



Figure S8 Duplicated genomic regions associated with the NRPSs PTRG_11818 & 11836. Putative proteins are designated as arrows. Proteins with similar function are coded with similar color. AT - acyltransferase; MT - methyltransferase. Protein coding regions and spacing between are to scale. Numbers above boxes indicate locus position in the reference. Numbers beneath boxes represent the % identity of the proteins relative to PTRG_11817 through 11822. Triangles represent transposons or transposon remnants.