b) Percentage QRNA scores > shuffled scores (x10^3)

Subtract Distributions

-5 0 5

-5 0 5

Percentage of z counts > shuffled counts (x10^3)

-5 0 5

-5 0 5

zMfold z-Score

Evofold Score (human-chimp alignments)

5' UTR

Coding

3' UTR

Intronic

Intergenic

Random

Percentage of z counts > shuffled counts (x10^3)

-5 0 5

-5 0 5

Percentage QRNA scores > shuffled scores (x10^3)

-5 0 5

-5 0 5

zMfold z-Score

Evofold Score (human-chimp alignments)