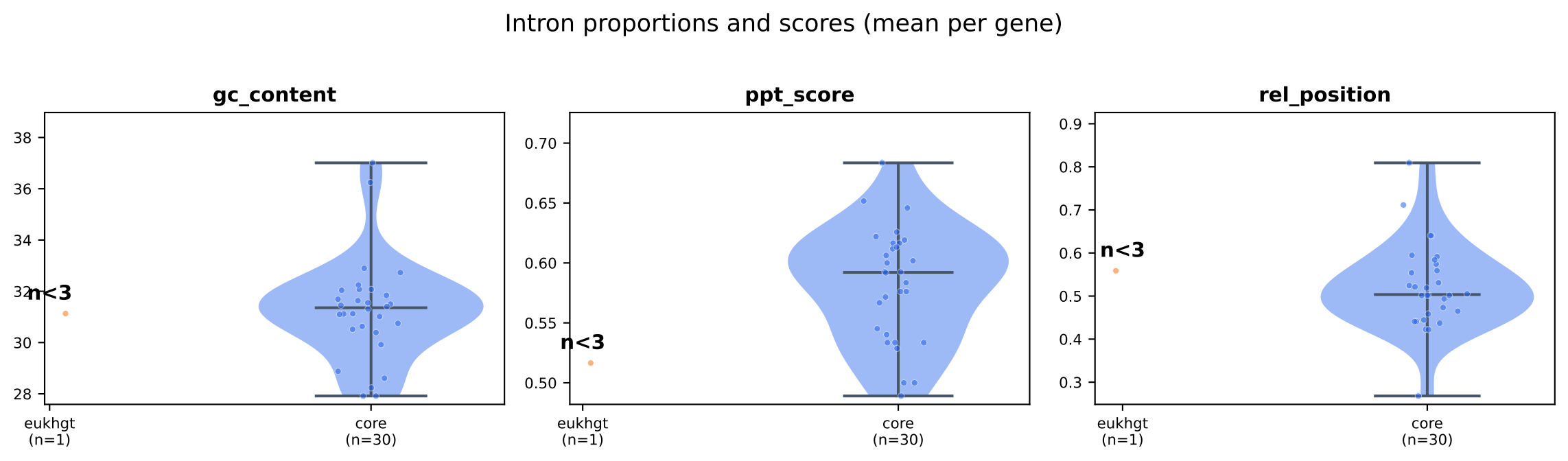

Intron proportions and scores (mean per gene)
gc_content
ppt_score
rel_position
n<3
n<3
n<3
38
36
34
32
30
28
0.70
0.65
0.60
0.55
0.50
0.9
0.8
0.7
0.6
0.5
0.4
0.3
eukhgt
(n=1)
core
(n=30)
eukhgt
(n=1)
core
(n=30)
eukhgt
(n=1)
core
(n=30)