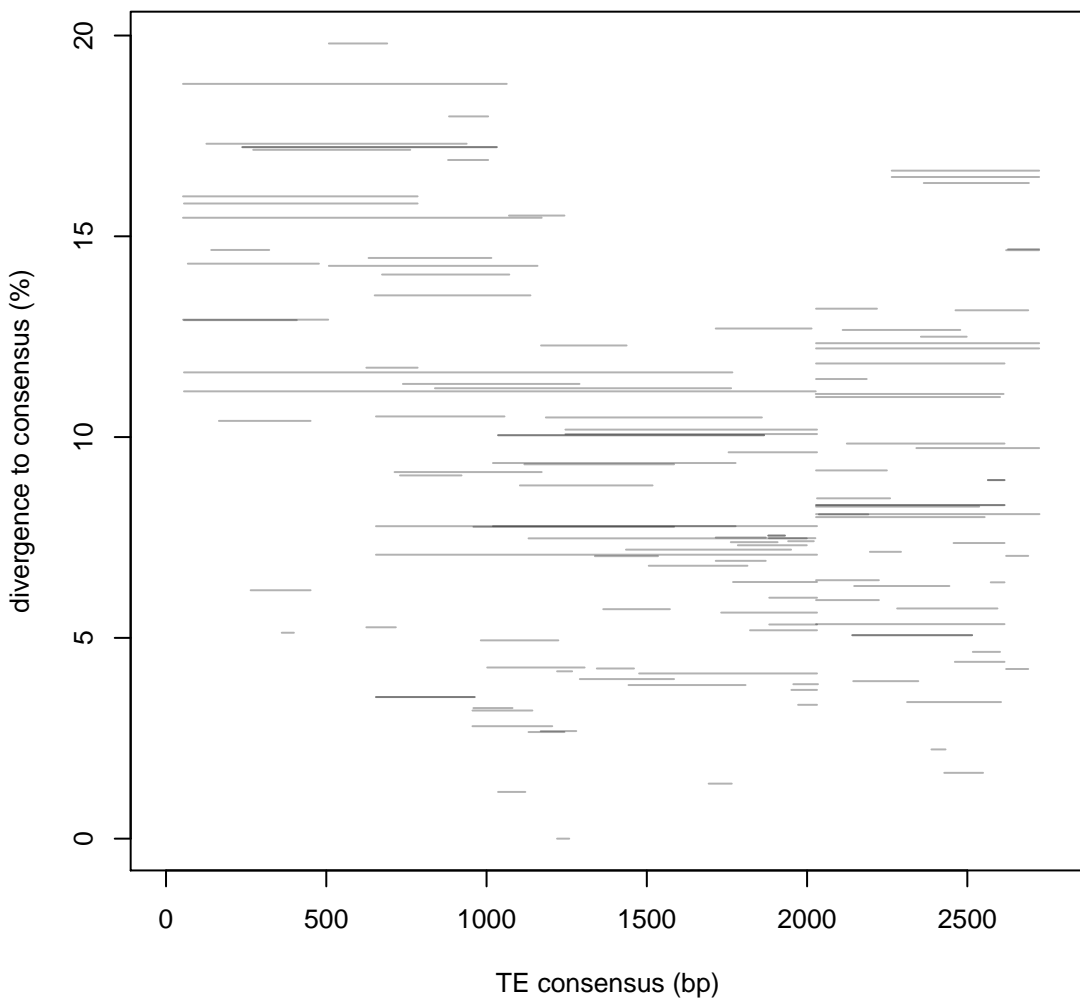
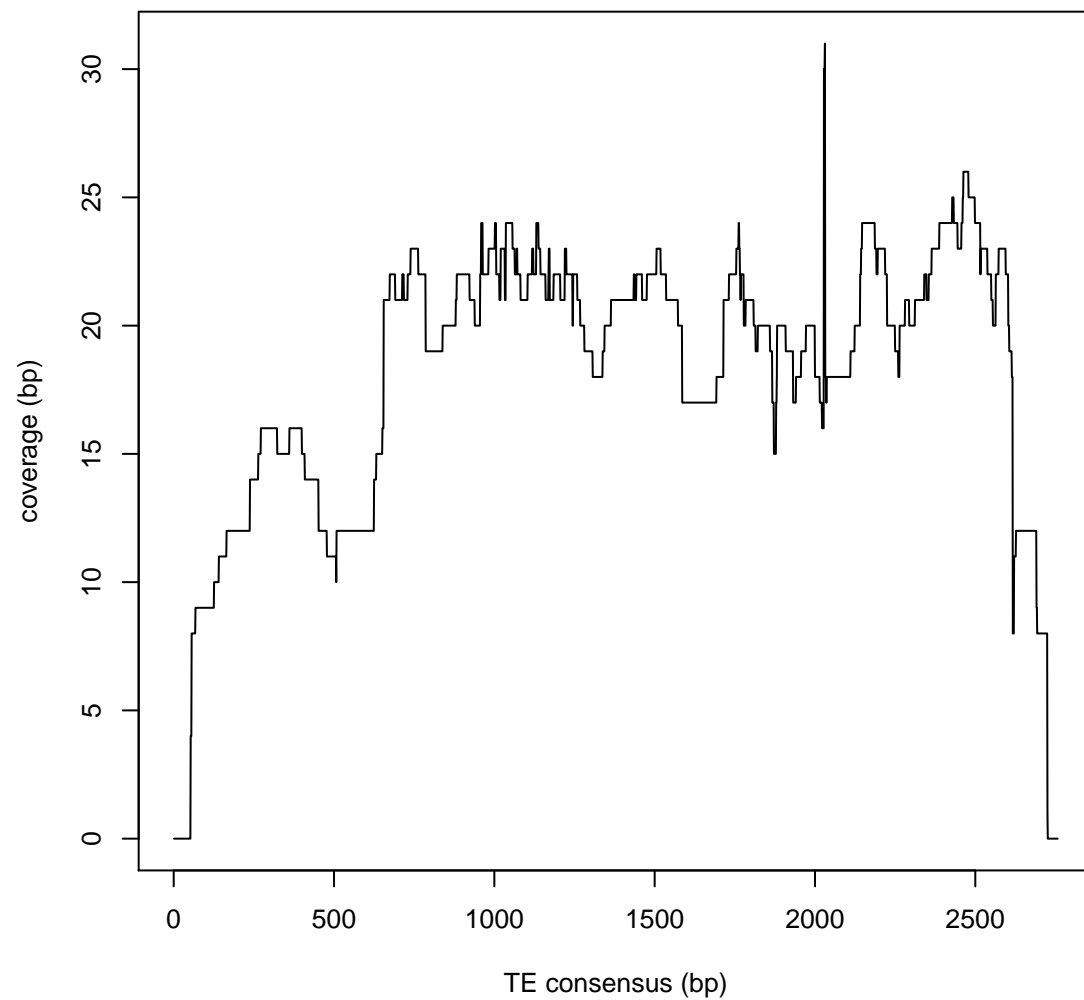


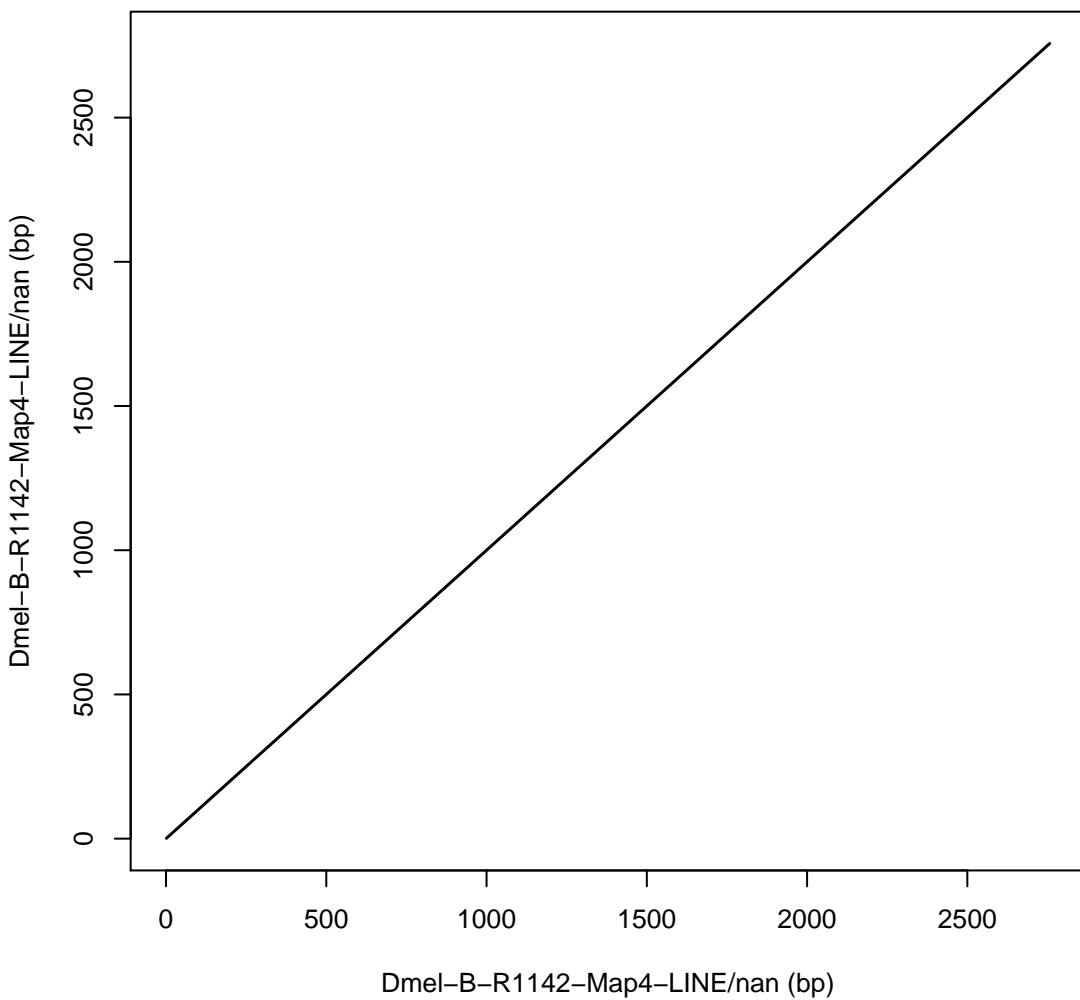
TE: Dmel-B-R1142-Map4-LINE/nan  
consensus size: 2757bp; fragments: 130; full length: 0 (>=2481.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

