

count

-1.0

-0.5

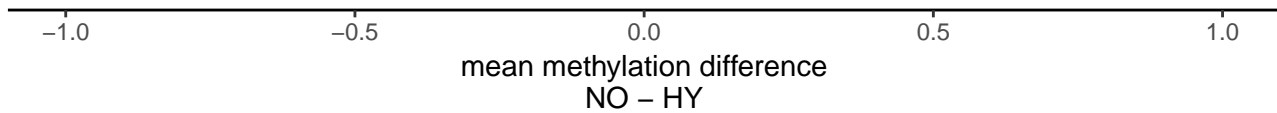
0.0

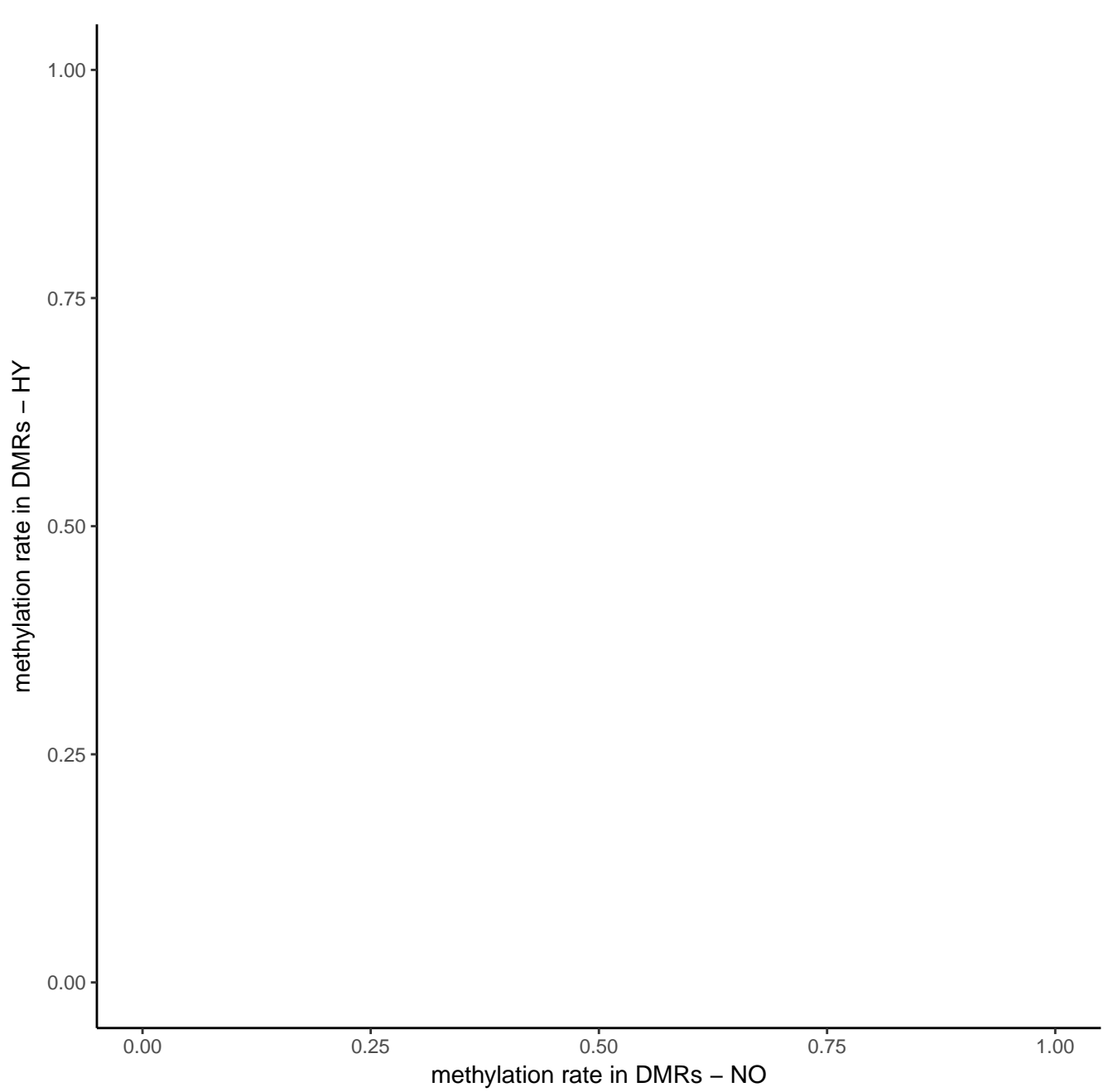
0.5

1.0

mean methylation difference
NO - HY

q-value





density

DMR length [nt]

density

DMR length [Cs]

DMR length [Cs]

DMR length [nt]