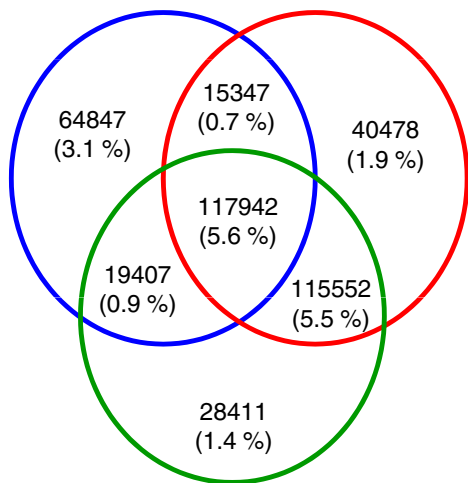


(a)

— RELARELA
 — RELAp50
 — RELAp52



number of distinct 11-mers enriched
 during EMSA-Seq
 from a starting pool
 of 2,097,152 sequences

(b)

Binding models generated using the
 top affinity binders from EMSA-Seq

