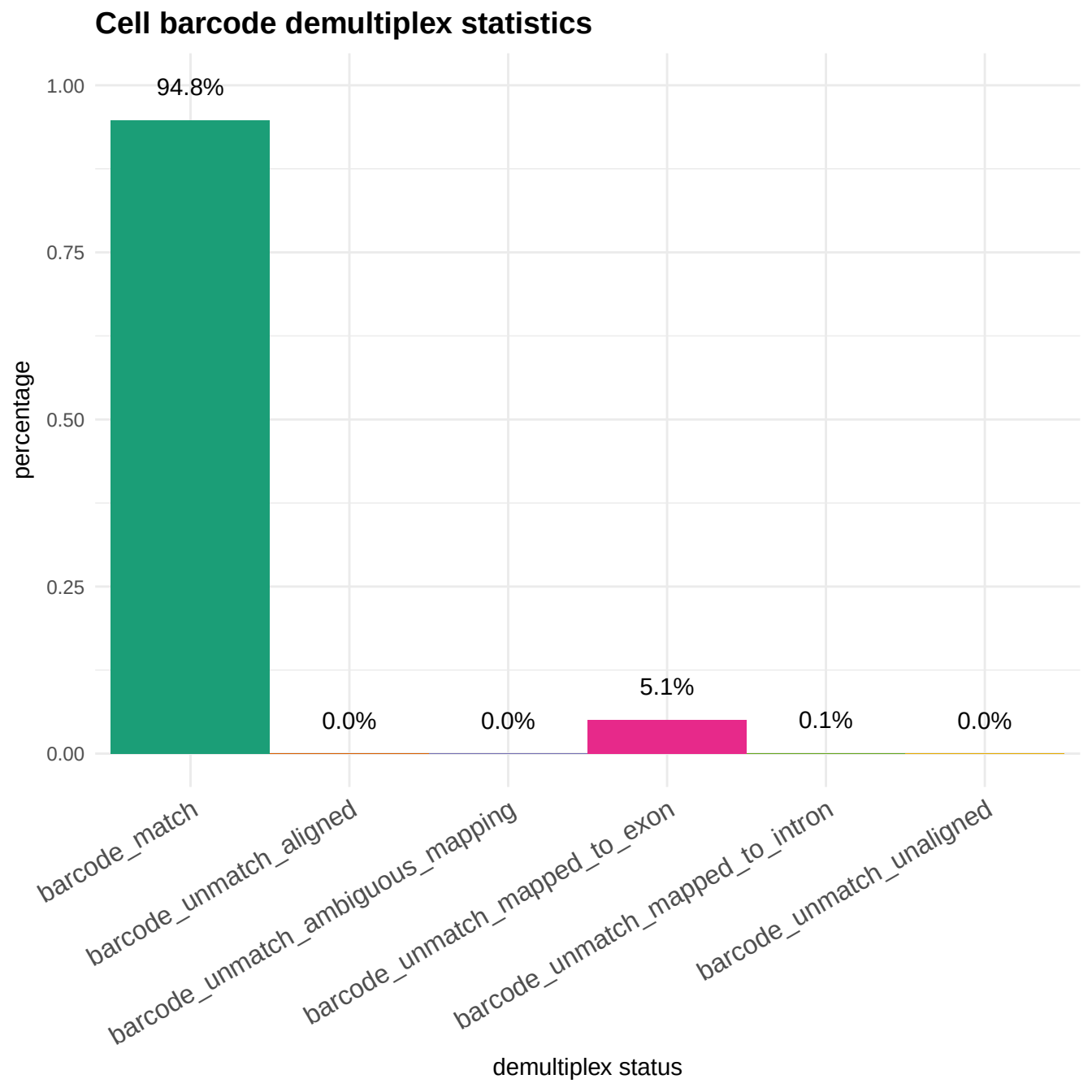
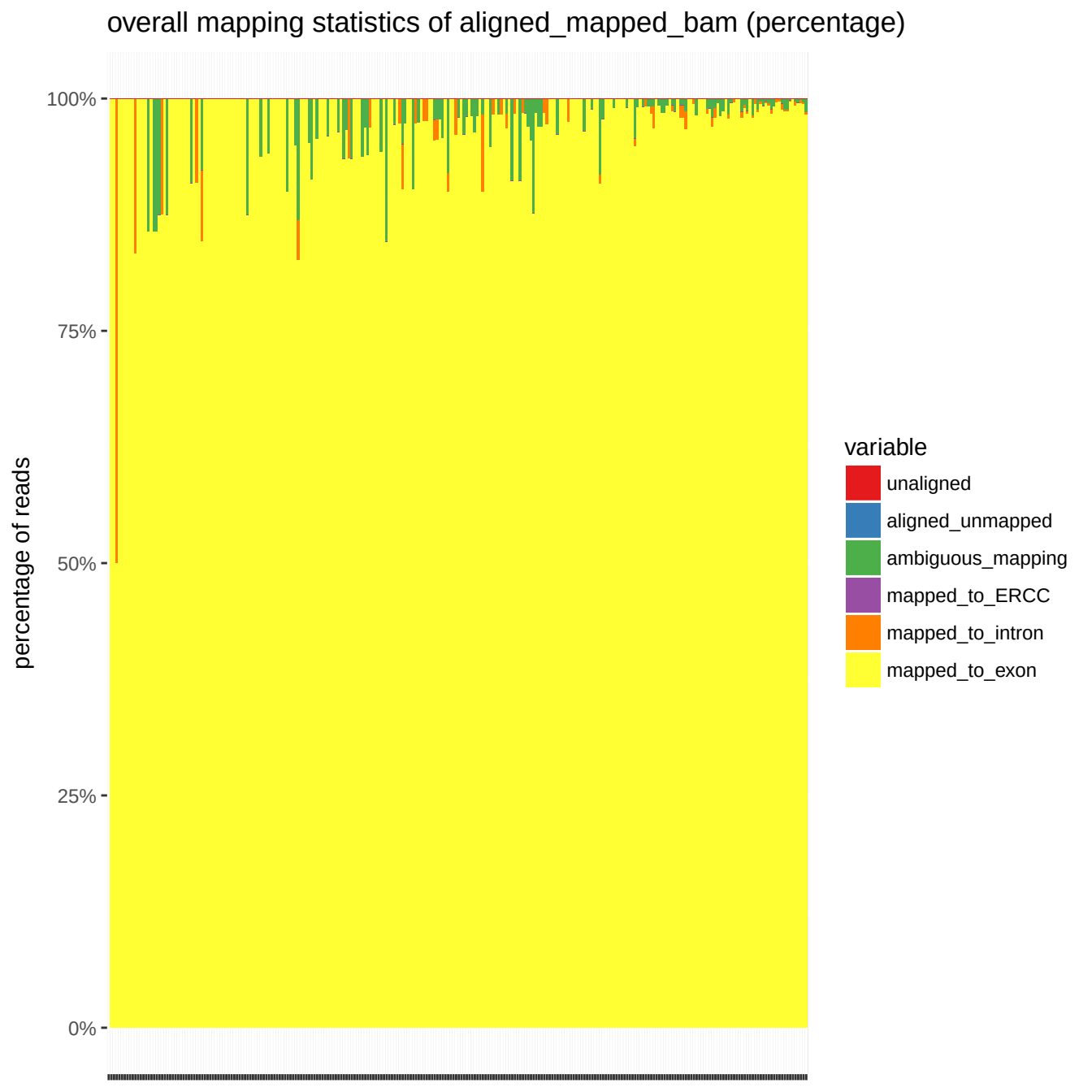
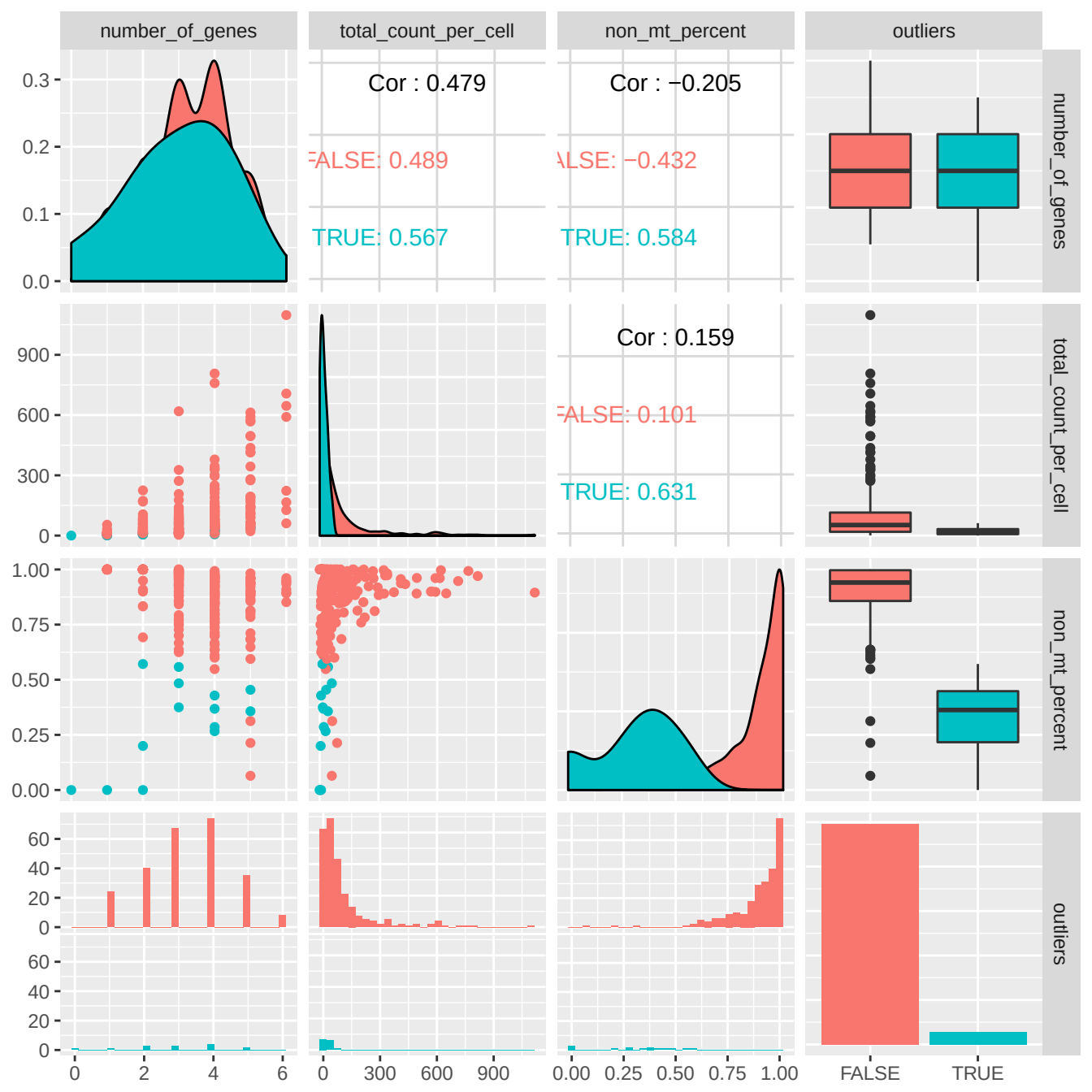

Cell barcode demultiplex statistics
1.00
0.75
0.50
0.25
0.00
percentage
94.8%
0.0%
0.0%
5.1%
0.1%
0.0%
barcode_match
barcode_unmatch_aligned
barcode_unmatch_ambiguous_mapping
barcode_unmatch_mapped_to_exon
barcode_unmatch_mapped_to_intron
barcode_unmatch_unaligned
demultiplex status


overall mapping statistics of aligned_mapped_bam (percentage)
100%
75%
50%
25%
0%
percentage of reads
variable
unaligned
aligned_unmapped
ambiguous_mapping
mapped_to_ERCC
mapped_to_intron
mapped_to_exon


number_of_genes
total_count_per_cell
non_mt_percent
outliers
Cor : 0.479
FALSE: 0.489
TRUE: 0.567
Cor : −0.205
ALSE: −0.432
TRUE: 0.584
Cor : 0.159
FALSE: 0.101
TRUE: 0.631
0.3
0.2
0.1
0.0
900
600
300
0
1.00
0.75
0.50
0.25
0.00
60
40
20
0
0
2
4
6
0
300
600
900
0.00
0.25
0.50
0.75
1.00
FALSE
TRUE