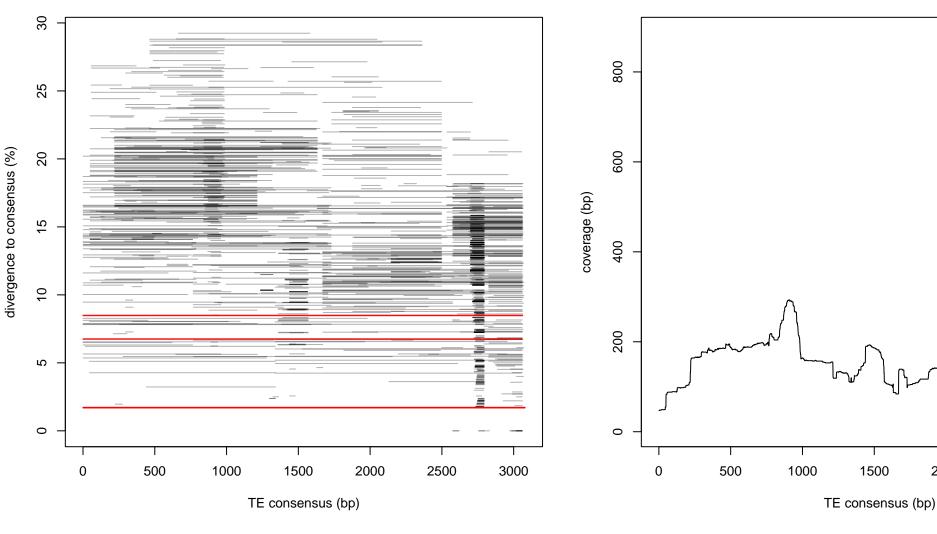
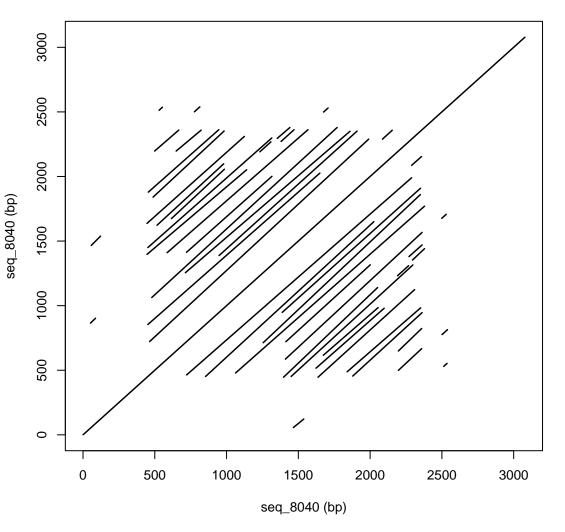
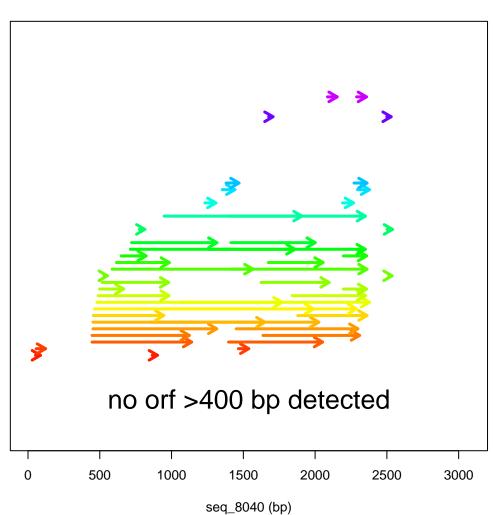
TE consensus genomic coverage



TE consensus self dotplot (blastn)

TE consensus structure and protein hits





1500

2000

2500

3000