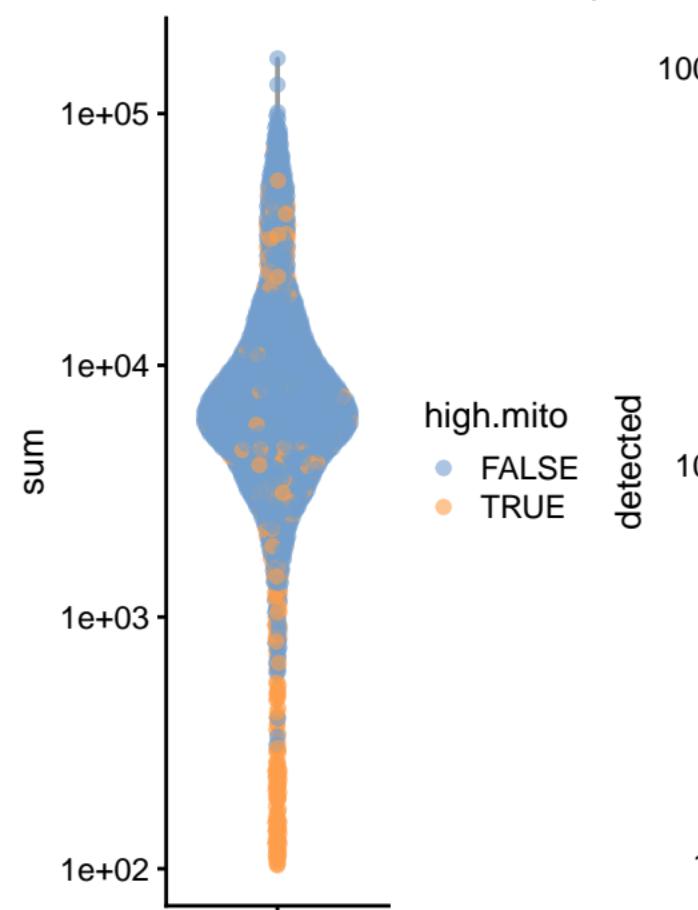
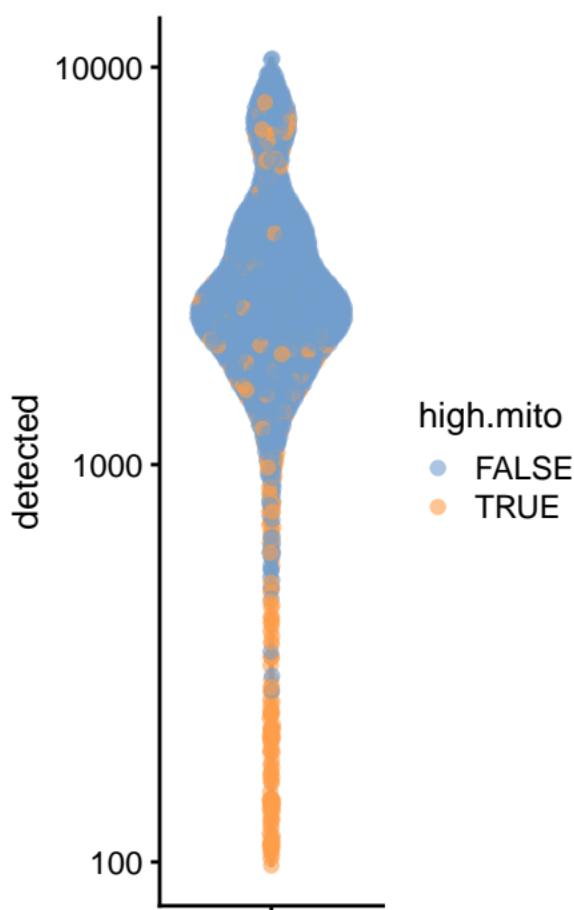


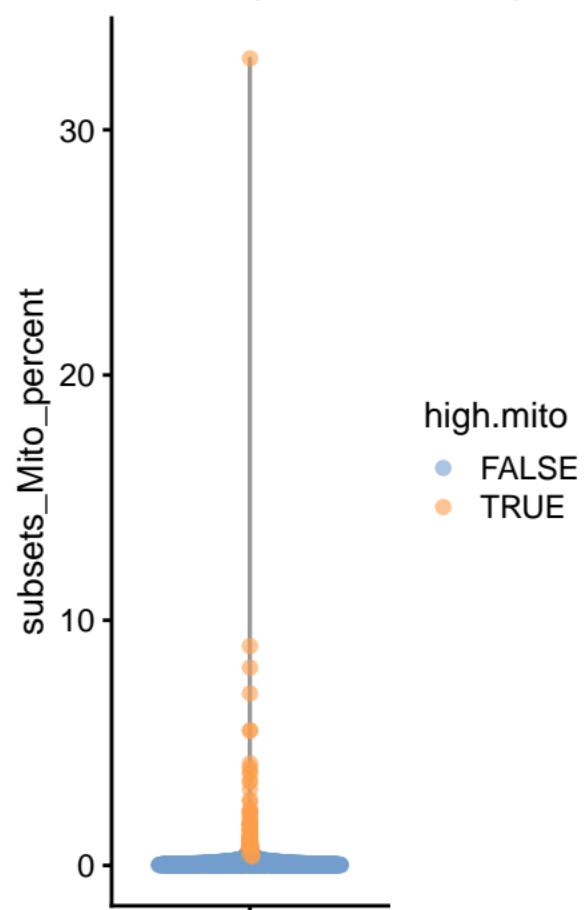
Total count: br5161.amy



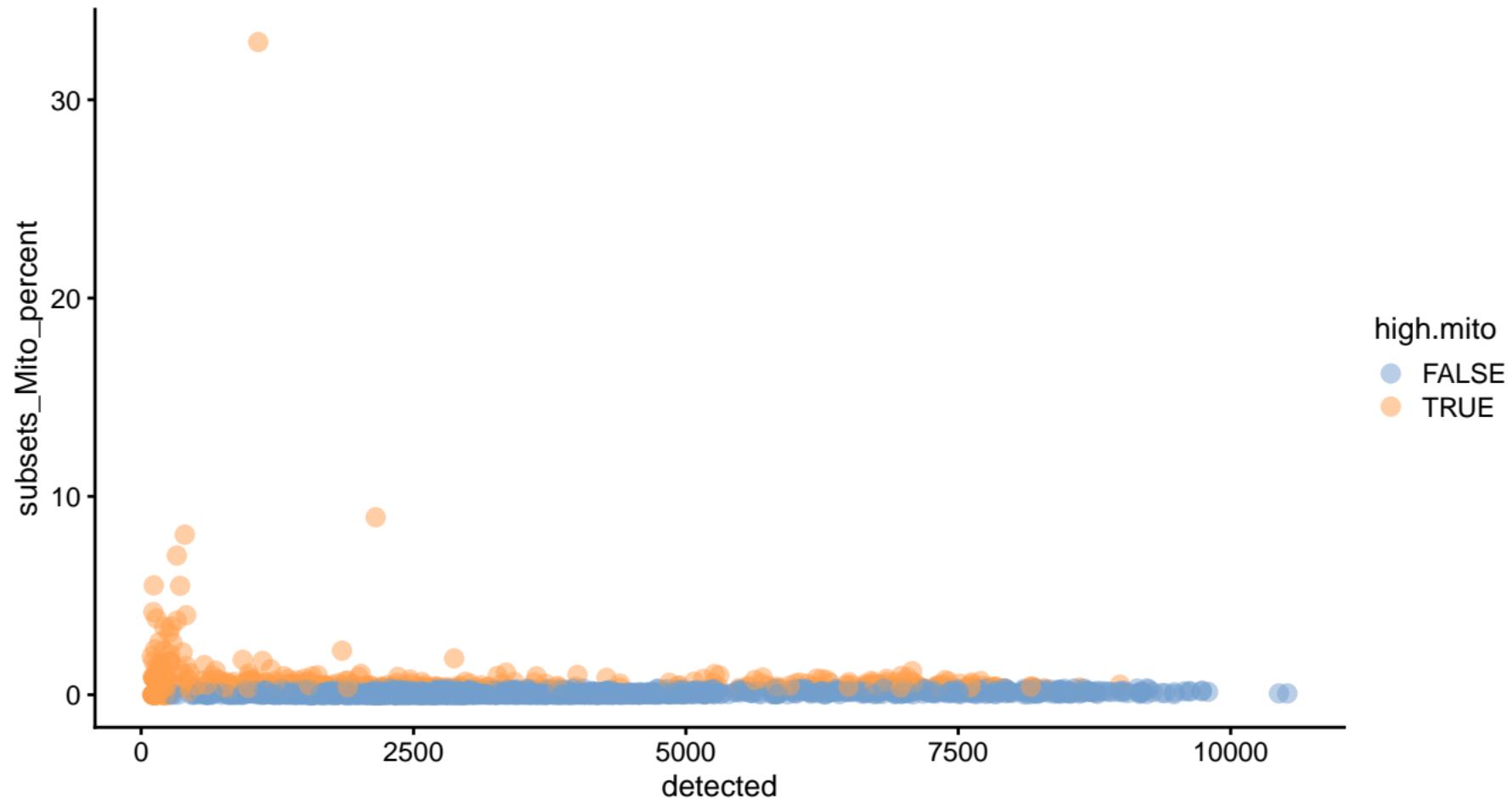
Detected features



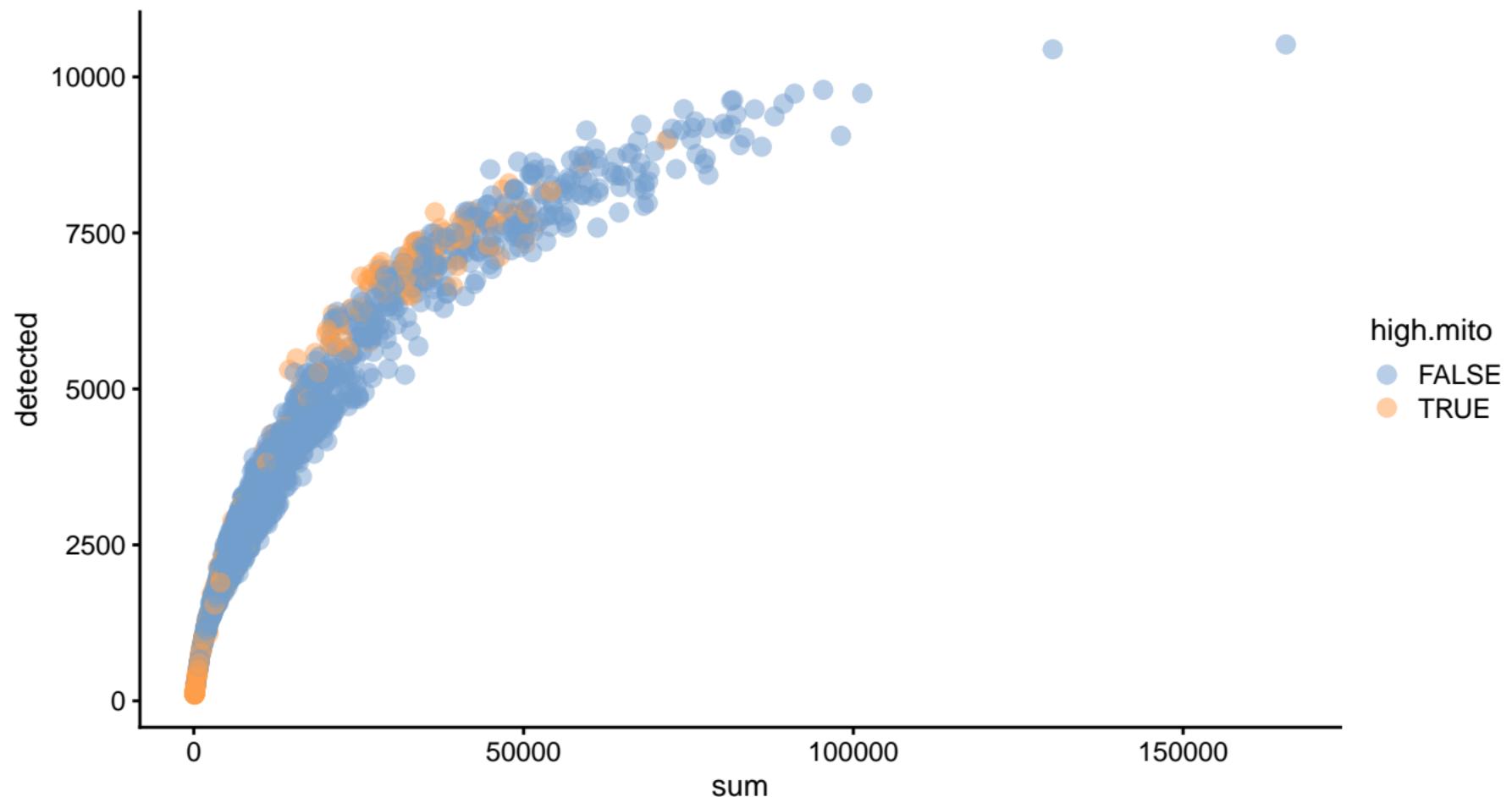
Mito % (cutoff = 0.321)



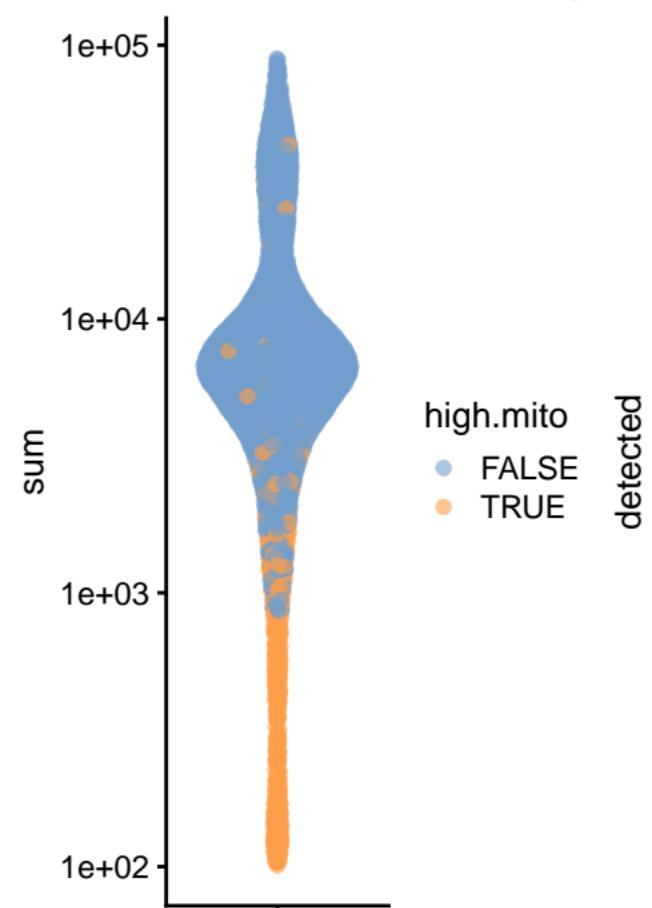
Sample: br5161.amy; pre-QC nNuclei: 3687; nNuclei kept: 3294 (0.89%)



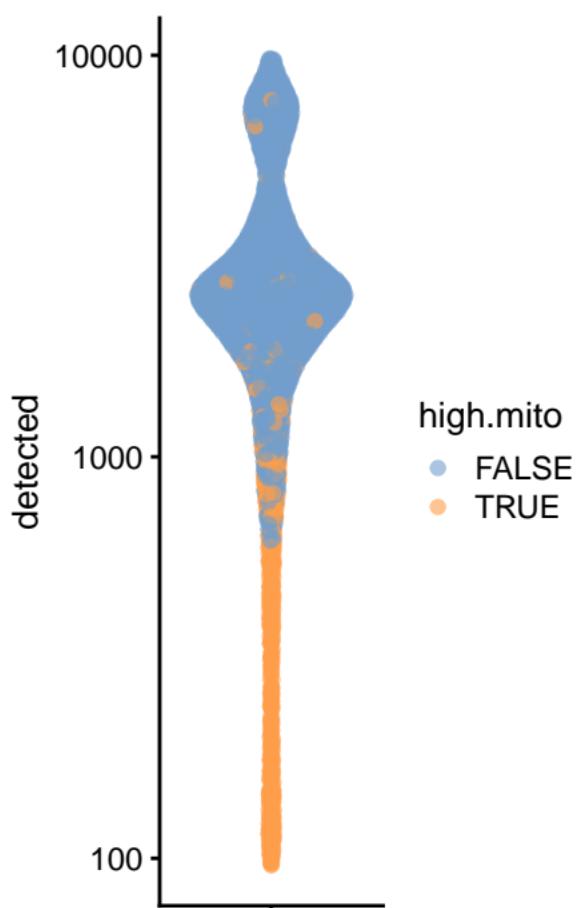
Sample: br5161.amy; pre-QC nNuclei: 3687; nNuclei kept: 3294 (0.89%)



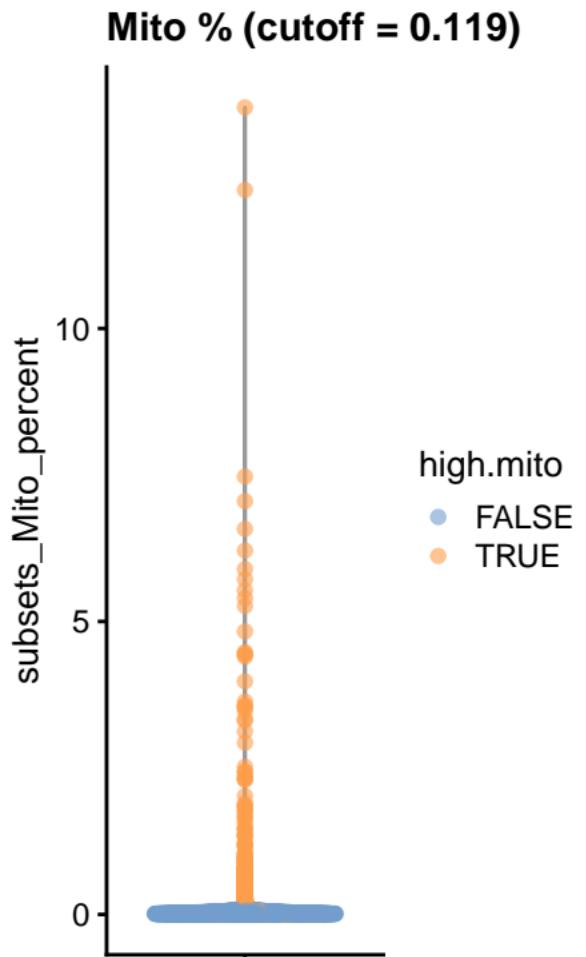
Total count: br5161.dlpfc



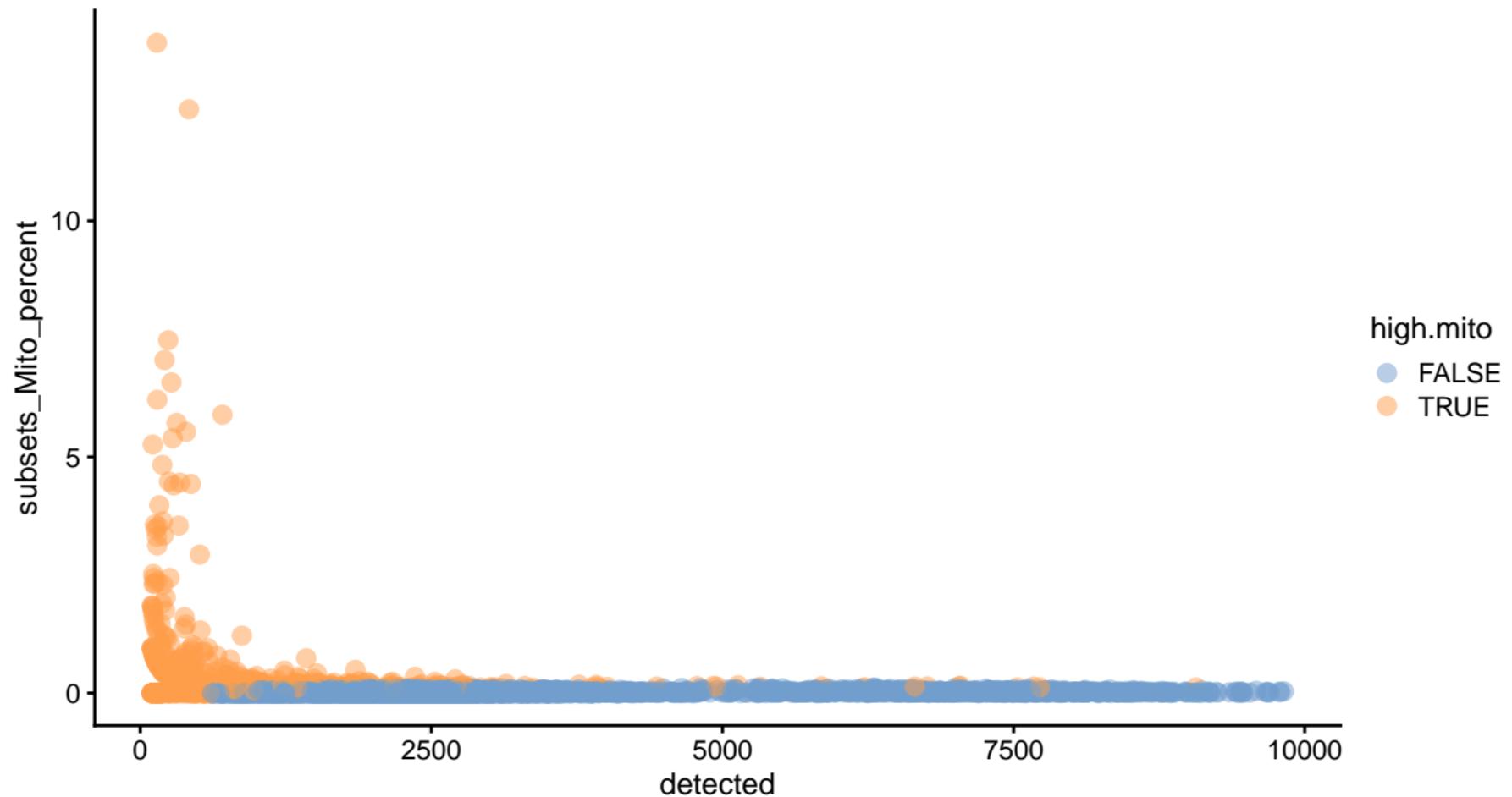
Detected features



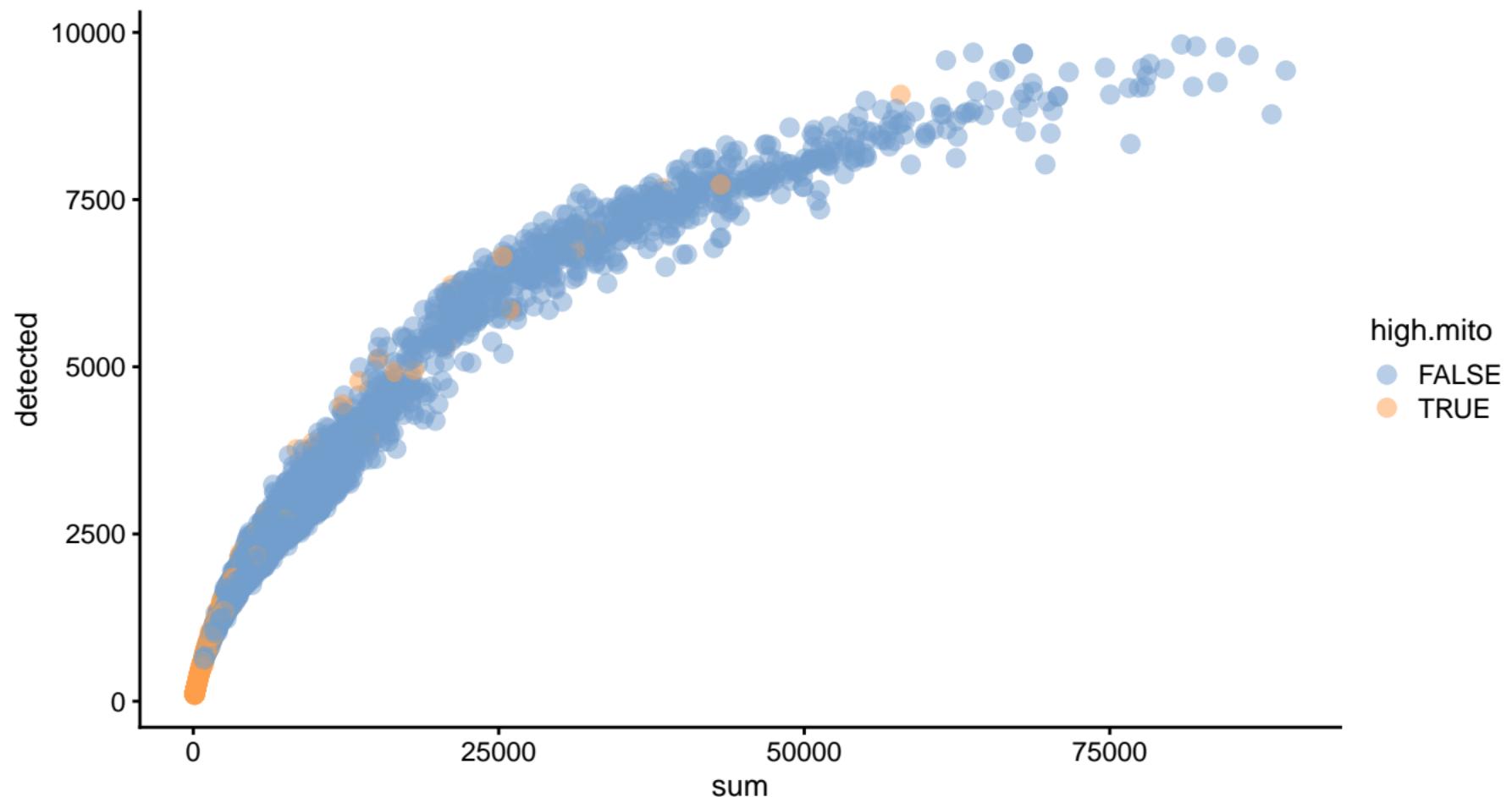
Mito % (cutoff = 0.119)



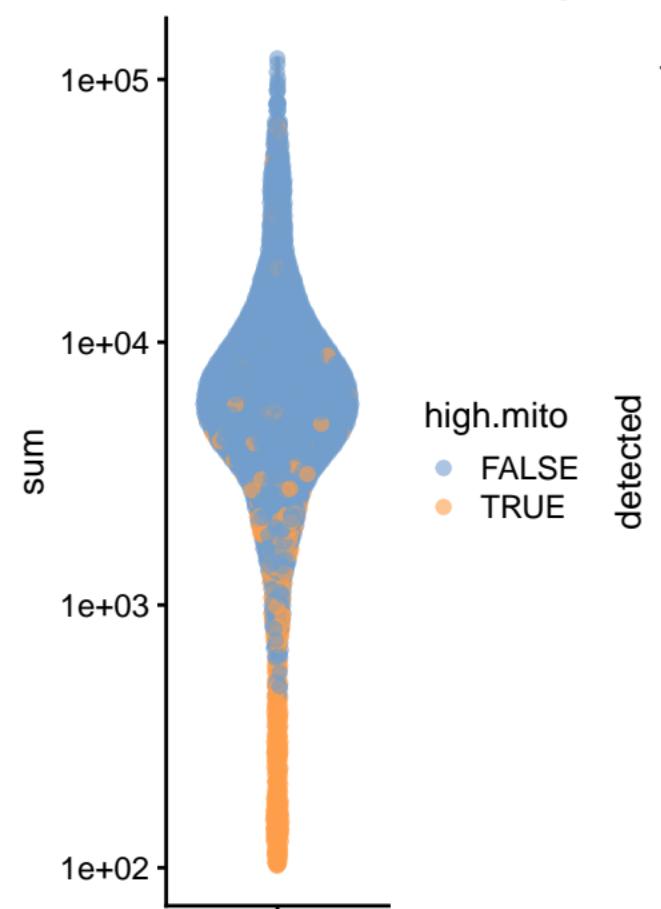
Sample: br5161.dlpfc; pre-QC nNuclei: 4788; nNuclei kept: 4215 (0.88%)



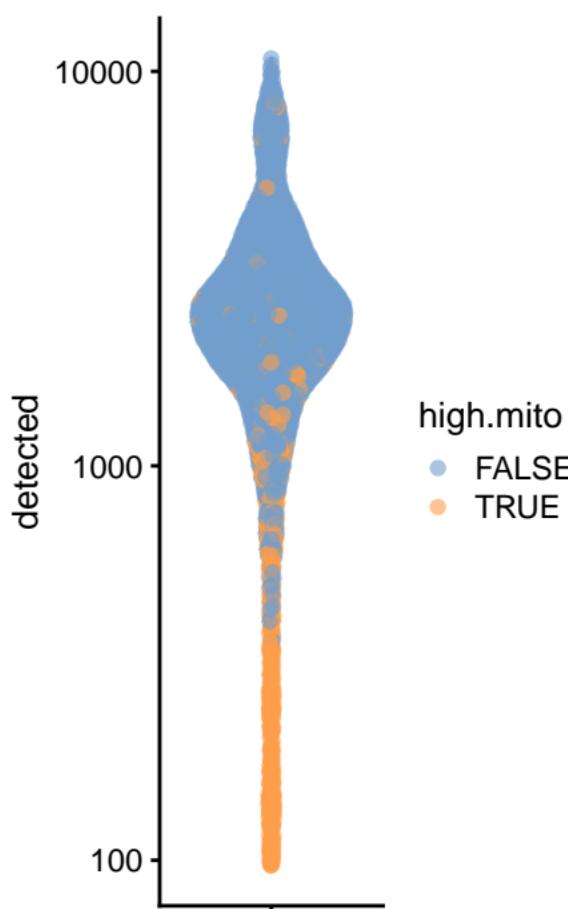
Sample: br5161.dlpfc; pre-QC nNuclei: 4788; nNuclei kept: 4215 (0.88%)



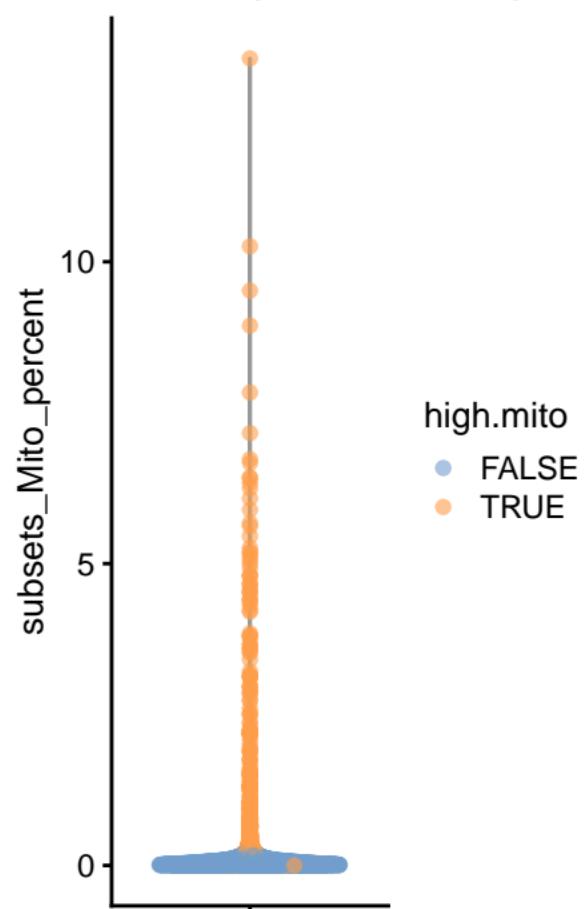
Total count: br5161.hpc



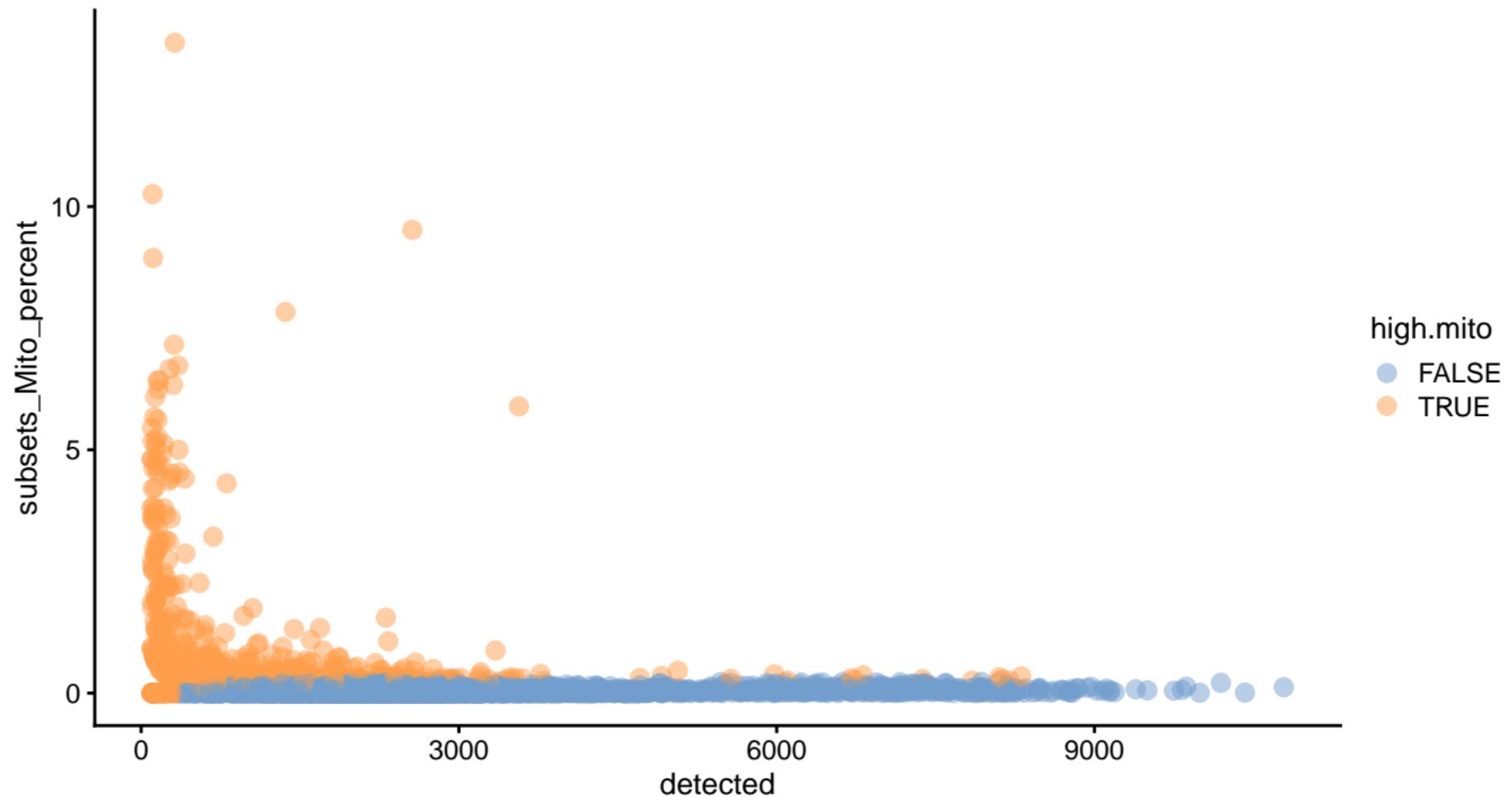
Detected features



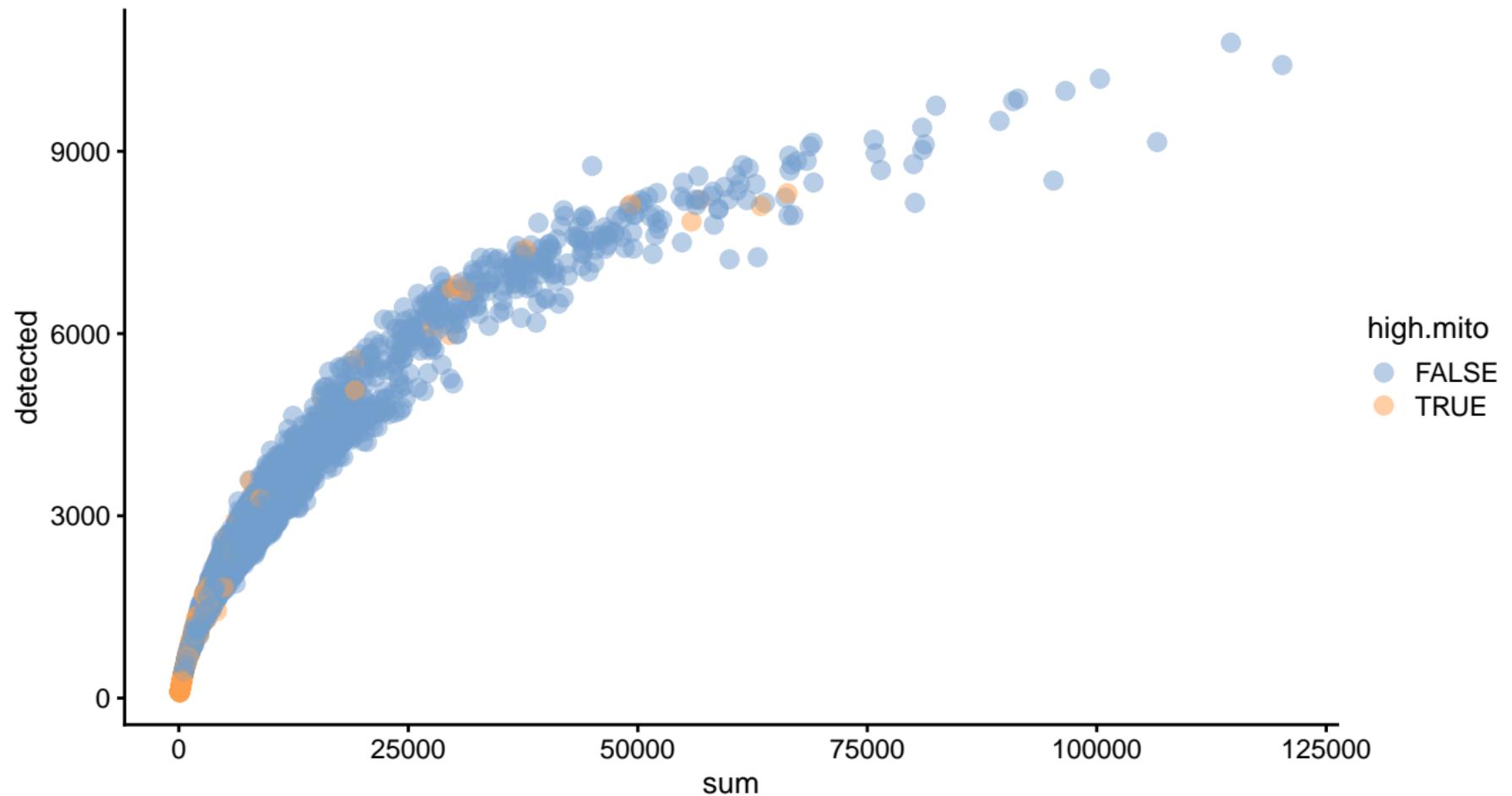
Mito % (cutoff = 0.236)



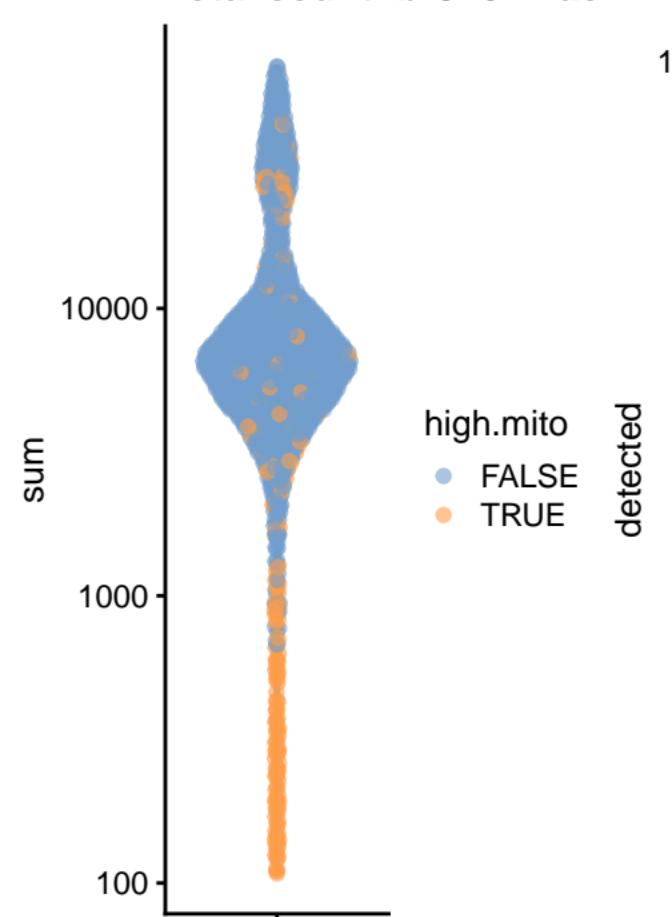
Sample: br5161.hpc; pre-QC nNuclei: 5042; nNuclei kept: 4421 (0.88%)



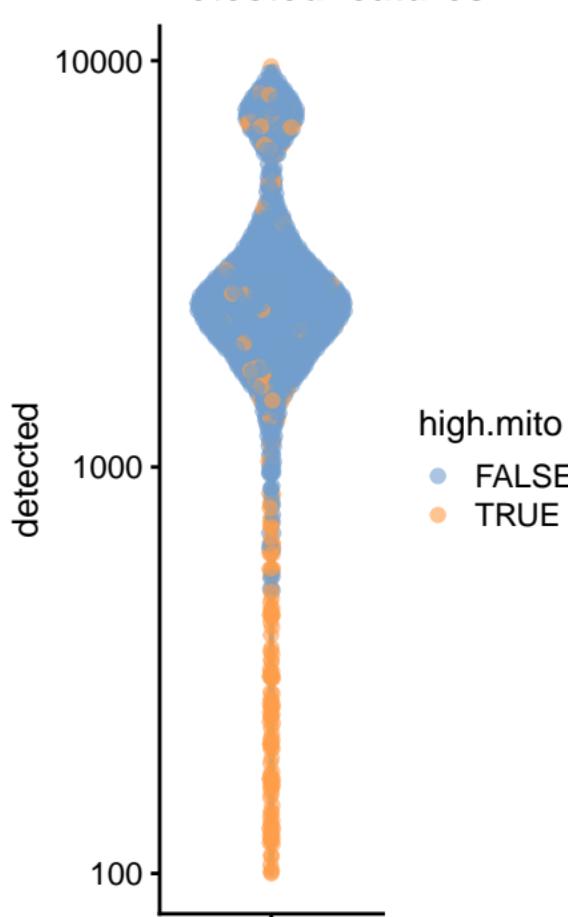
Sample: br5161.hpc; pre-QC nNuclei: 5042; nNuclei kept: 4421 (0.88%)



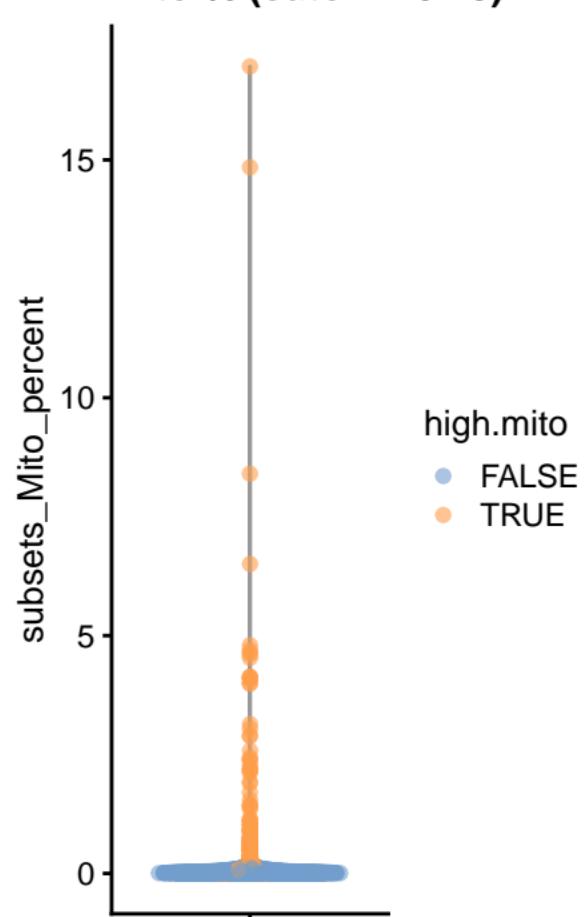
Total count: br5161.nac



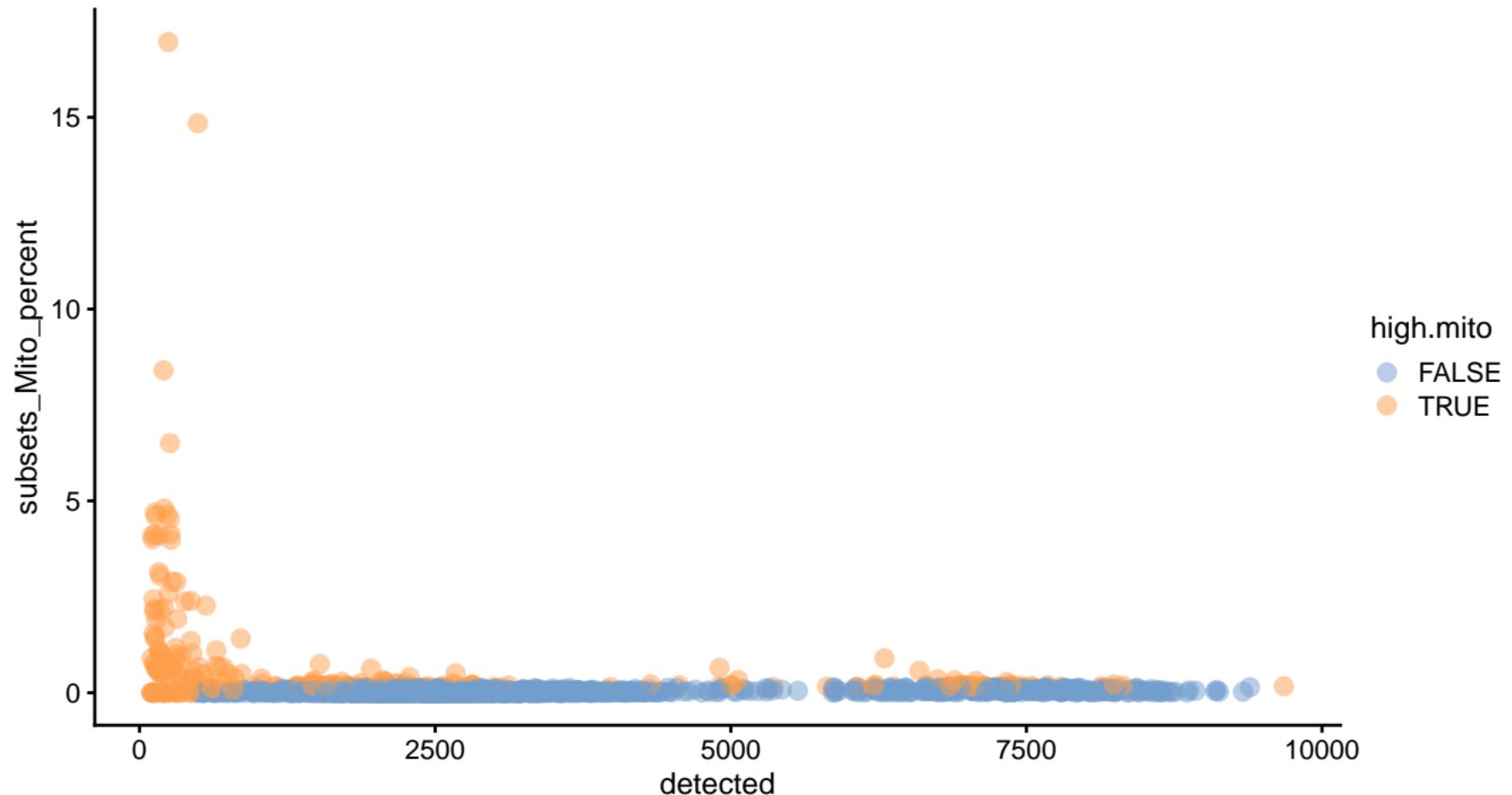
Detected features



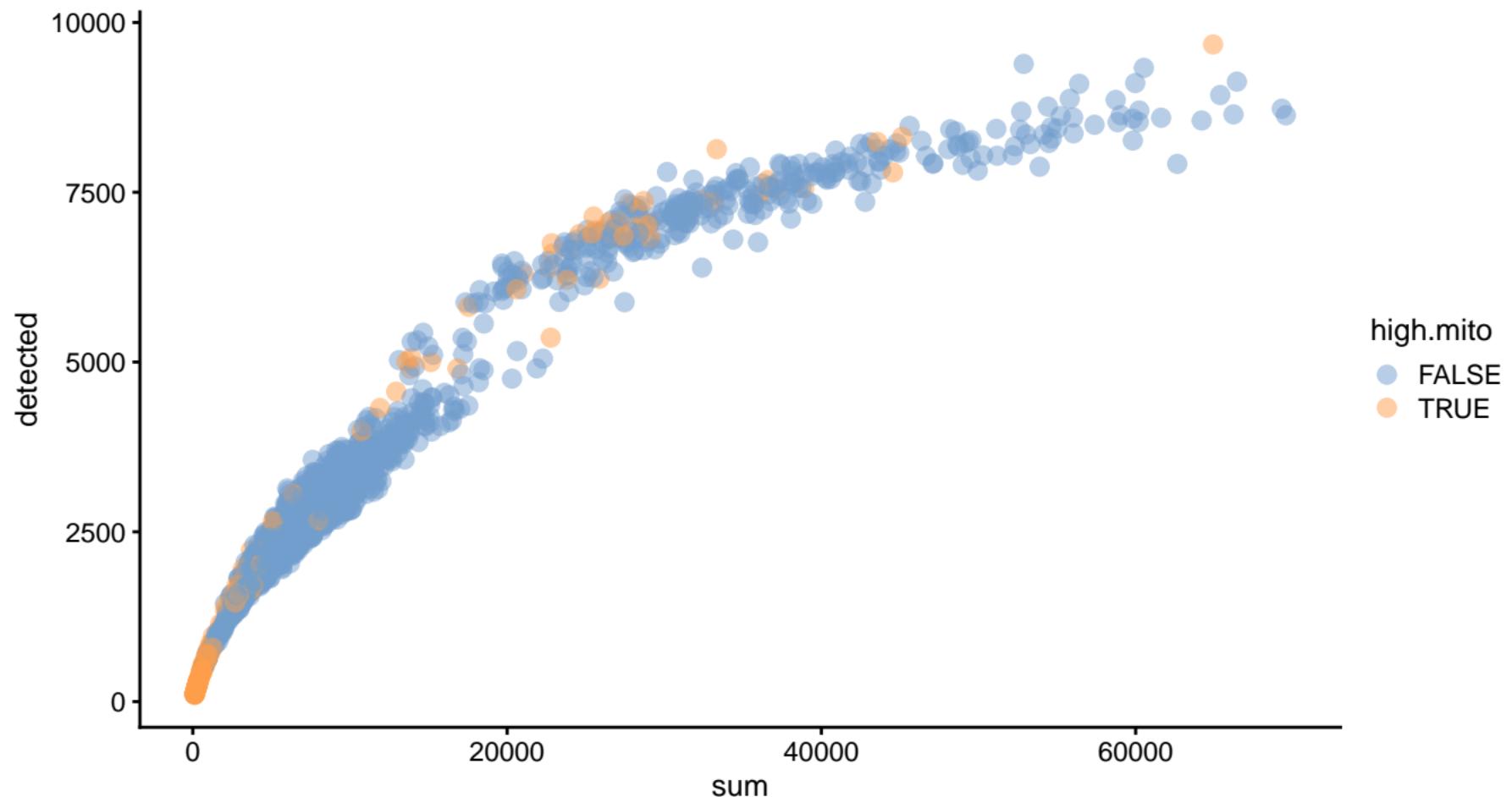
Mito % (cutoff = 0.15)

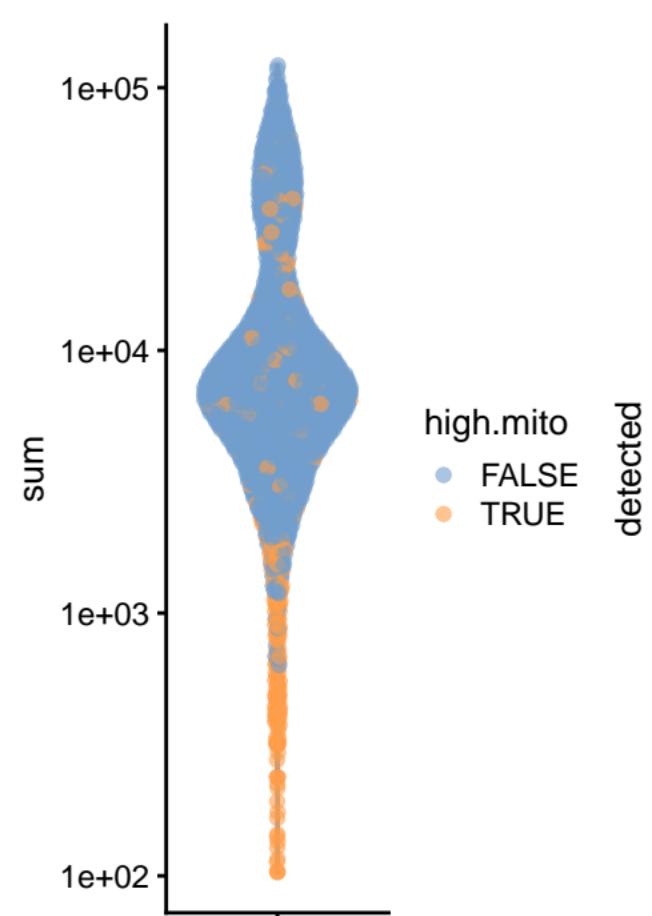
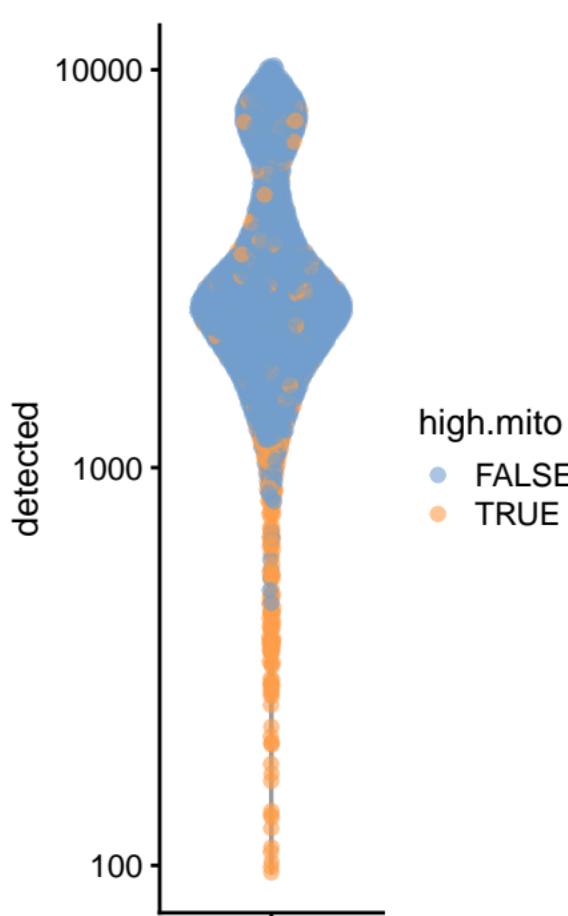
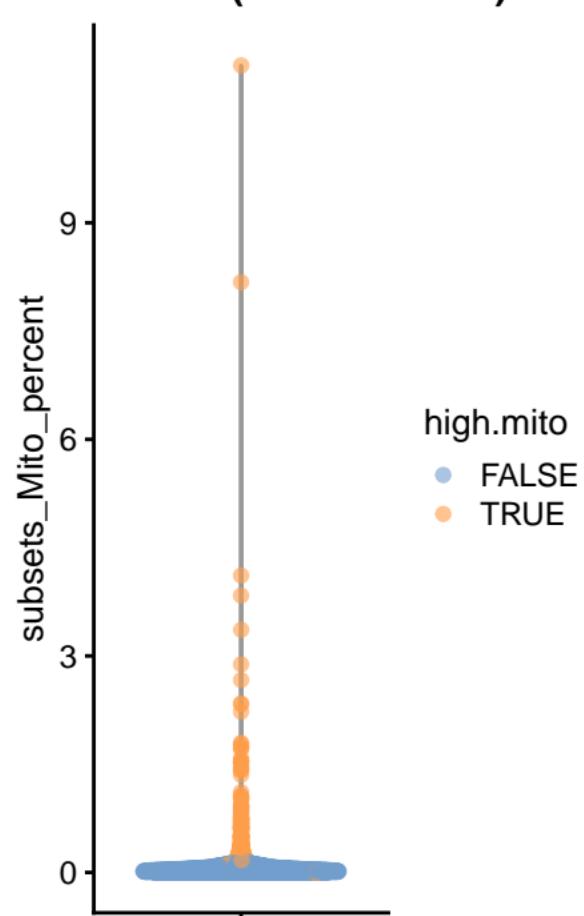


Sample: br5161.nac; pre-QC nNuclei: 2294; nNuclei kept: 2055 (0.9%)

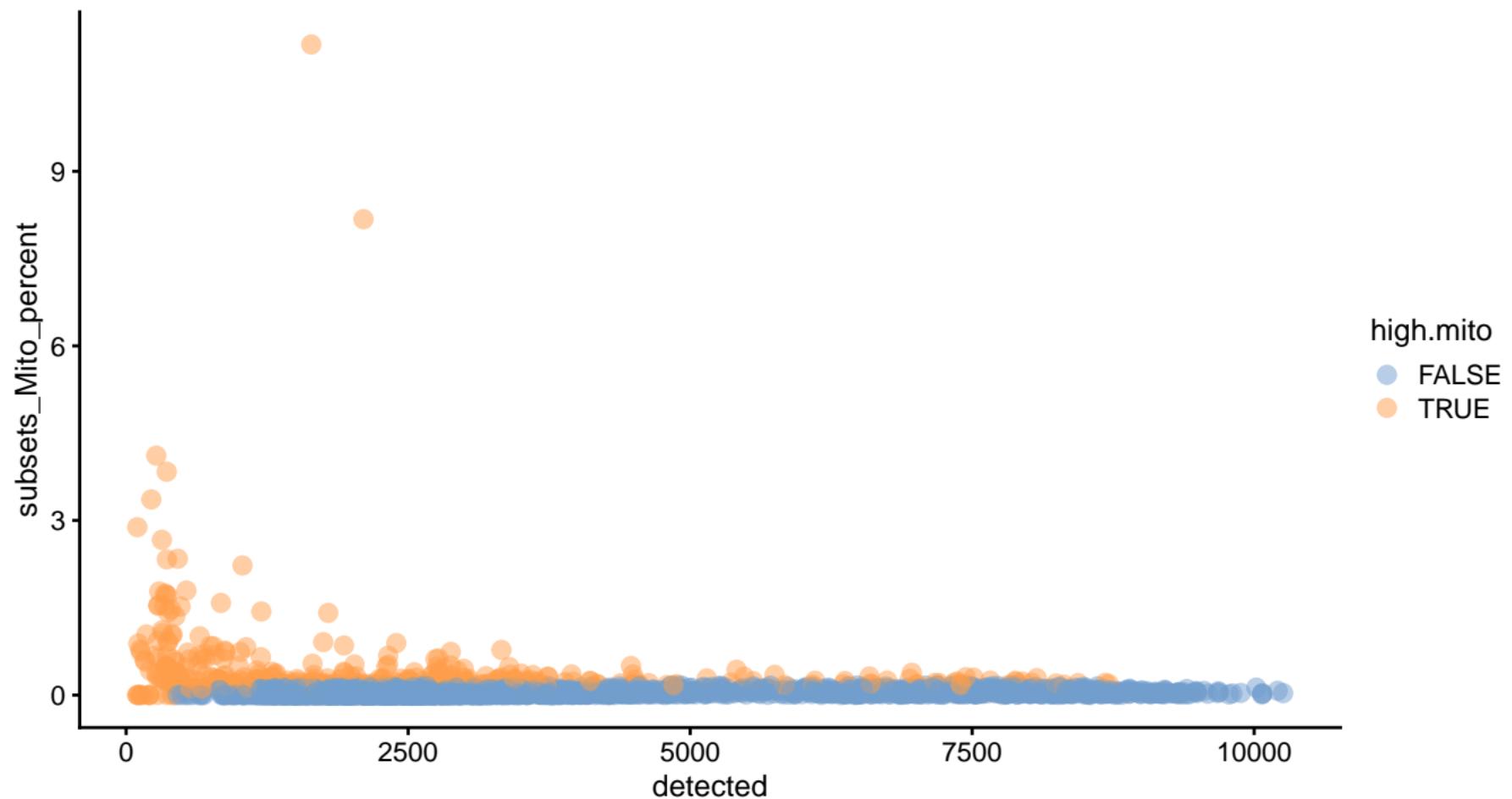


Sample: br5161.nac; pre-QC nNuclei: 2294; nNuclei kept: 2055 (0.9%)

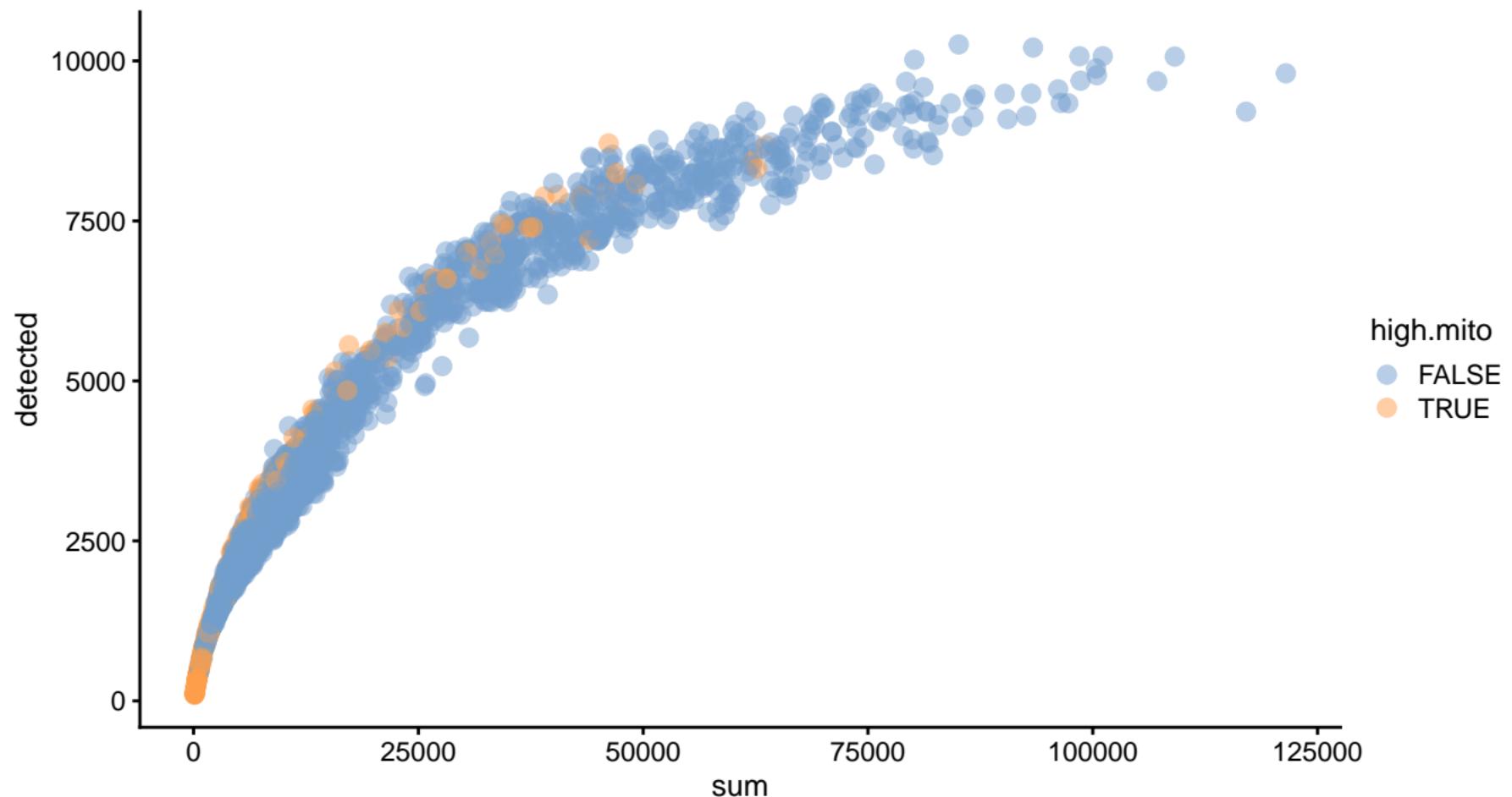


Total count: br5161.sacc**Detected features****Mito % (cutoff = 0.169)**

Sample: br5161.sacc; pre-QC nNuclei: 3547; nNuclei kept: 3174 (0.89%)



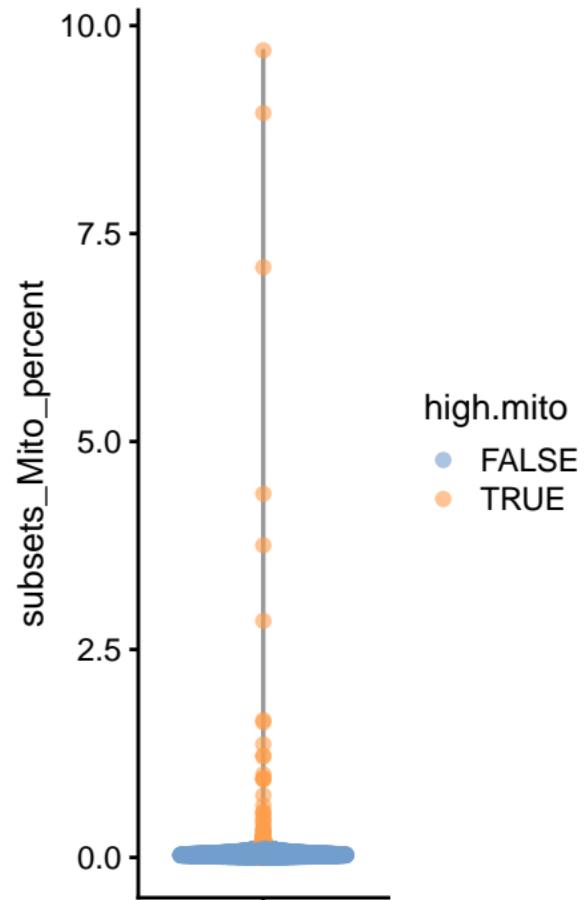
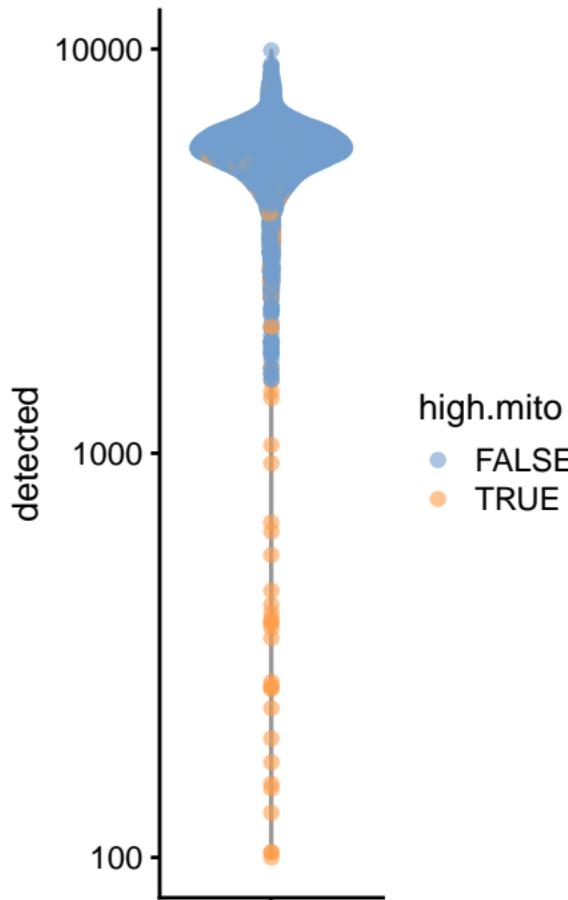
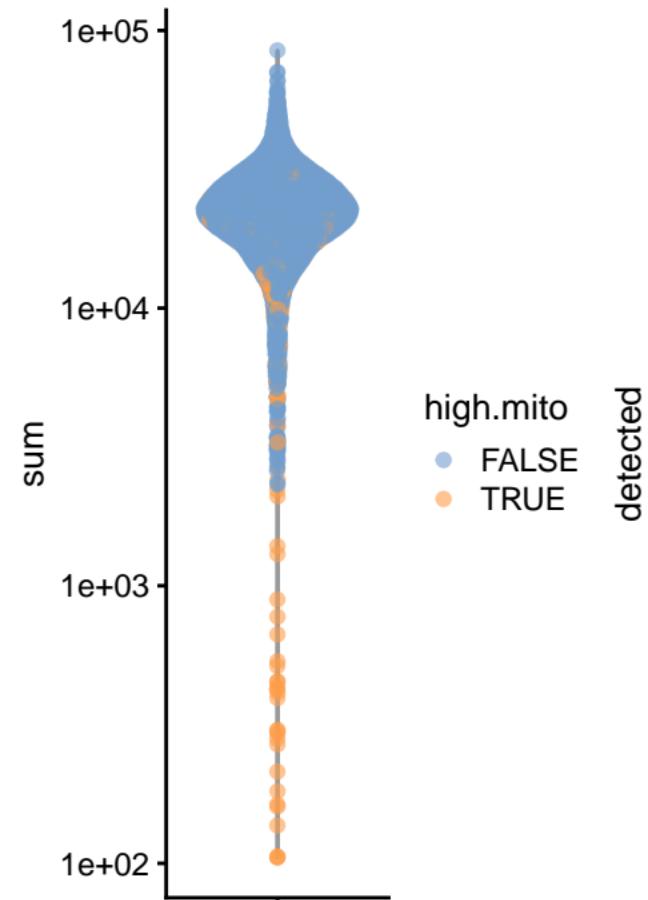
Sample: br5161.sacc; pre-QC nNuclei: 3547; nNuclei kept: 3174 (0.89%)



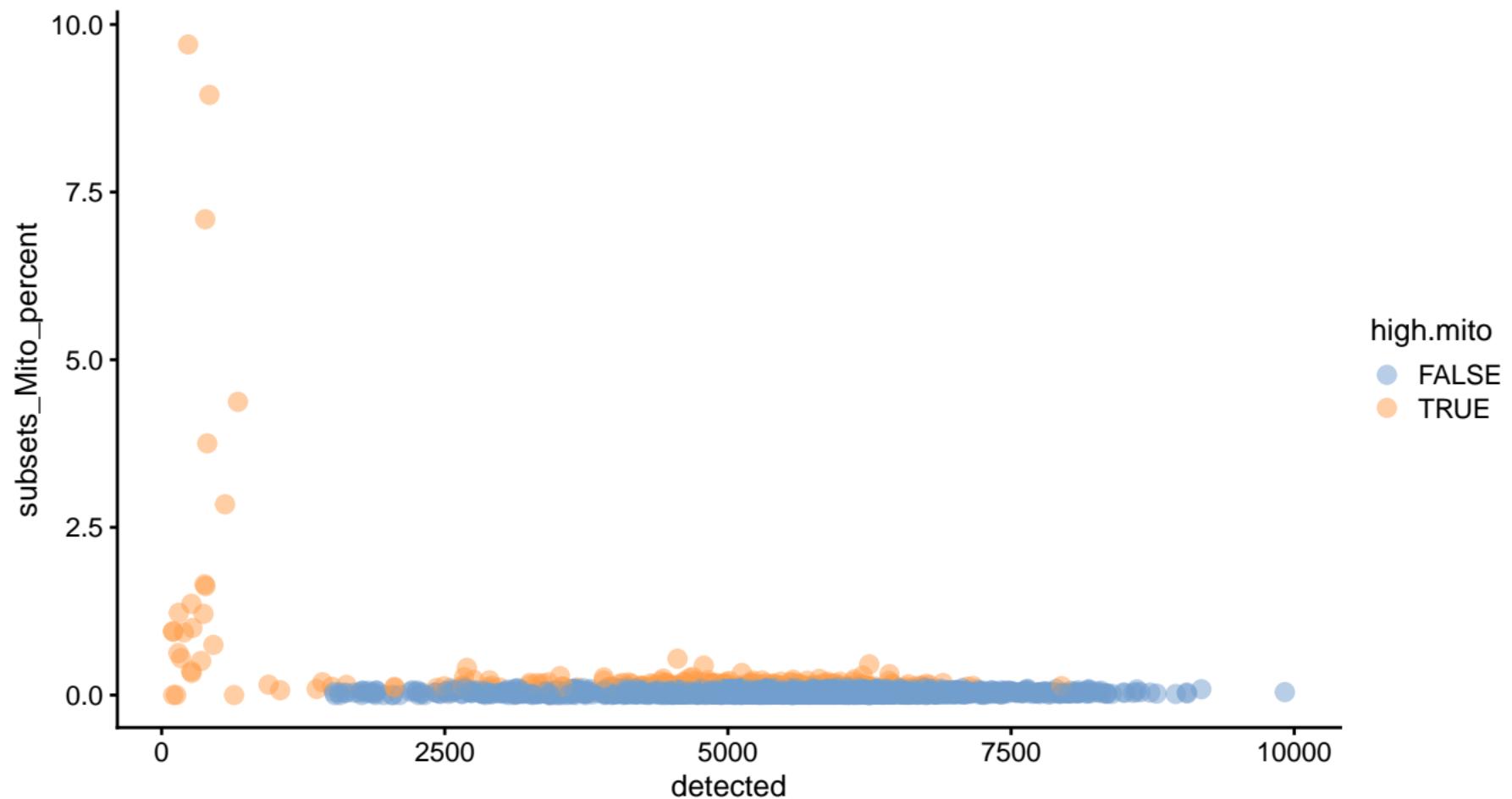
Total count: br5207.nac.neun

Detected features

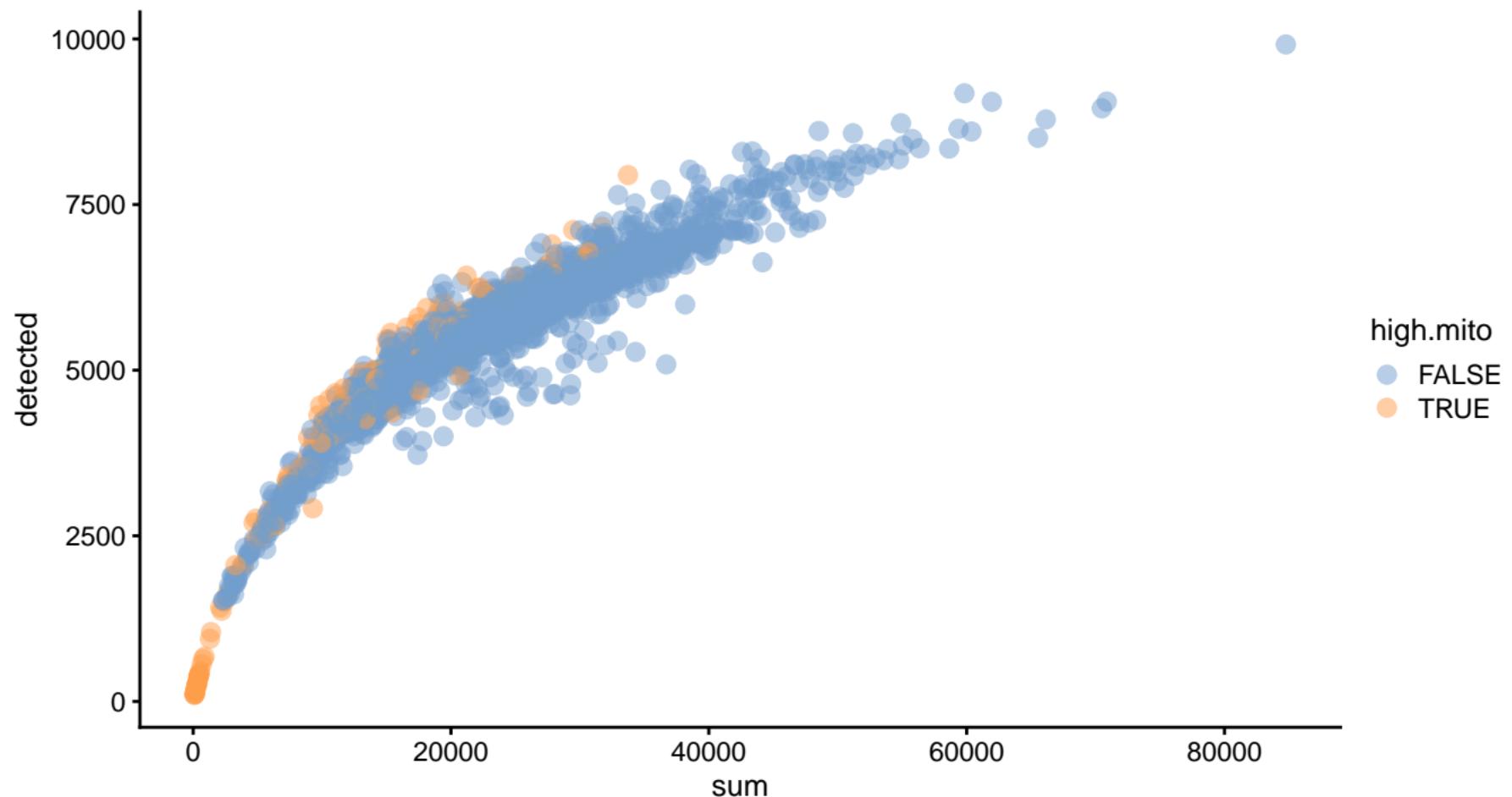
Mito % (cutoff = 0.119)



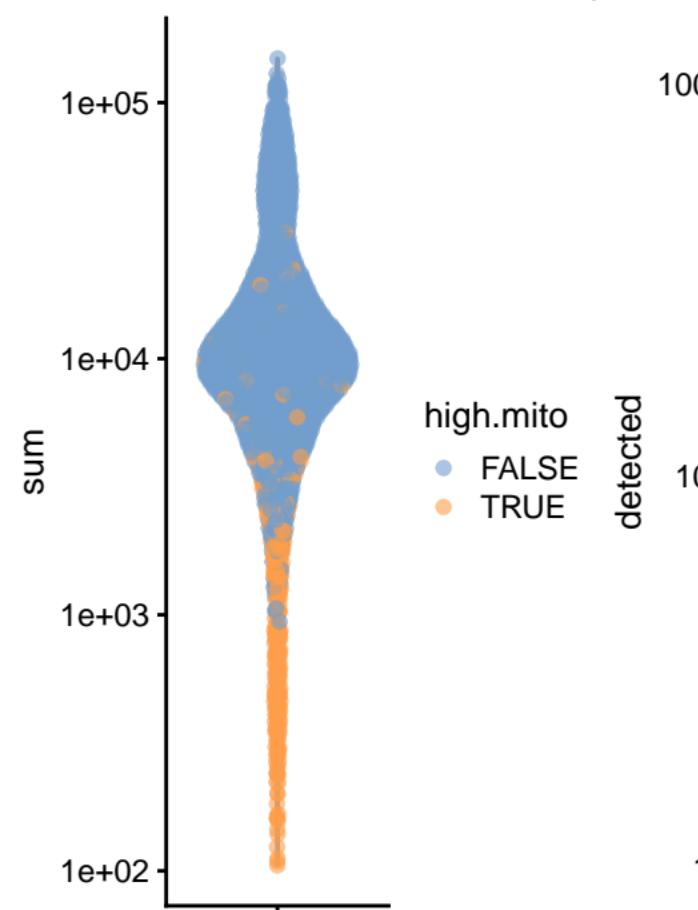
Sample: br5207.nac.neun; pre-QC nNuclei: 4682; nNuclei kept: 4425 (0.95%)



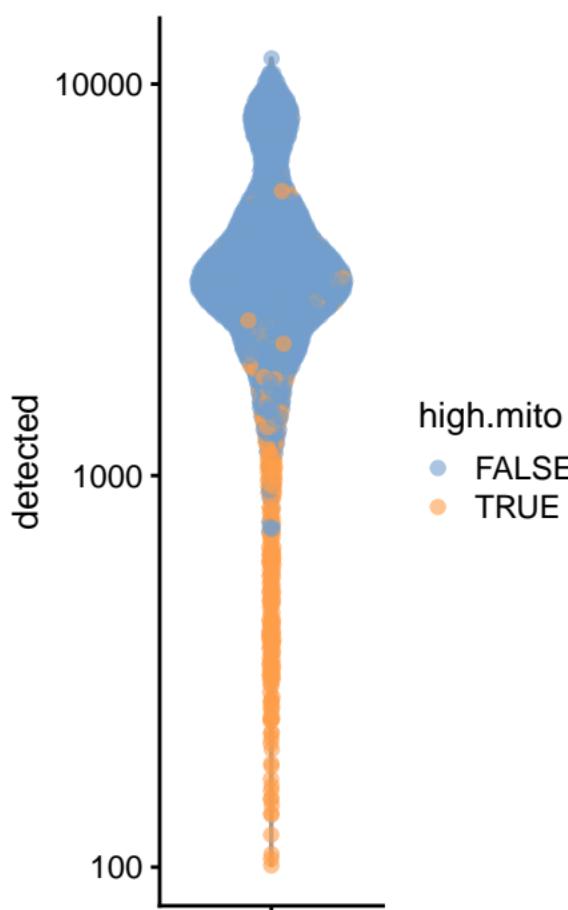
Sample: br5207.nac.neun; pre-QC nNuclei: 4682; nNuclei kept: 4425 (0.95%)



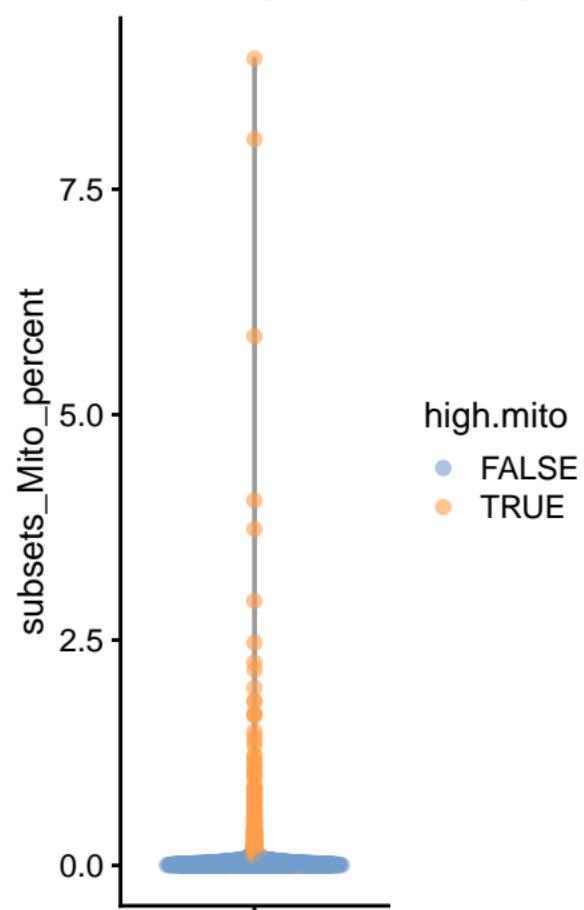
Total count: br5212.amy



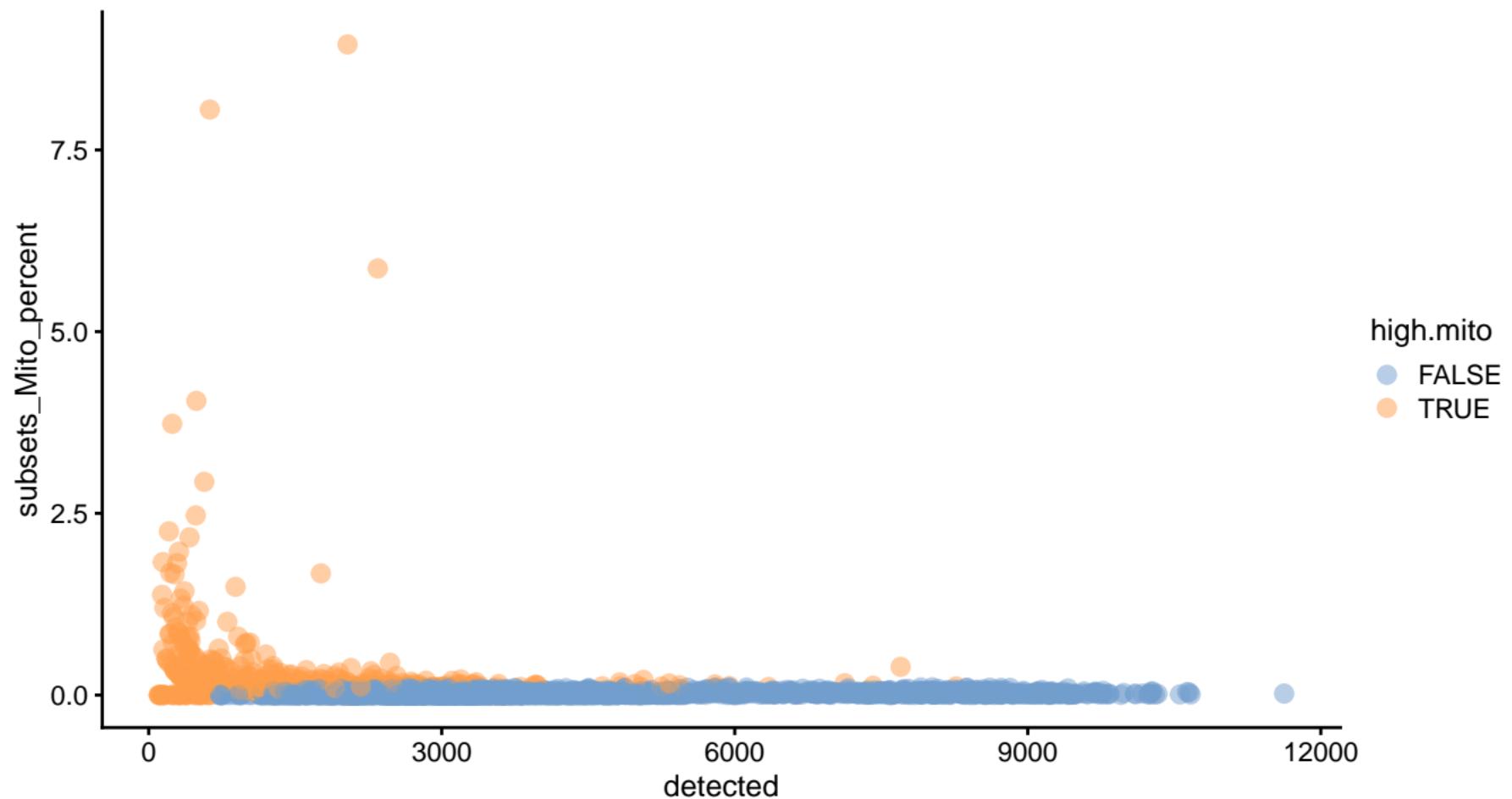
Detected features



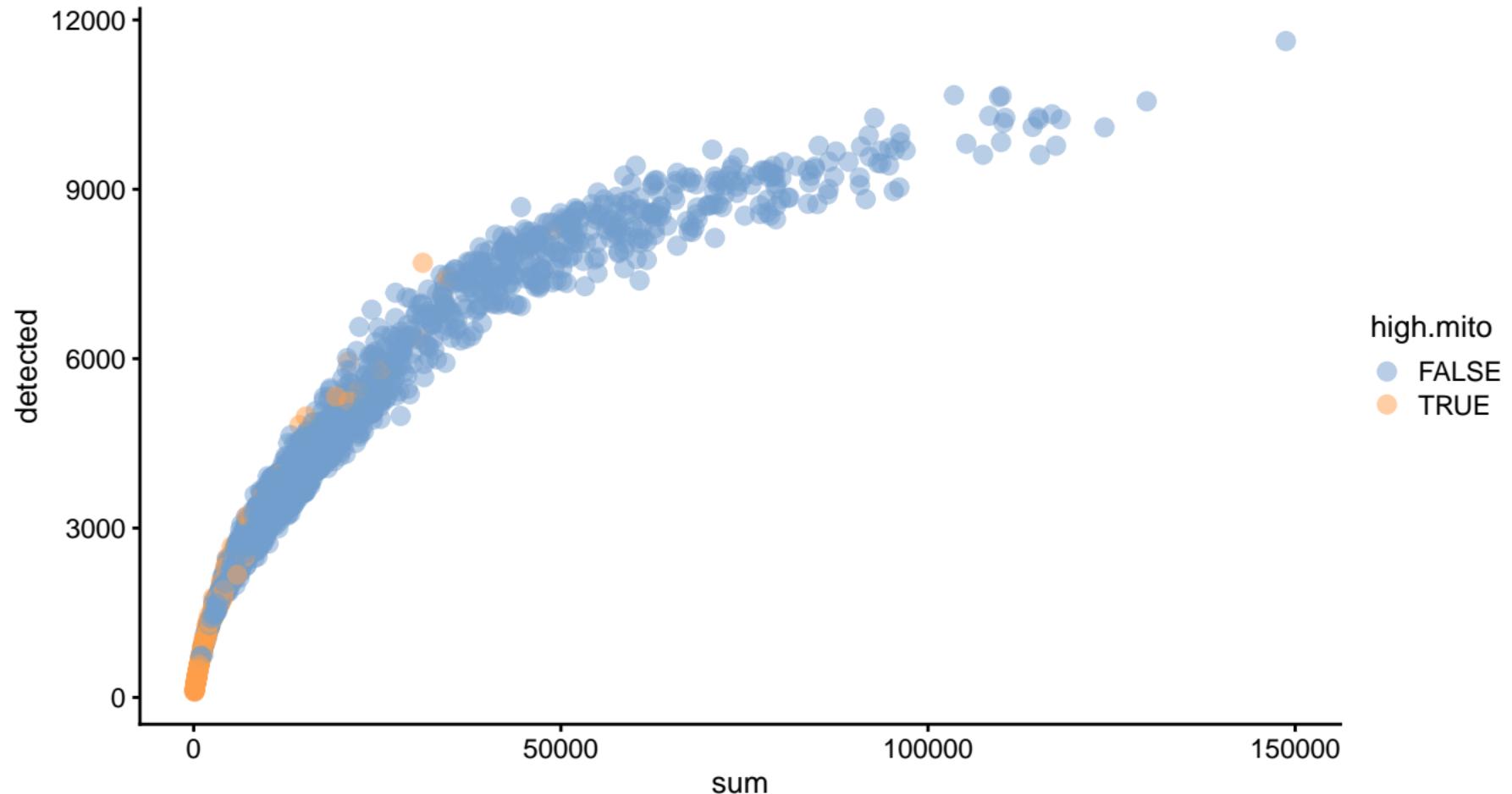
Mito % (cutoff = 0.111)



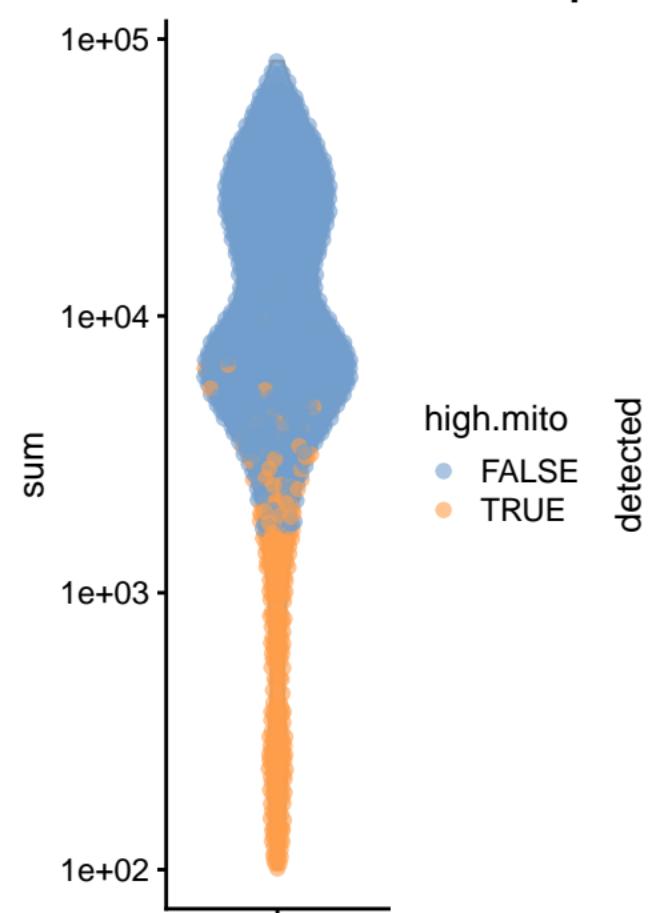
Sample: br5212.amy; pre-QC nNuclei: 3661; nNuclei kept: 3259 (0.89%)



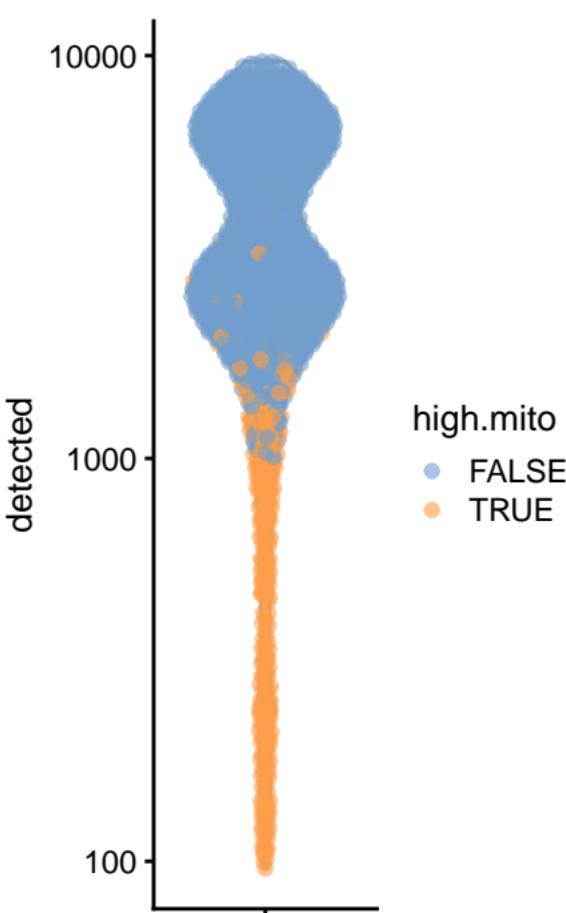
Sample: br5212.amy; pre-QC nNuclei: 3661; nNuclei kept: 3259 (0.89%)



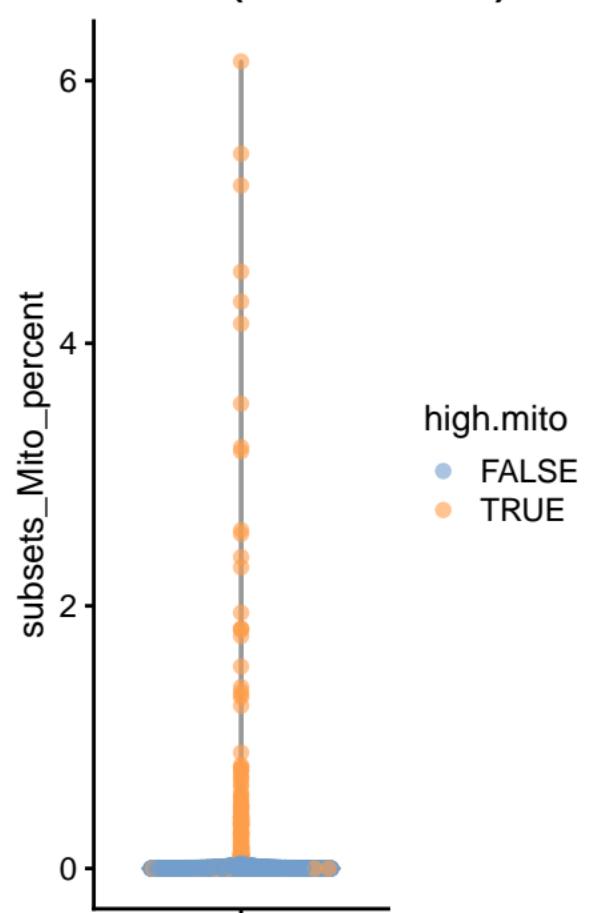
Total count: br5212.dlpfc



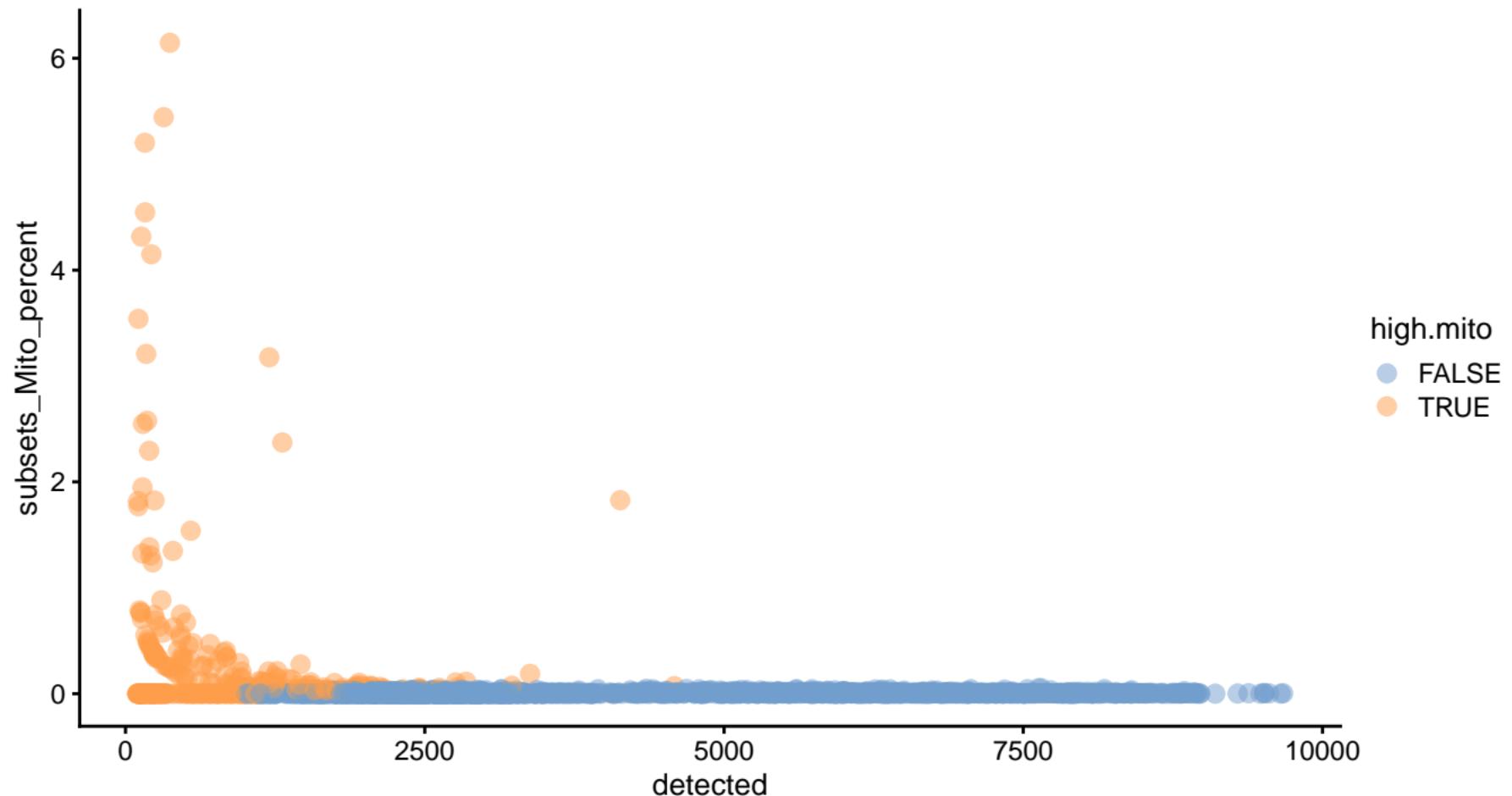
Detected features



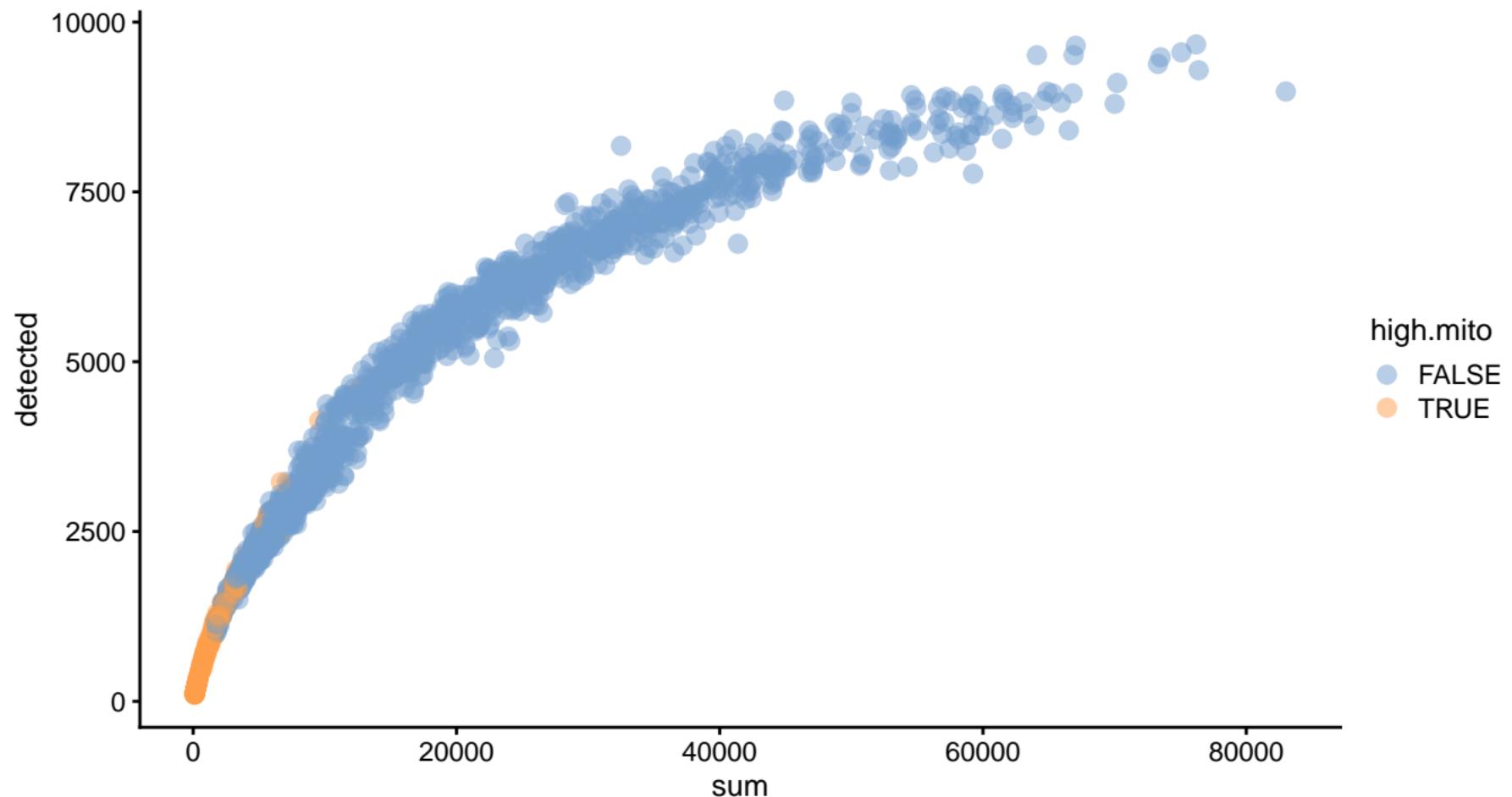
Mito % (cutoff = 0.061)



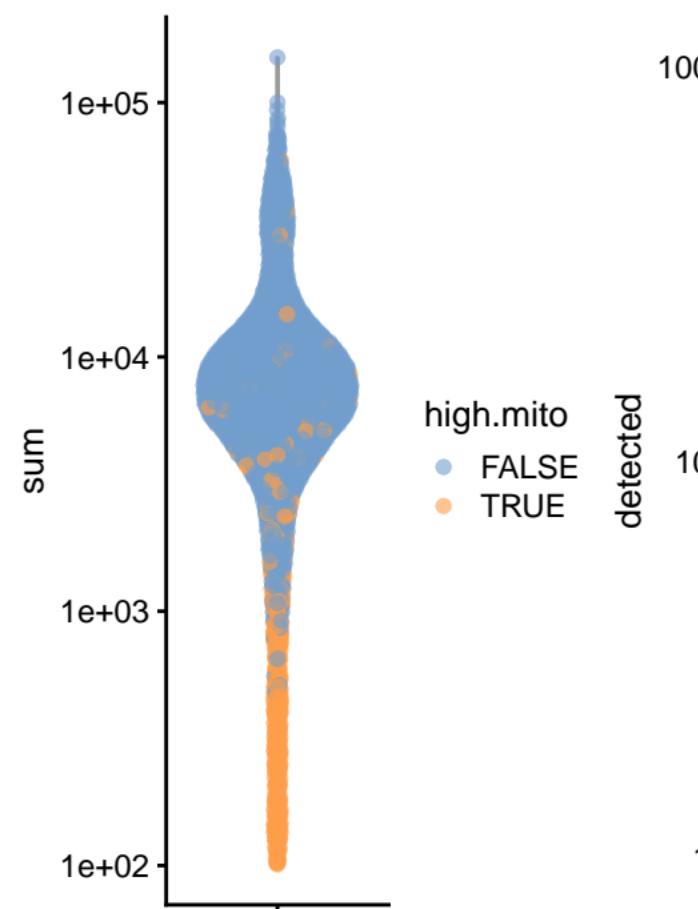
Sample: br5212.dlpfc; pre-QC nNuclei: 2006; nNuclei kept: 1693 (0.84%)



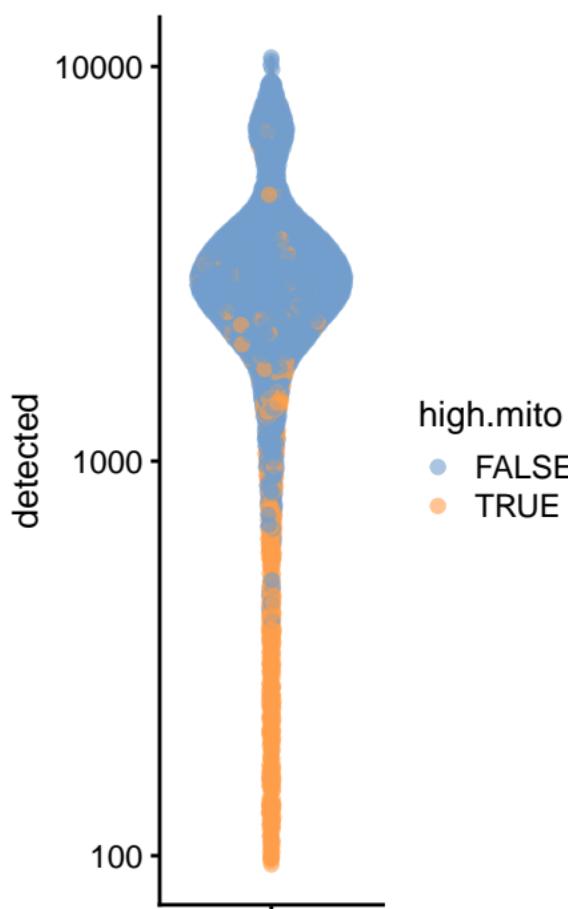
Sample: br5212.dlpfc; pre-QC nNuclei: 2006; nNuclei kept: 1693 (0.84%)



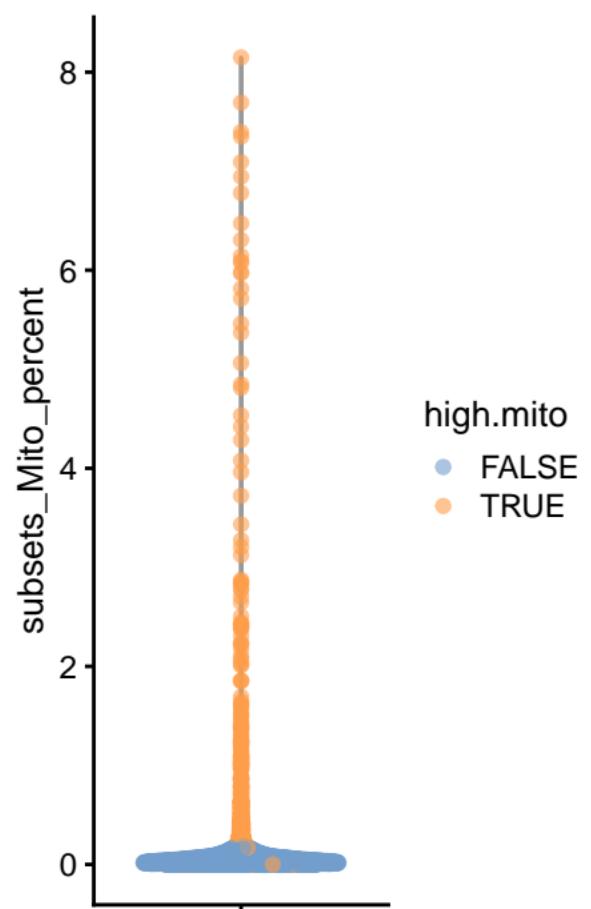
Total count: br5212.hpc



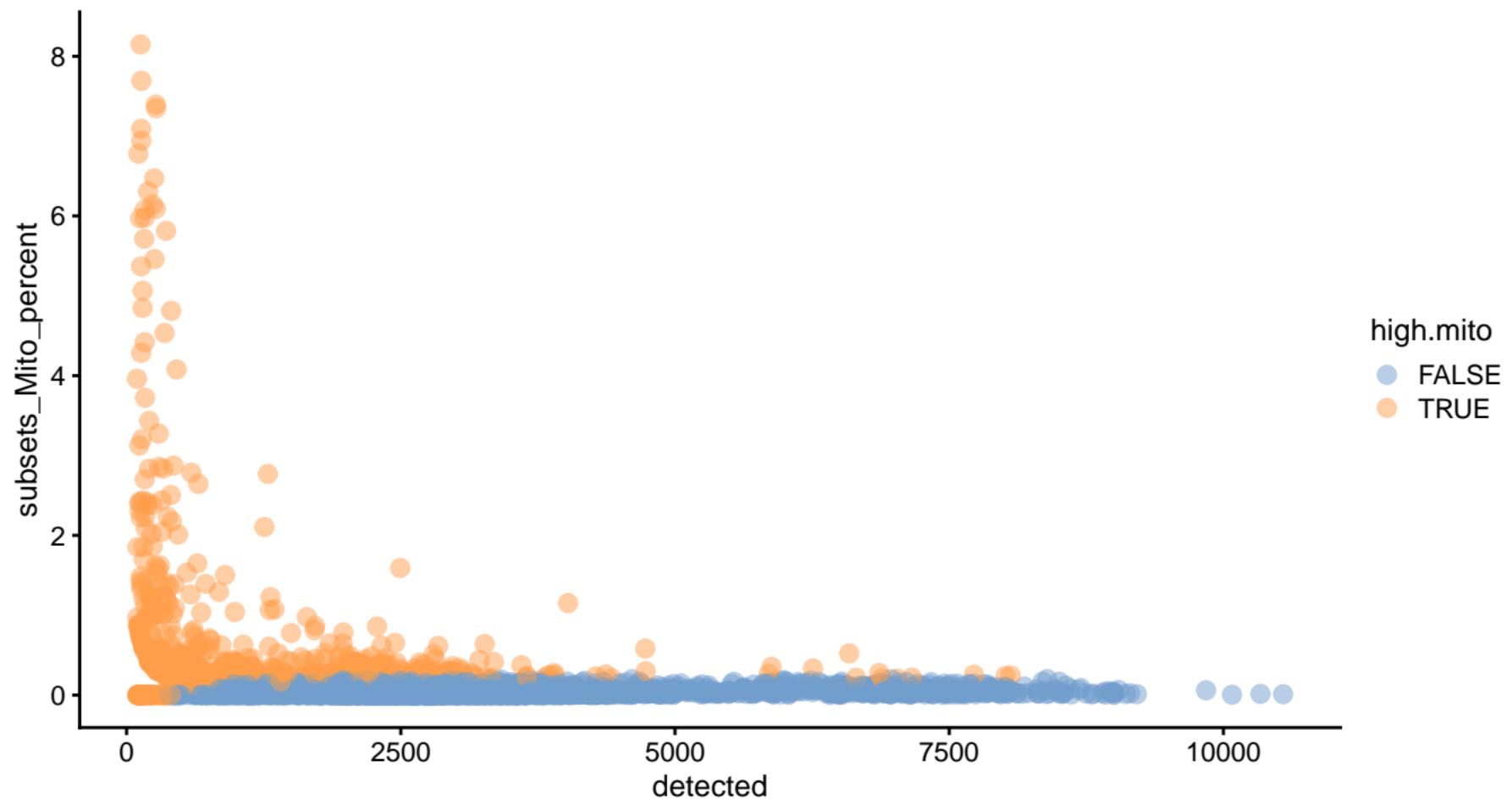
Detected features



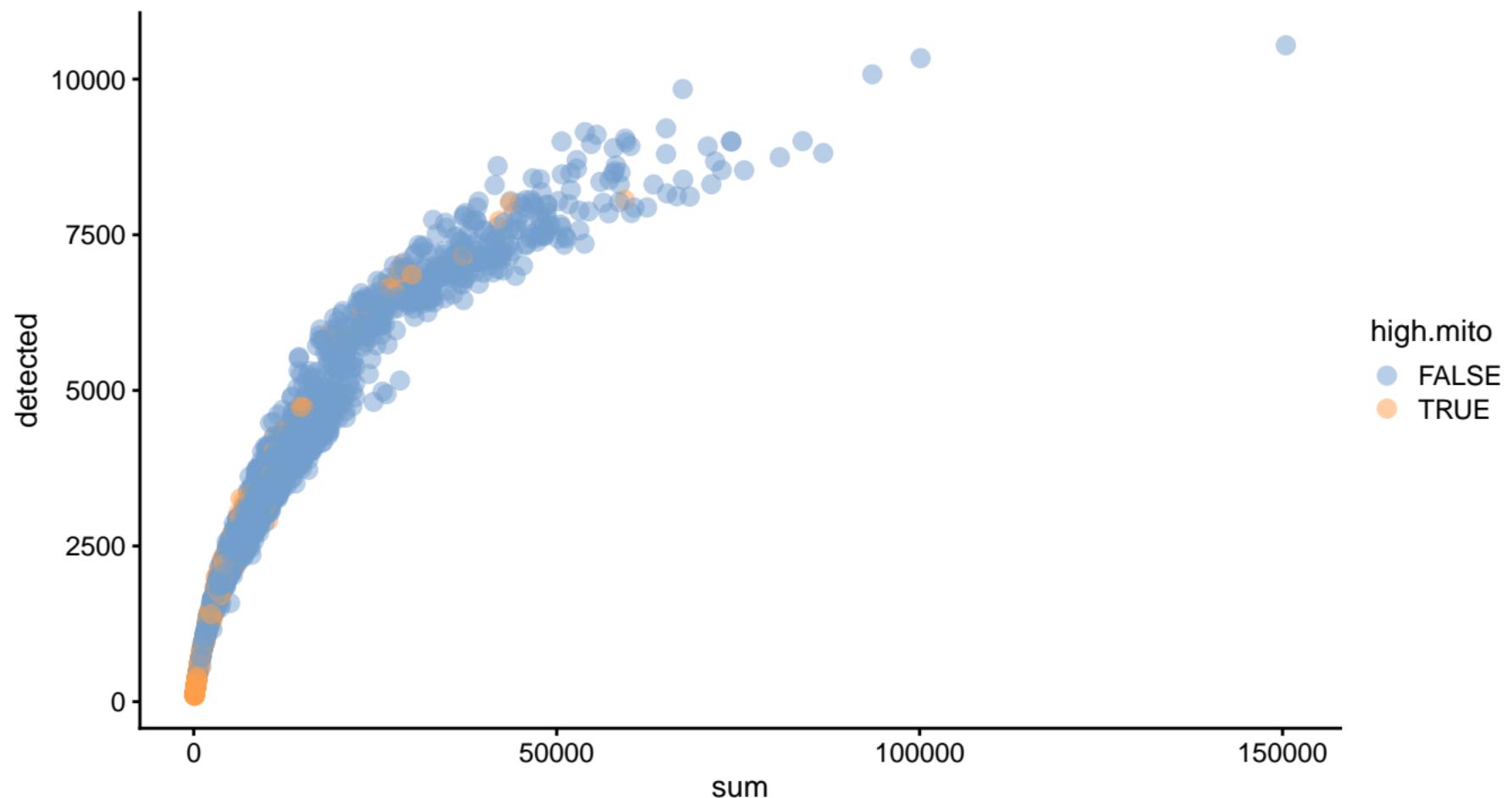
Mito % (cutoff = 0.207)



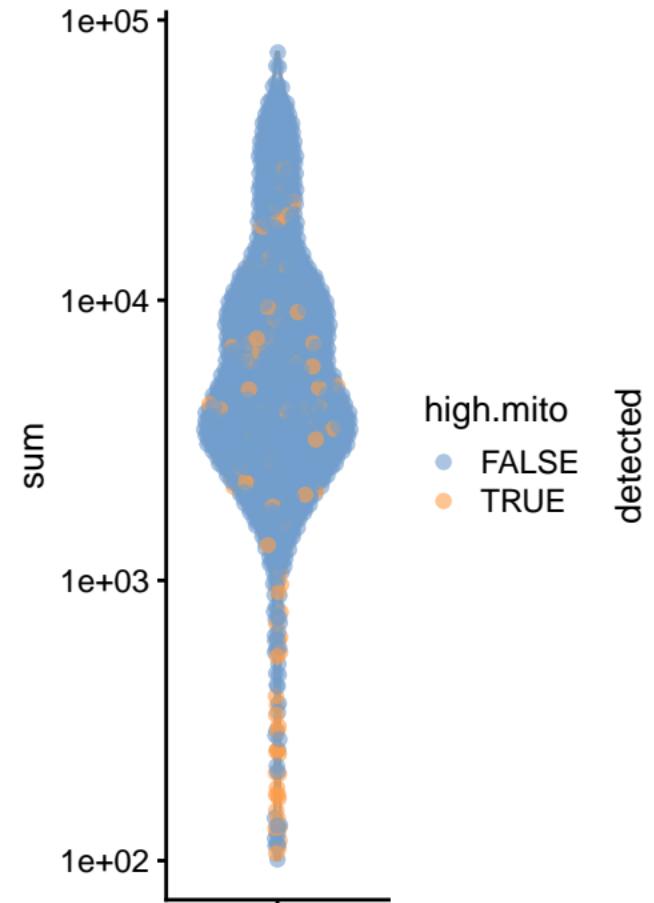
Sample: br5212.hpc; pre-QC nNuclei: 4538; nNuclei kept: 3977 (0.88%)



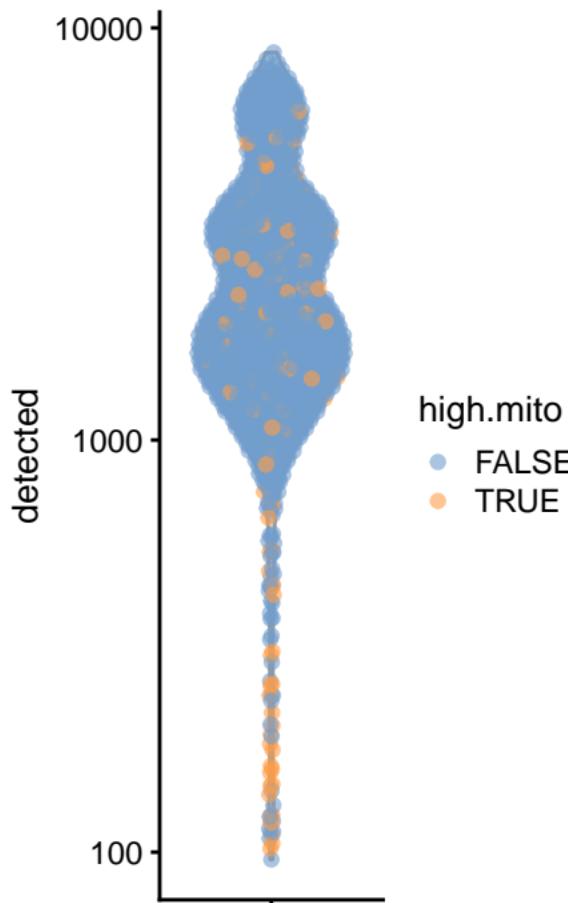
Sample: br5212.hpc; pre-QC nNuclei: 4538; nNuclei kept: 3977 (0.88%)



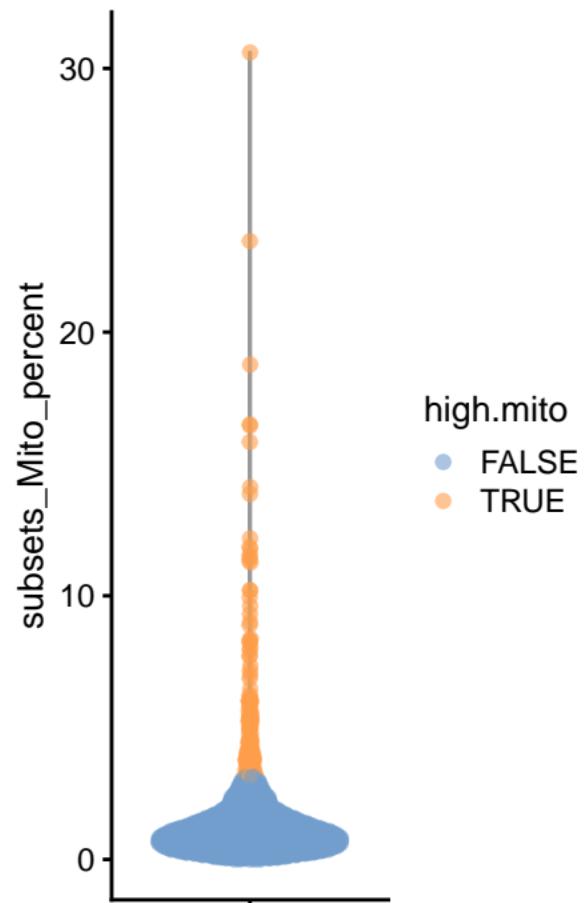
Total count: br5212.nac



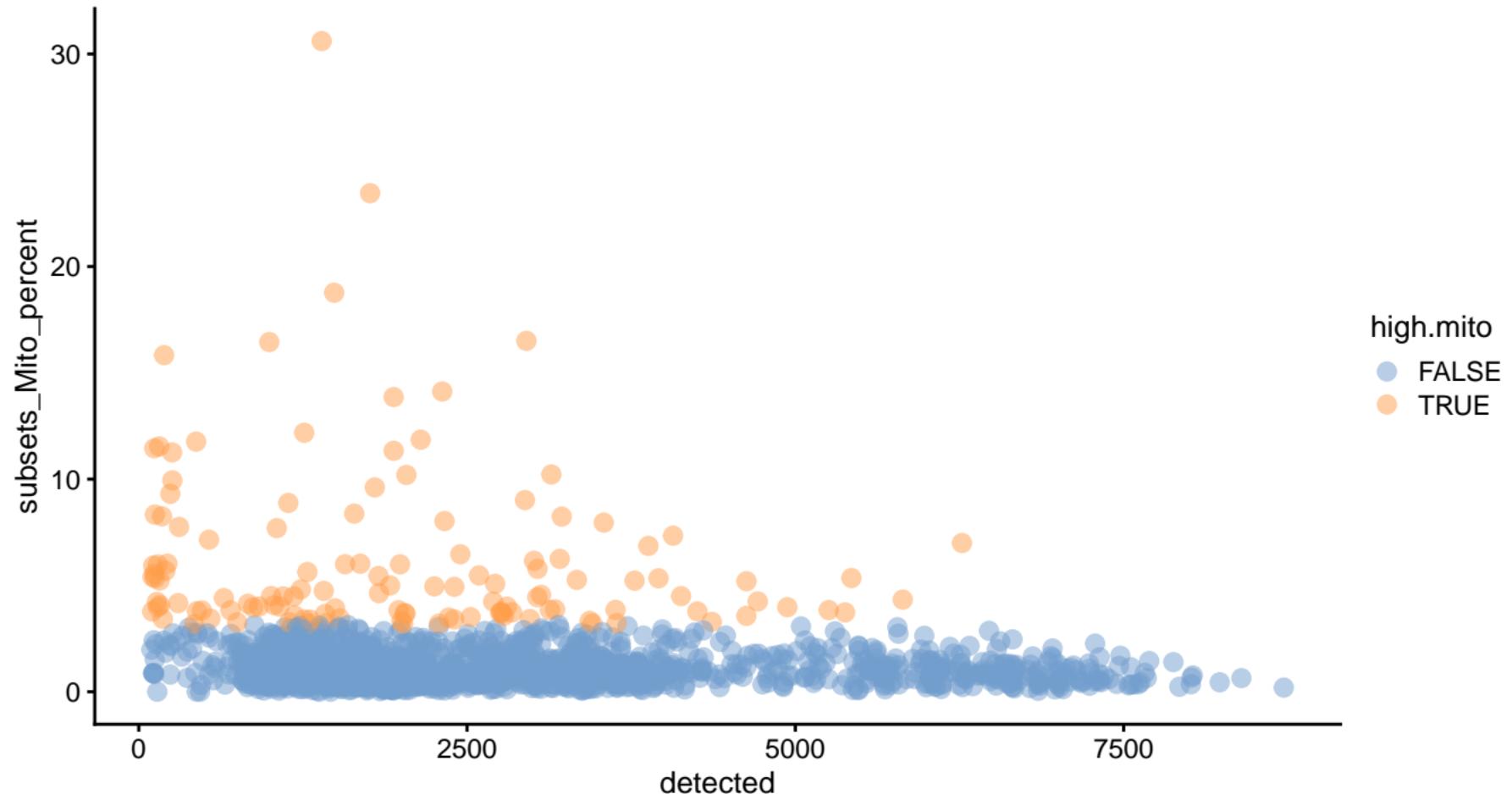
Detected features



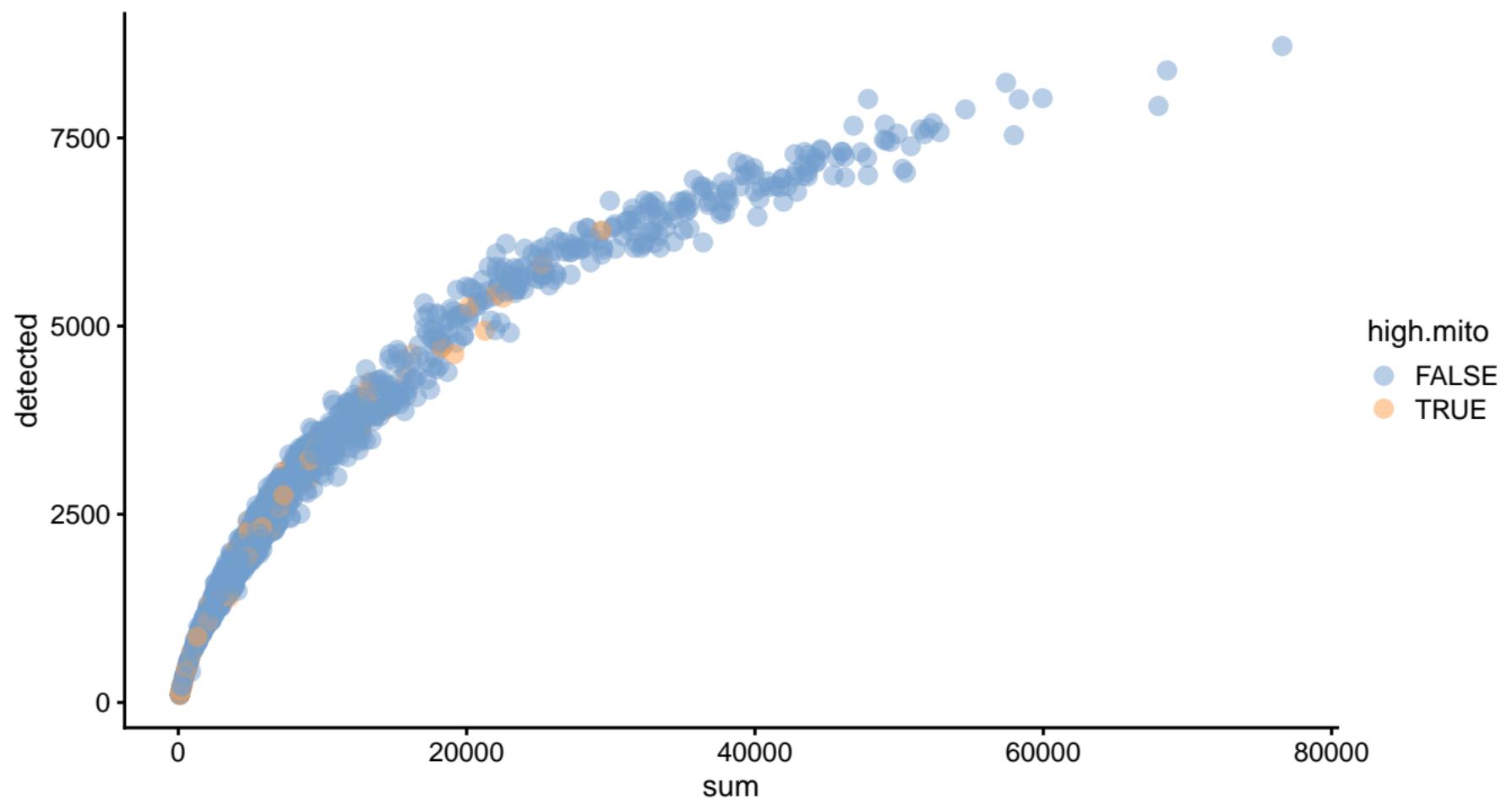
Mito % (cutoff = 3.216)

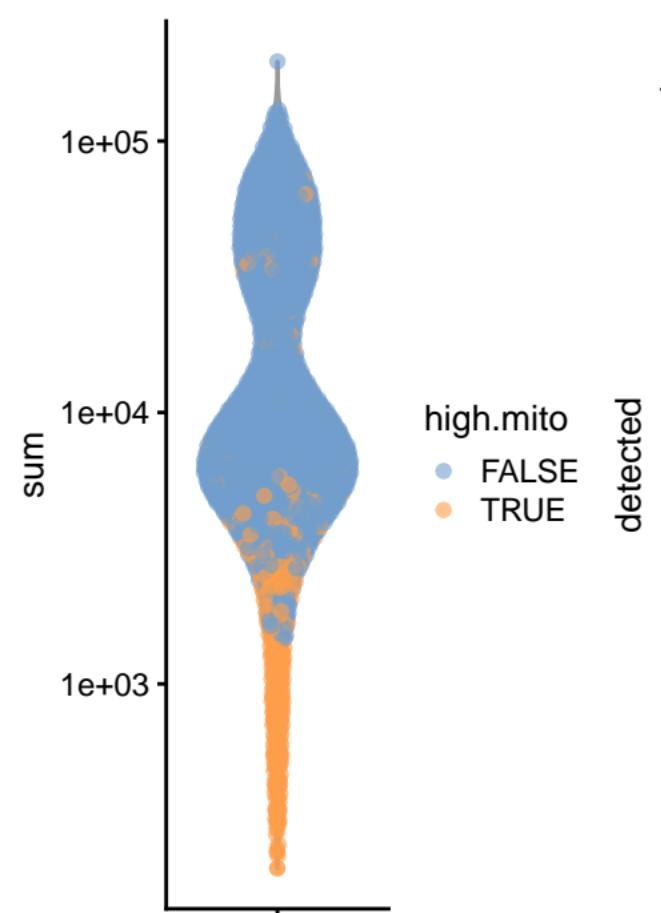
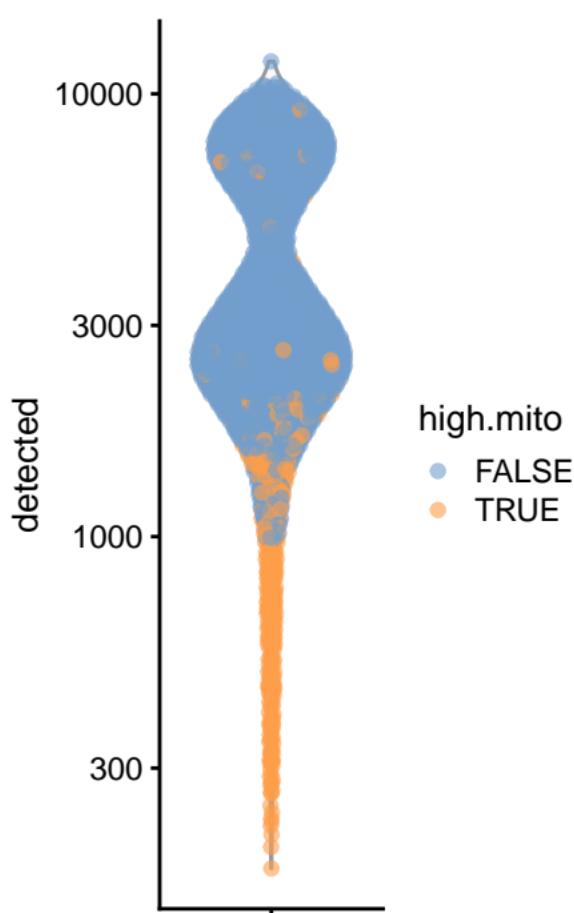
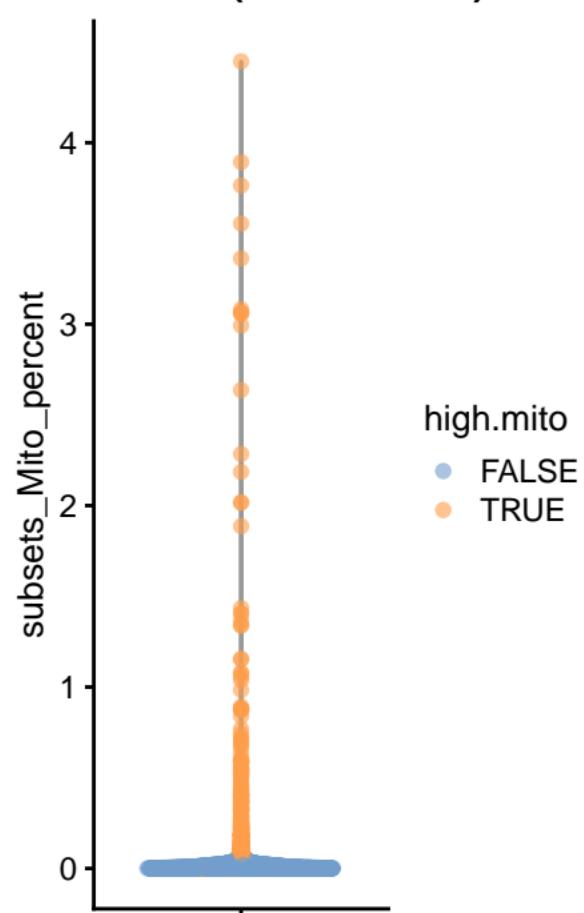


Sample: br5212.nac; pre-QC nNuclei: 1901; nNuclei kept: 1773 (0.93%)

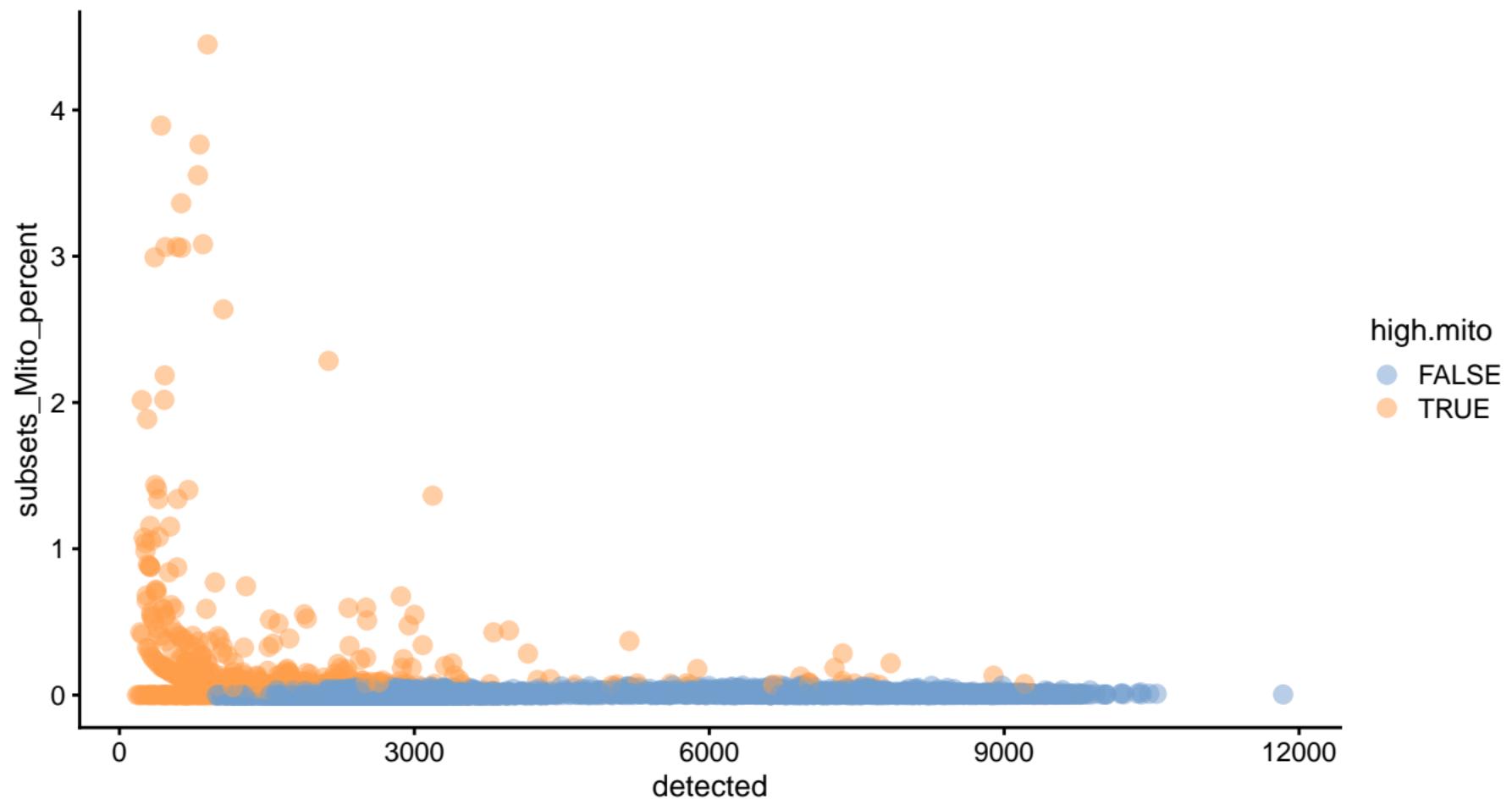


Sample: br5212.nac; pre-QC nNuclei: 1901; nNuclei kept: 1773 (0.93%)

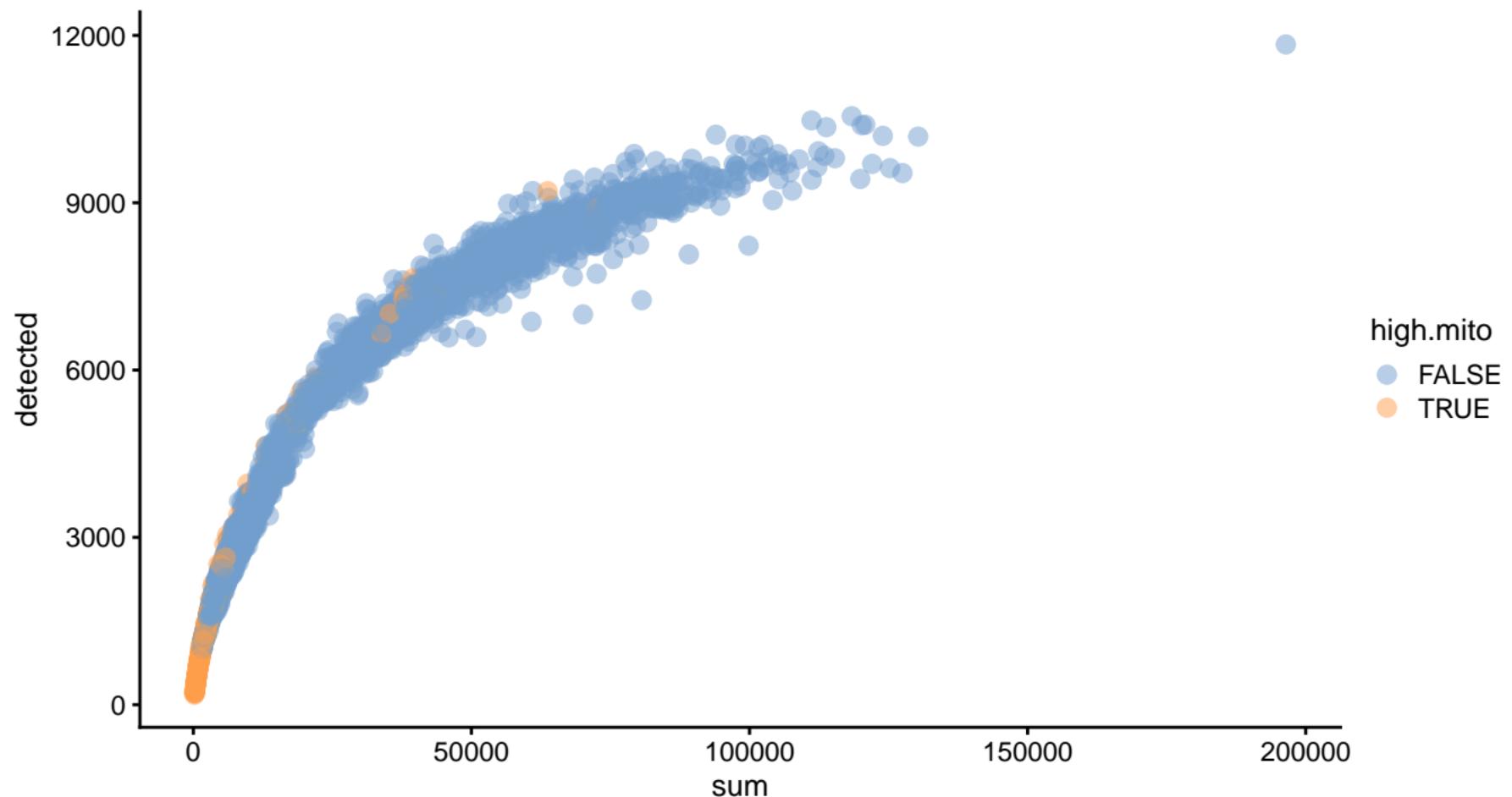


Total count: br5212.sacc**Detected features****Mito % (cutoff = 0.07)**

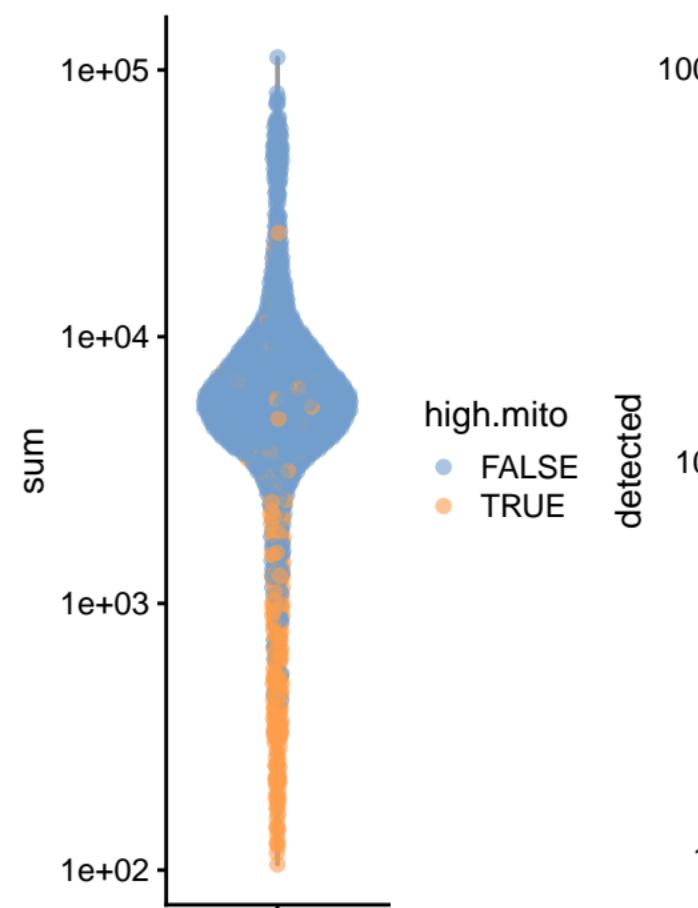
Sample: br5212.sacc; pre-QC nNuclei: 4398; nNuclei kept: 3880 (0.88%)



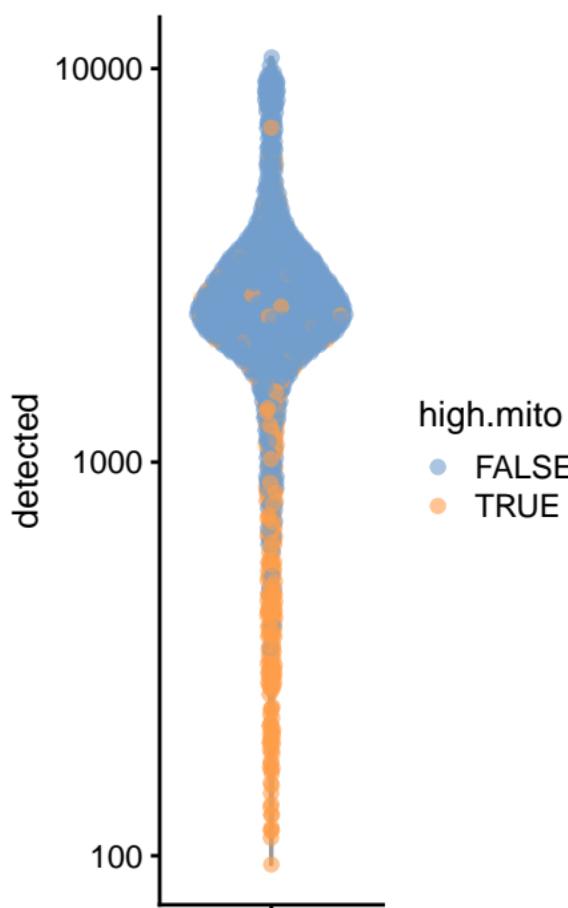
Sample: br5212.sacc; pre-QC nNuclei: 4398; nNuclei kept: 3880 (0.88%)



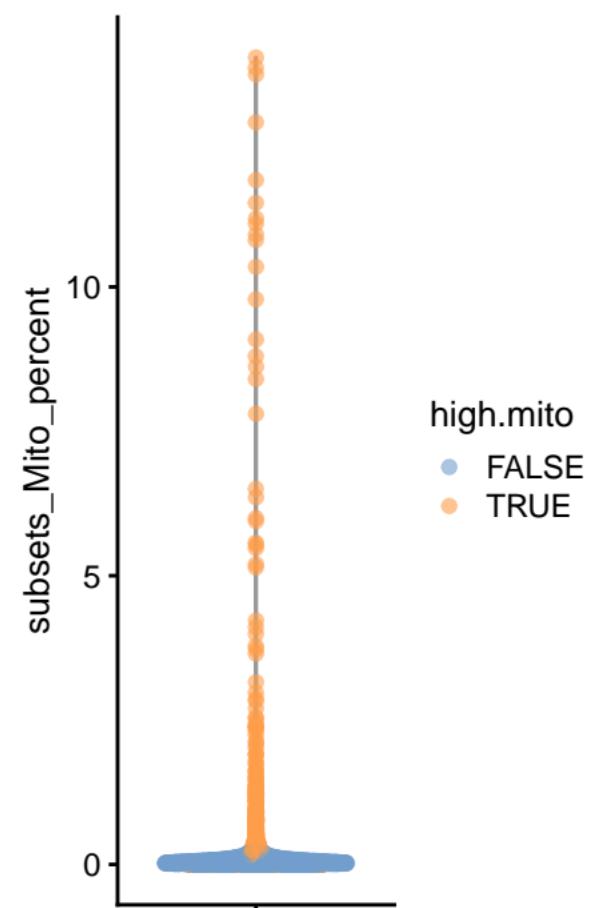
Total count: br5287.hpc



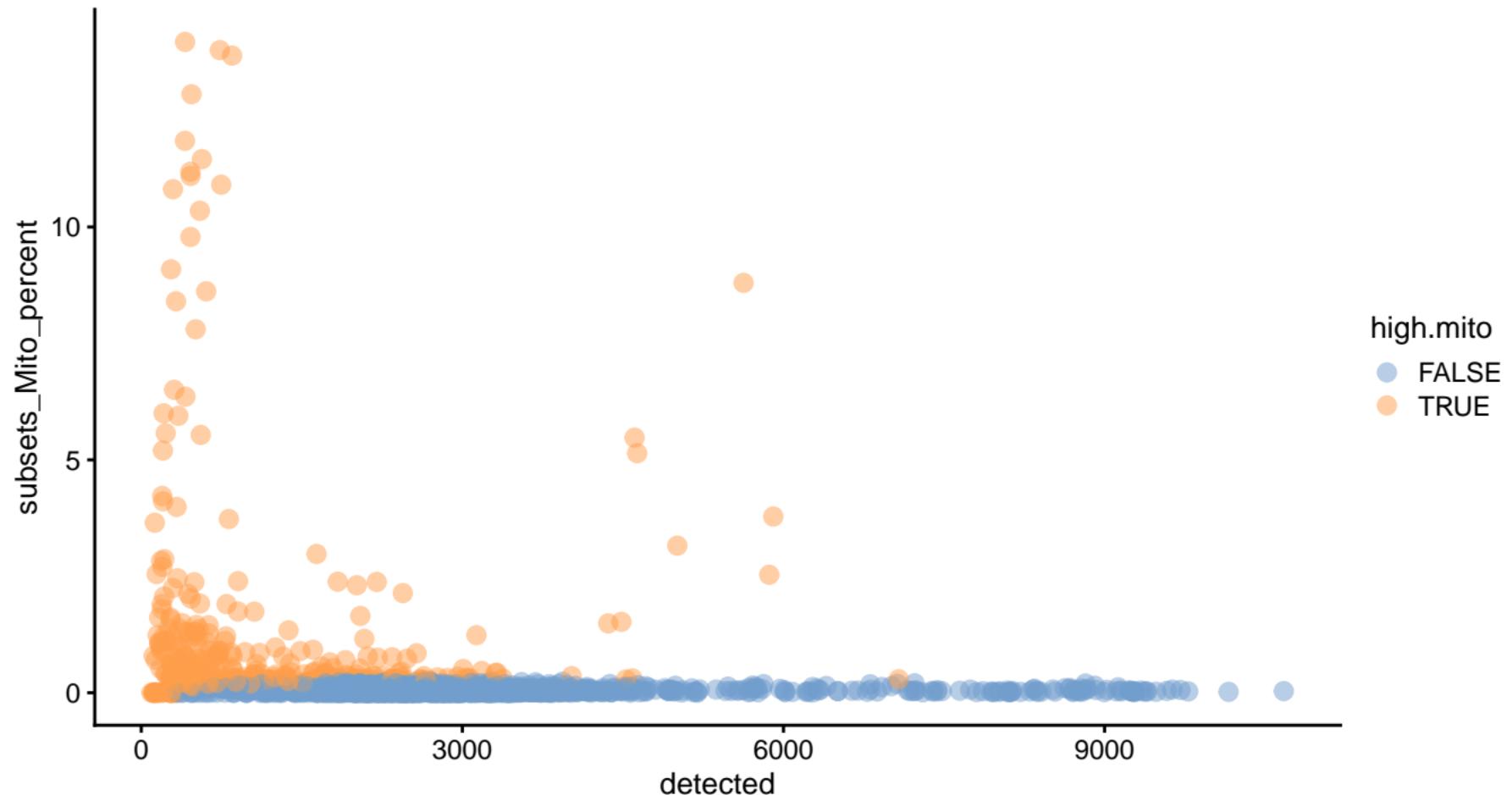
Detected features



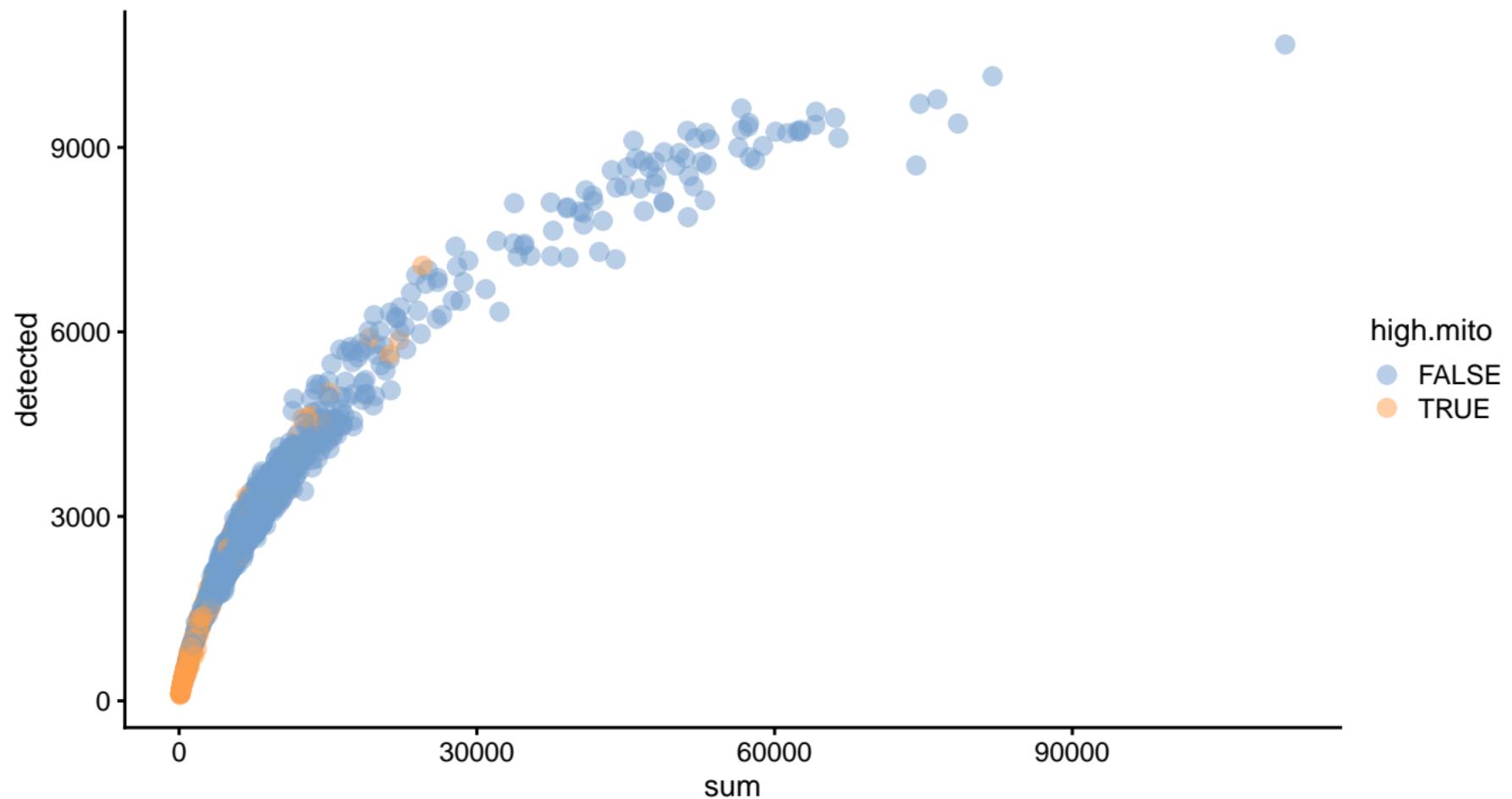
Mito % (cutoff = 0.253)



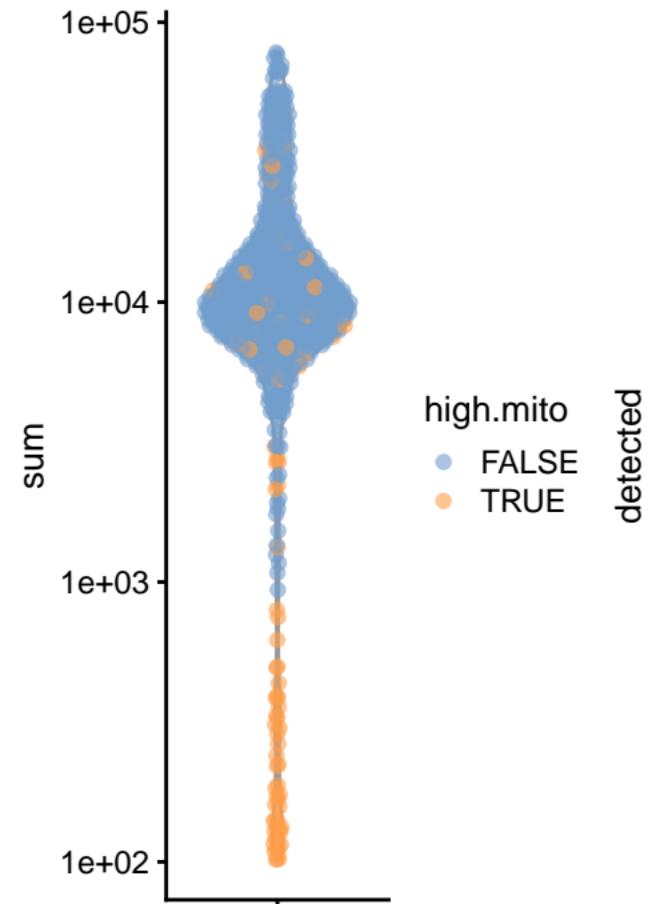
Sample: br5287.hpc; pre-QC nNuclei: 2189; nNuclei kept: 1870 (0.85%)



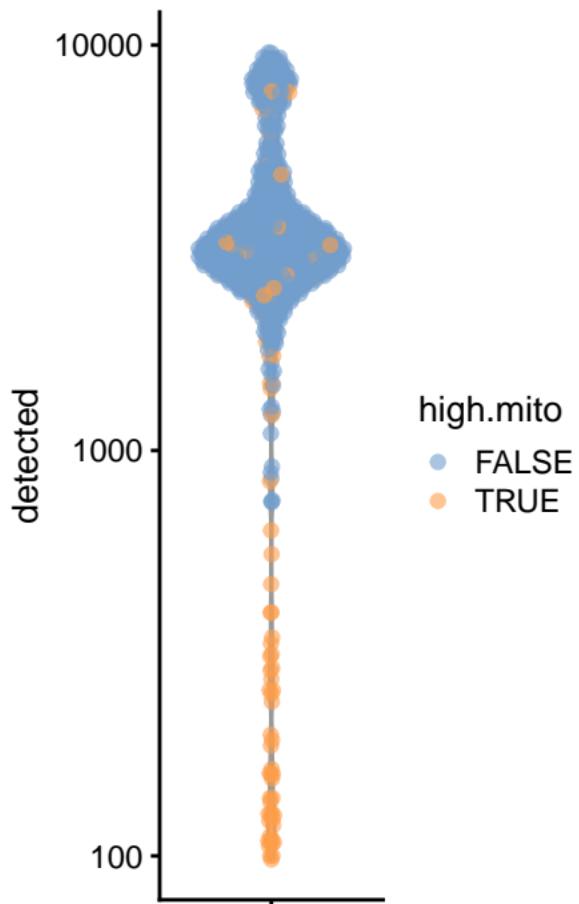
Sample: br5287.hpc; pre-QC nNuclei: 2189; nNuclei kept: 1870 (0.85%)



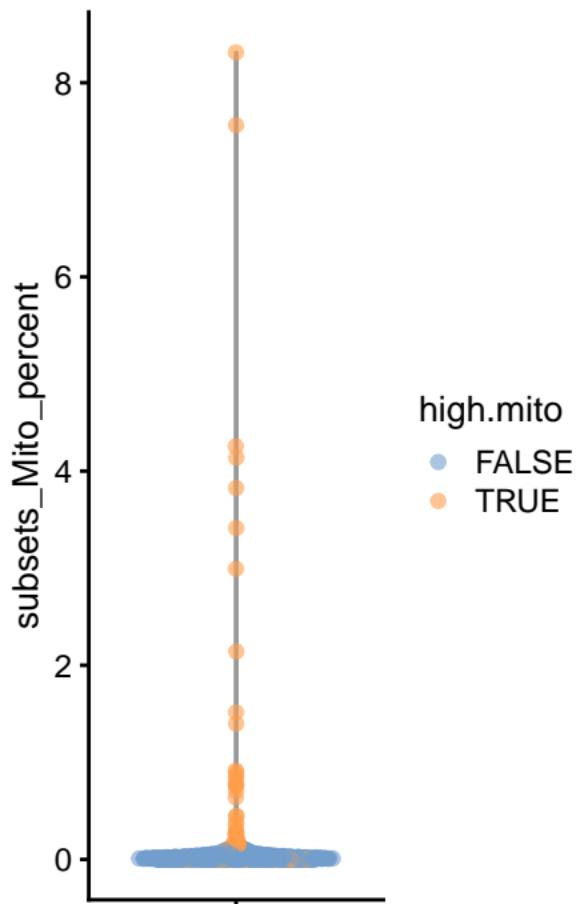
Total count: br5287.nac



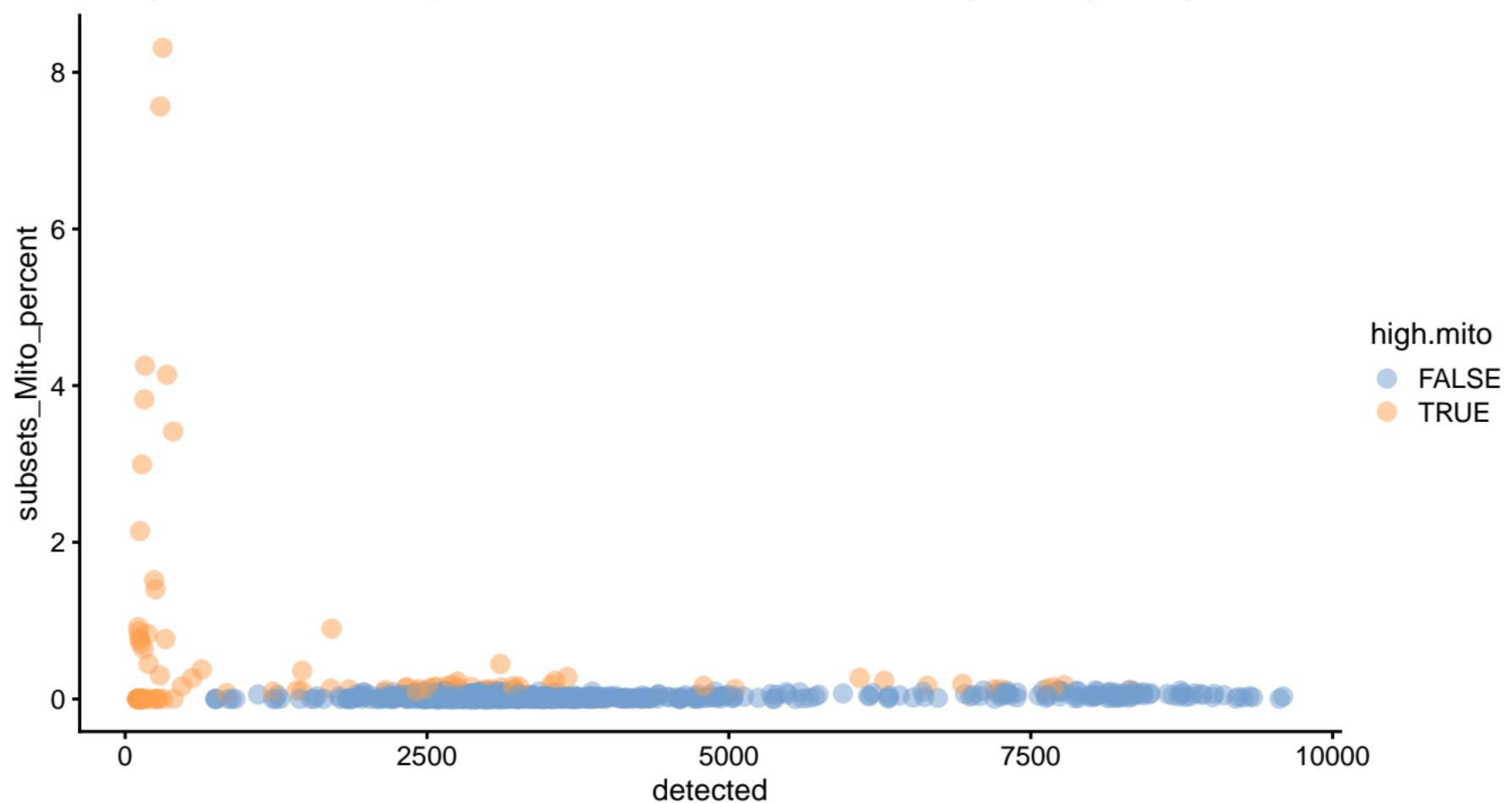
Detected features



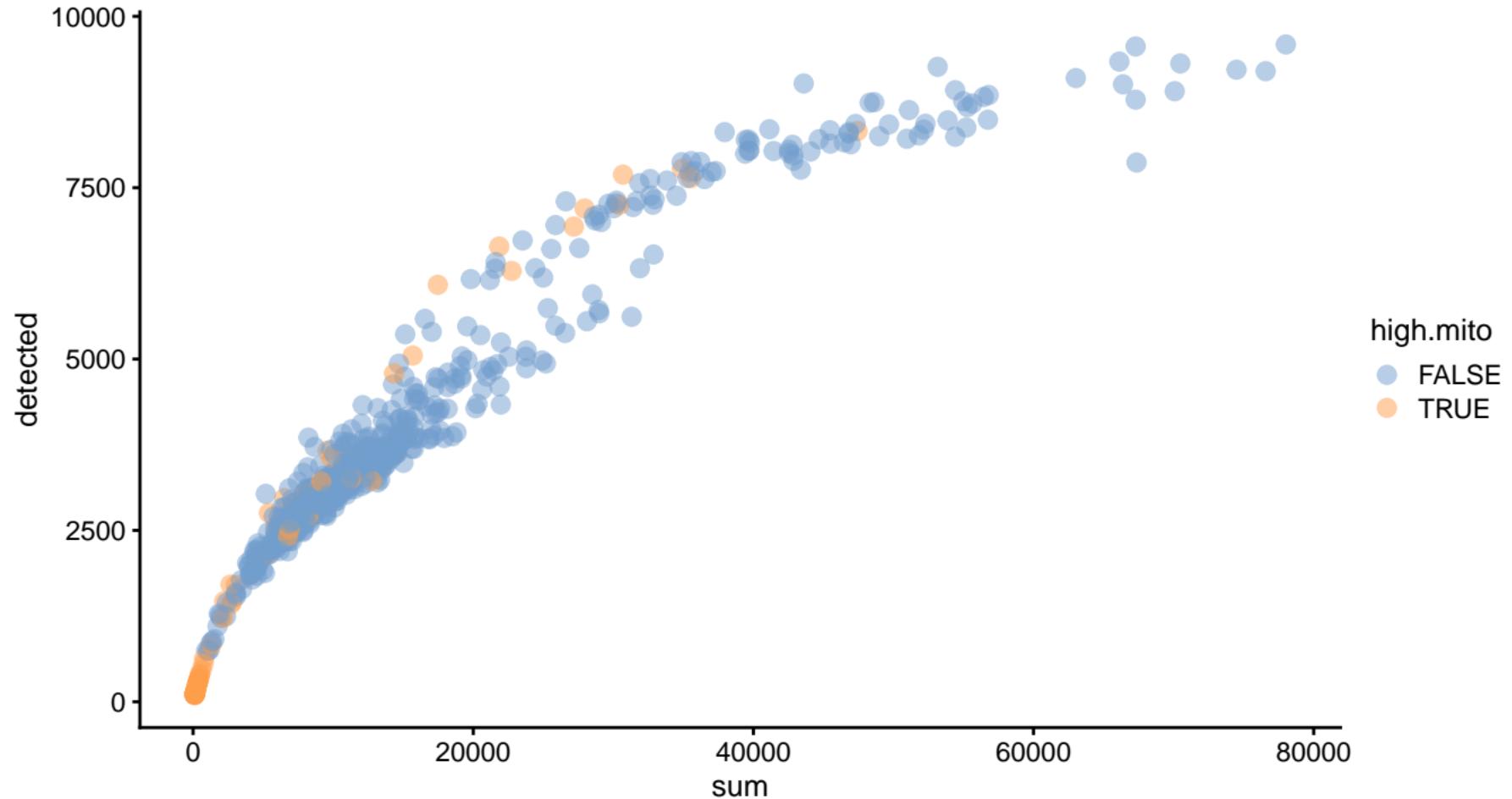
Mito % (cutoff = 0.116)

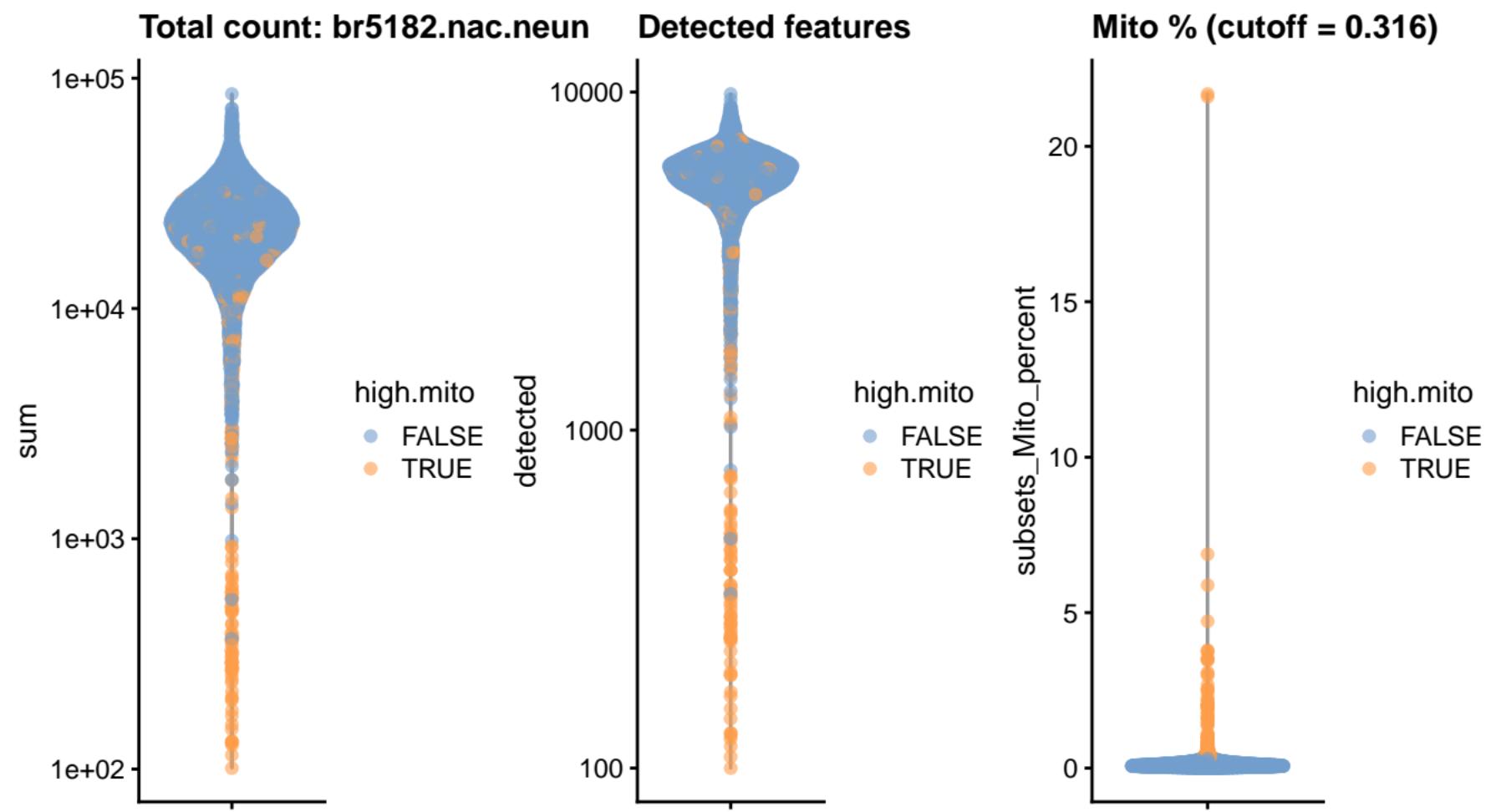


Sample: br5287.nac; pre-QC nNuclei: 770; nNuclei kept: 681 (0.88%)

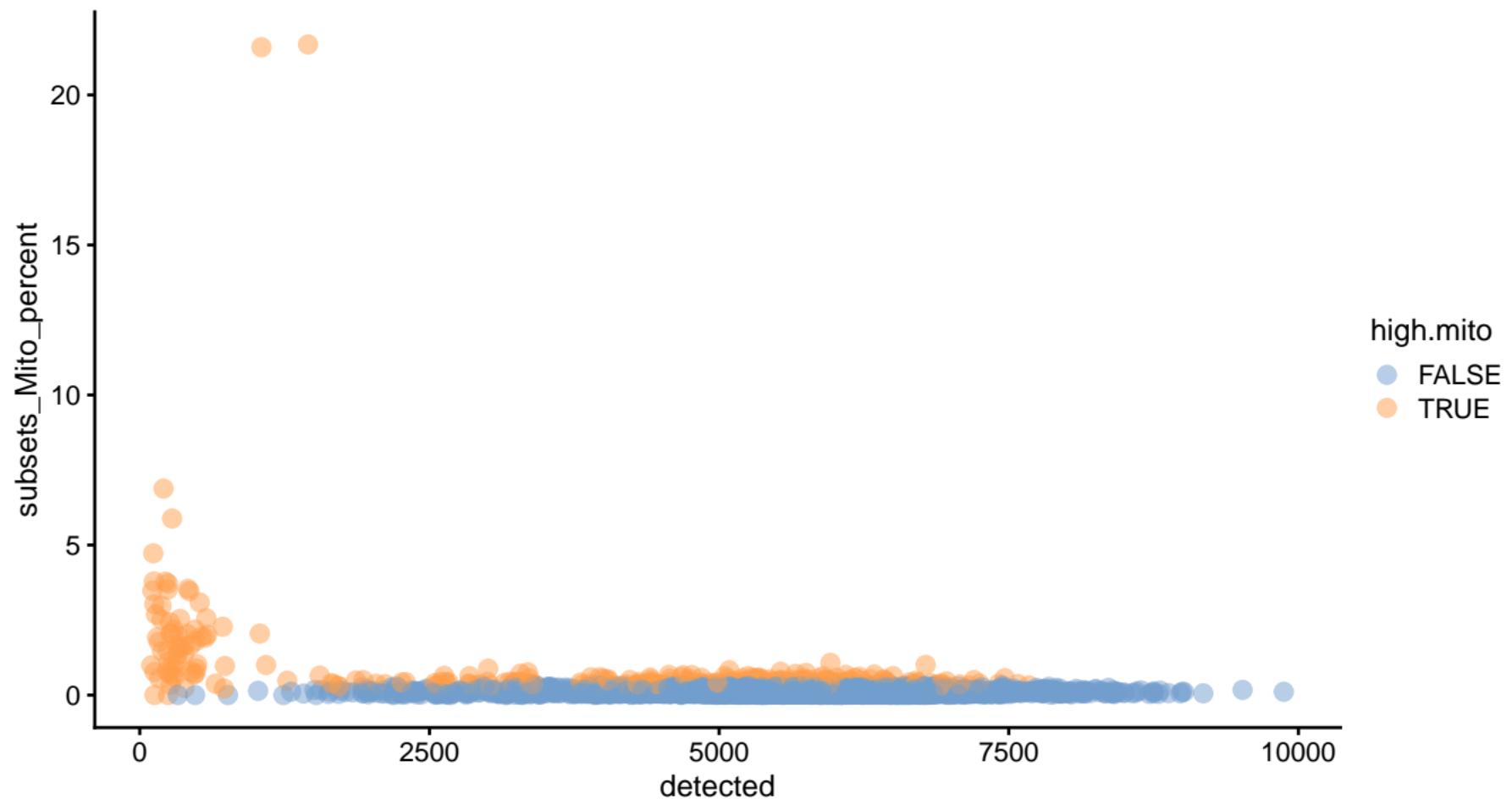


Sample: br5287.nac; pre-QC nNuclei: 770; nNuclei kept: 681 (0.88%)



Total count: br5182.nac.neun

Sample: br5182.nac.neun; pre-QC nNuclei: 4602; nNuclei kept: 4265 (0.93%)



Sample: br5182.nac.neun; pre-QC nNuclei: 4602; nNuclei kept: 4265 (0.93%)

