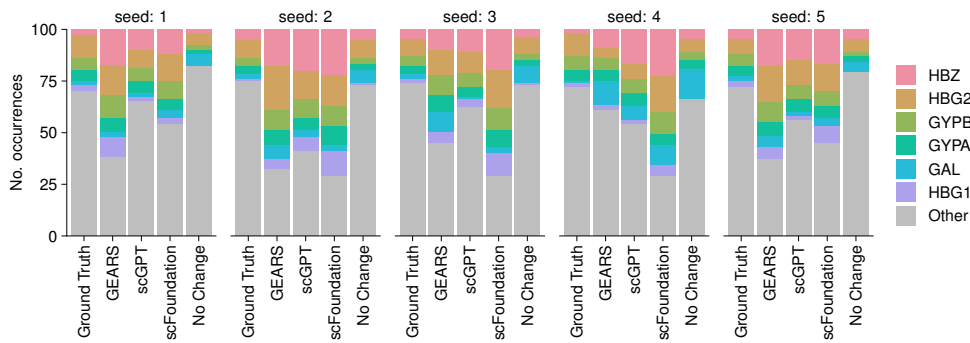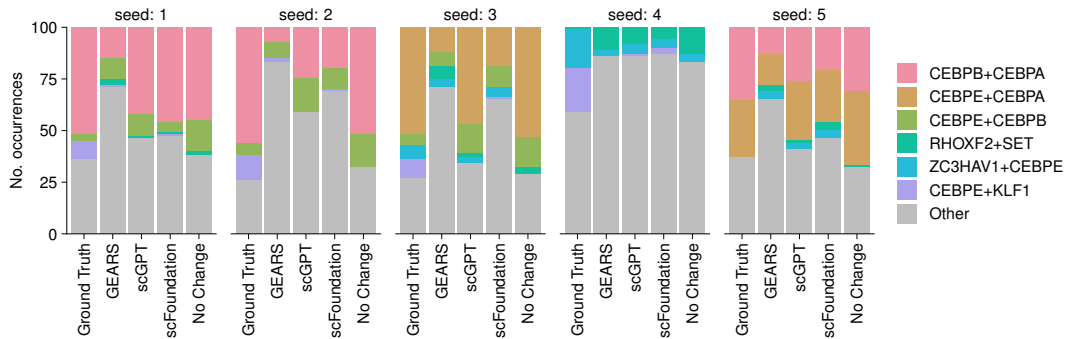**(A) Reoccurring genes among 100 top synergistic predictions**

(B) Reoccurring perturbations among 100 top synergistic predictions

(C) Expression predictions vs observed and additive model