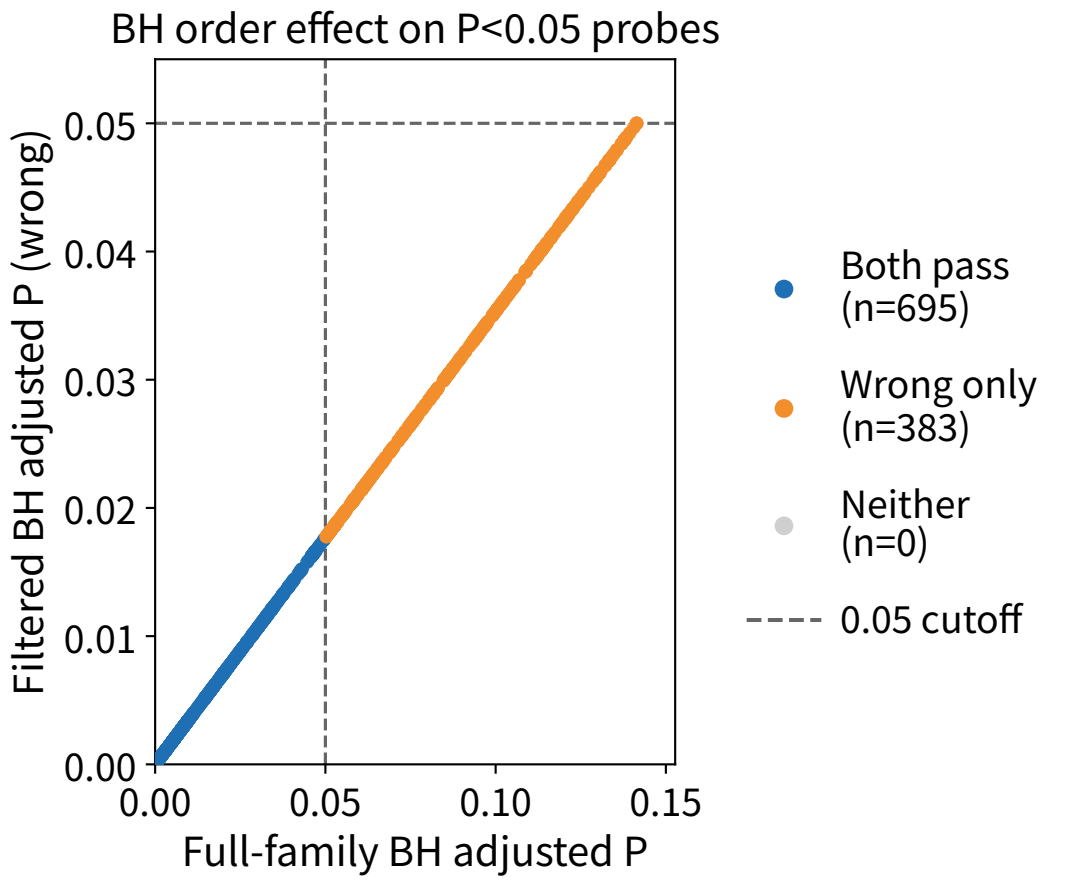
BH order effect on P<0.05 probes
Filtered BH adjusted P (wrong)
Full-family BH adjusted P
0.00
0.01
0.02
0.03
0.04
0.05
0.00
0.05
0.10
0.15
Both pass
(n=695)
Wrong only
(n=383)
Neither
(n=0)
0.05 cutoff