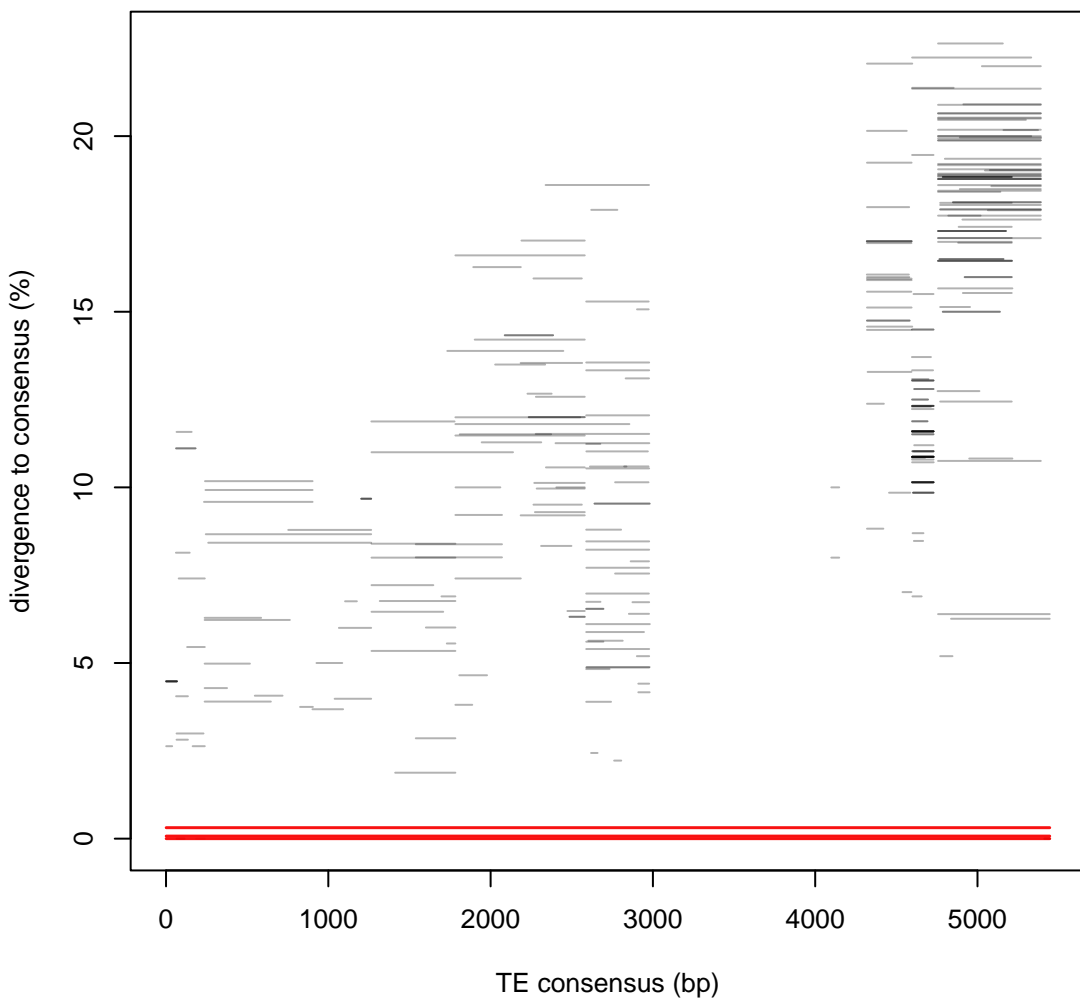
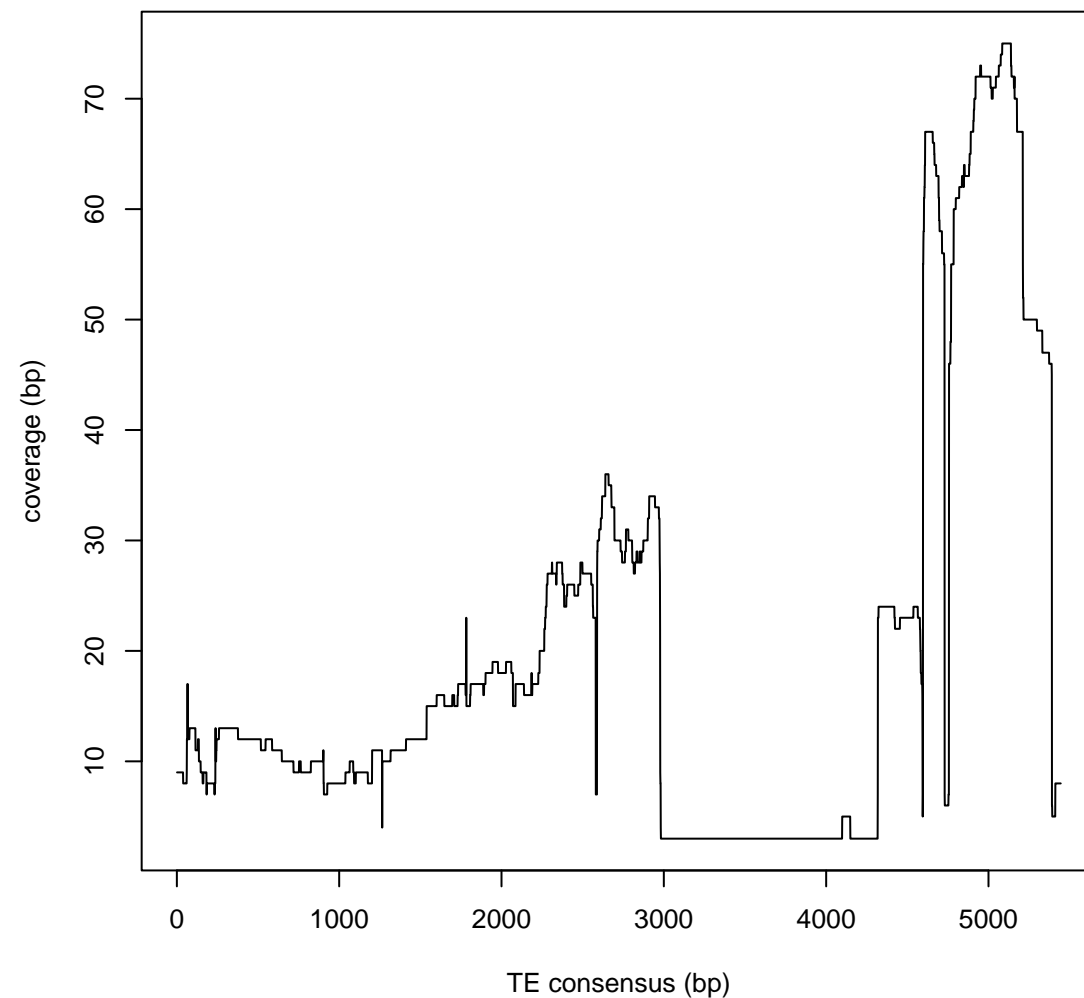


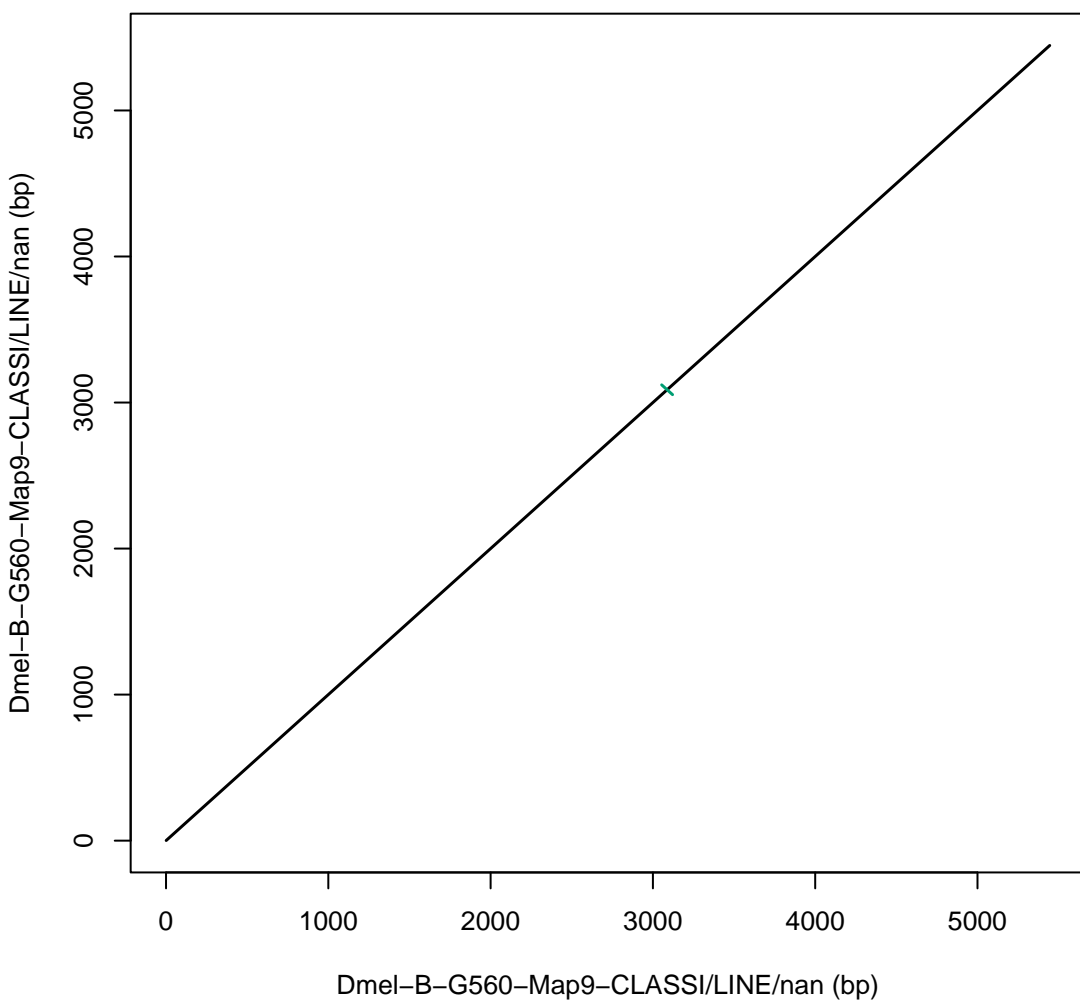
TE: Dmel-B-G560-Map9-CLASSI/LINE/nan
consensus size: 5445bp; fragments: 304; full length: 3 (>=4900.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

