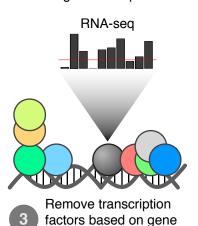
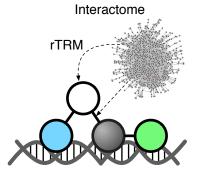


Add computational prediction of binding for the query transcription factors



expression



Identifying TRMs from PPI networks