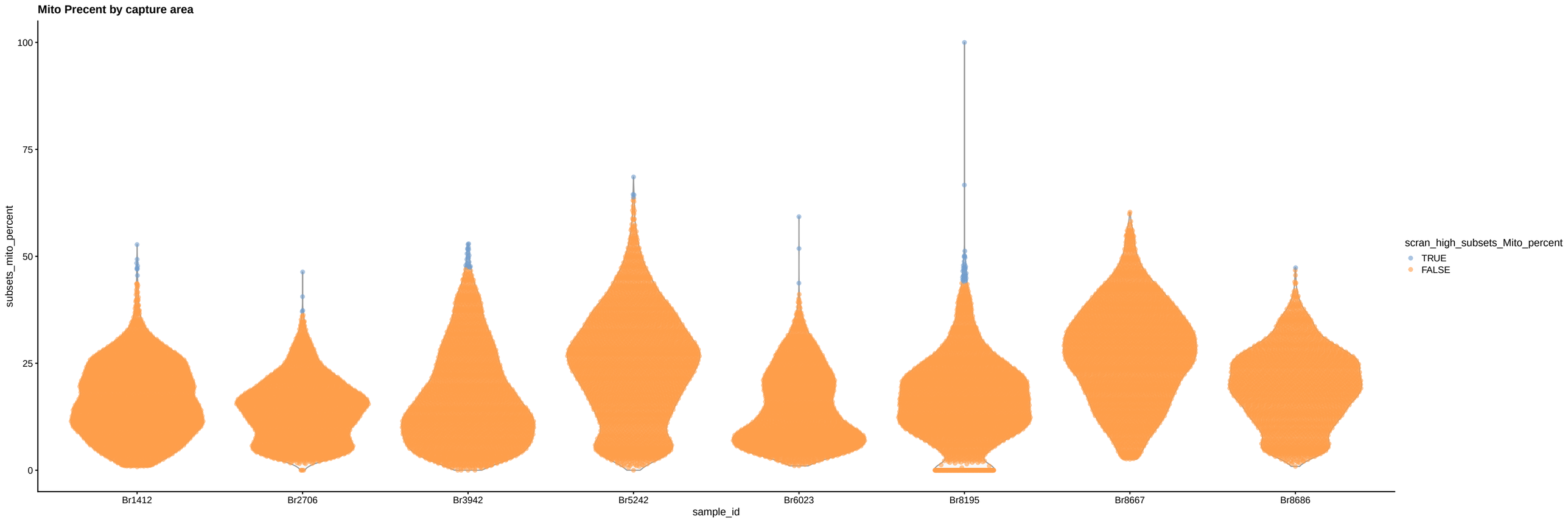
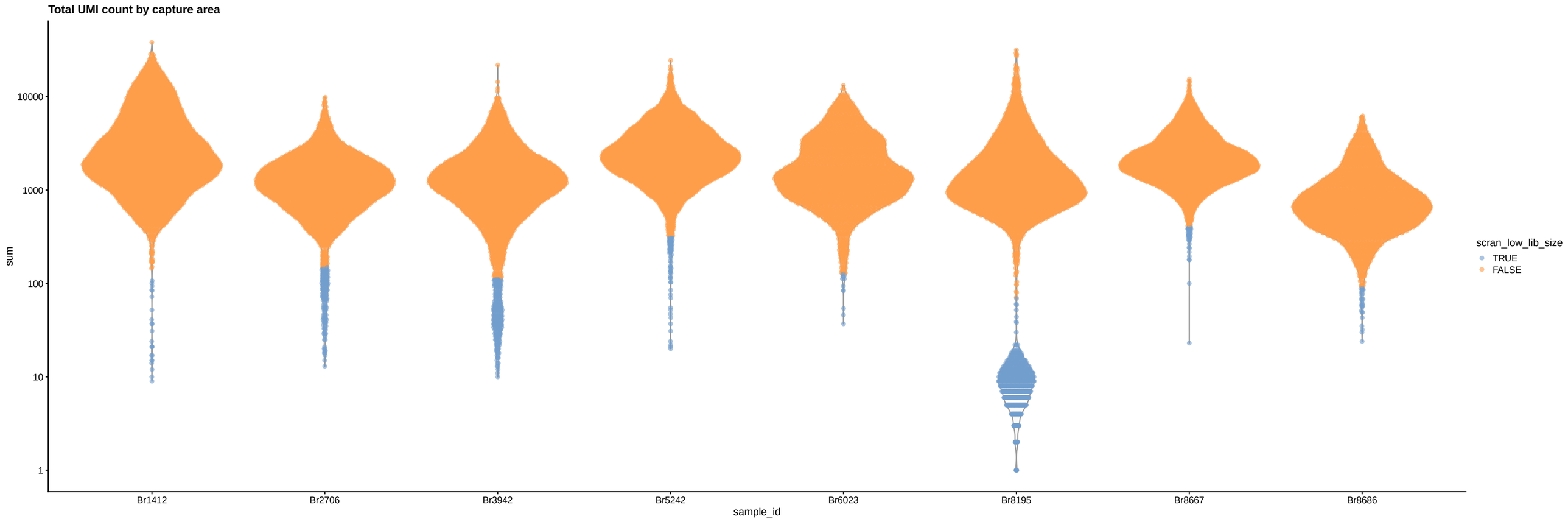
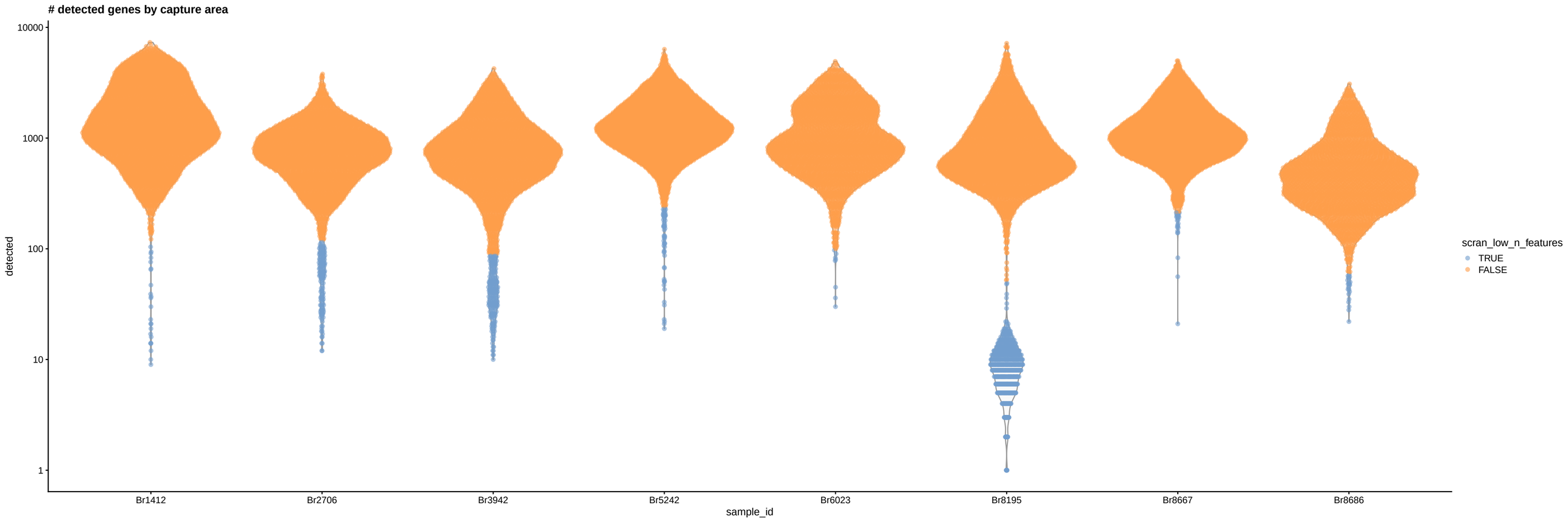
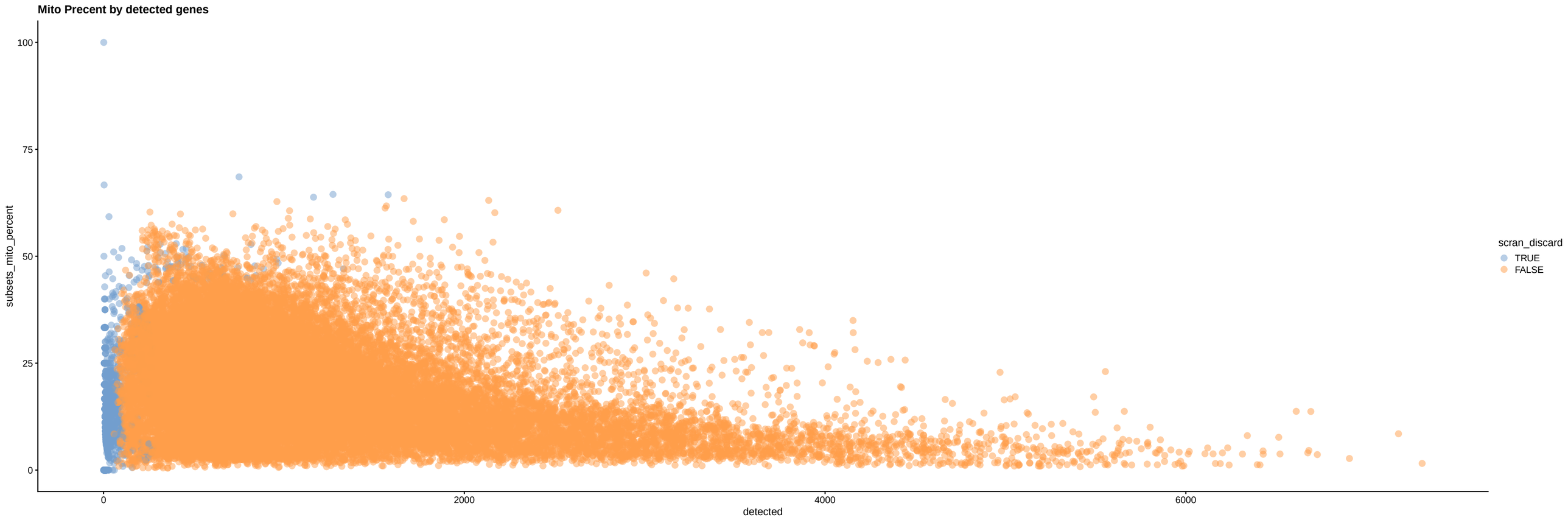

Mito Precent by capture area
subsets_mito_percent
100
75
50
25
0
Br1412
Br2706
Br3942
Br5242
Br6023
Br8195
Br8667
Br8686
sample_id
scran_high_subsets_Mito_percent
TRUE
FALSE


Total UMI count by capture area
sum
10000
1000
100
10
1
Br1412
Br2706
Br3942
Br5242
Br6023
Br8195
Br8667
Br8686
sample_id
scran_low_lib_size
TRUE
FALSE


detected genes by capture area
10000
1000
100
10
1
detected
Br1412
Br2706
Br3942
Br5242
Br6023
Br8195
Br8667
Br8686
sample_id
scran_low_n_features
TRUE
FALSE


Mito Precent by detected genes
subsets_mito_percent
100
75
50
25
0
0
2000
4000
6000
detected
scran_discard
TRUE
FALSE