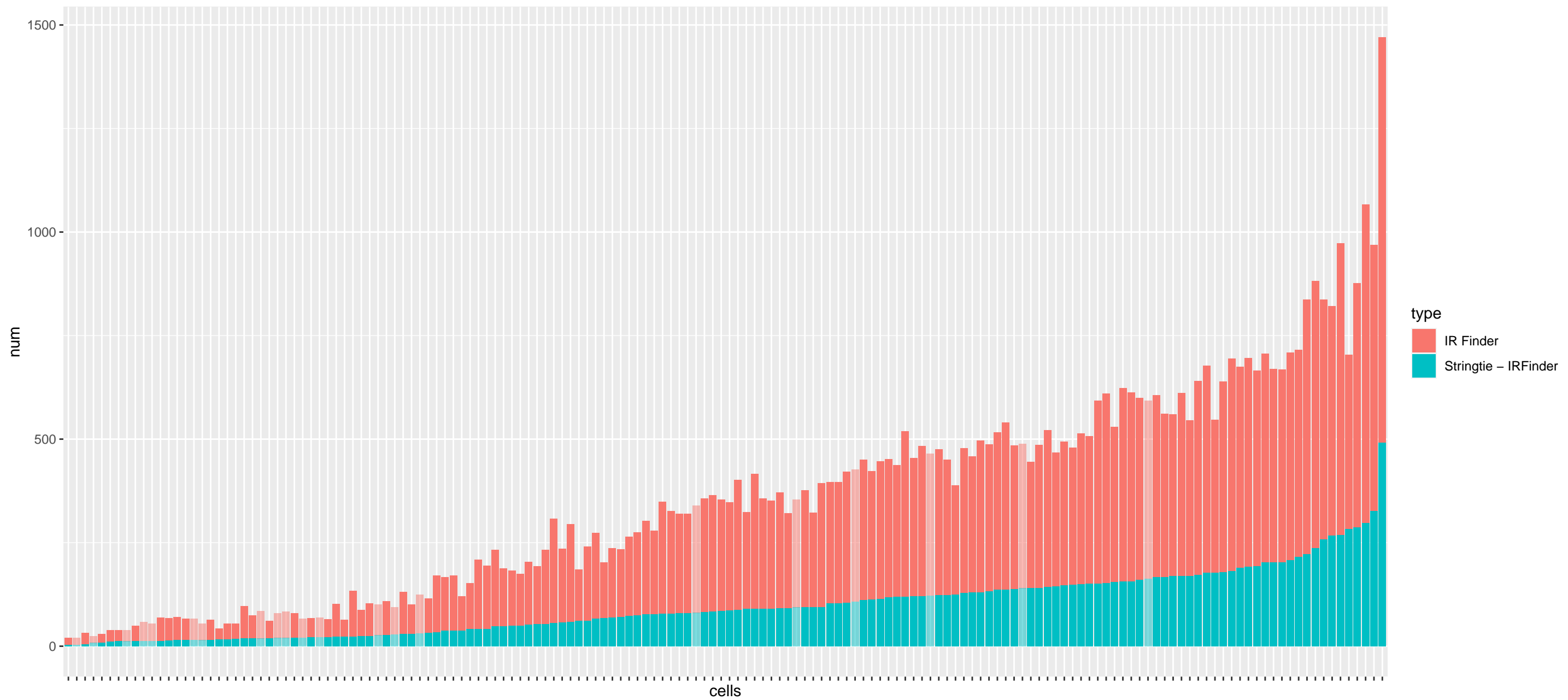
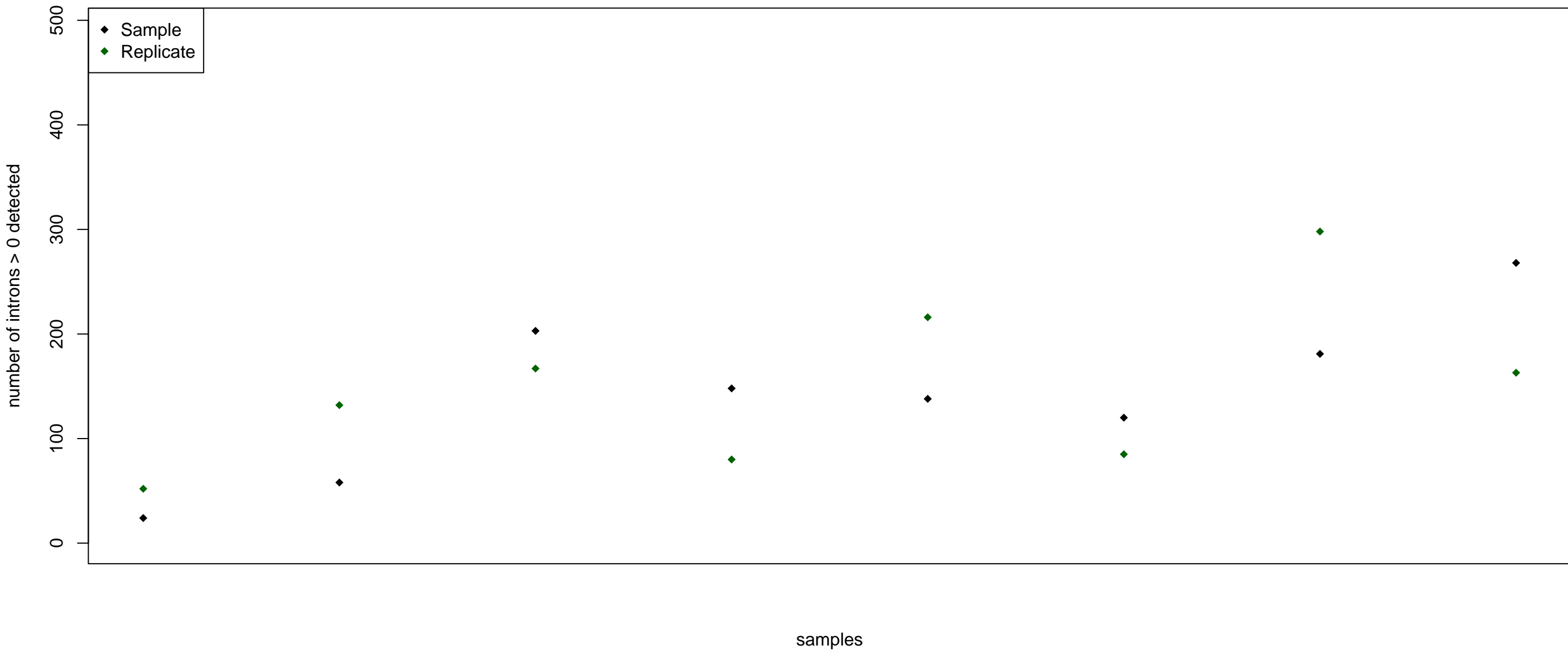


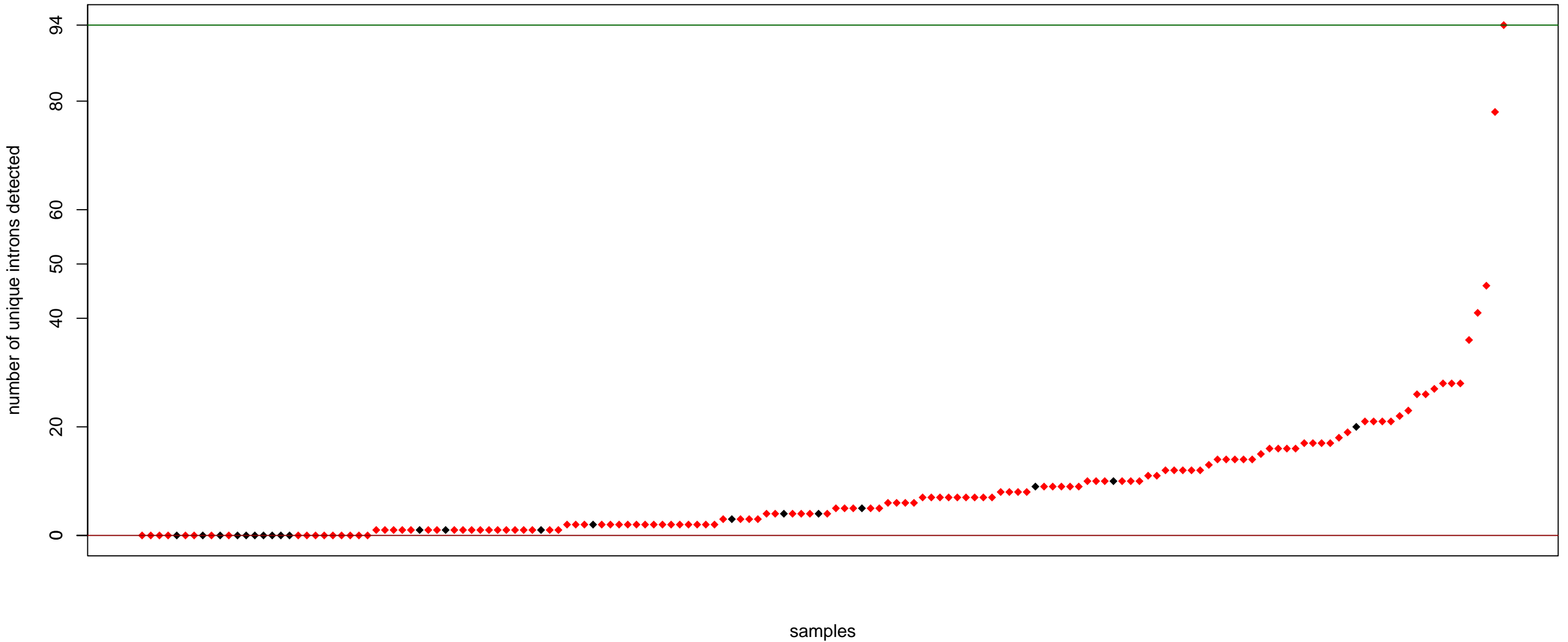
Intron retention events with at least 0.2 coverage



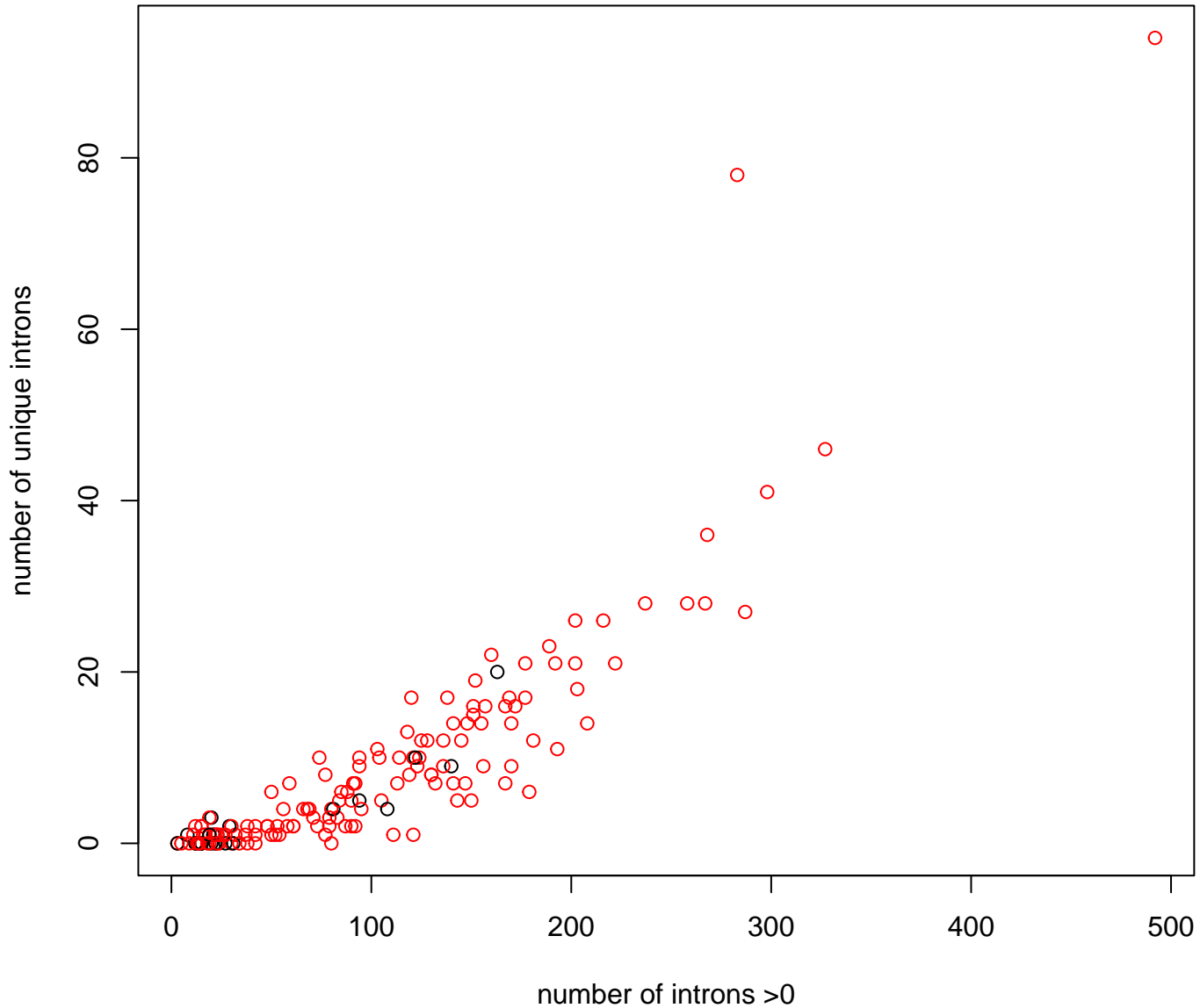
# Intersected introns



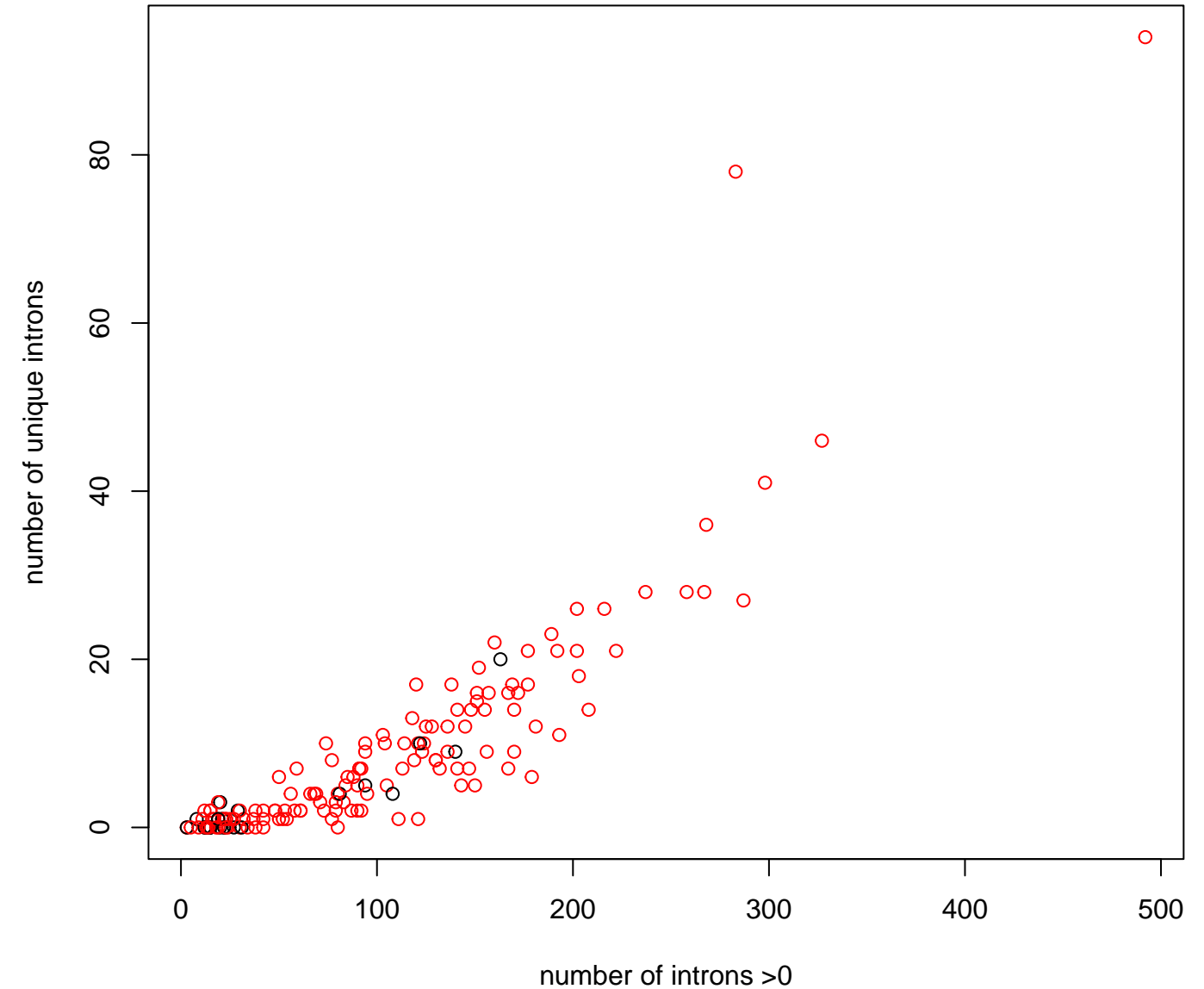
Total: 1334 introns and 1274 for CRIS classified



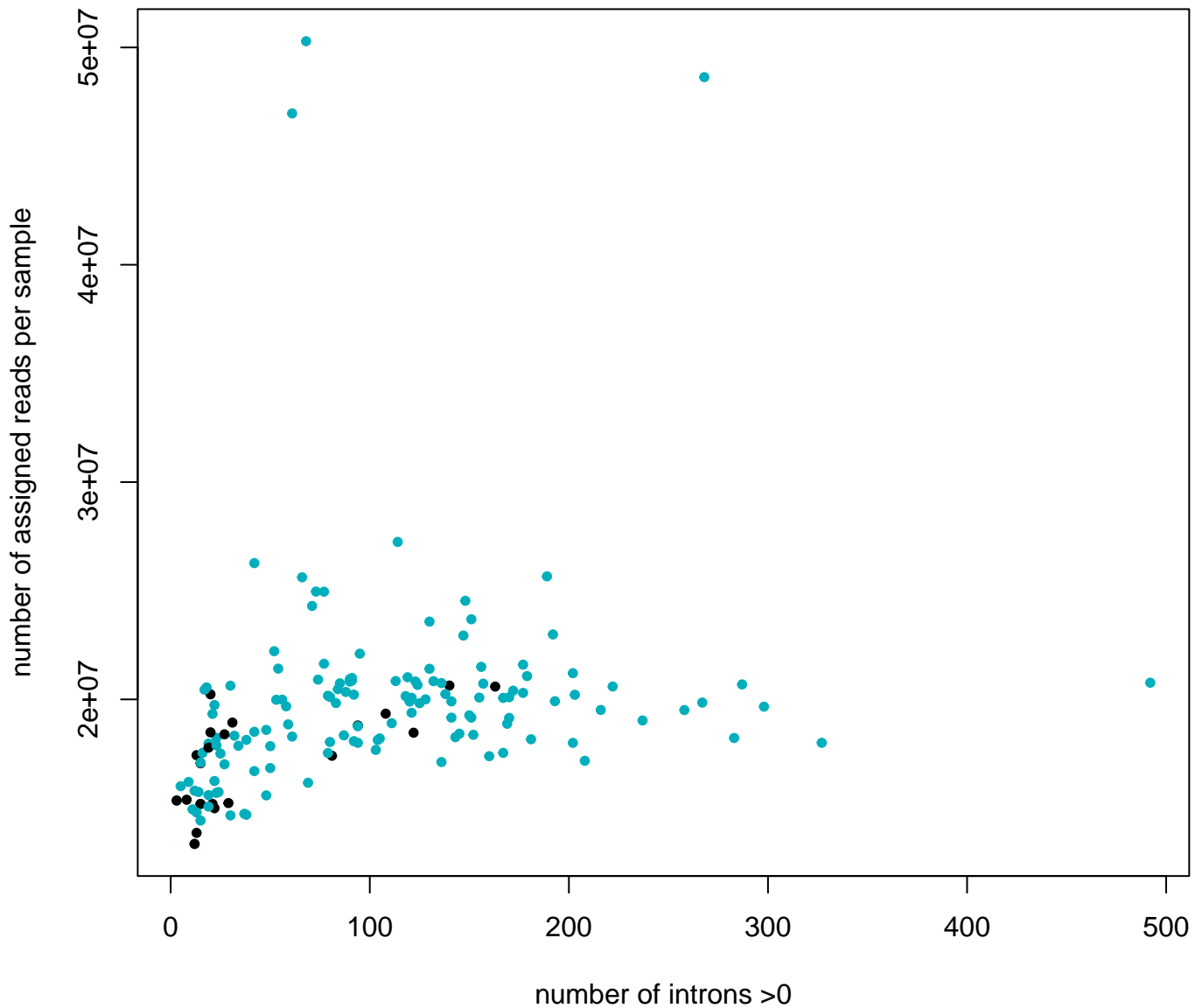
**Pearson correlation between number of total introns and unique introns per sample: 0.87593685620235 and 0.875716396095812 for those classified for CRIS**



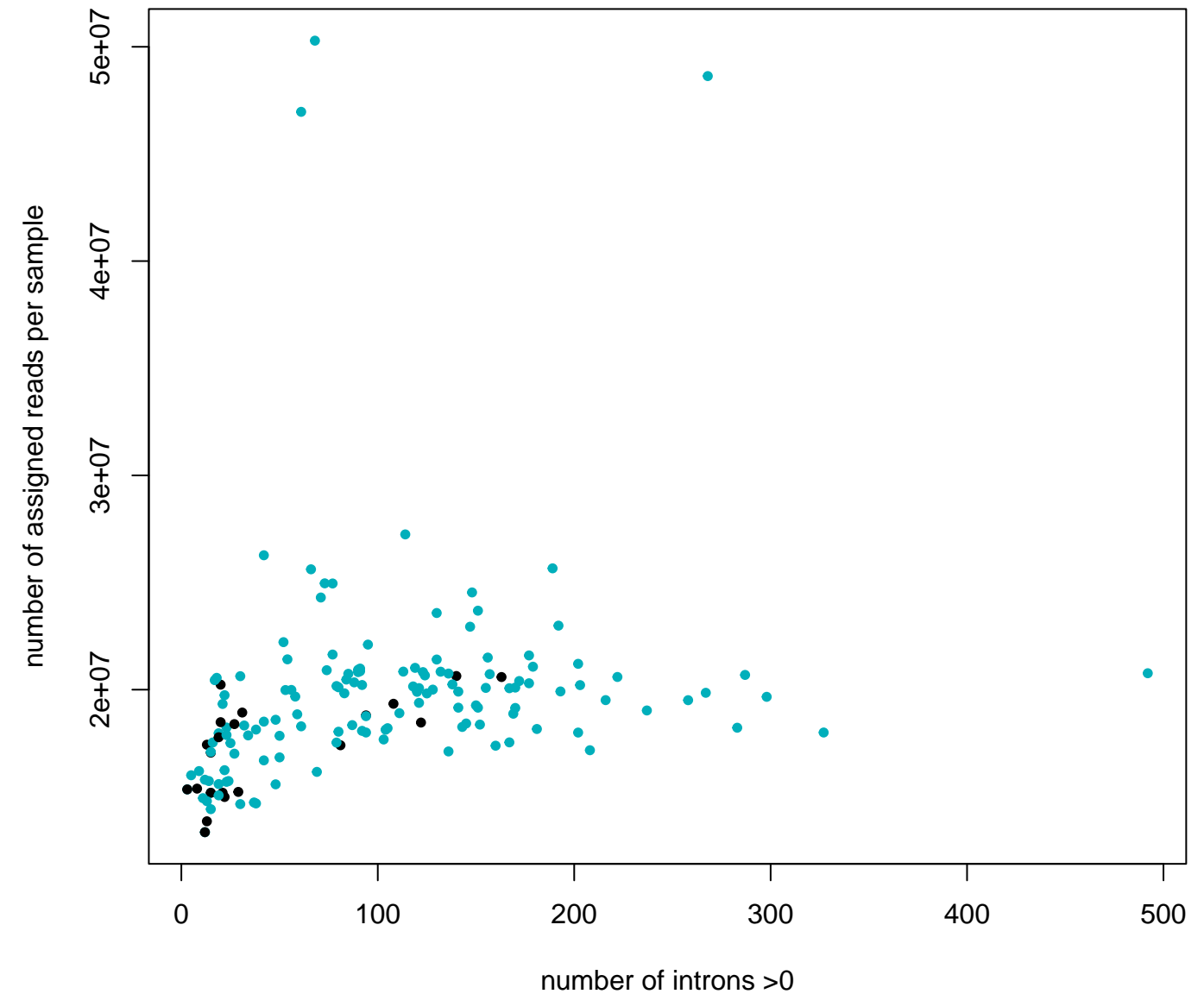
**Spearman correlation between number of total introns and unique introns per sample: 0.90439457627603 and 0.901513566000073 for those classified for CRIS**



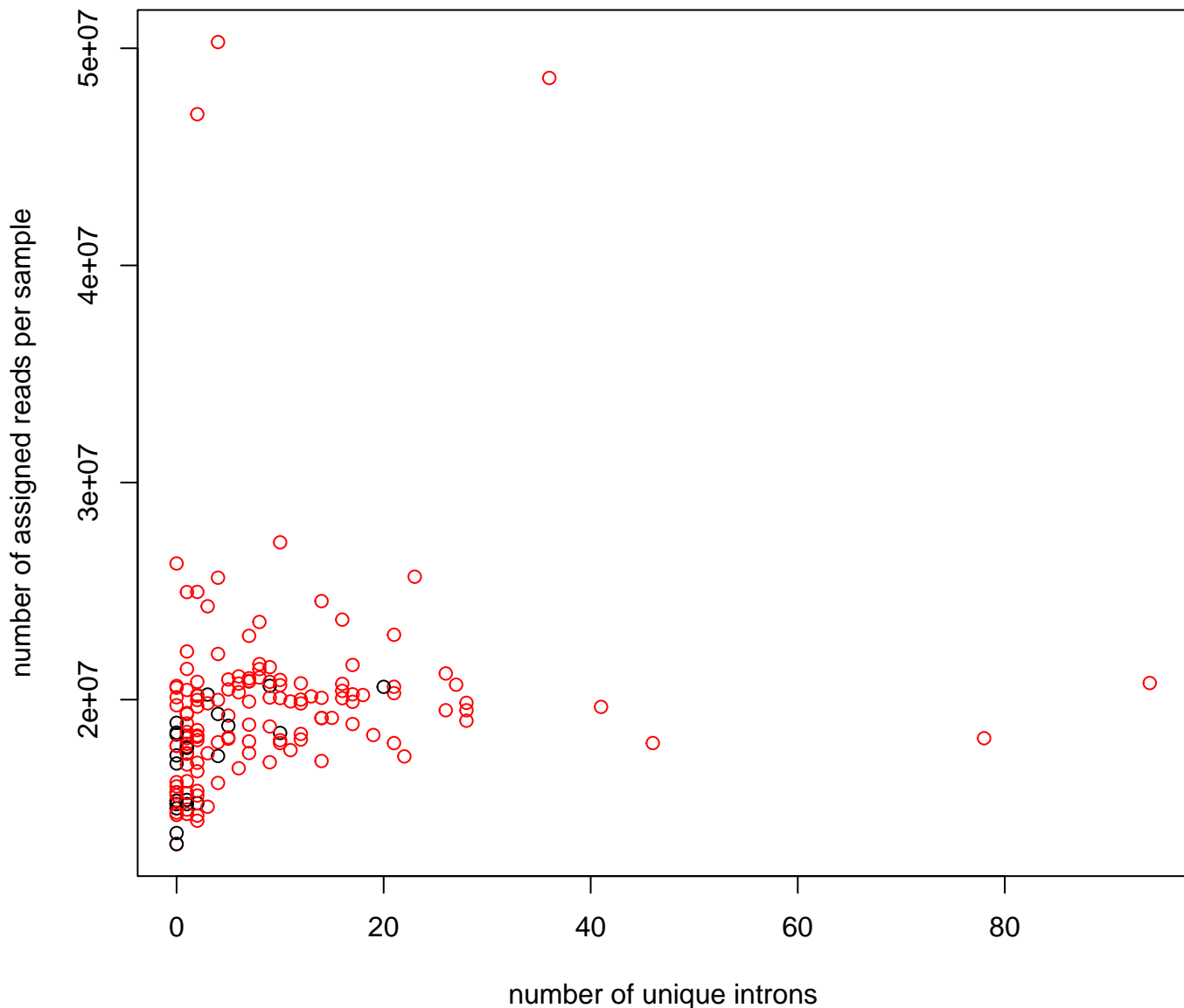
**Pearson correlation between number  
of total introns and assigned reads: 0.26619054982758 and  
0.21189521335656 for those classified for CRIS**



**Spearman correlation between number  
of total introns and assigned reads: 0.527646796133711 and  
0.437052015952506 for those classified for CRIS**



**Pearson correlation between number of total introns and assigned reads: 0.175512188784208 and 0.135737750770721 for those classified for CRIS**



**Spearman correlation between number of total introns and assigned reads: 0.450913691467083 and 0.360214291585262 for those classified for CRIS**

