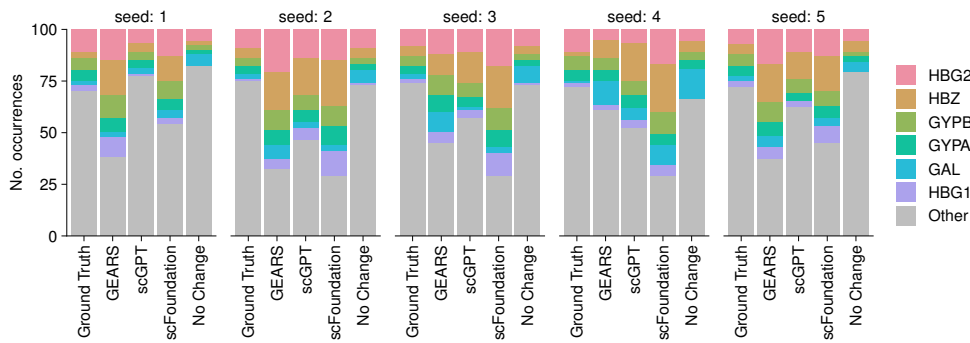
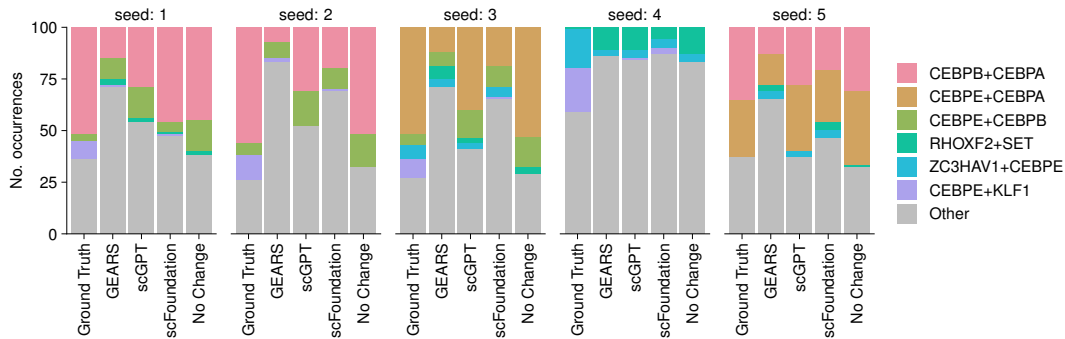


**(A) Reoccurring genes among 100 top synergistic predictions****(B) Reoccurring perturbations among 100 top synergistic predictions****(C) Expression predictions vs observed and additive model**