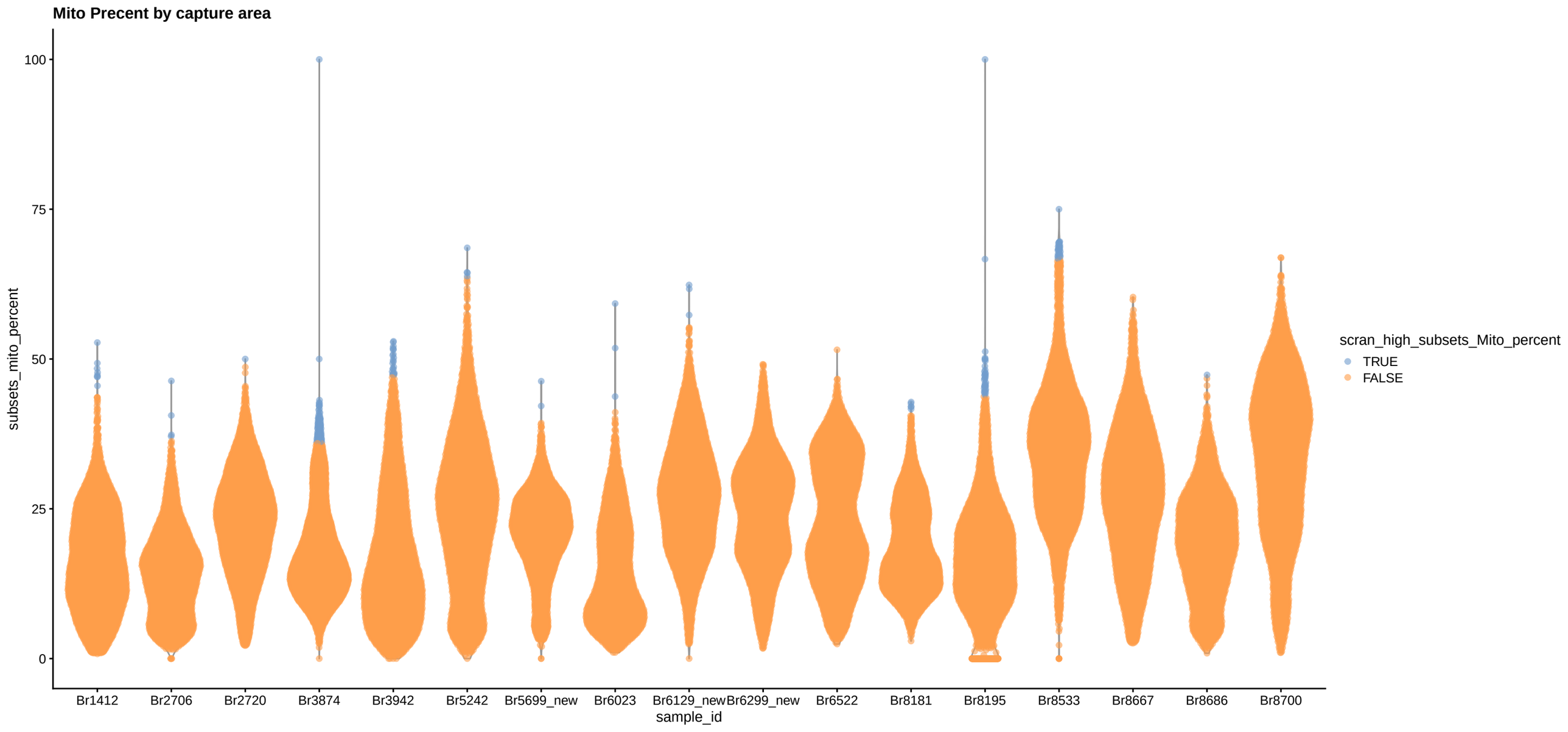
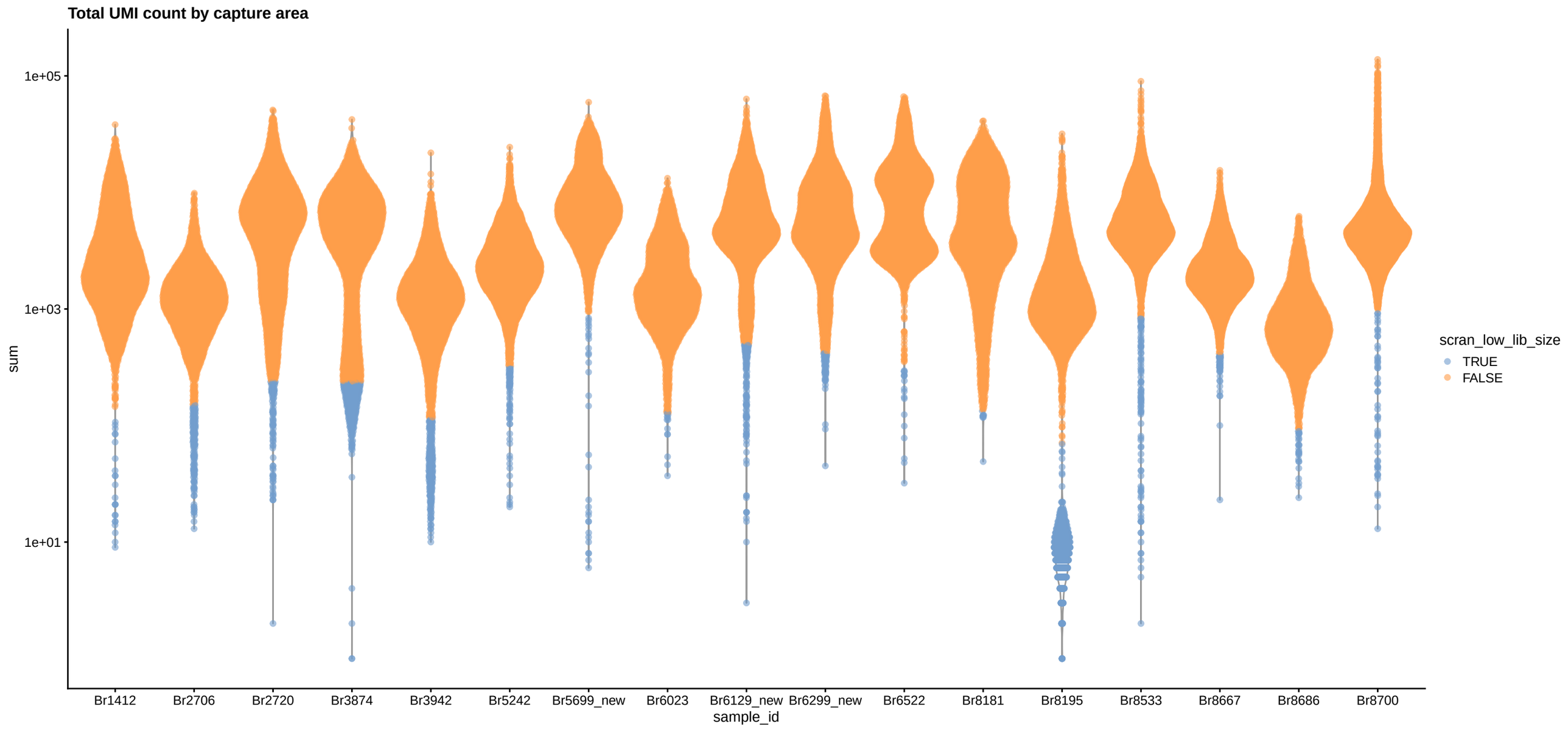
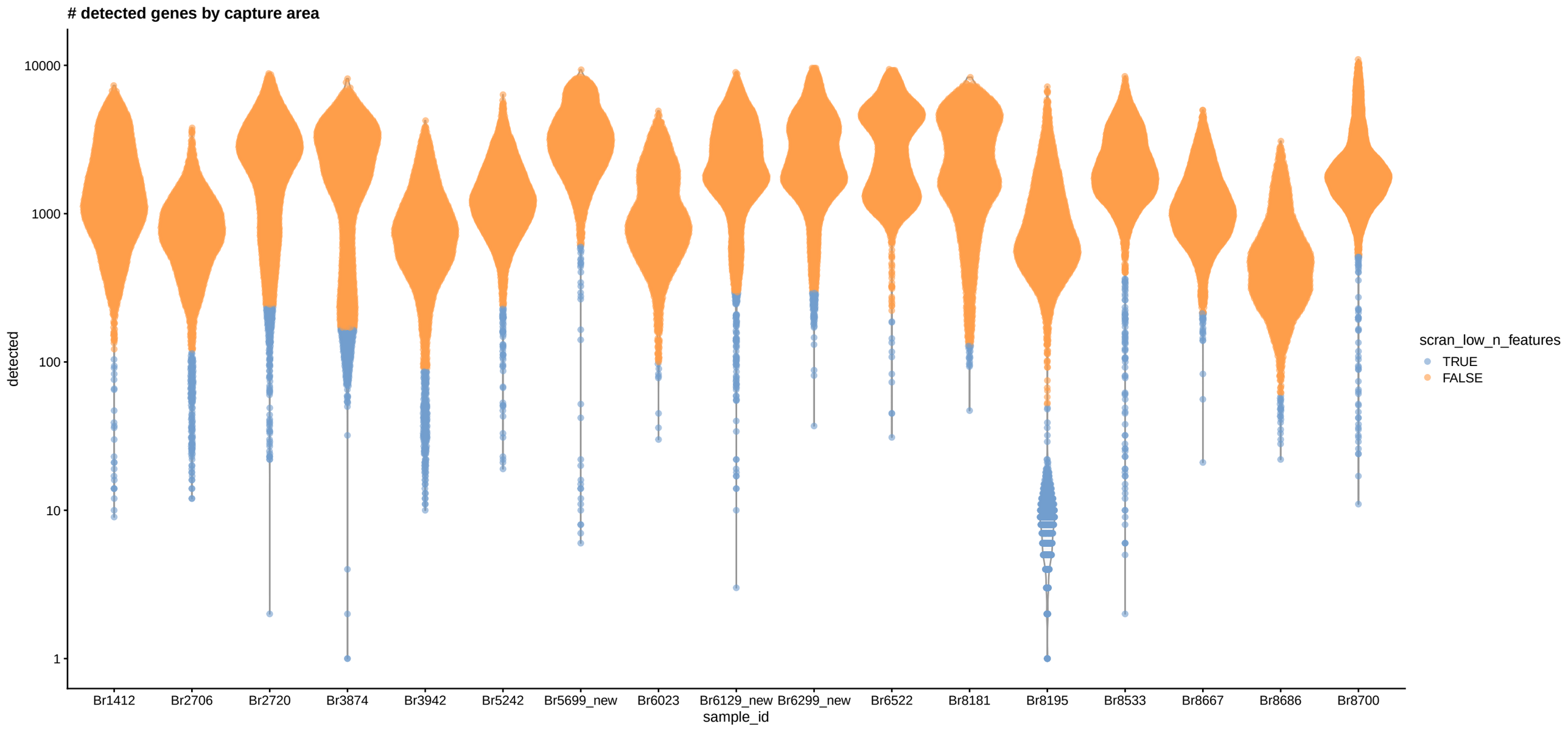
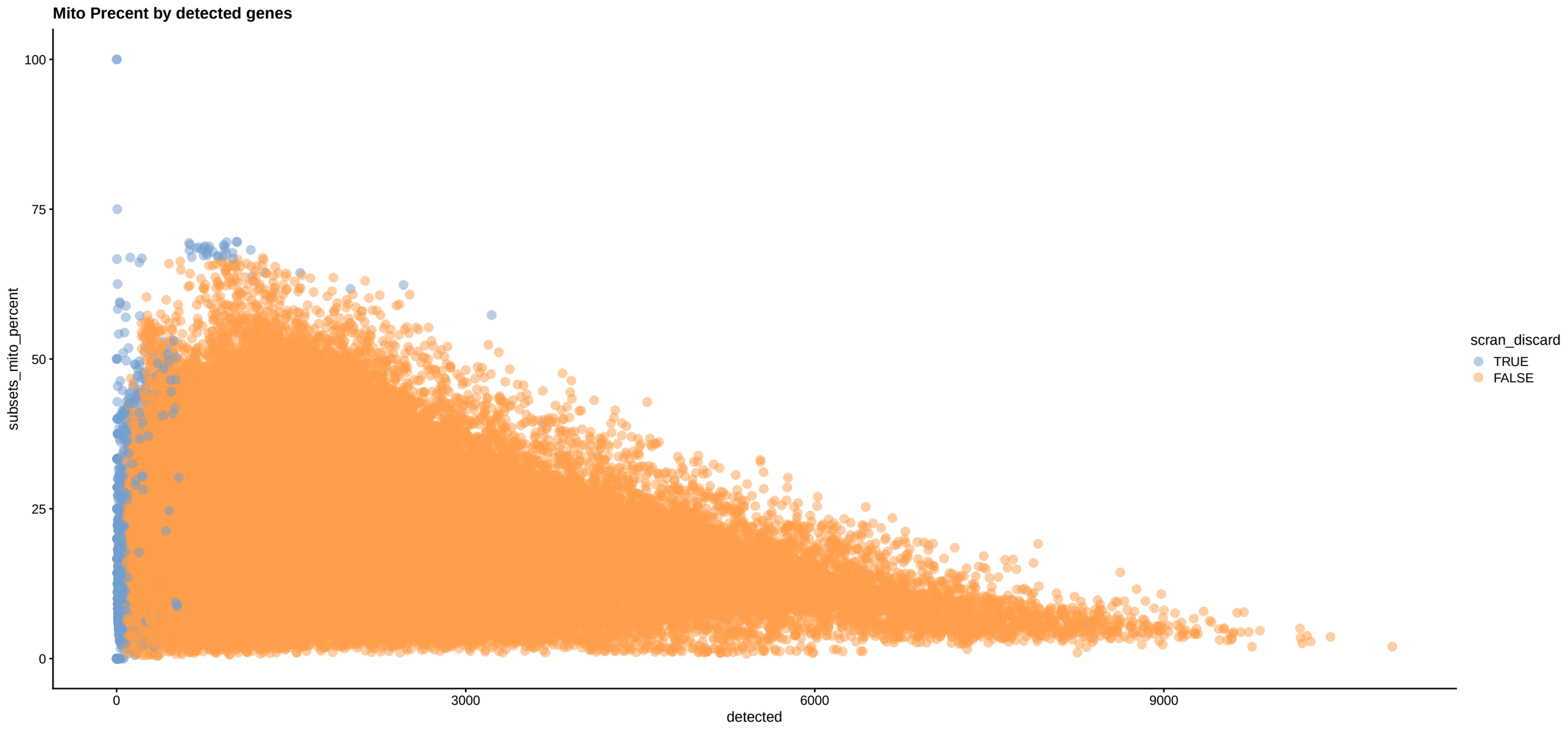

Mito Precent by capture area
subsets_mito_percent
100
75
50
25
0
Br1412
Br2706
Br2720
Br3874
Br3942
Br5242
Br5699_new
Br6023
Br6129_new
Br6299_new
Br6522
Br8181
Br8195
Br8533
Br8667
Br8686
Br8700
sample_id
scran_high_subsets_Mito_percent
TRUE
FALSE


Total UMI count by capture area
1e+05
1e+03
1e+01
sum
Br1412
Br2706
Br2720
Br3874
Br3942
Br5242
Br5699_new
Br6023
Br6129_new
Br6299_new
Br6522
Br8181
Br8195
Br8533
Br8667
Br8686
Br8700
sample_id
scran_low_lib_size
TRUE
FALSE


detected genes by capture area
detected
10000
1000
100
10
1
Br1412
Br2706
Br2720
Br3874
Br3942
Br5242
Br5699_new
Br6023
Br6129_new
Br6299_new
Br6522
Br8181
Br8195
Br8533
Br8667
Br8686
Br8700
sample_id
scran_low_n_features
TRUE
FALSE


Mito Precent by detected genes
subsets_mito_percent
100
75
50
25
0
0
3000
6000
9000
detected
scran_discard
TRUE
FALSE