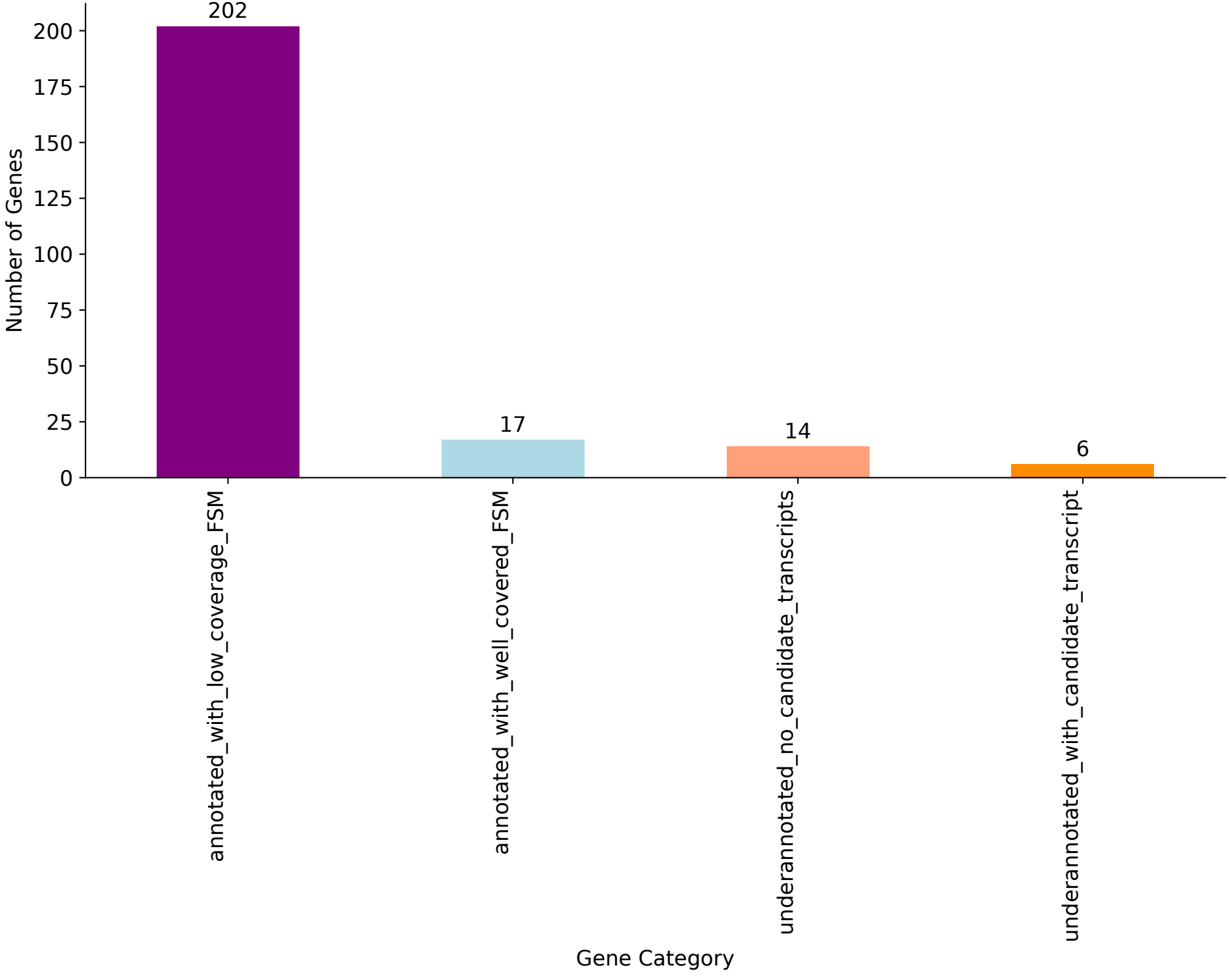
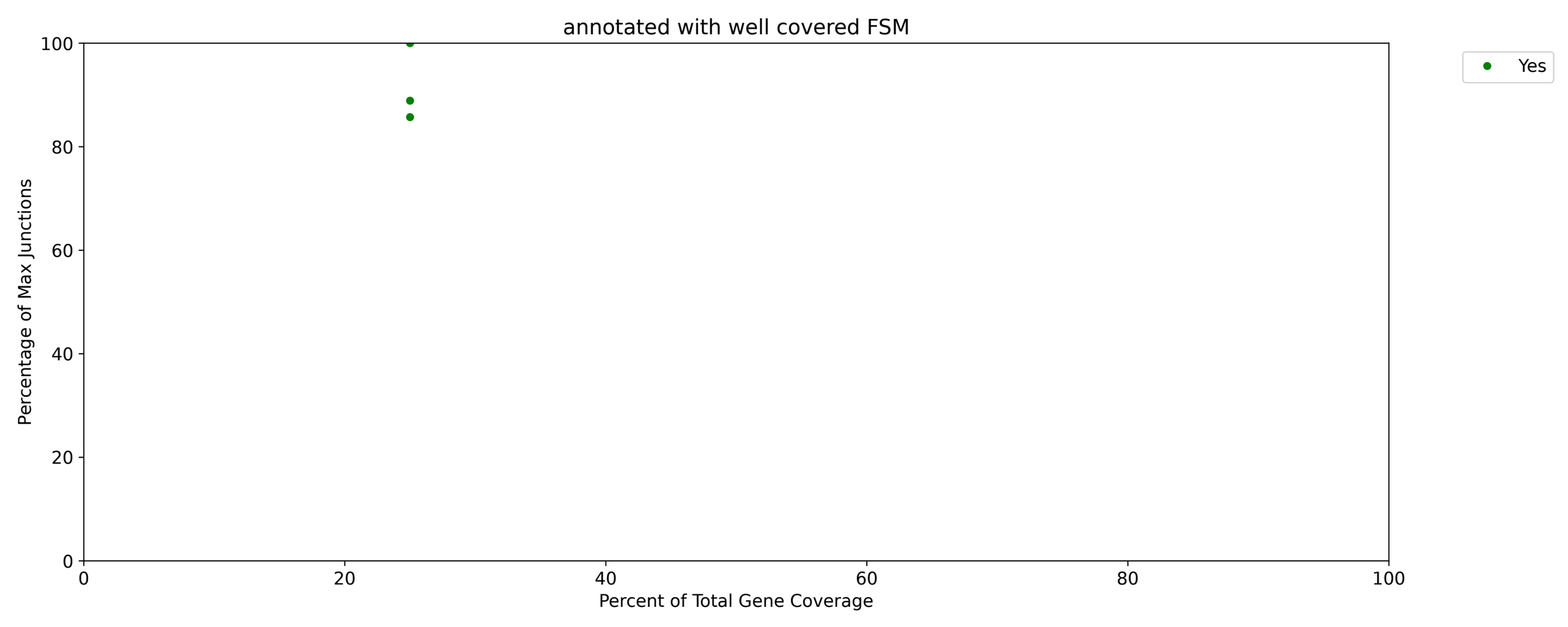
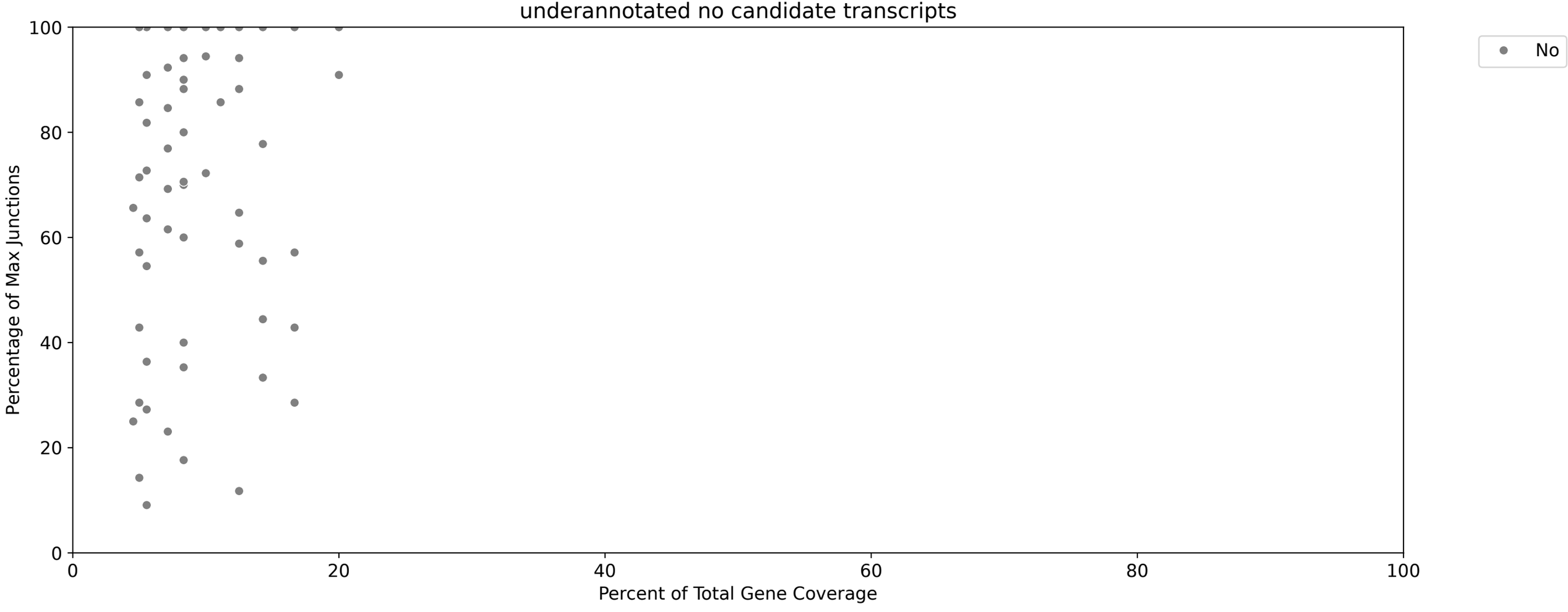
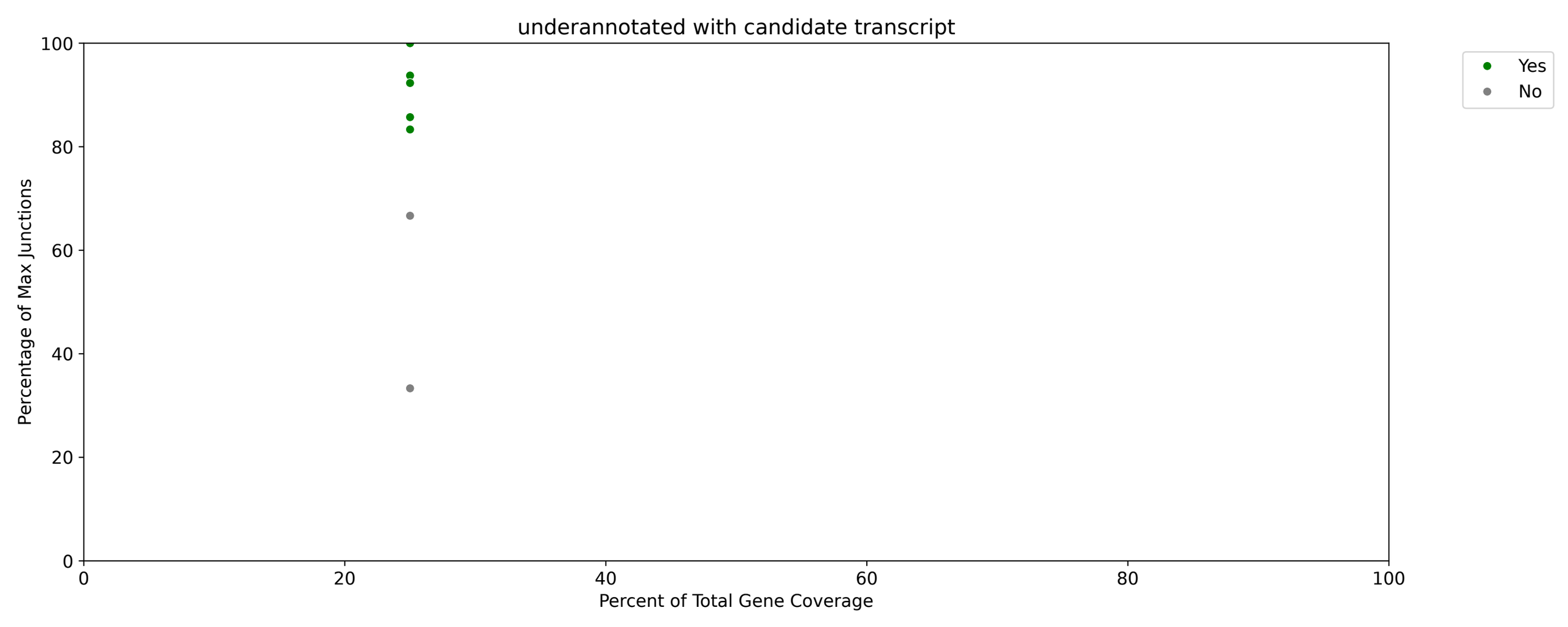
SQANTI-reads annotation report


202
17
14
6
Number of Genes
0
25
50
75
100
125
150
175
200
annotated_with_low_coverage_FSM
annotated_with_well_covered_FSM
underannotated_no_candidate_transcripts
underannotated_with_candidate_transcript
Gene Category


annotated with well covered FSM
Yes
Percentage of Max Junctions
Percent of Total Gene Coverage
0
20
40
60
80
100


underannotated no candidate transcripts
Percentage of Max Junctions
Percent of Total Gene Coverage
No
0
20
40
60
80
100


underannotated with candidate transcript
Yes
No
Percentage of Max Junctions
Percent of Total Gene Coverage
0
20
40
60
80
100