

is_1.2_order_magnitude



bigger



no



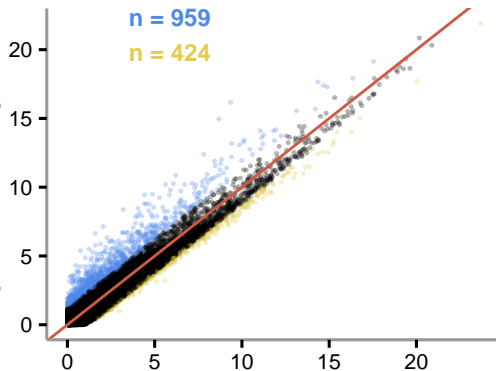
smaller

log10 p-value
(transcriptome diversity controlled)

PValue_transcriptome_diversity_tpm

n = 959

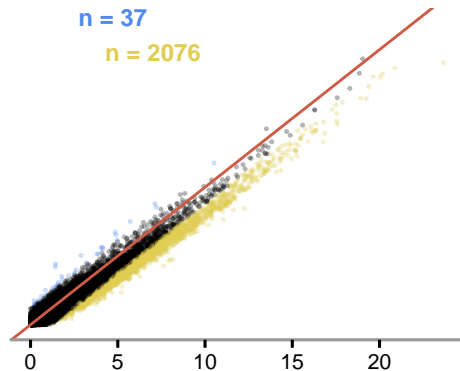
n = 424



PValue_transcriptome_diversity_tpm_scrambl

n = 37

n = 2076



log10 p-value
(original)