TE: rnd-1_family-4-TIR/P TE consensus genomic coverage consensus size: 7752bp; fragments: 6909; full length: 7 (>=6976.8bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits rnd-1_family-4-TIR/P (bp) trofit4 LTR/Copia Retrofit4 LTR/Copia Retrofit4 LTR/Copia rnd-1_family-4-TIR/P (bp) rnd-1_family-4-TIR/P (bp)