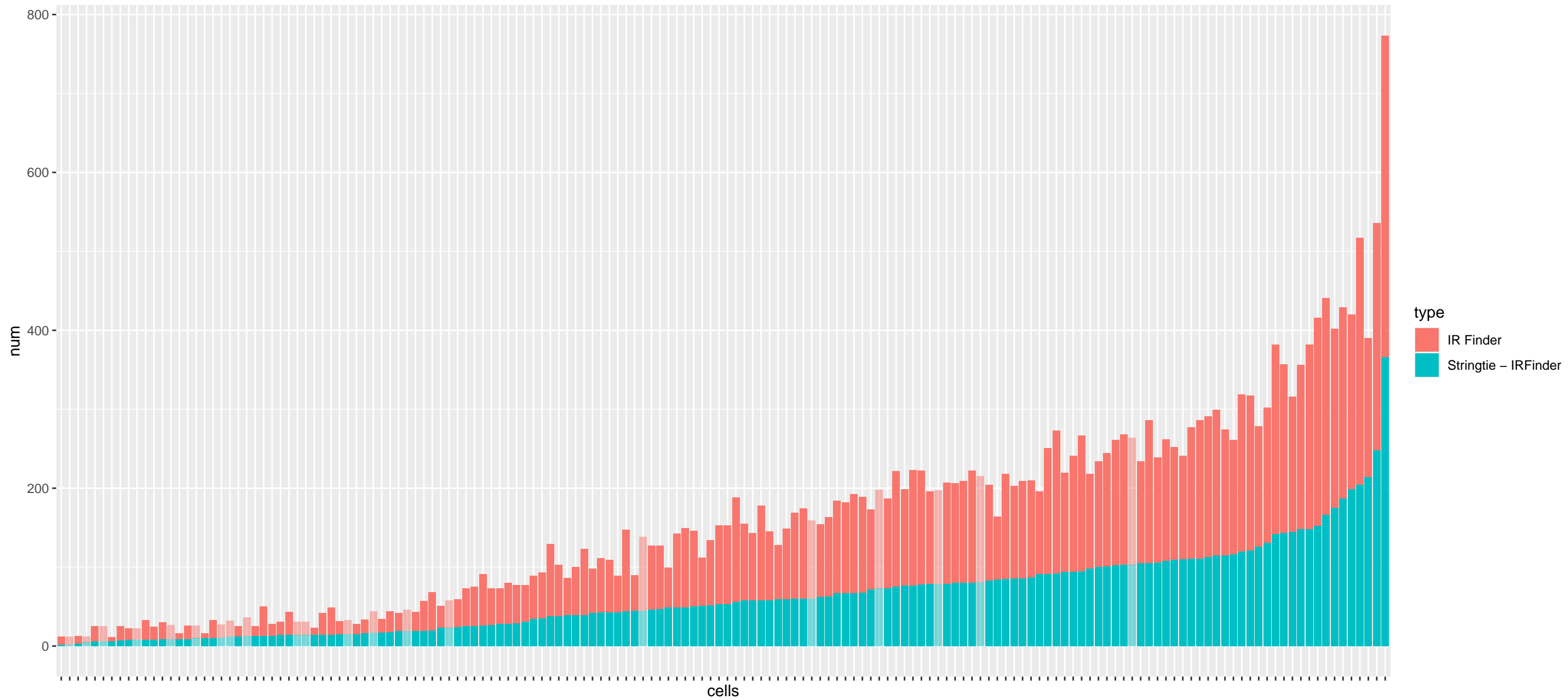
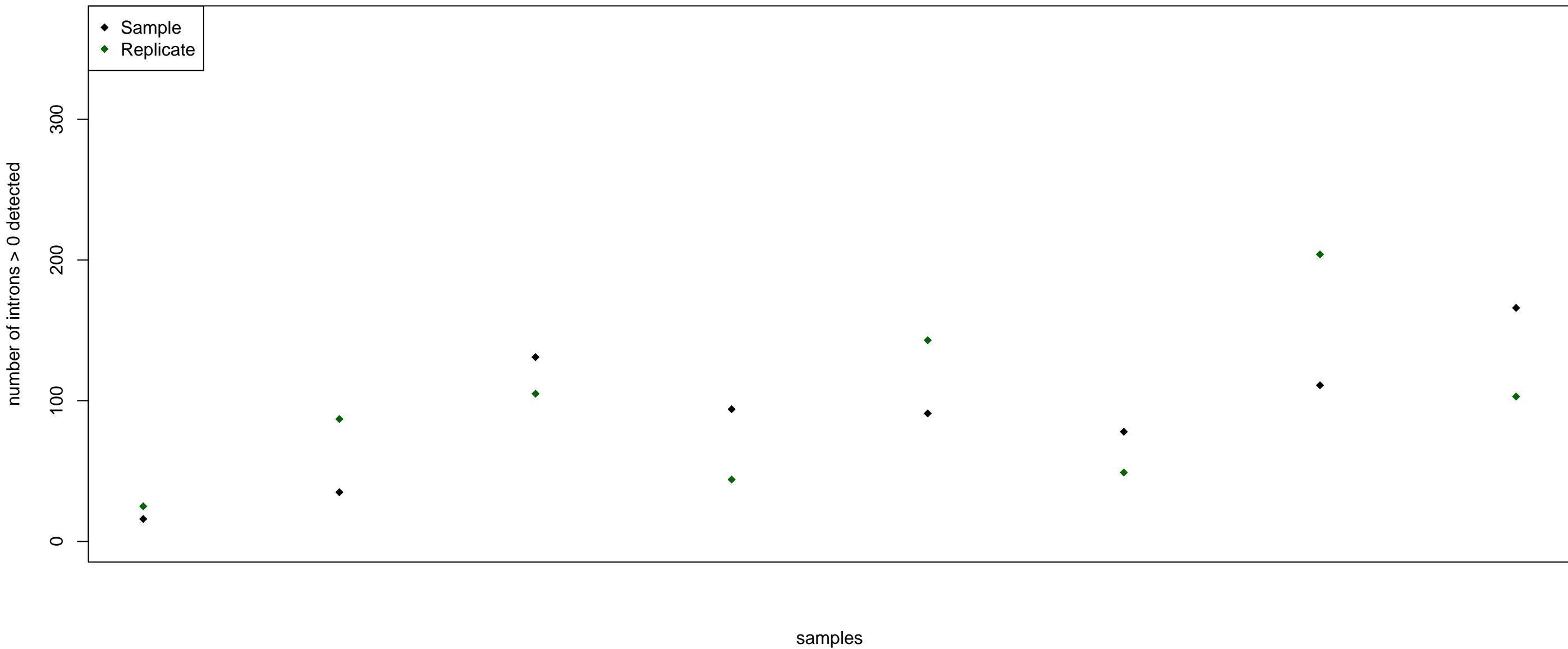


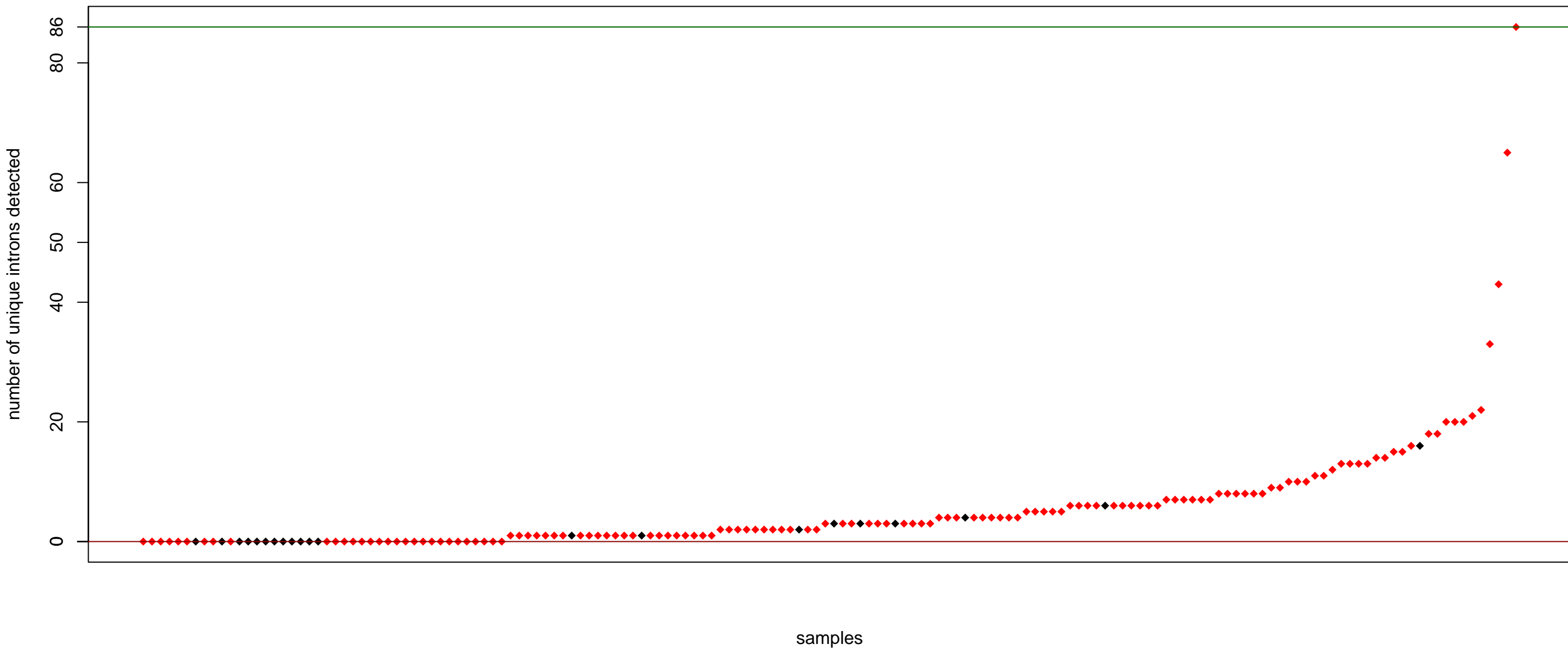
Intron retention events with at least 0.3 coverage



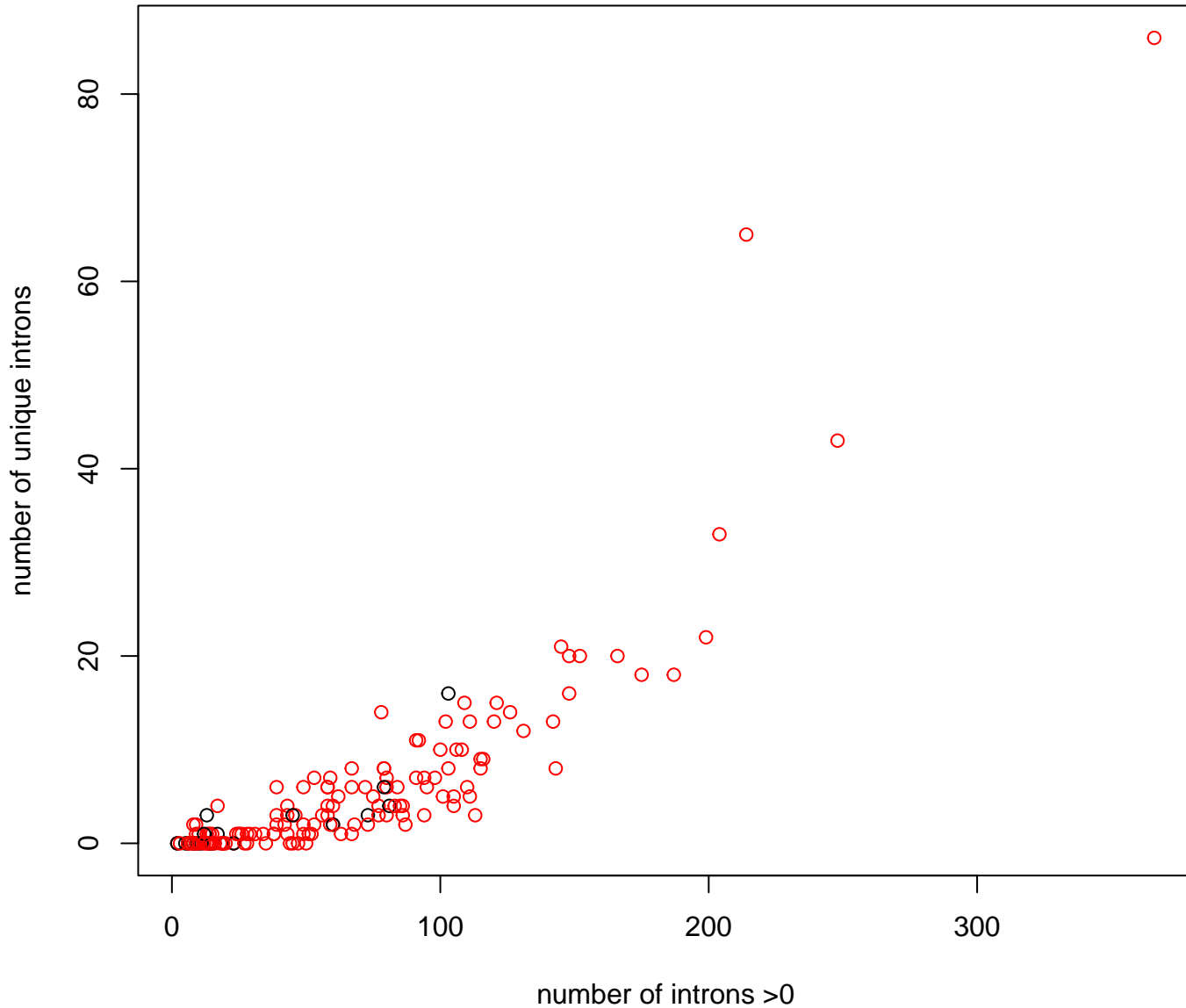
Intersected introns



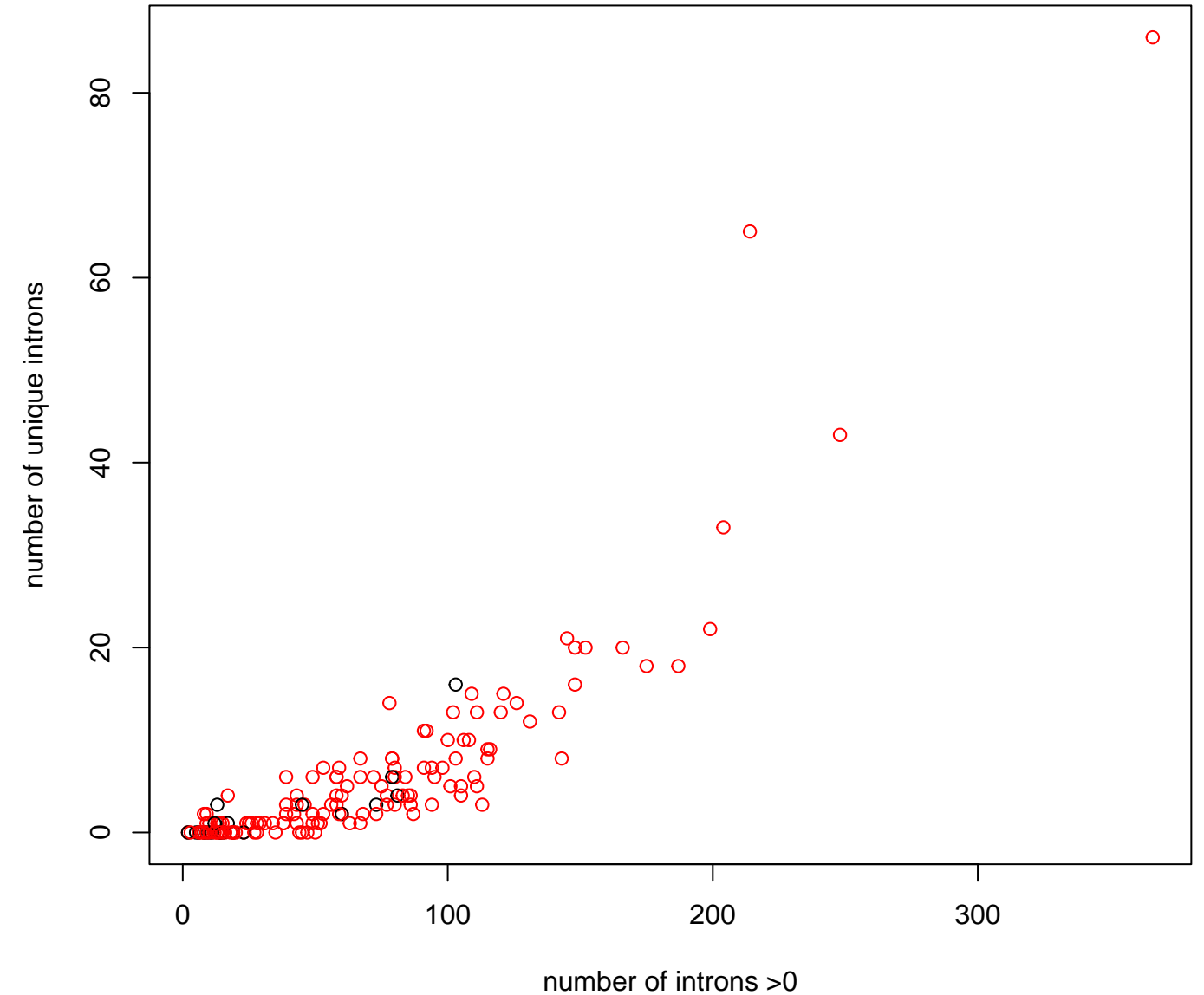
Total: 898 introns and 859 for CRIS classified



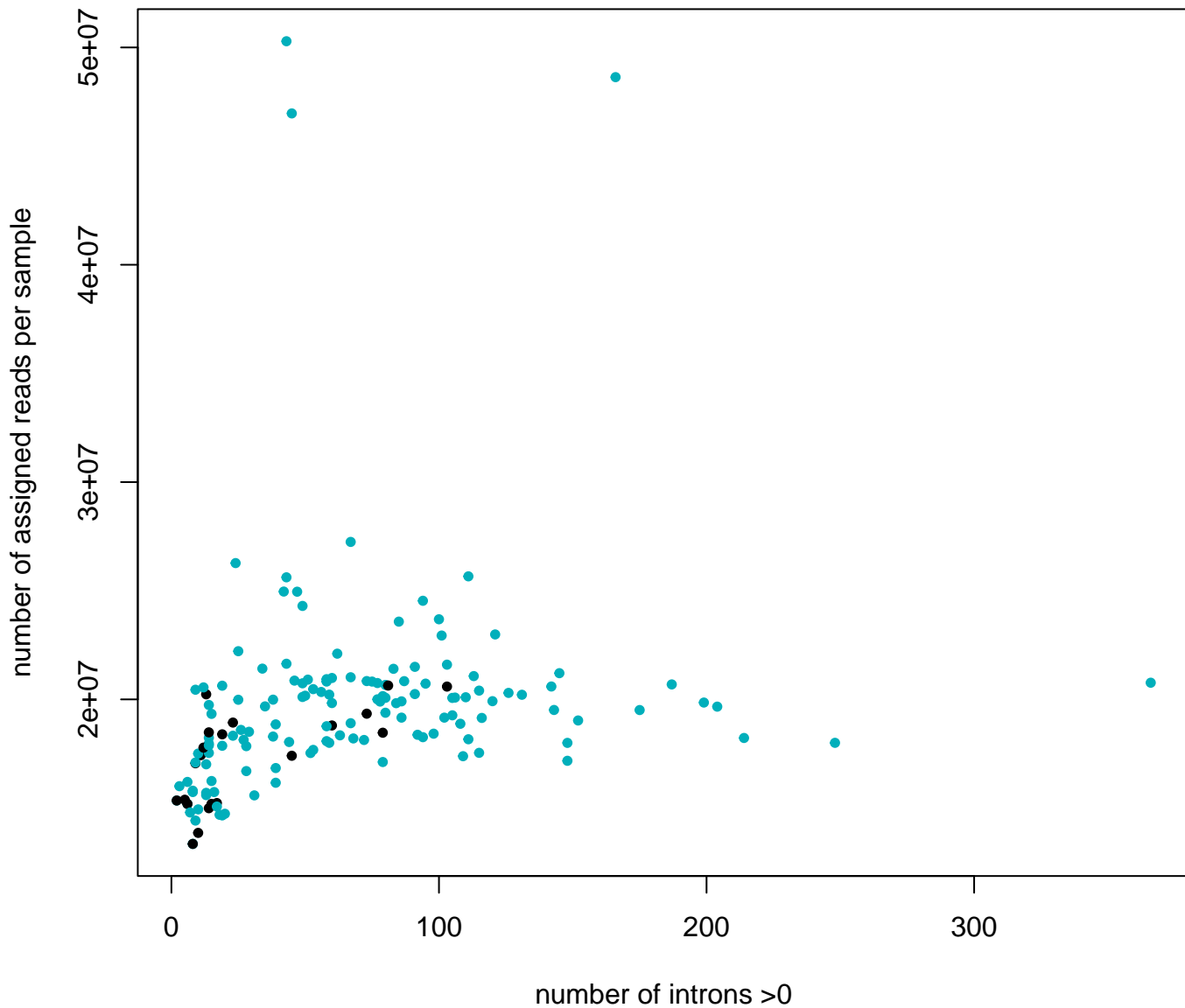
**Pearson correlation between number of total introns
and unique introns per sample: 0.859209135013674 and
0.862252861094247 for those classified for CRIS**



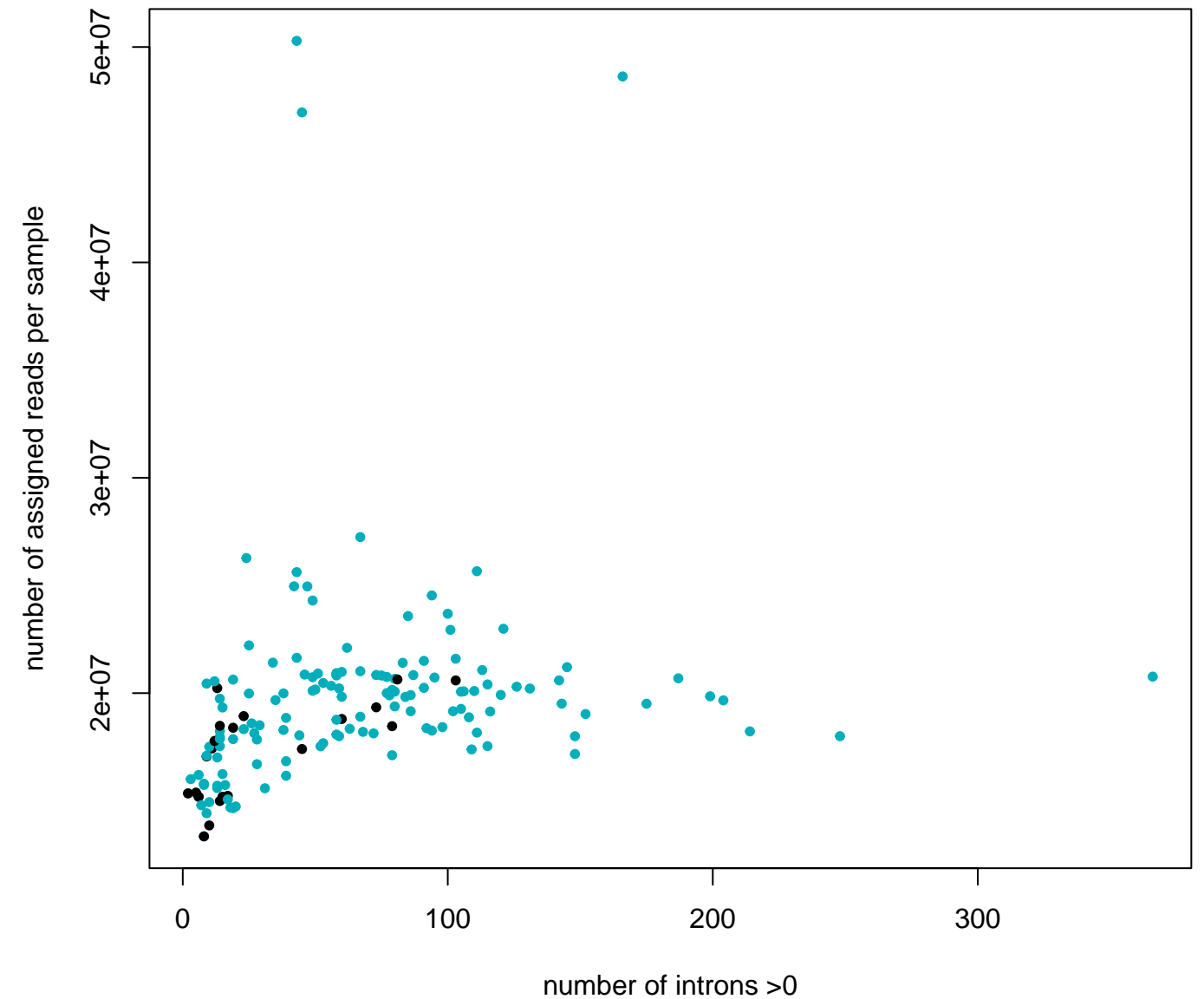
**Spearman correlation between number of total introns
and unique introns per sample: 0.878378448852275 and
0.878857284234974 for those classified for CRIS**



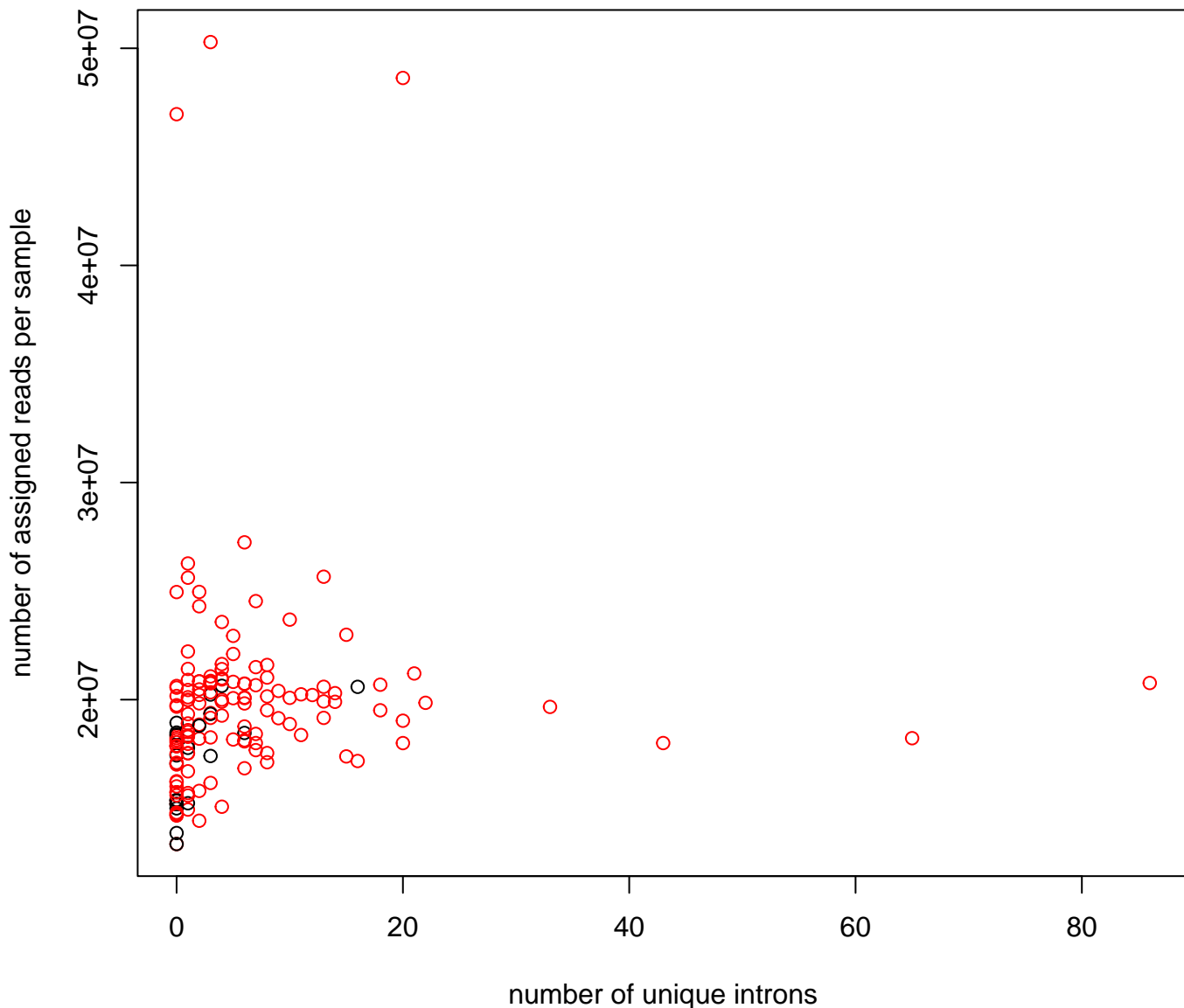
**Pearson correlation between number
of total introns and assigned reads: 0.238639036848374 and
0.185016229519923 for those classified for CRIS**



**Spearman correlation between number
of total introns and assigned reads: 0.520370108119765 and
0.426703759862017 for those classified for CRIS**



**Pearson correlation between number of total
introns and assigned reads: 0.111646712329477 and
0.0758240912388996 for those classified for CRIS**



**Spearman correlation between number of total
introns and assigned reads: 0.429508406702596 and
0.333590679604399 for those classified for CRIS**

