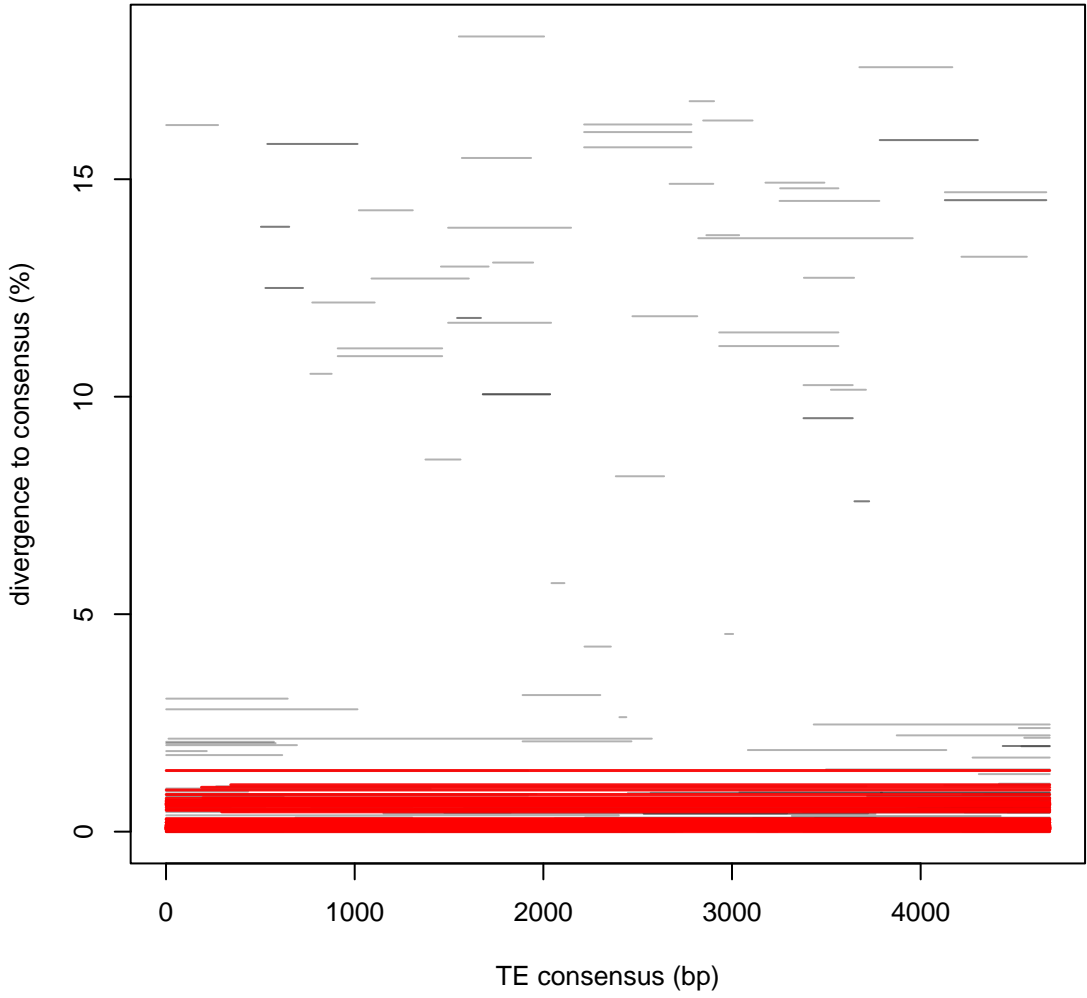
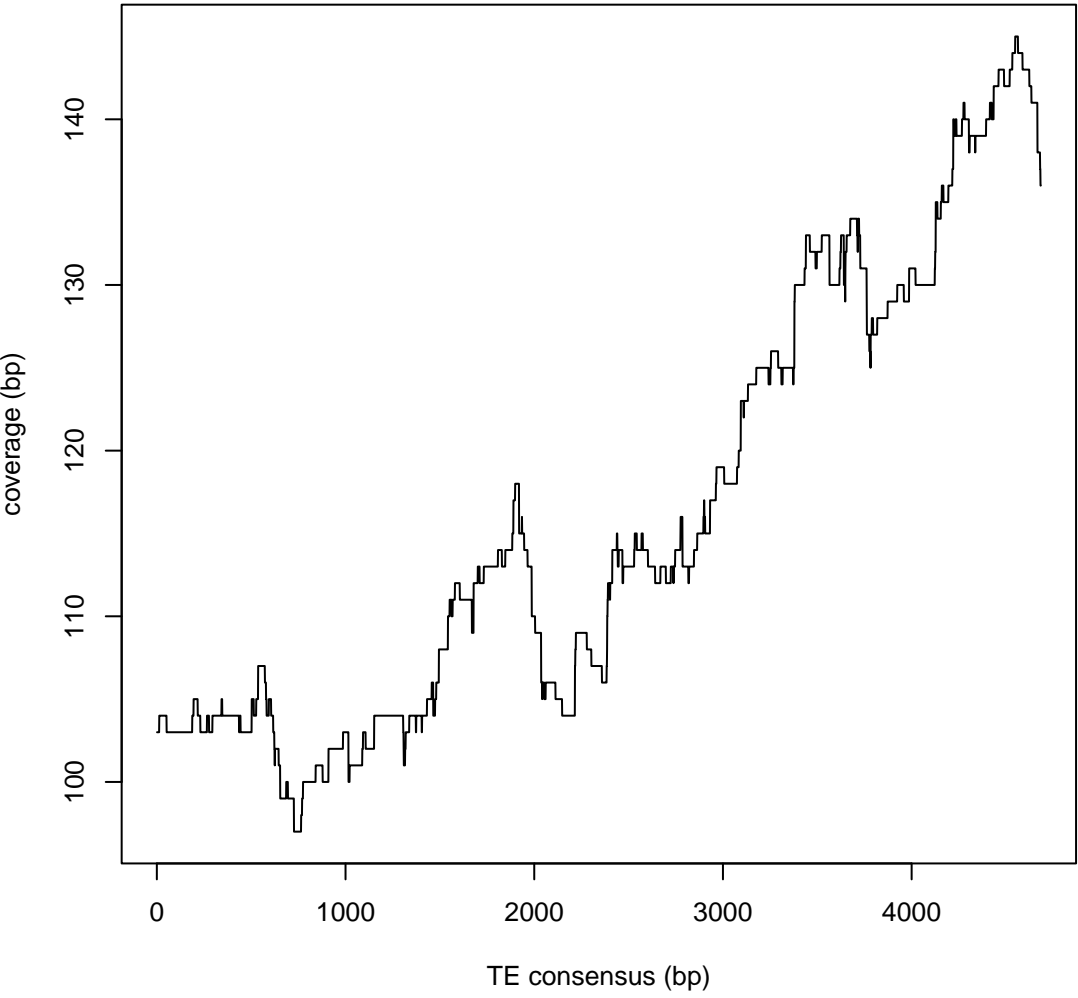


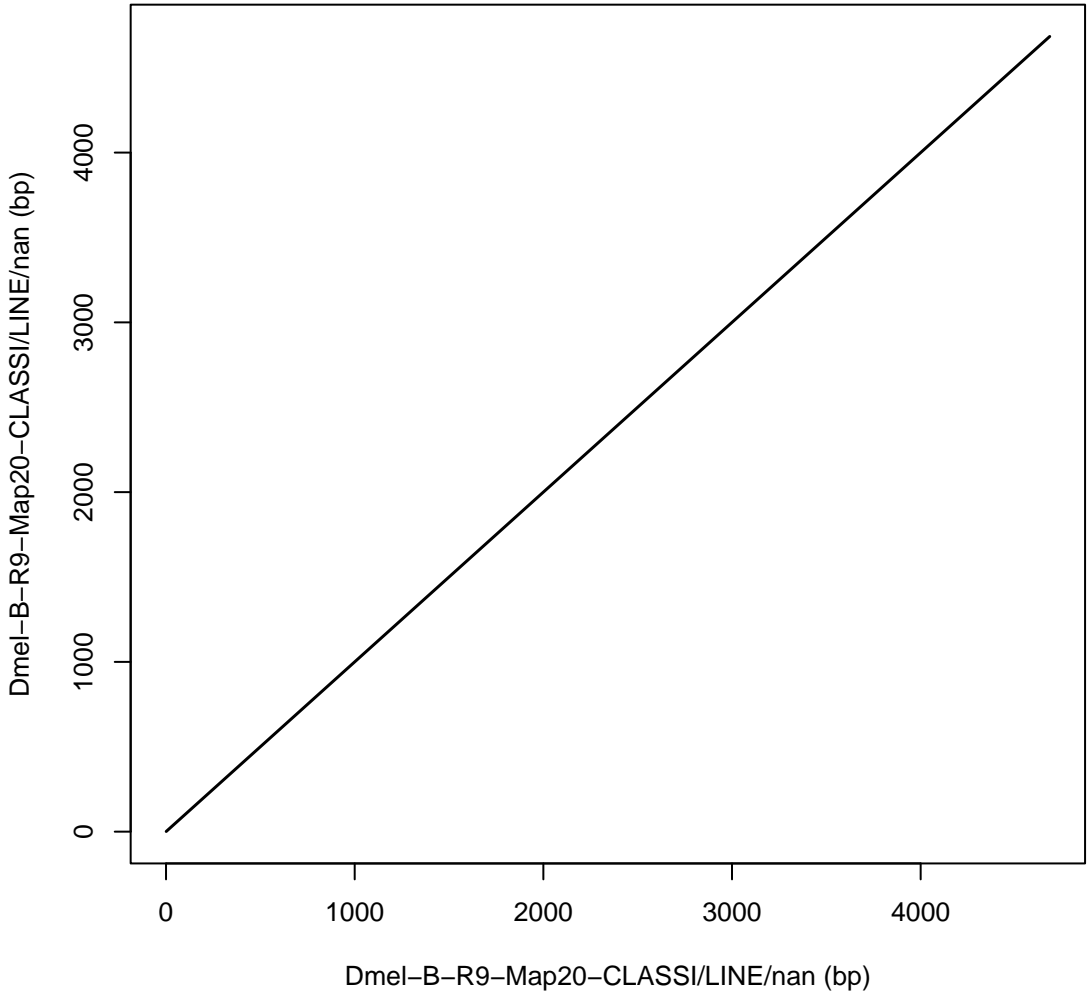
TE: Dmel-B-R9-Map20-CLASSI/LINE/nan  
consensus size: 4684bp; fragments: 262; full length: 75 (>=4215.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

