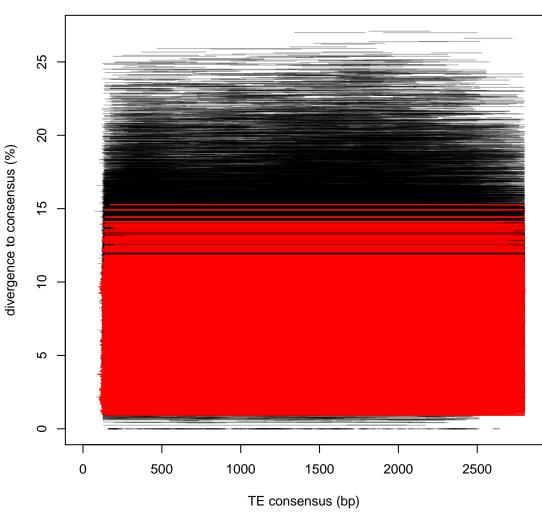
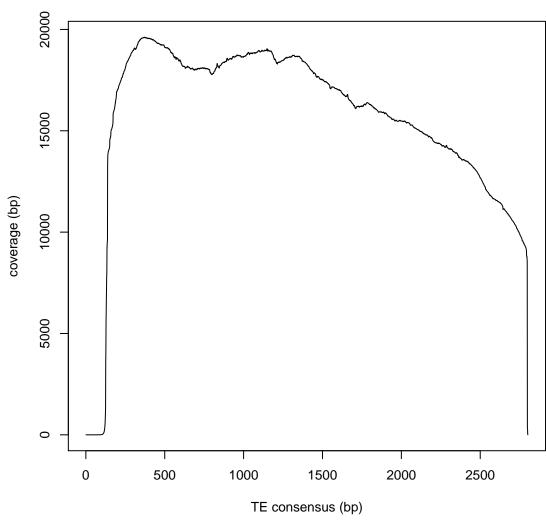
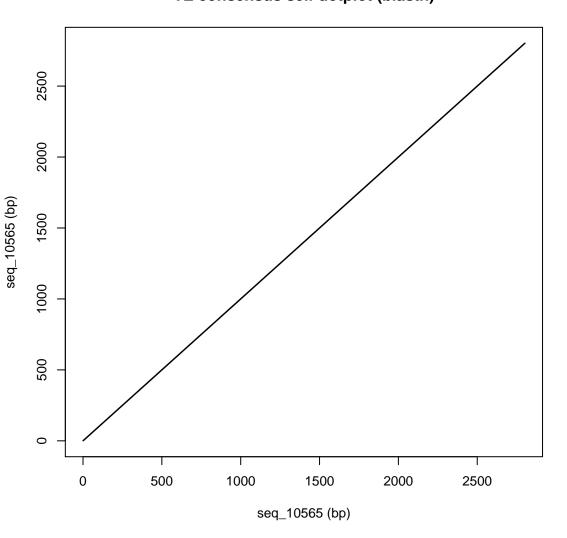
TE: seq_10565 consensus size: 2802bp; fragments: 53758; full length: 3280 (>=2521.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

