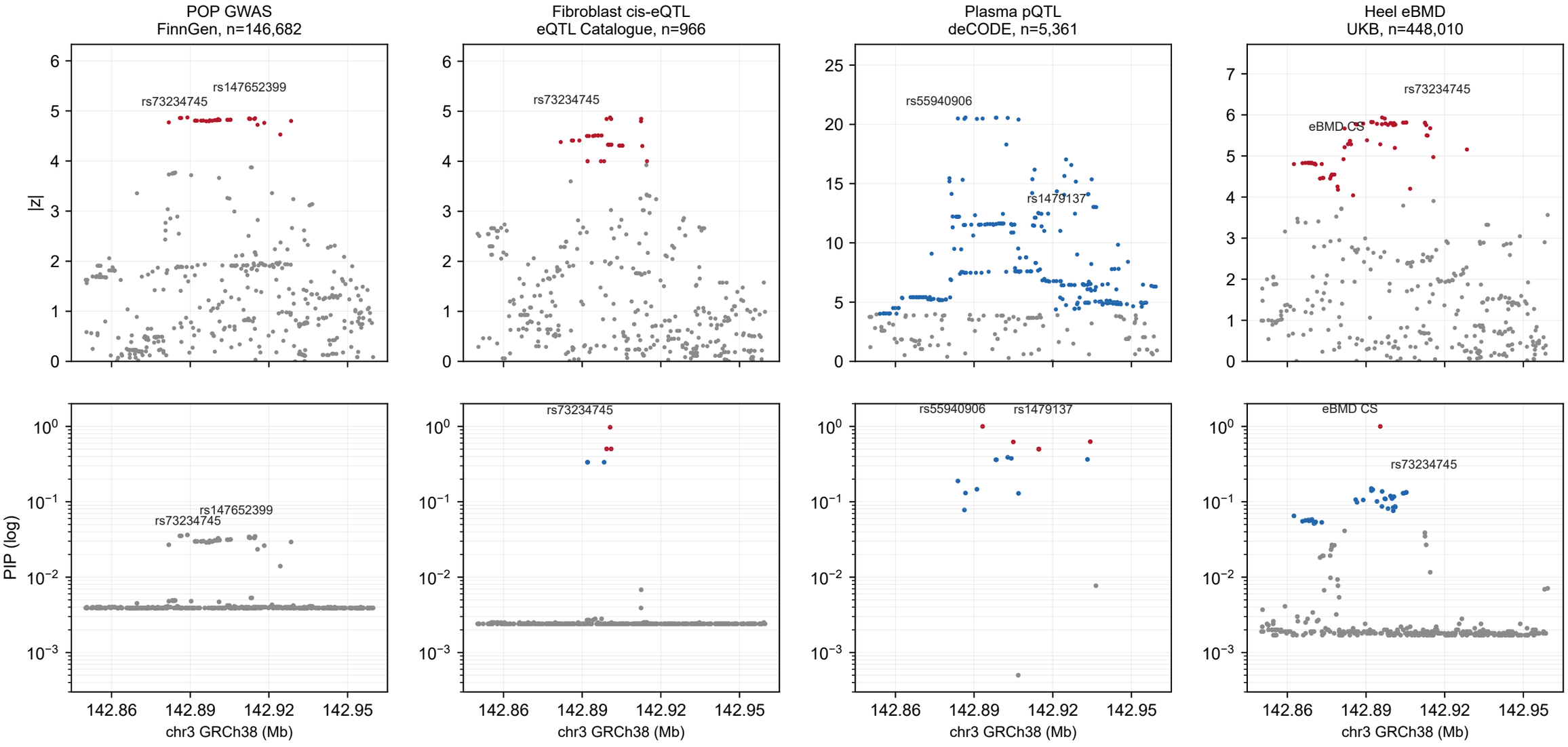


Fibroblast eQTL: 3 tight CS (purity ≥0.98); pQTL: 8 CS (purity ≥0.90); eBMD: 1 single-variant CS + one 1,112-variant CS (purity = 0, D'≈1 block); POP: no CS (max PIP 0.04).
Credible sets are phenotype-specific and mutually non-overlapping (all pairwise Jaccard ≤0.011).