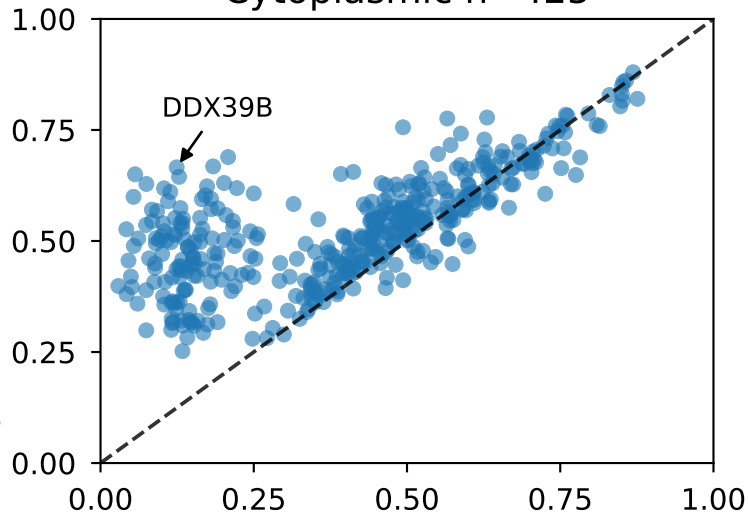


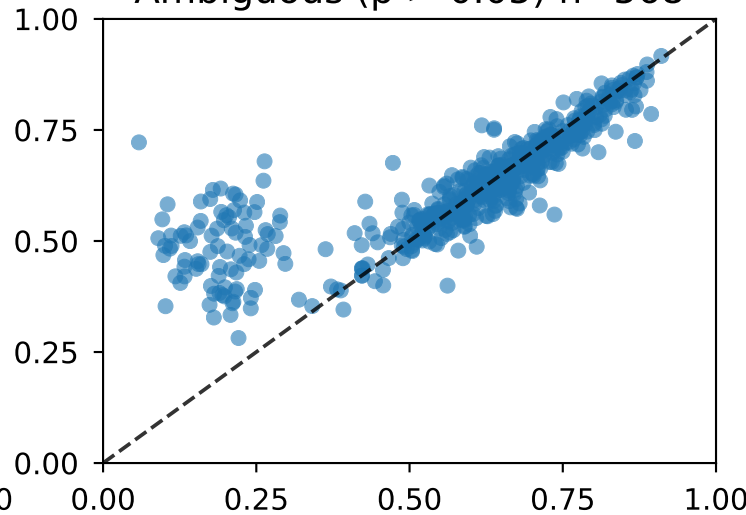
Nuclear predictions with retained intron

Cytoplasmic n=425



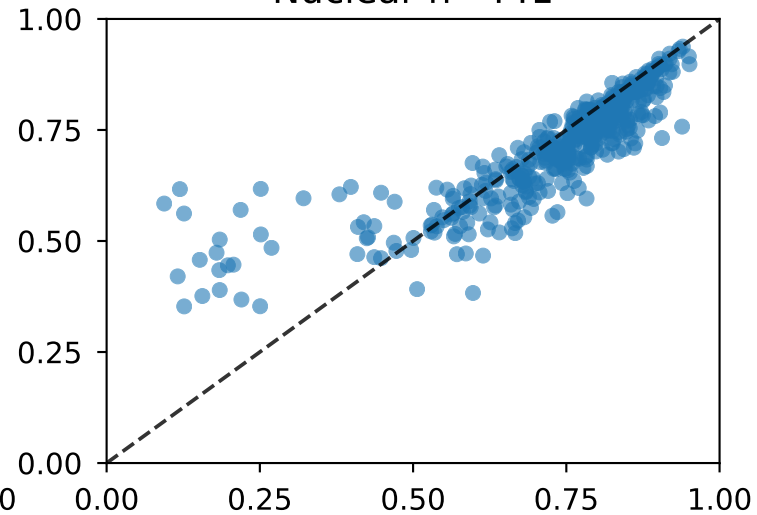
Nuclear predictions without retained intron

Ambiguous ($p > 0.05$) n=568



Nuclear predictions without retained intron

Nuclear n=441



Nuclear predictions without retained intron