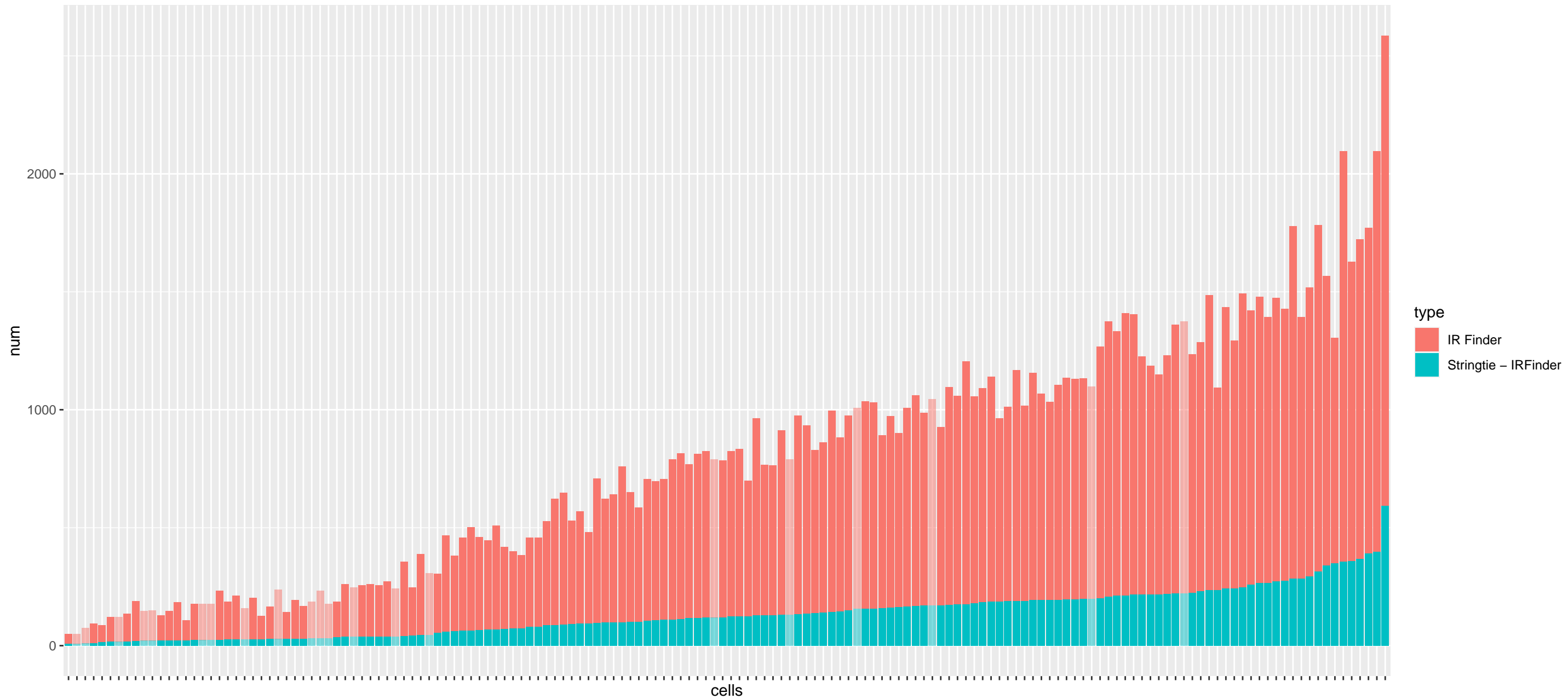
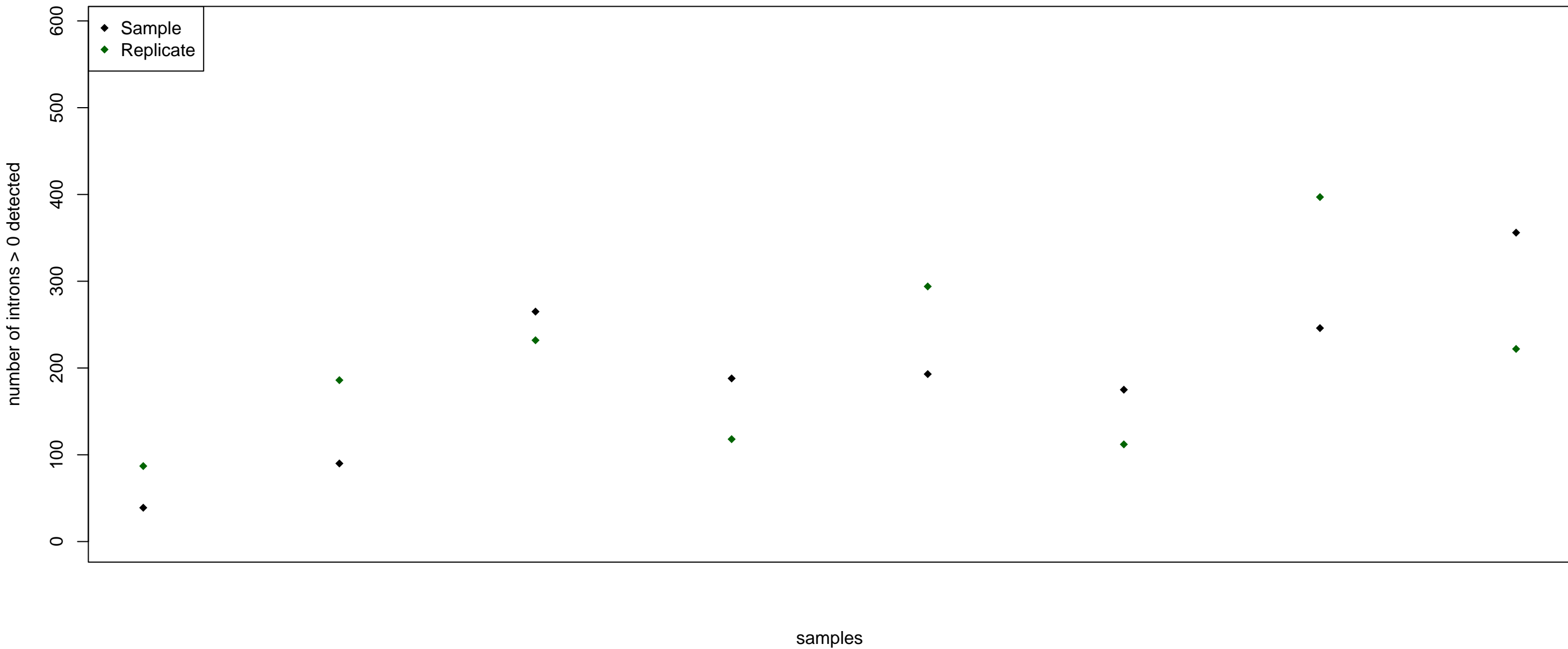


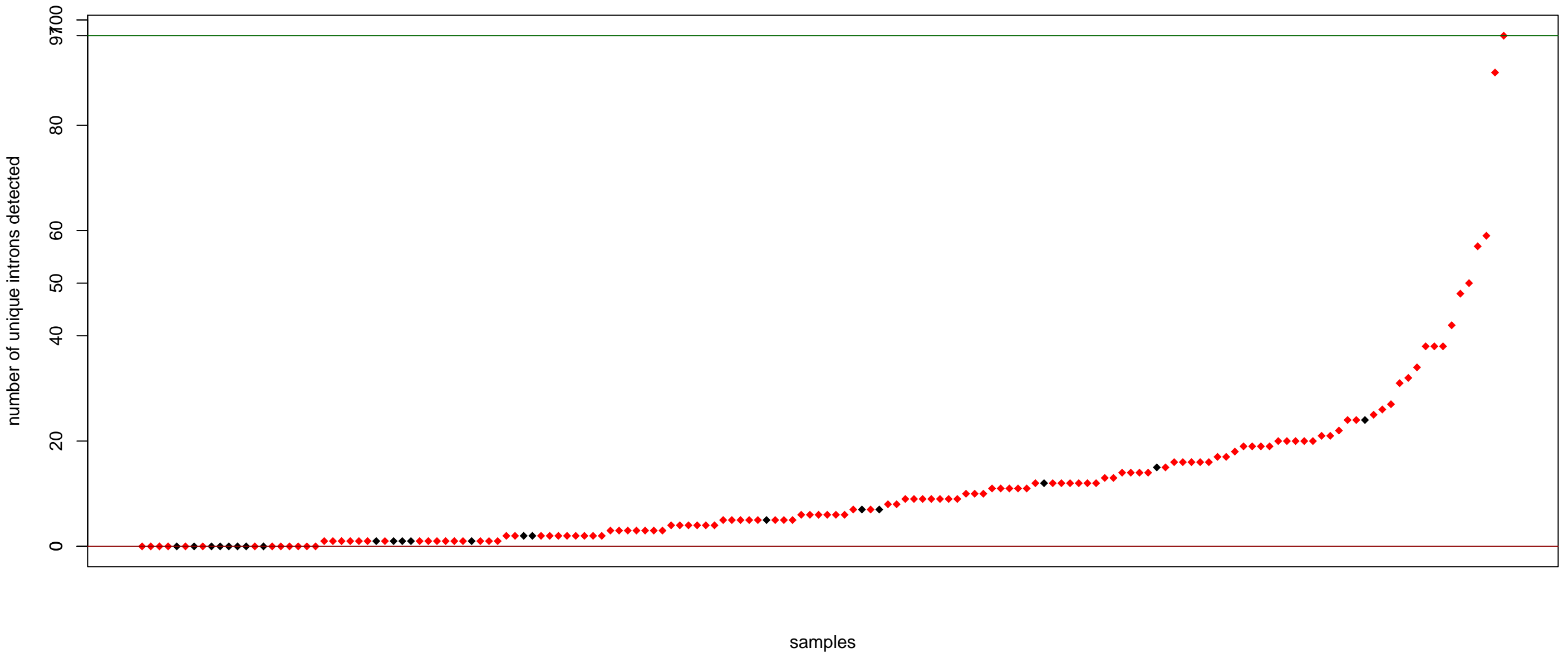
Intron retention events with at least 0.1 coverage



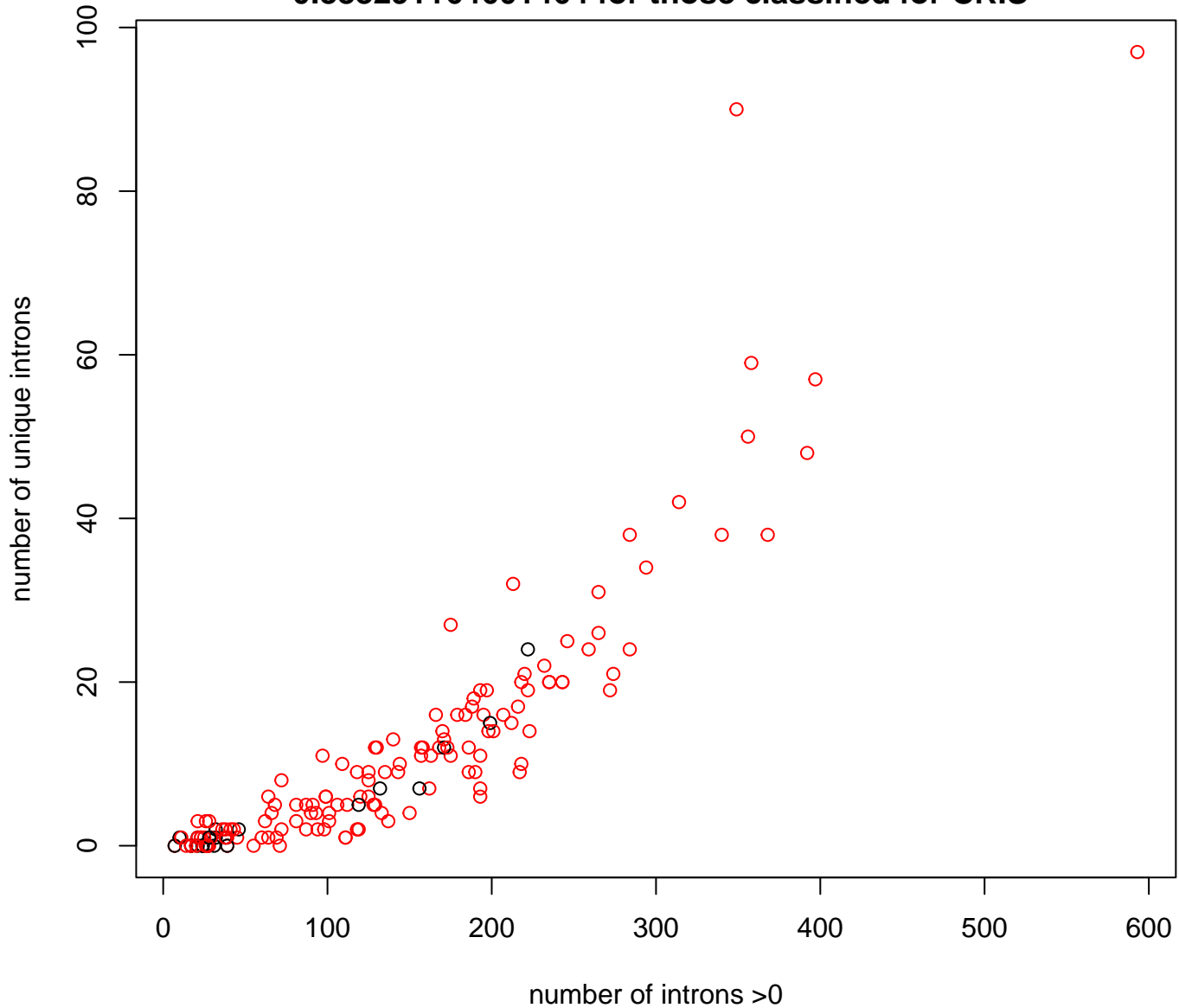
# Intersected introns



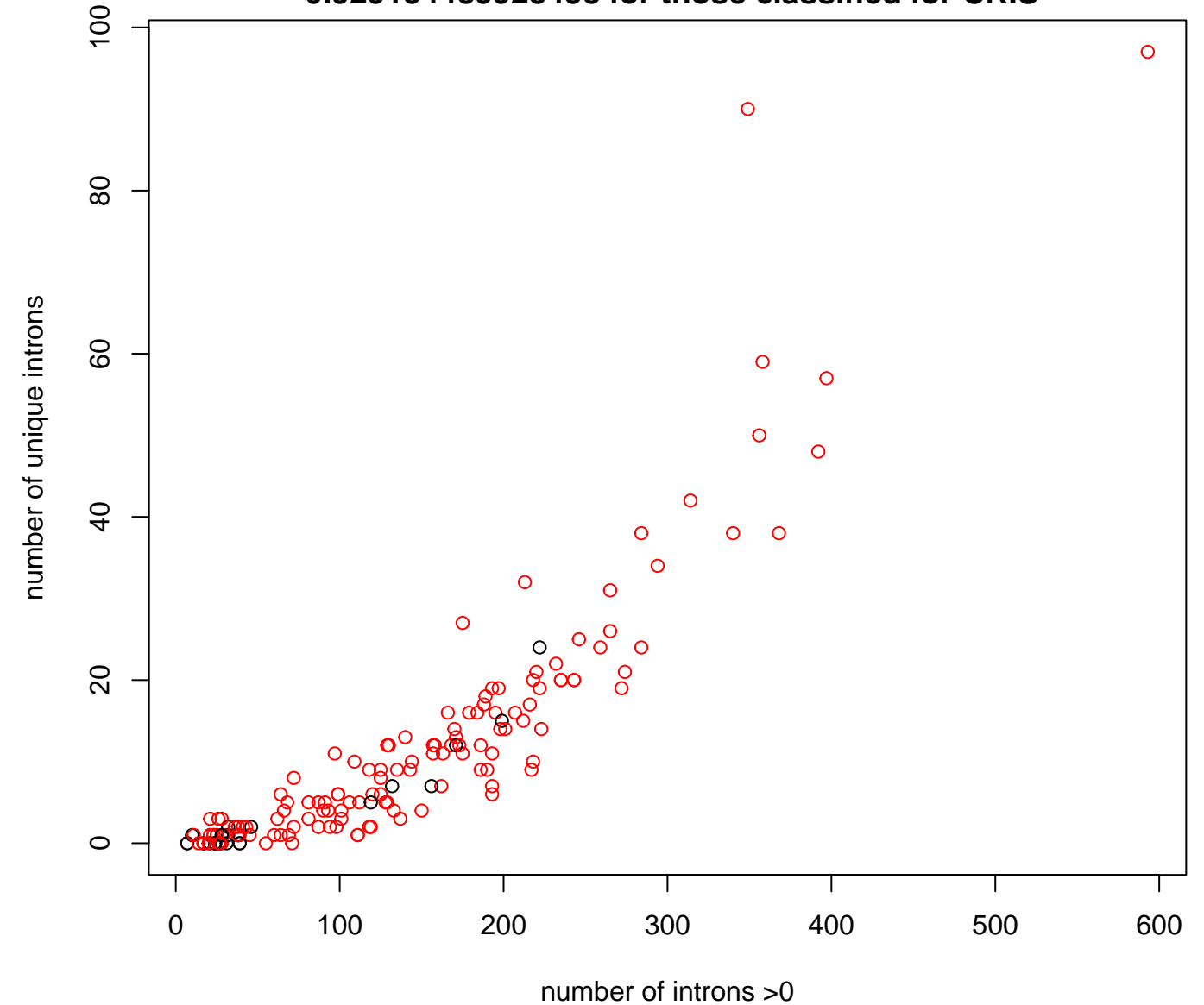
Total: 1747 introns and 1668 for CRIS classified



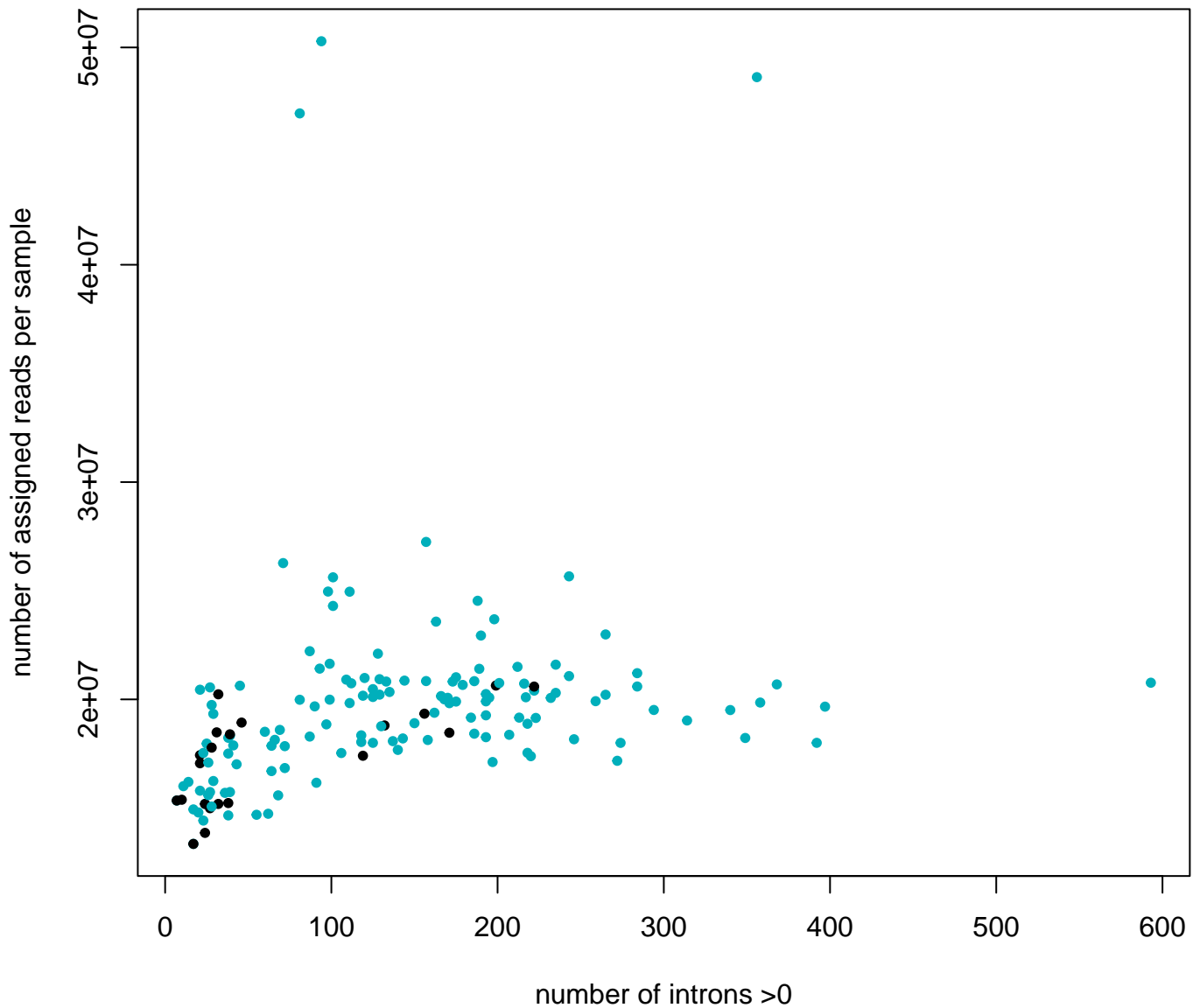
**Pearson correlation between number of total introns  
and unique introns per sample: 0.88854685985107 and  
0.888291164601464 for those classified for CRIS**



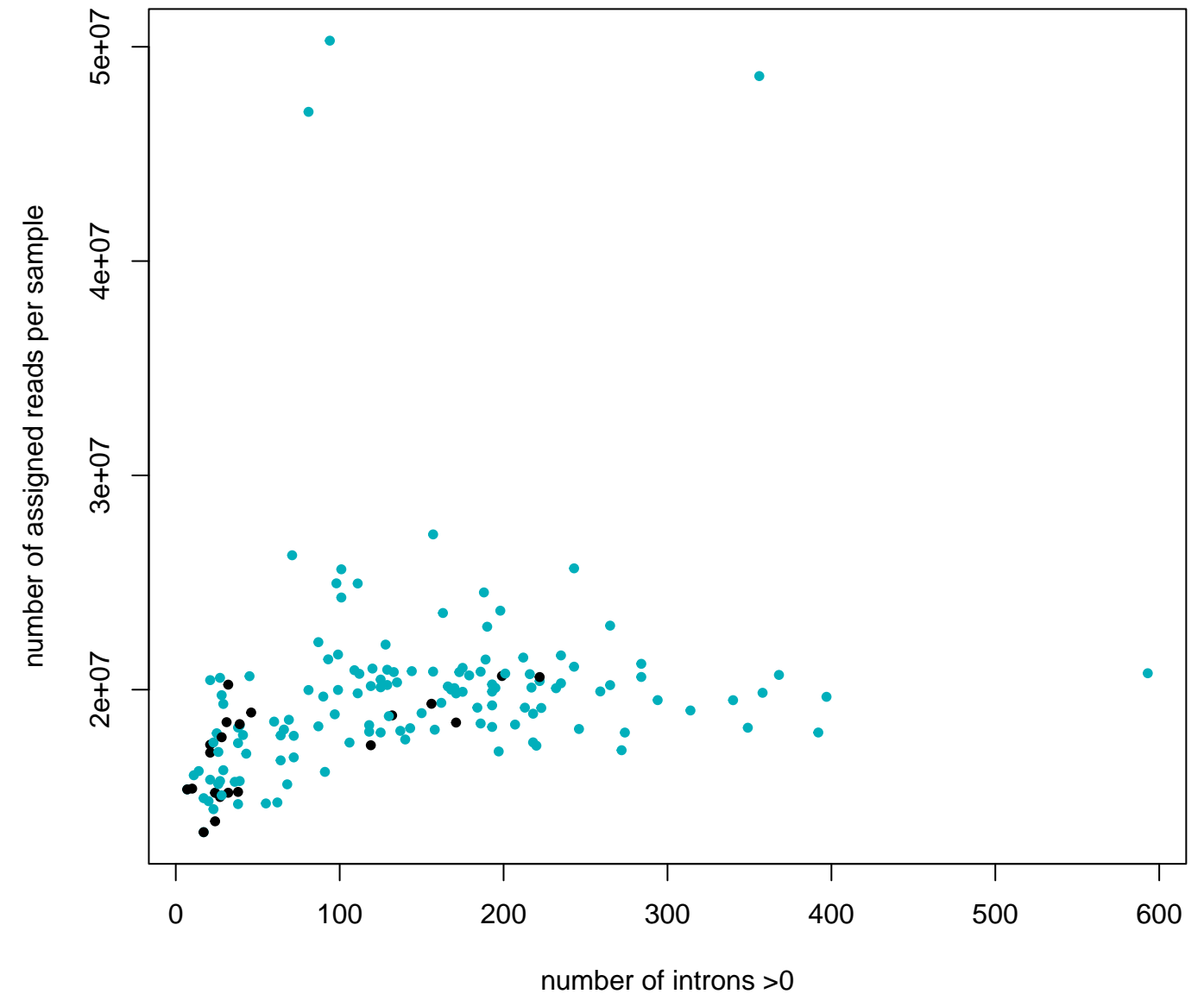
**Spearman correlation between number of total introns  
and unique introns per sample: 0.934014167677062 and  
0.929134489928458 for those classified for CRIS**



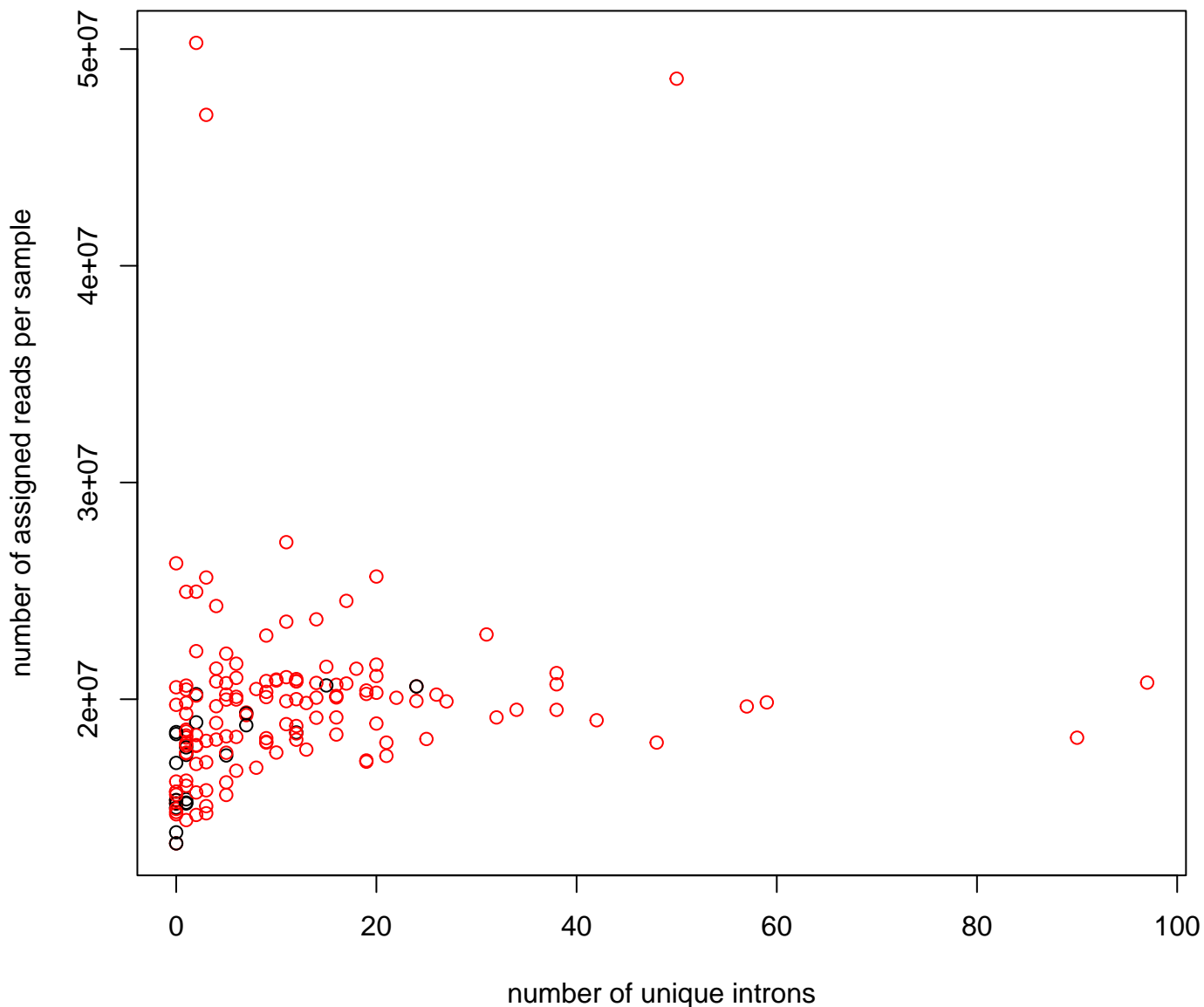
**Pearson correlation between number  
of total introns and assigned reads: 0.276785157834385 and  
0.220807395958334 for those classified for CRIS**



**Spearman correlation between number  
of total introns and assigned reads: 0.530549902152014 and  
0.438813622961837 for those classified for CRIS**



**Pearson correlation between number of total  
introns and assigned reads: 0.183576571247453 and  
0.139816516921745 for those classified for CRIS**



**Spearman correlation between number of total  
introns and assigned reads: 0.460221877994785 and  
0.356600175561678 for those classified for CRIS**

