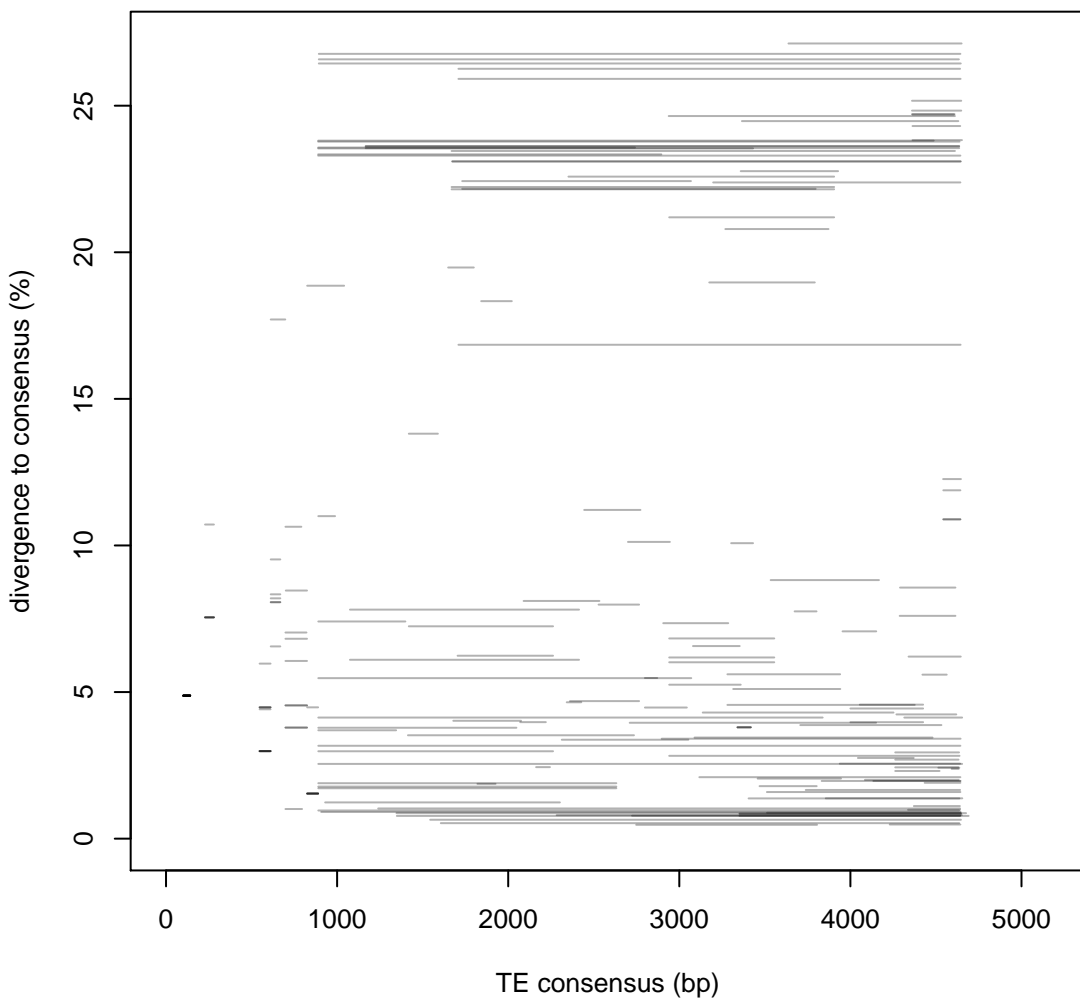
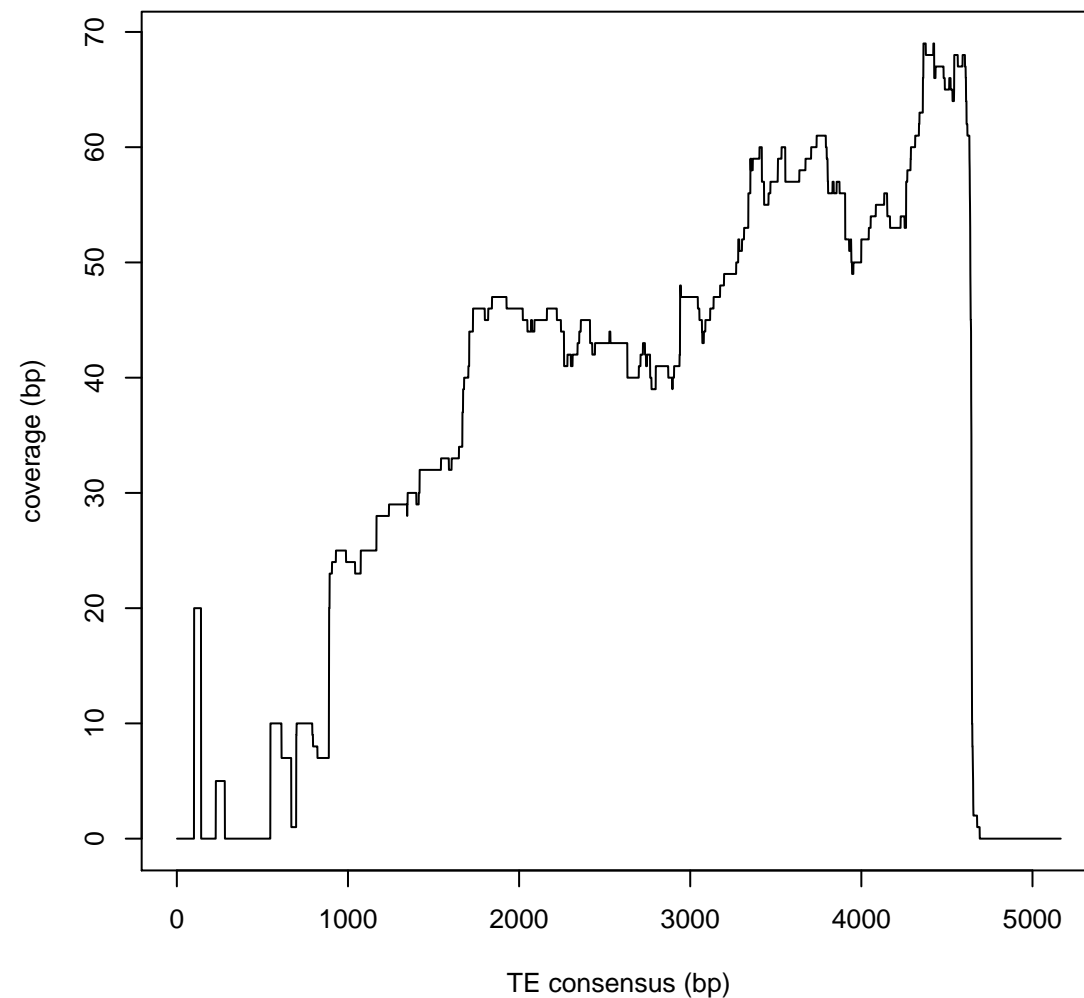


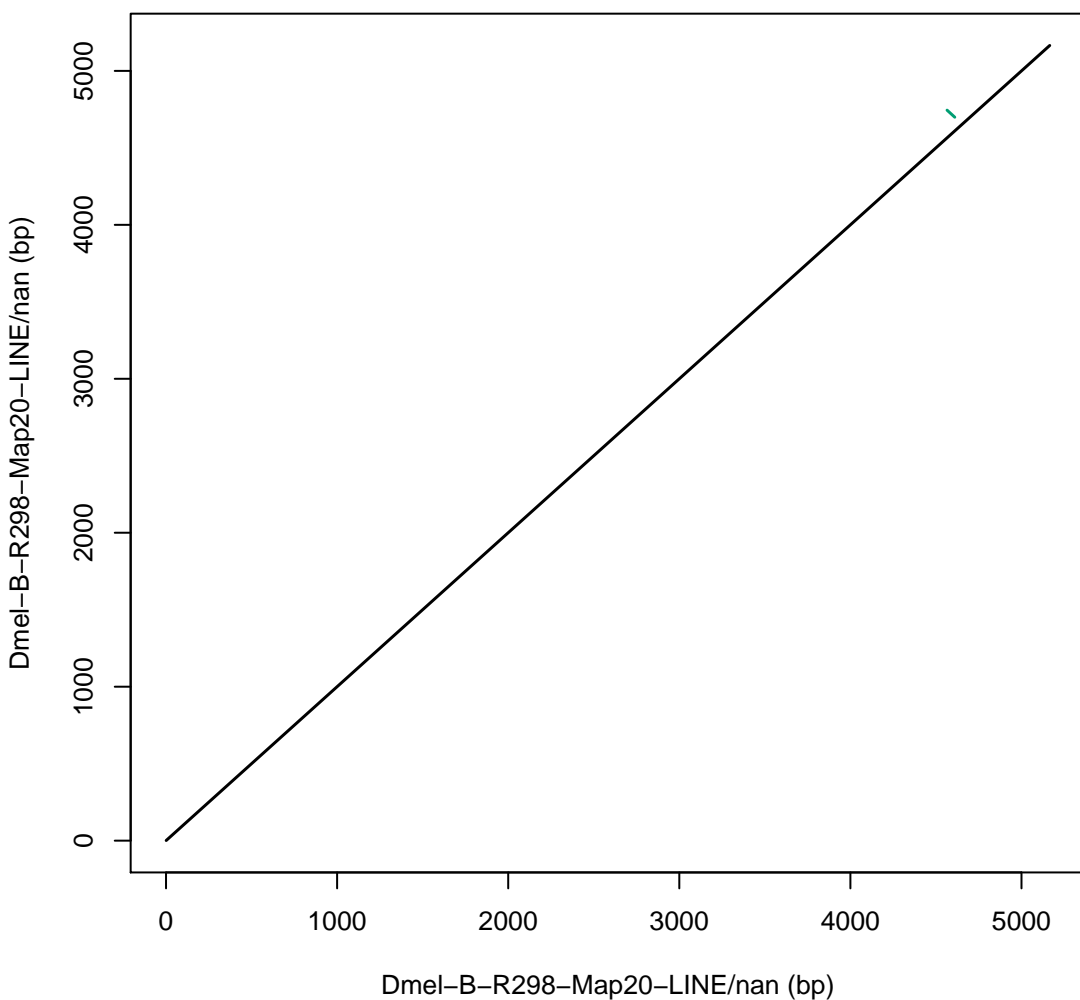
TE: Dmel-B-R298-Map20-LINE/nan  
consensus size: 5165bp; fragments: 201; full length: 0 (>=4648.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

