

Supplementary Materials: Acontia, a Specialised Defensive Structure, Has Low Venom Complexity in *Calliactis polypus*

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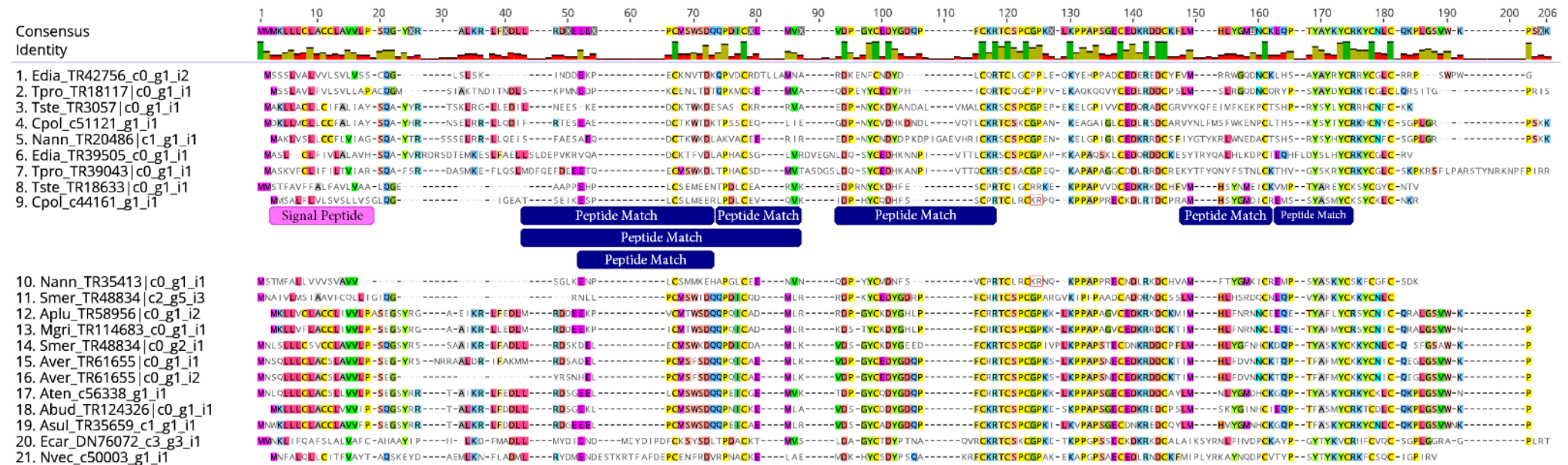


Figure S1. Alignment of amino acid sequences for the Unknown 12C candidate genes and the consensus sequence and identity percentage above. Annotation of the candidate identified in the *Calliactis polypus* acontia proteome is depicted with the signal peptide and peptide matches from the mass spectra of the SWATH-MS and DDA pooled acontia sample. Consensus identity percentages are as follows, green bar for 100 %, yellow bar for ≥ 50 % and < 100 %, and red bar for < 50 %.