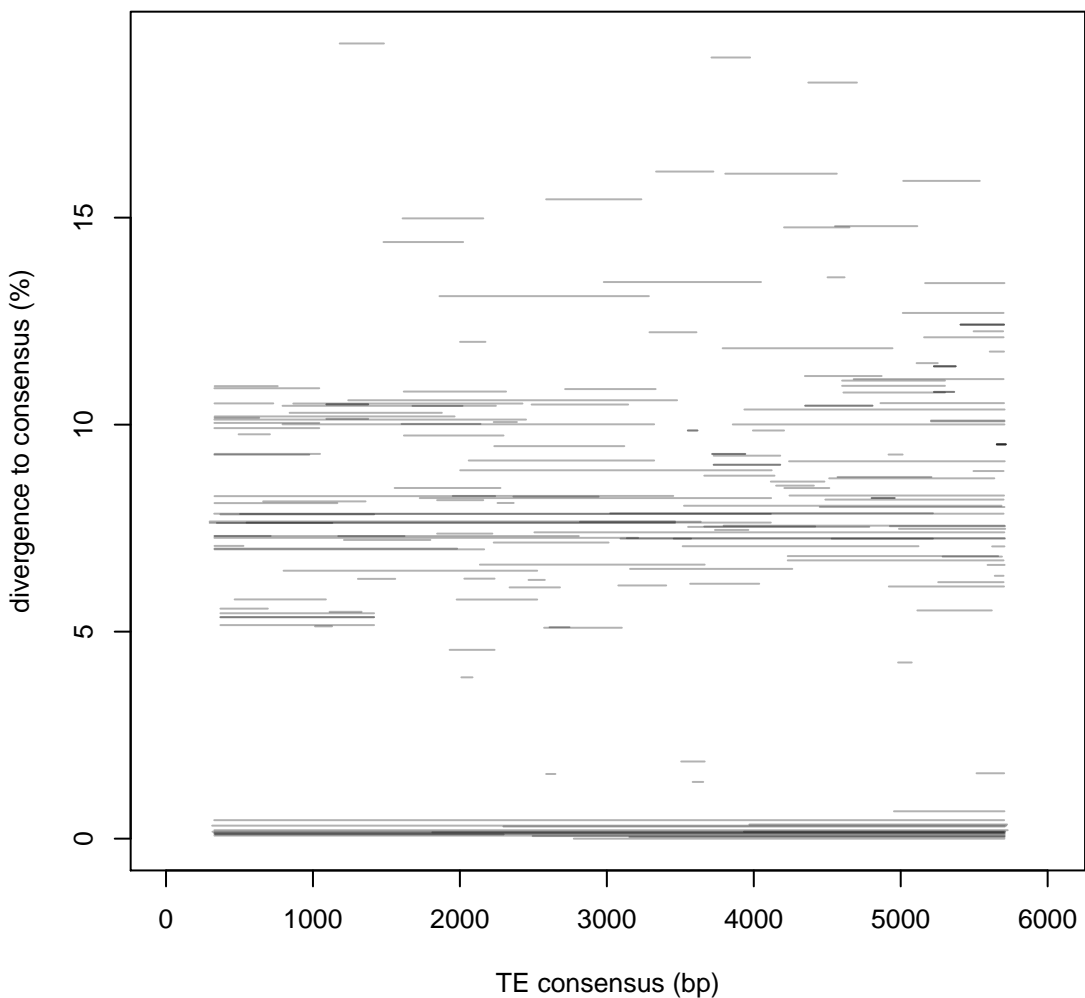
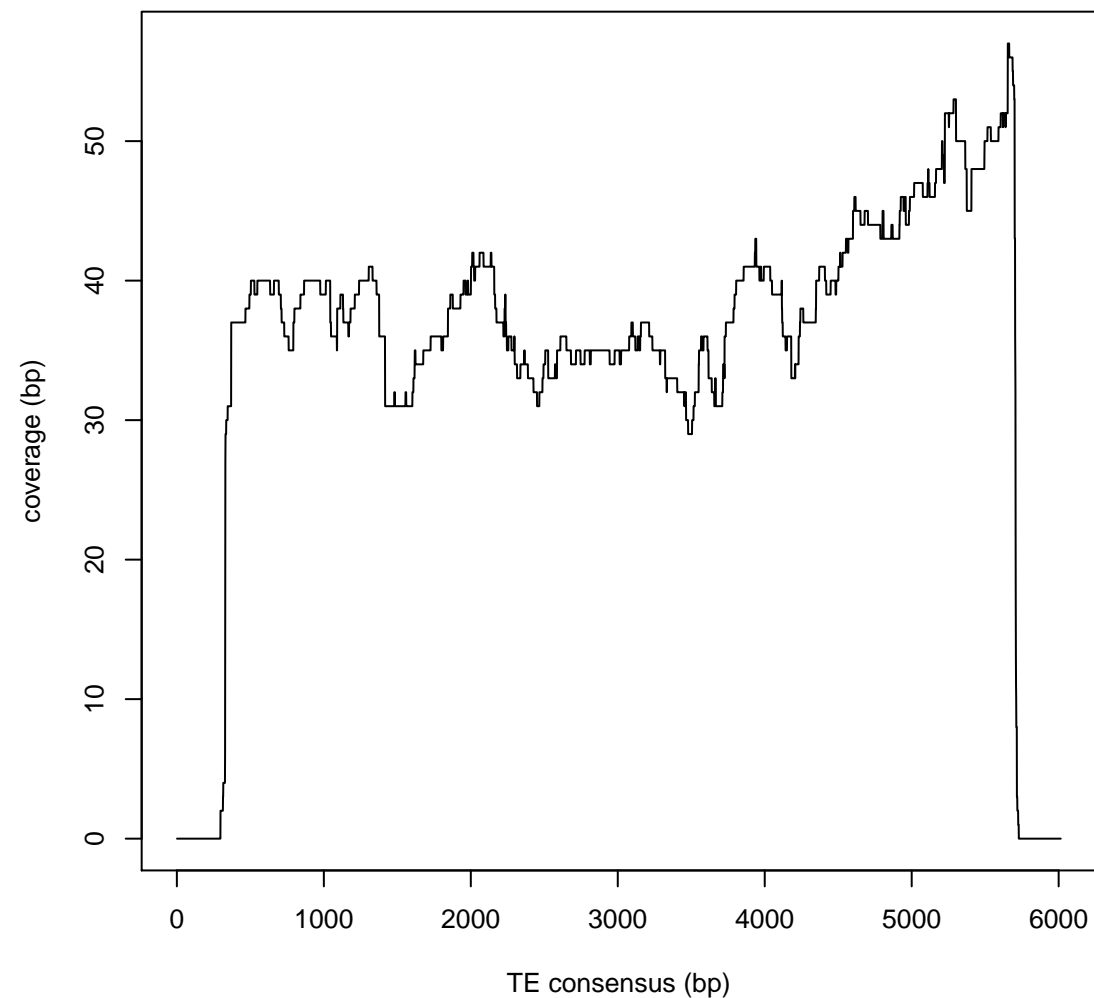


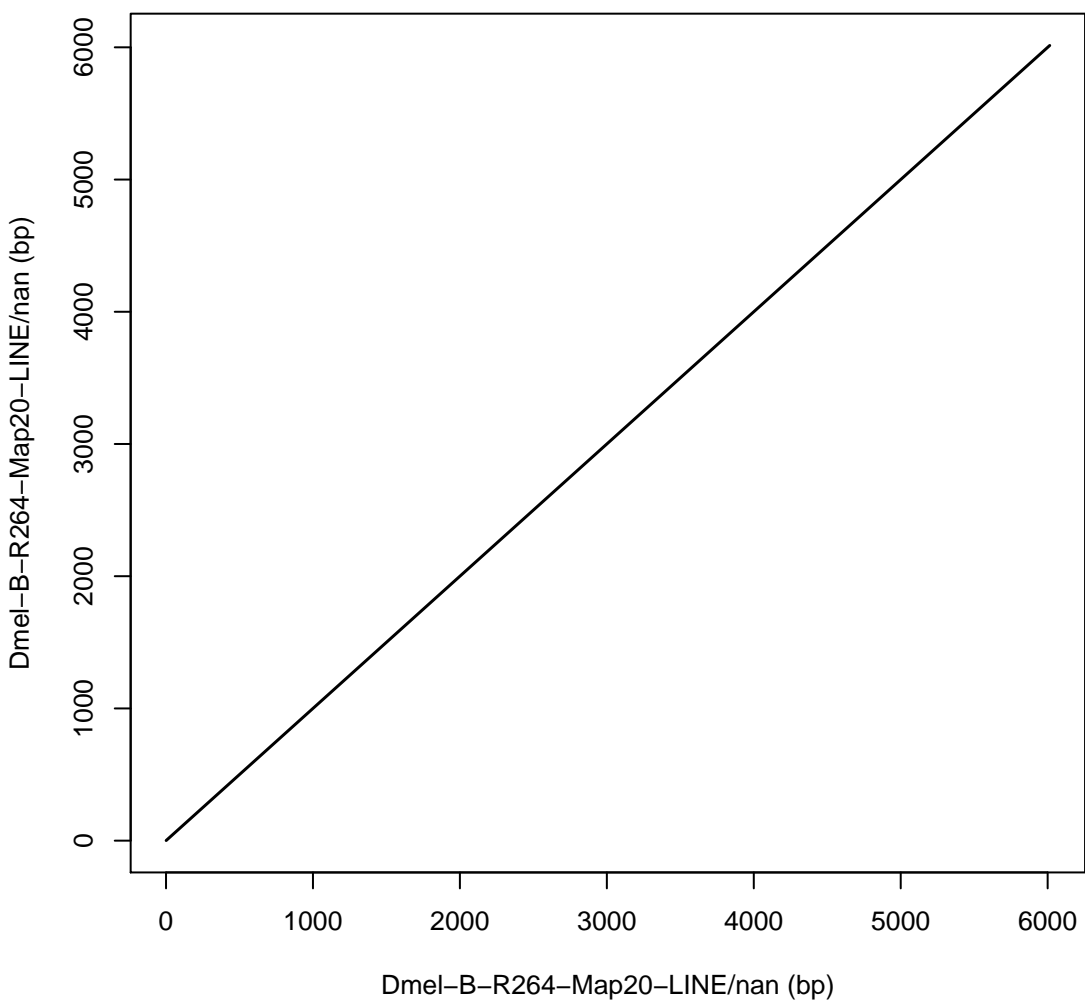
TE: Dmel-B-R264-Map20-LINE/nan  
consensus size: 6014bp; fragments: 193; full length: 0 (>=5412.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

