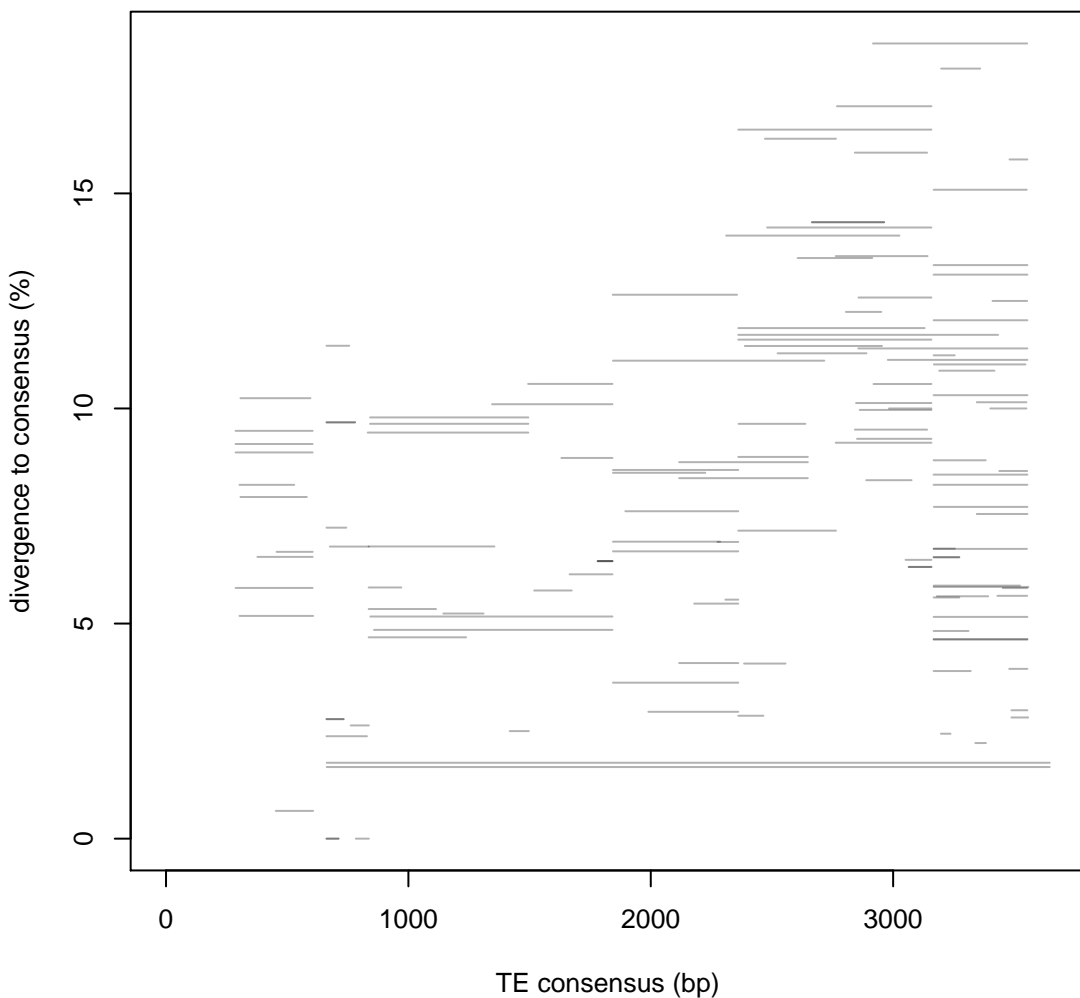
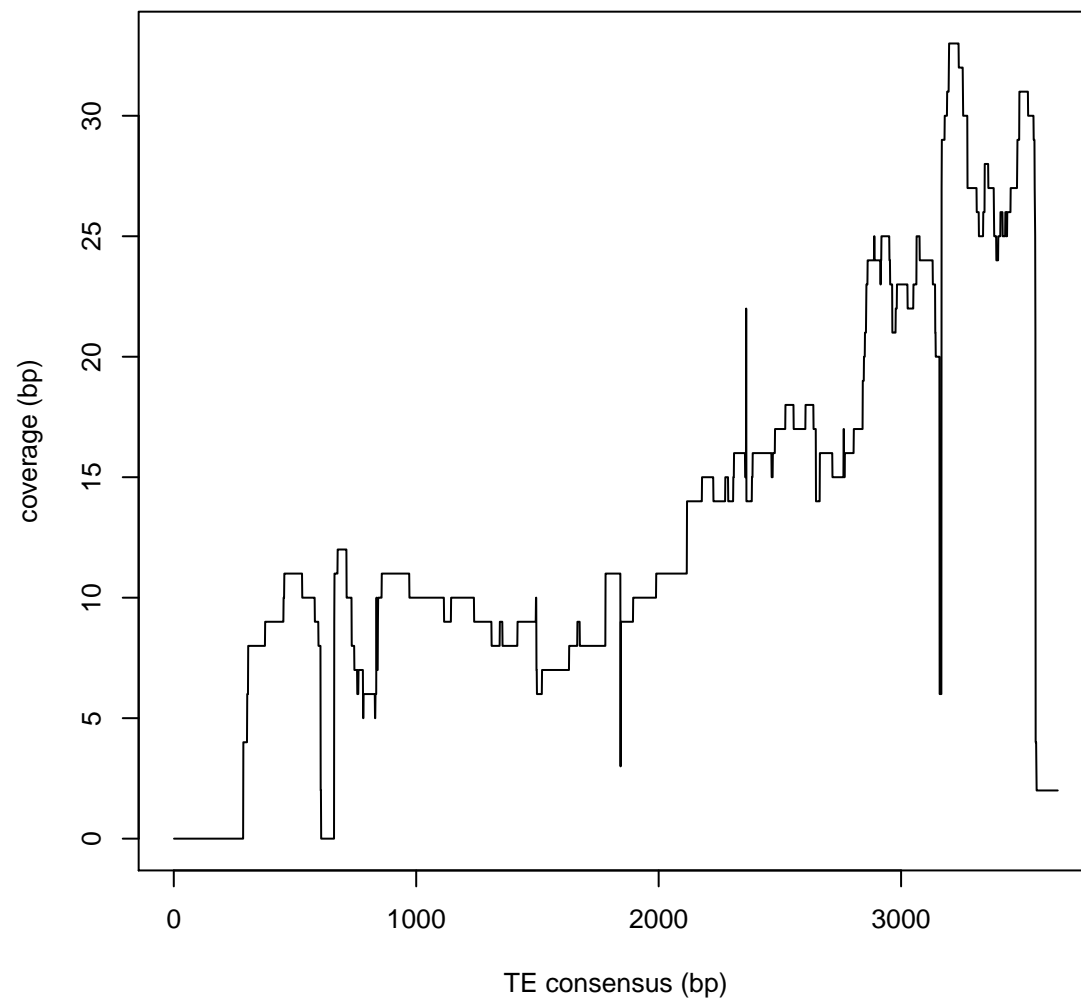


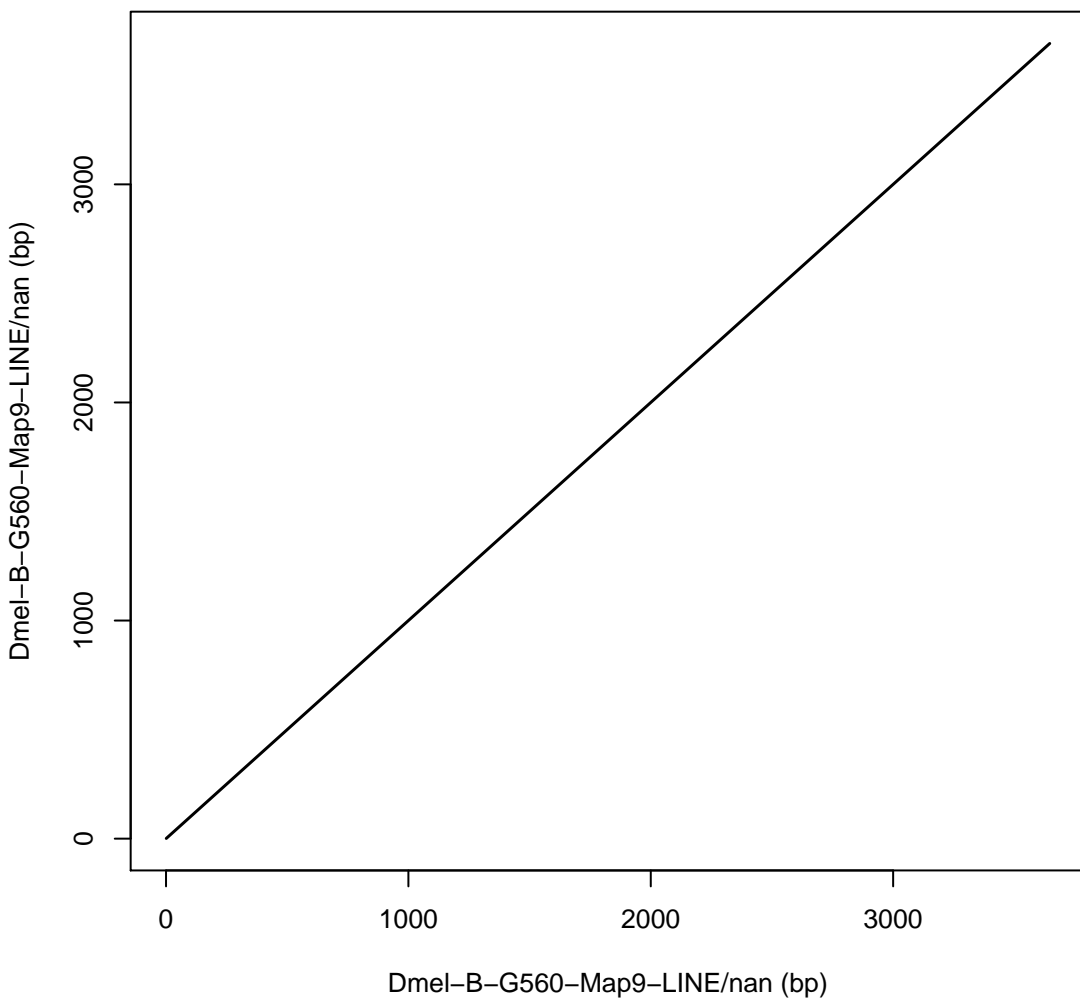
TE: Dmel-B-G560-Map9-LINE/nan  
consensus size: 3646bp; fragments: 134; full length: 0 (>=3281.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

