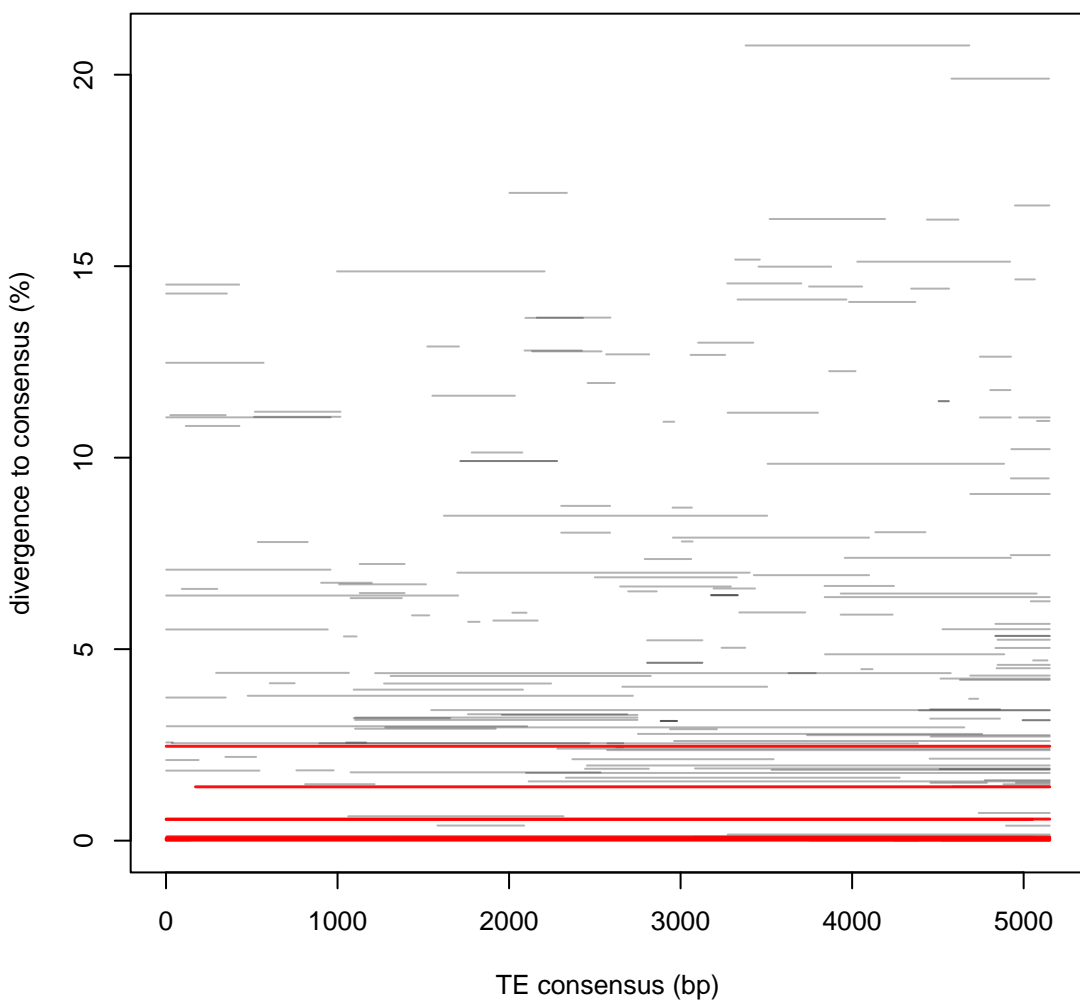
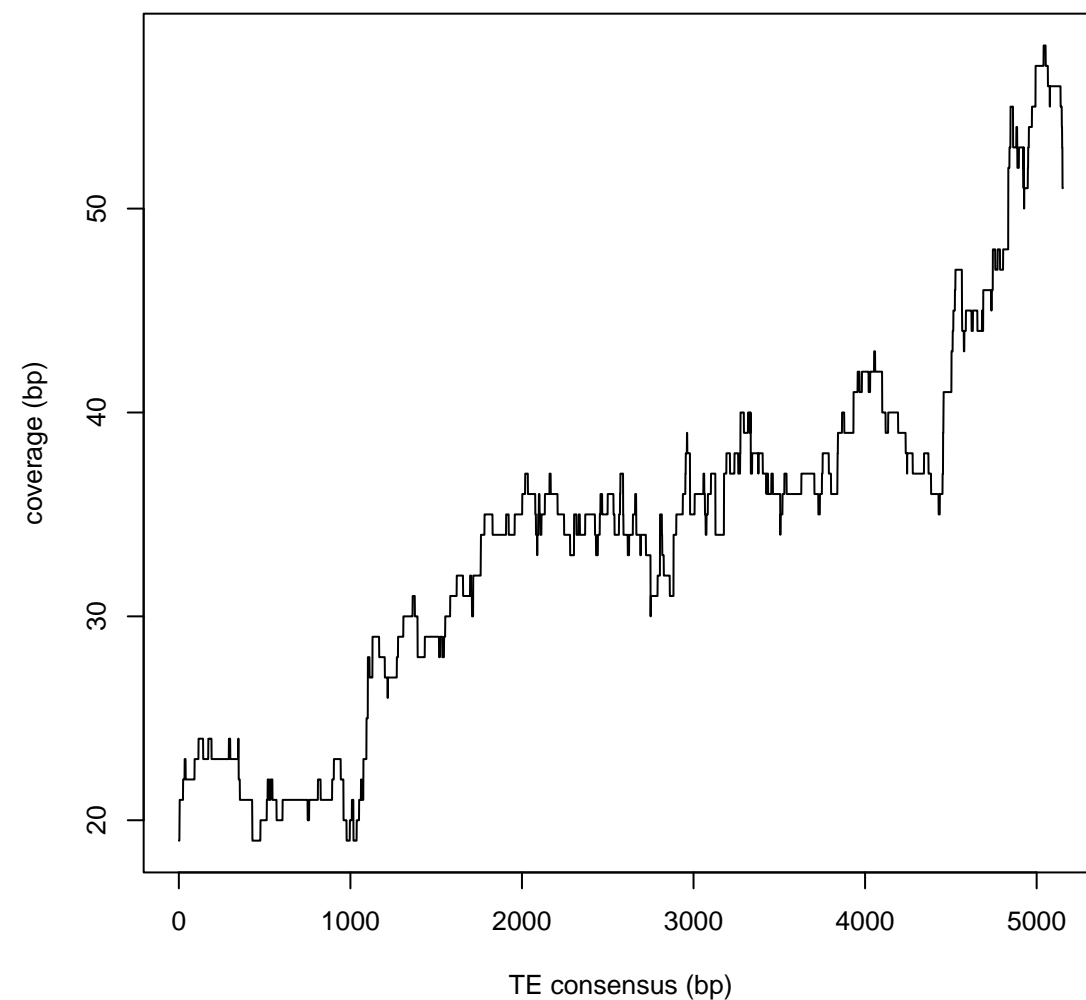


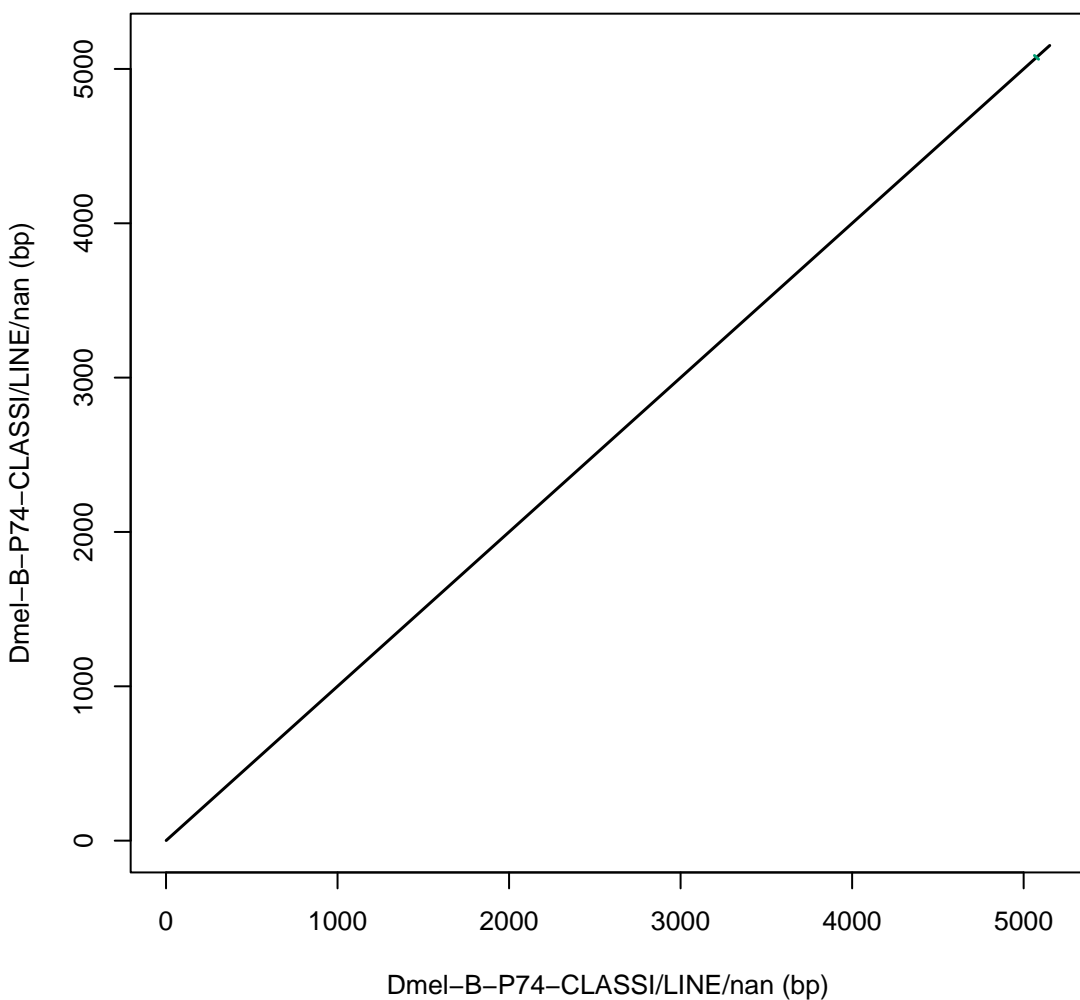
TE: Dmel-B-P74-CLASSI/LINE/nan  
consensus size: 5152bp; fragments: 191; full length: 9 (>=4636.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

