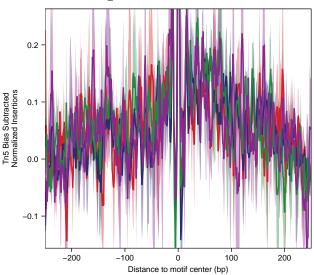
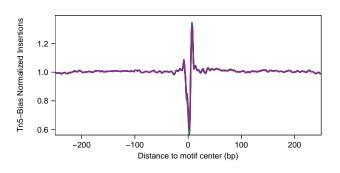
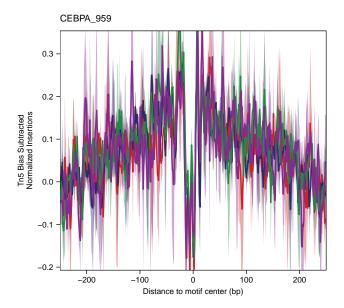
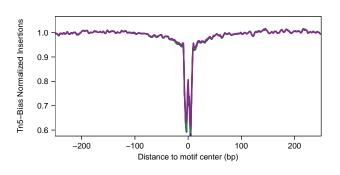


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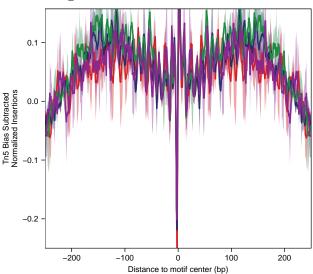


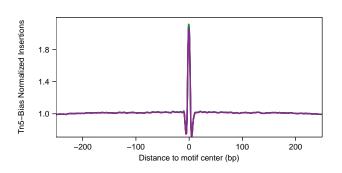




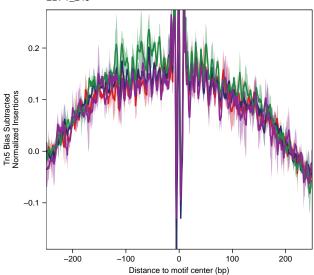


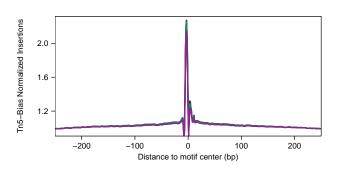




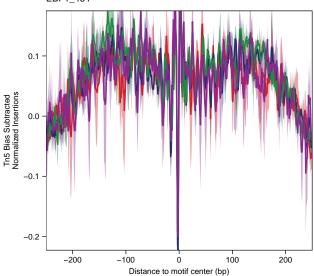


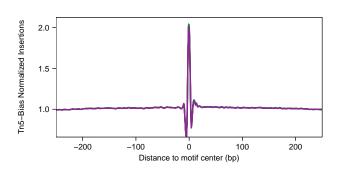


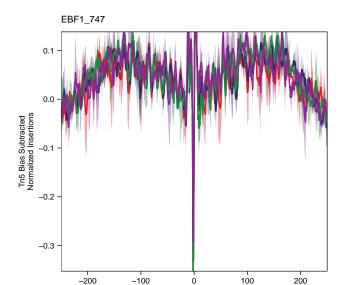


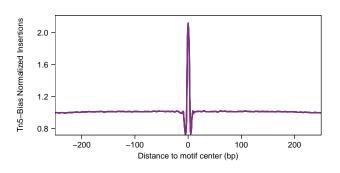






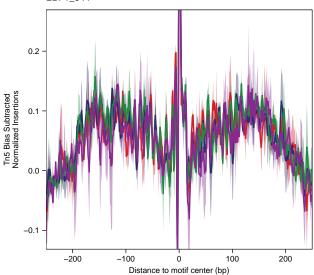


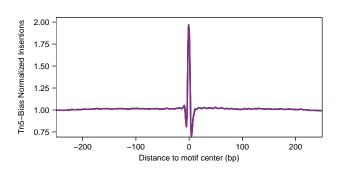




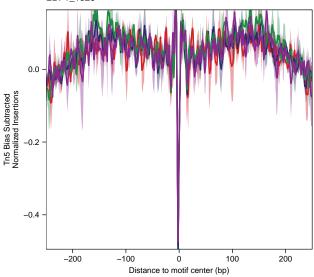
Distance to motif center (bp)

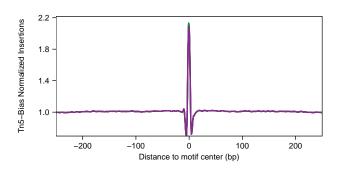


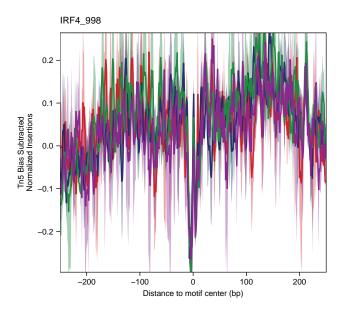


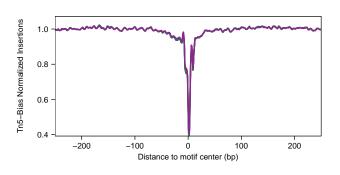


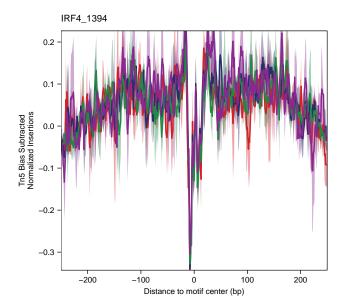


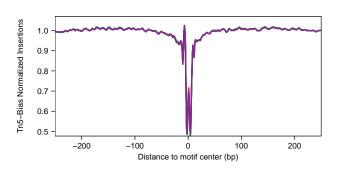




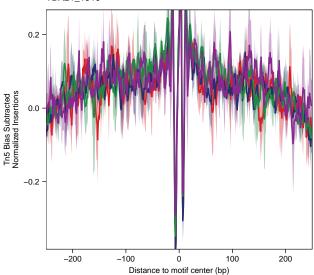


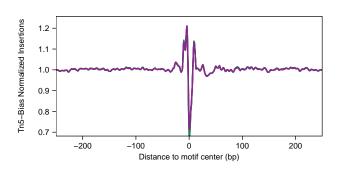


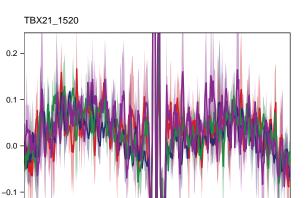












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Distance to motif center (bp)

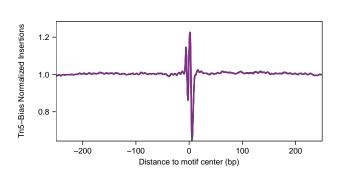
100

200

Tn5 Bias Subtracted Normalized Insertions

-0.2

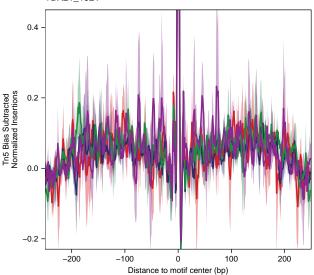
-200

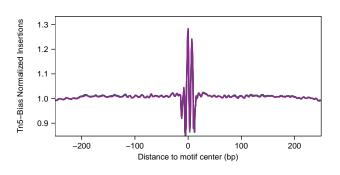


group CFA_3DDA CFA_ApAP CFA_Veh Saline_Veh

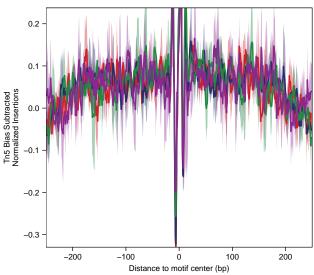
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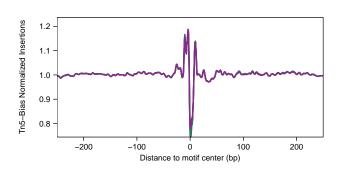


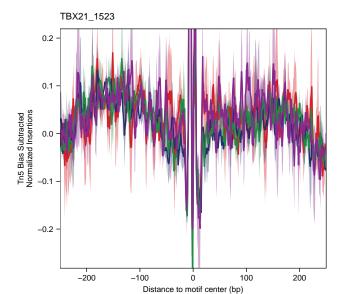


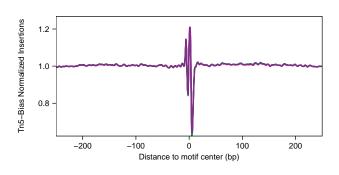




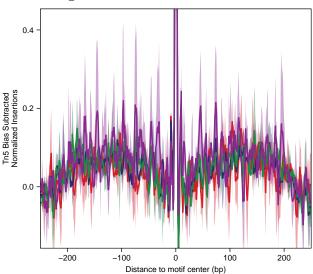


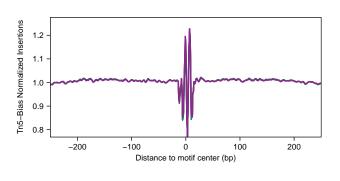


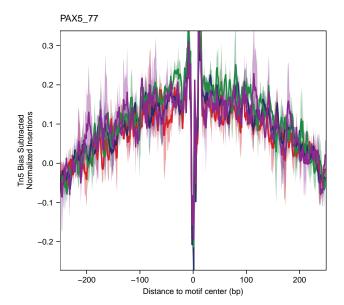


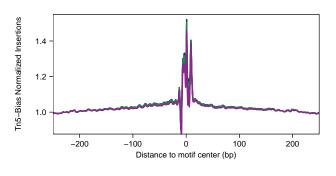




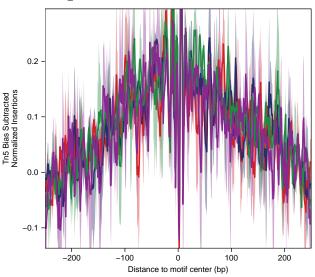


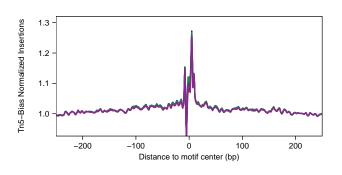


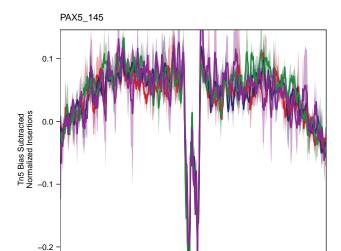


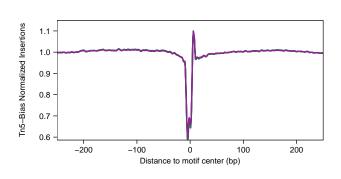












Distance to motif center (bp)

-200

-100

100

200



