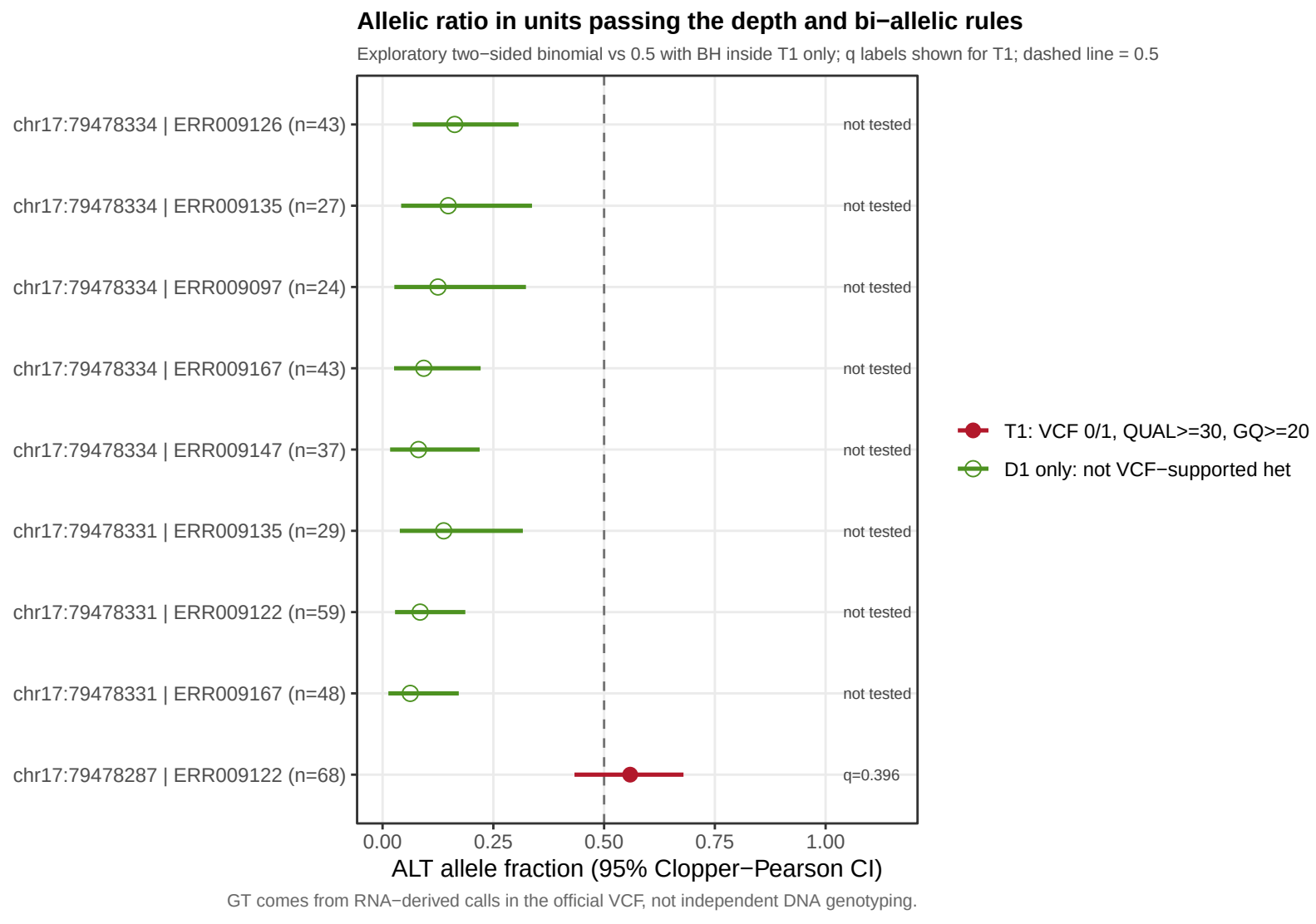

Allelic ratio in units passing the depth and bi−allelic rules
Exploratory two−sided binomial vs 0.5 with BH inside T1 only; q labels shown for T1; dashed line = 0.5
chr17:79478334 | ERR009126 (n=43)
chr17:79478334 | ERR009135 (n=27)
chr17:79478334 | ERR009097 (n=24)
chr17:79478334 | ERR009167 (n=43)
chr17:79478334 | ERR009147 (n=37)
chr17:79478331 | ERR009135 (n=29)
chr17:79478331 | ERR009122 (n=59)
chr17:79478331 | ERR009167 (n=48)
chr17:79478287 | ERR009122 (n=68)
not tested
not tested
not tested
not tested
not tested
not tested
not tested
not tested
q=0.396
0.00
0.25
0.50
0.75
1.00
ALT allele fraction (95% Clopper−Pearson CI)
T1: VCF 0/1, QUAL>=30, GQ>=20
D1 only: not VCF−supported het
GT comes from RNA−derived calls in the official VCF, not independent DNA genotyping.