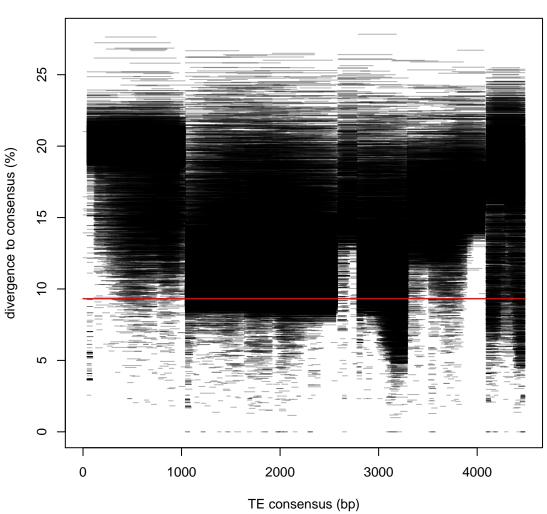
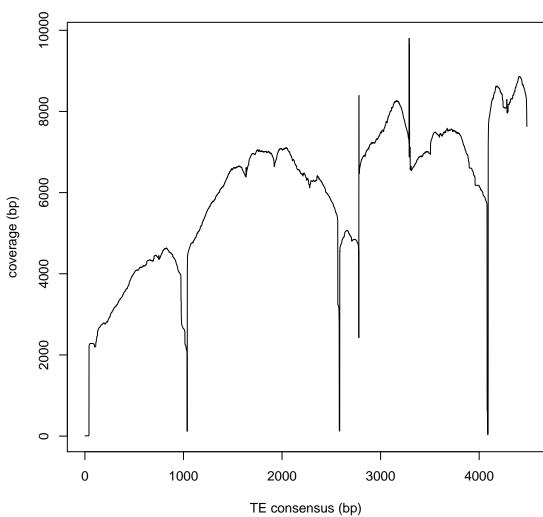
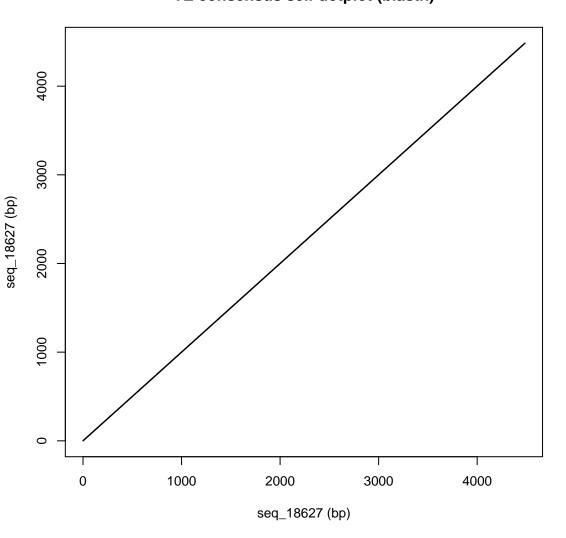
TE: seq_18627 consensus size: 4484bp; fragments: 54609; full length: 1 (>=4035.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

