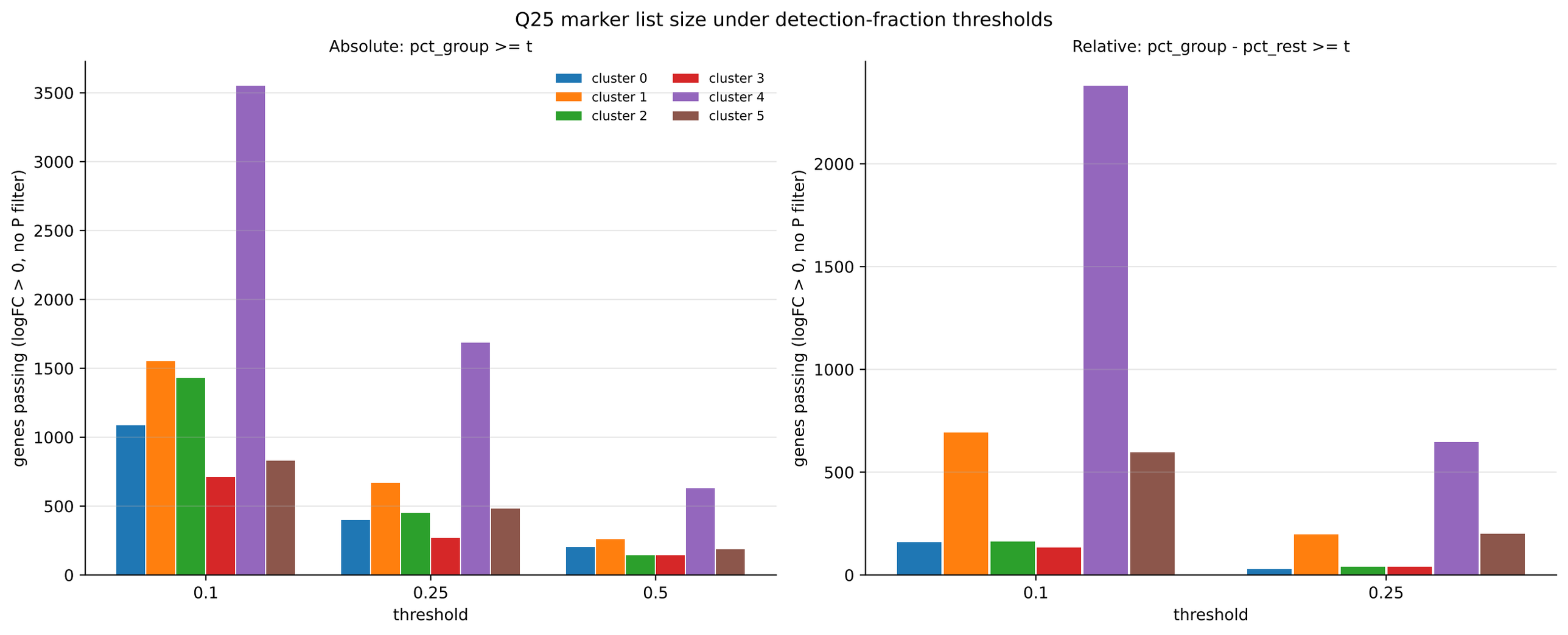

Q25 marker list size under detection-fraction thresholds
Absolute: pct_group >= t
Relative: pct_group - pct_rest >= t
cluster 0
cluster 1
cluster 2
cluster 3
cluster 4
cluster 5
genes passing (logFC > 0, no P filter)
threshold
0.1
0.25
0.5
0
500
1000
1500
2000
2500
3000
3500