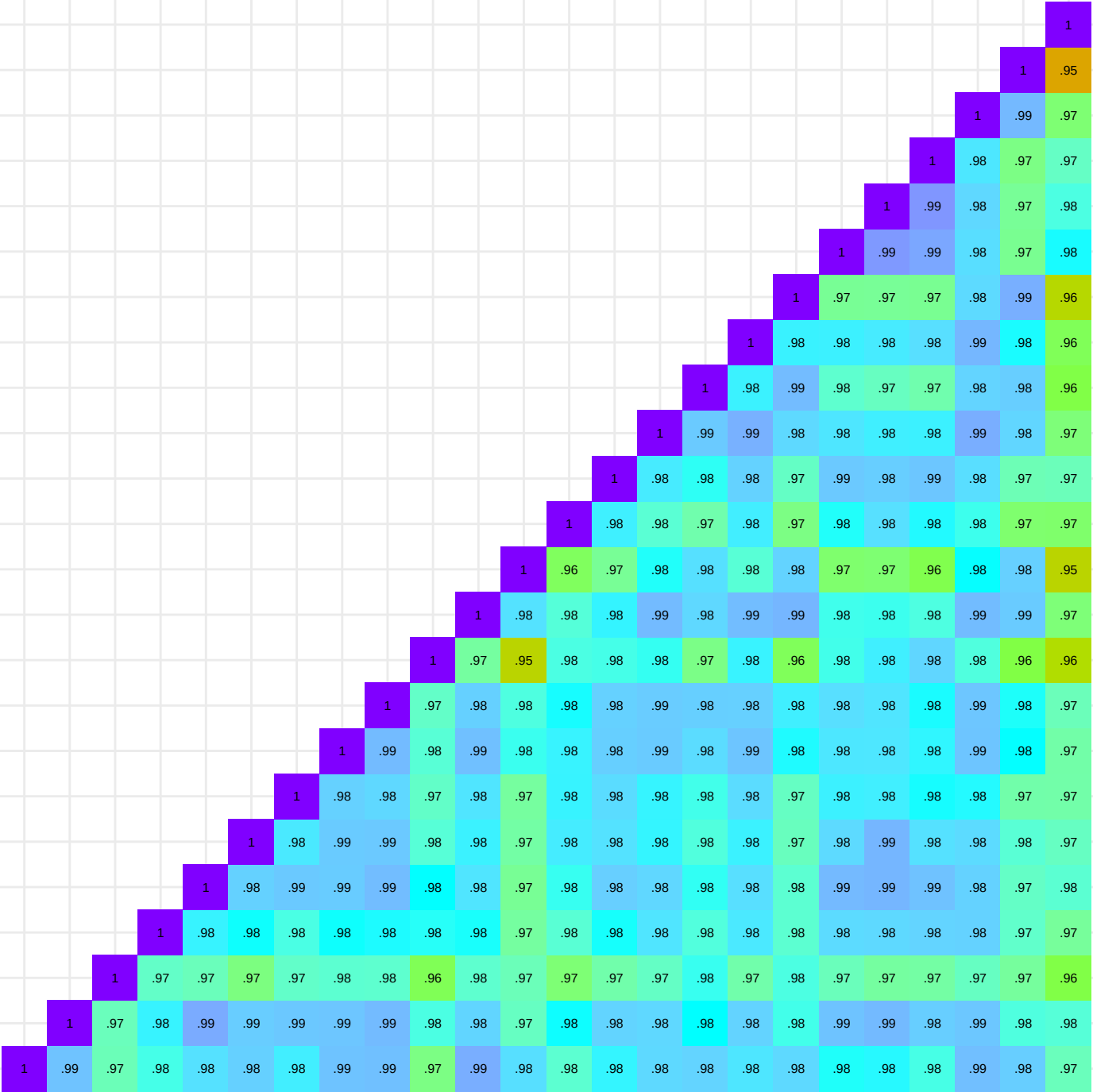
1.000
0.995
0.990
0.985

Var1

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

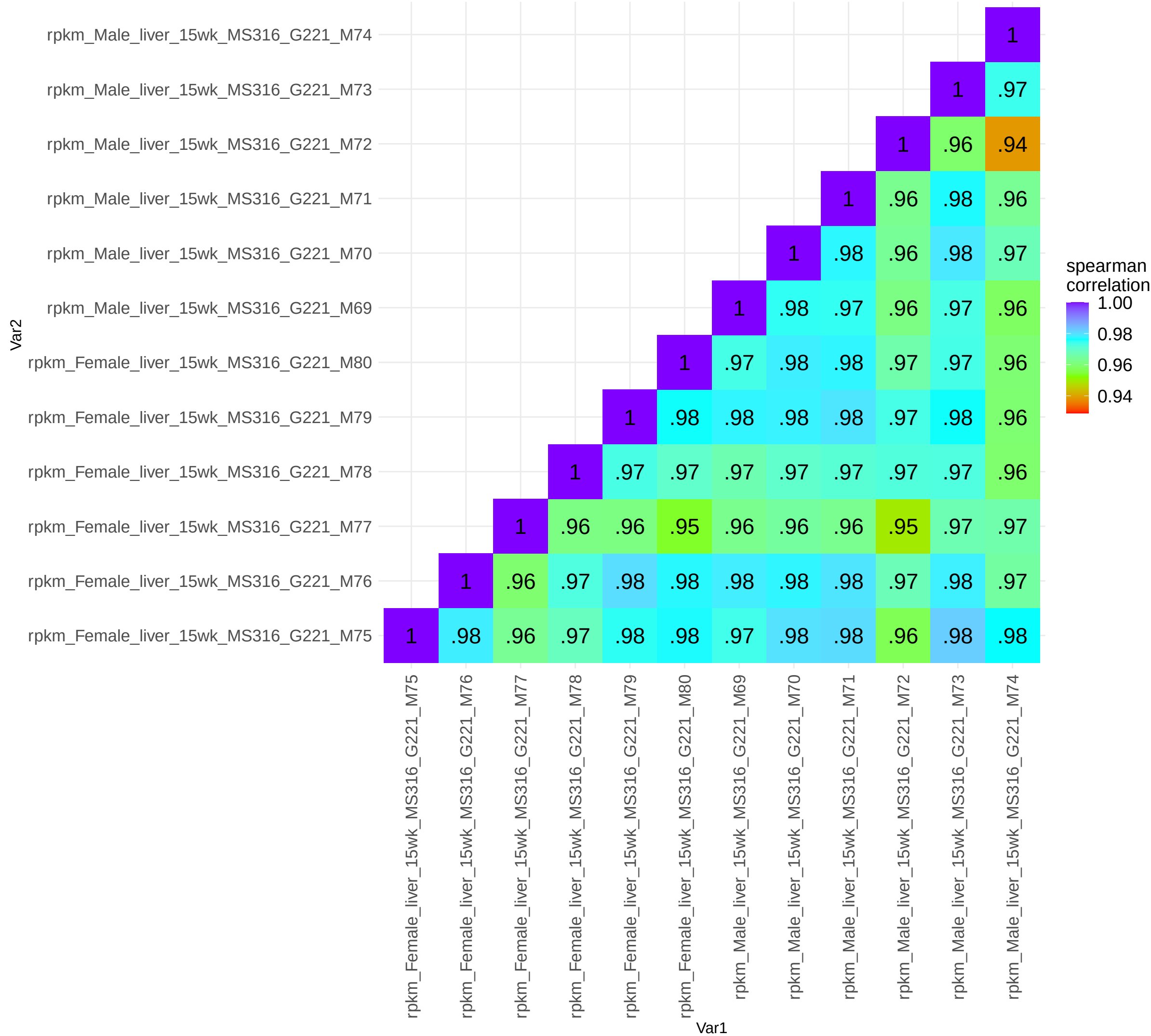
Var2

rpkm_Female_liver_8wk_G216_G216_M72
rpkm_Female_liver_8wk_G216_G216_M71
rpkm_Female_liver_8wk_G216_G216_M70
rpkm_Female_liver_8wk_G216_G216_M69
rpkm_Female_liver_8wk_G216_G216_M68
rpkm_Female_liver_8wk_G216_G216_M67
rpkm_Male_liver_8wk_G216_G216_M66
rpkm_Male_liver_8wk_G216_G216_M65
rpkm_Male_liver_8wk_G216_G216_M64
rpkm_Male_liver_8wk_G216_G216_M63
rpkm_Male_liver_8wk_G216_G216_M62
rpkm_Male_liver_8wk_G216_G216_M61
rpkm_Male_liver_15wk_MS316_G221_M74
rpkm_Male_liver_15wk_MS316_G221_M73
rpkm_Male_liver_15wk_MS316_G221_M72
rpkm_Male_liver_15wk_MS316_G221_M71
rpkm_Male_liver_15wk_MS316_G221_M70
rpkm_Male_liver_15wk_MS316_G221_M69
rpkm_Female_liver_15wk_MS316_G221_M80
rpkm_Female_liver_15wk_MS316_G221_M79
rpkm_Female_liver_15wk_MS316_G221_M78
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rpkm_Female_liver_15wk_MS316_G221_M76
rpkm_Female_liver_15wk_MS316_G221_M75

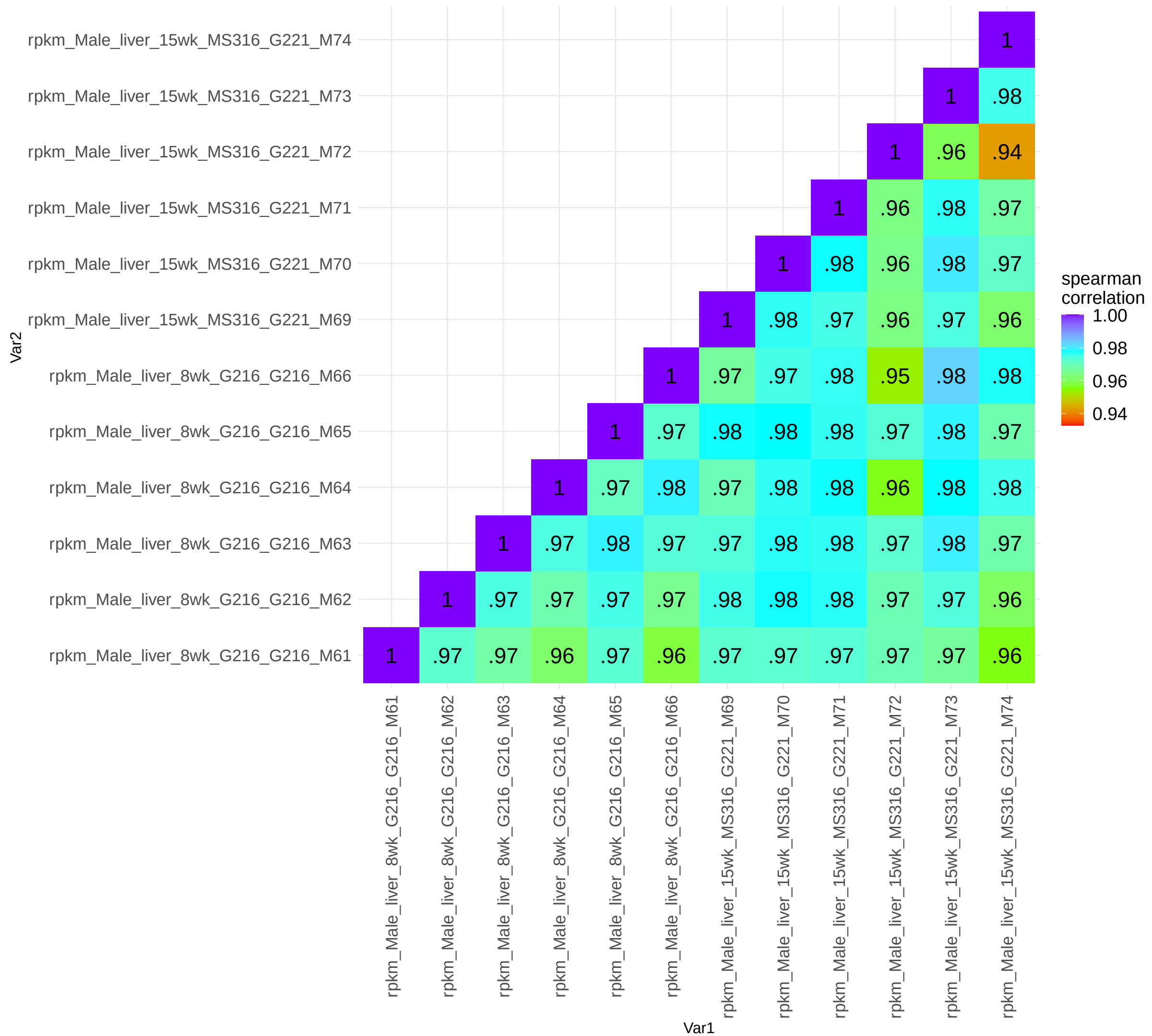


Var1

G221_RNASEQ_MS316, 1_Spearman_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_Spearman_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_Spearman_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

