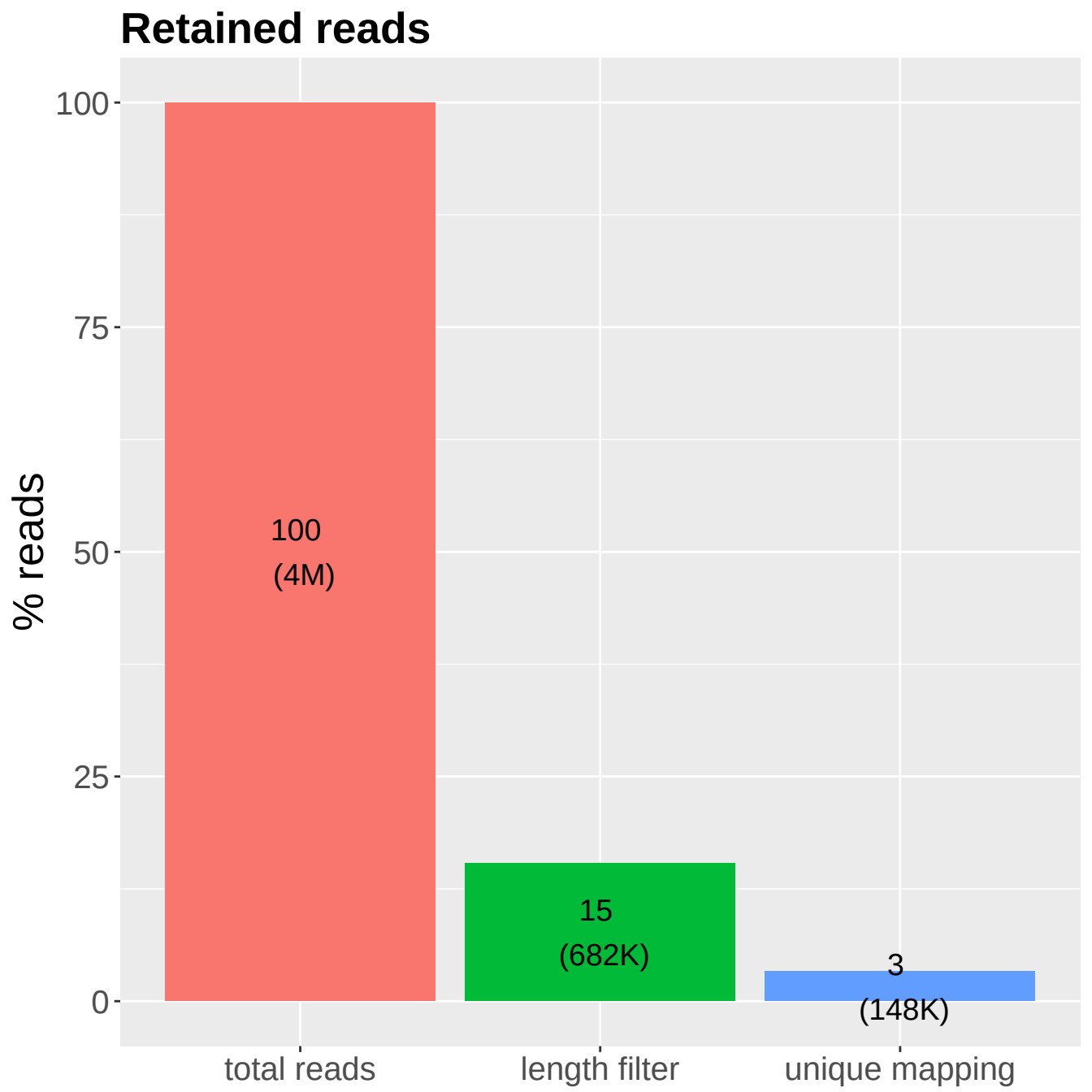
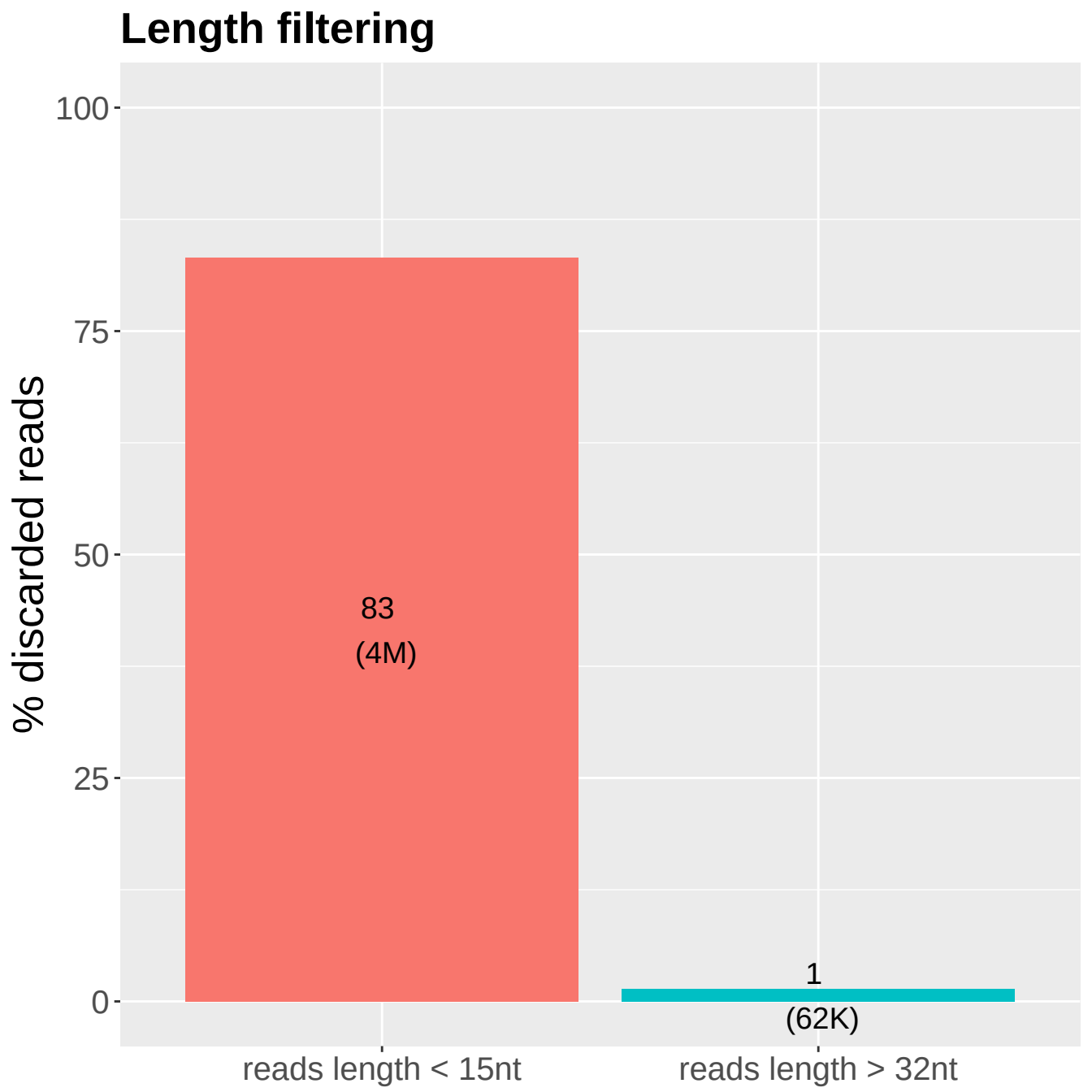
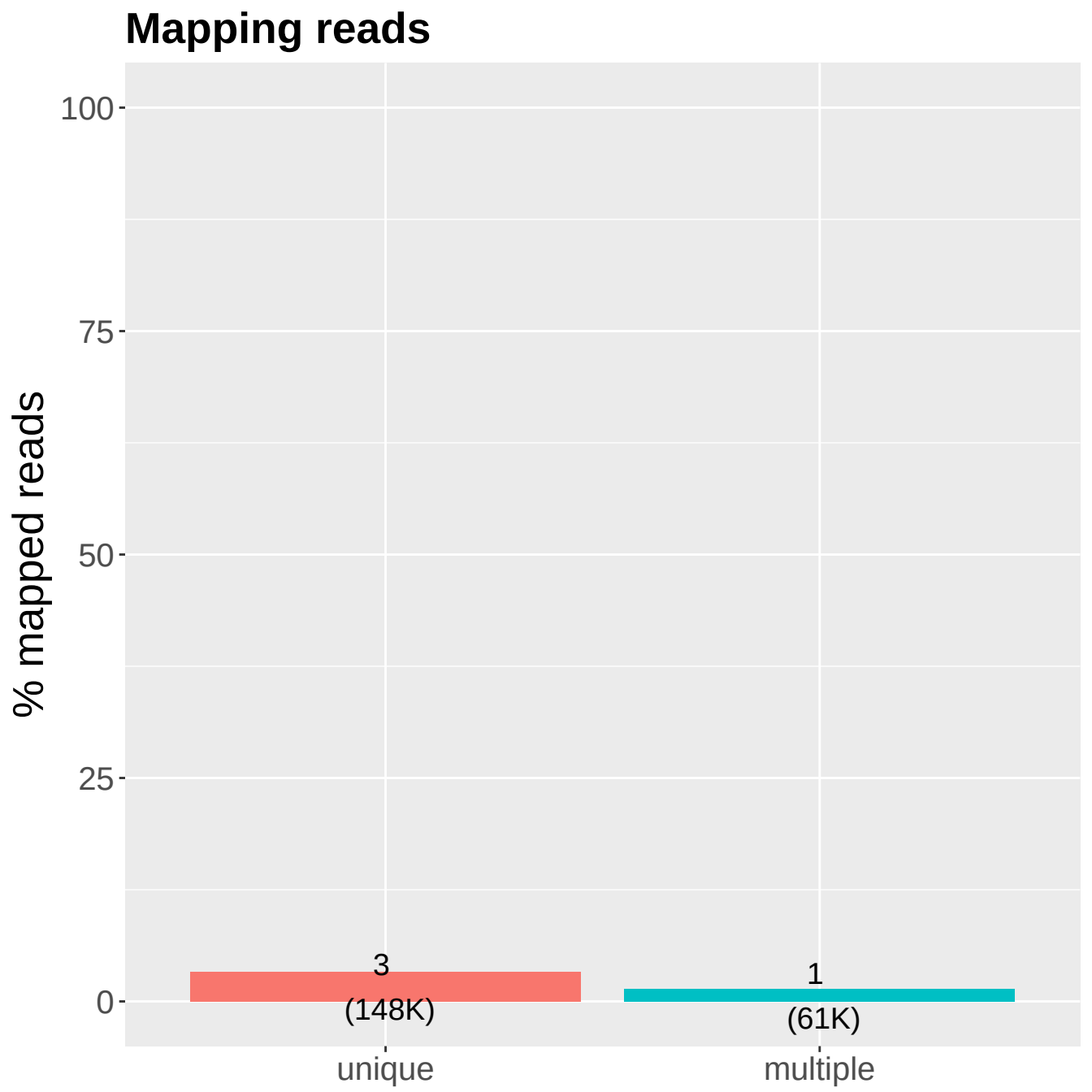
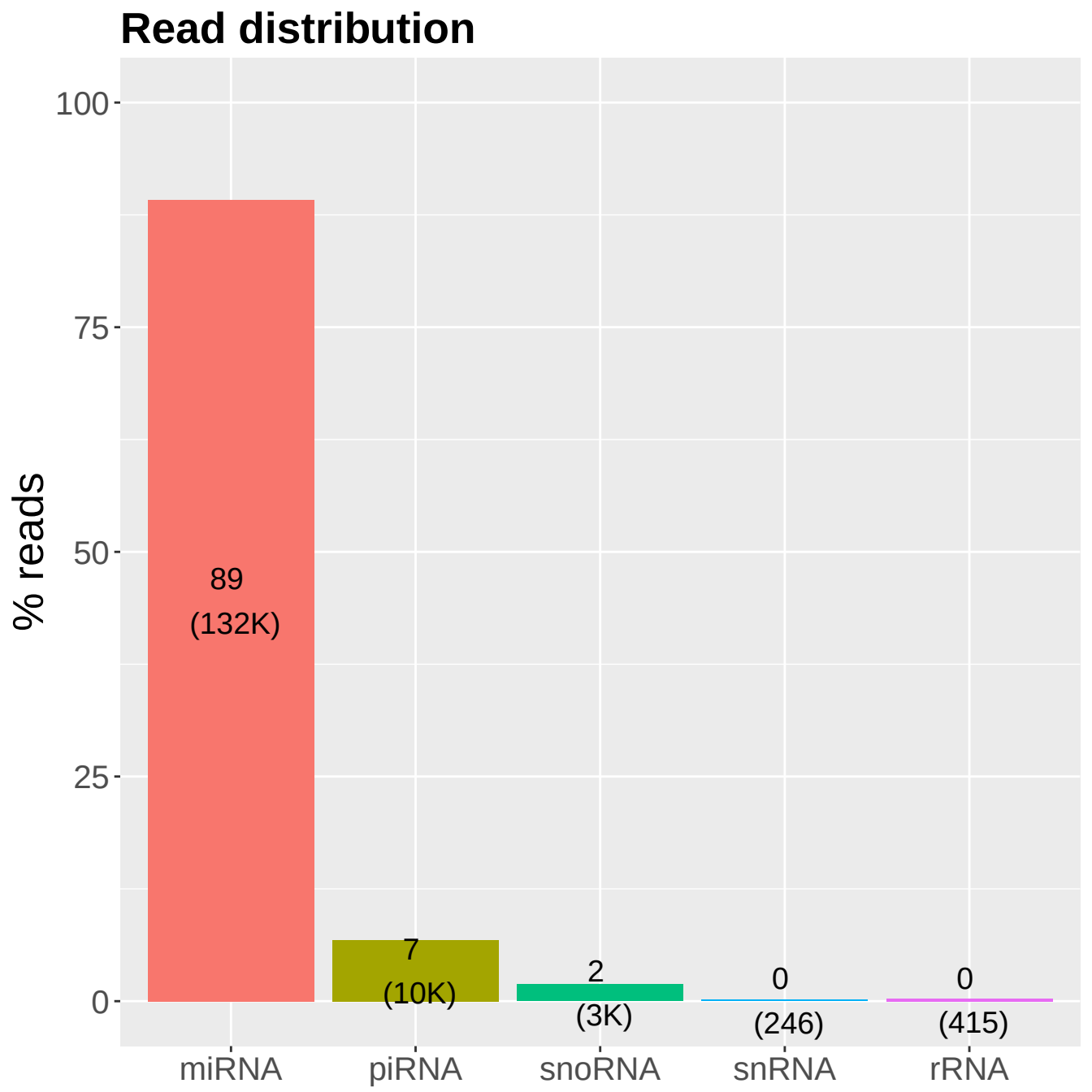

Retained reads
100
75
50
25
0
% reads
100
(4M)
15
(682K)
3
(148K)
total reads
length filter
unique mapping


Length filtering
100
75
50
25
0
% discarded reads
83
(4M)
1
(62K)
reads length < 15nt
reads length > 32nt


Mapping reads
100
75
50
25
0
% mapped reads
3
(148K)
1
(61K)
unique
multiple


Read distribution
100
75
50
25
0
% reads
89
(132K)
7
(10K)
2
(3K)
0
(246)
0
(415)
miRNA
piRNA
snoRNA
snRNA
rRNA