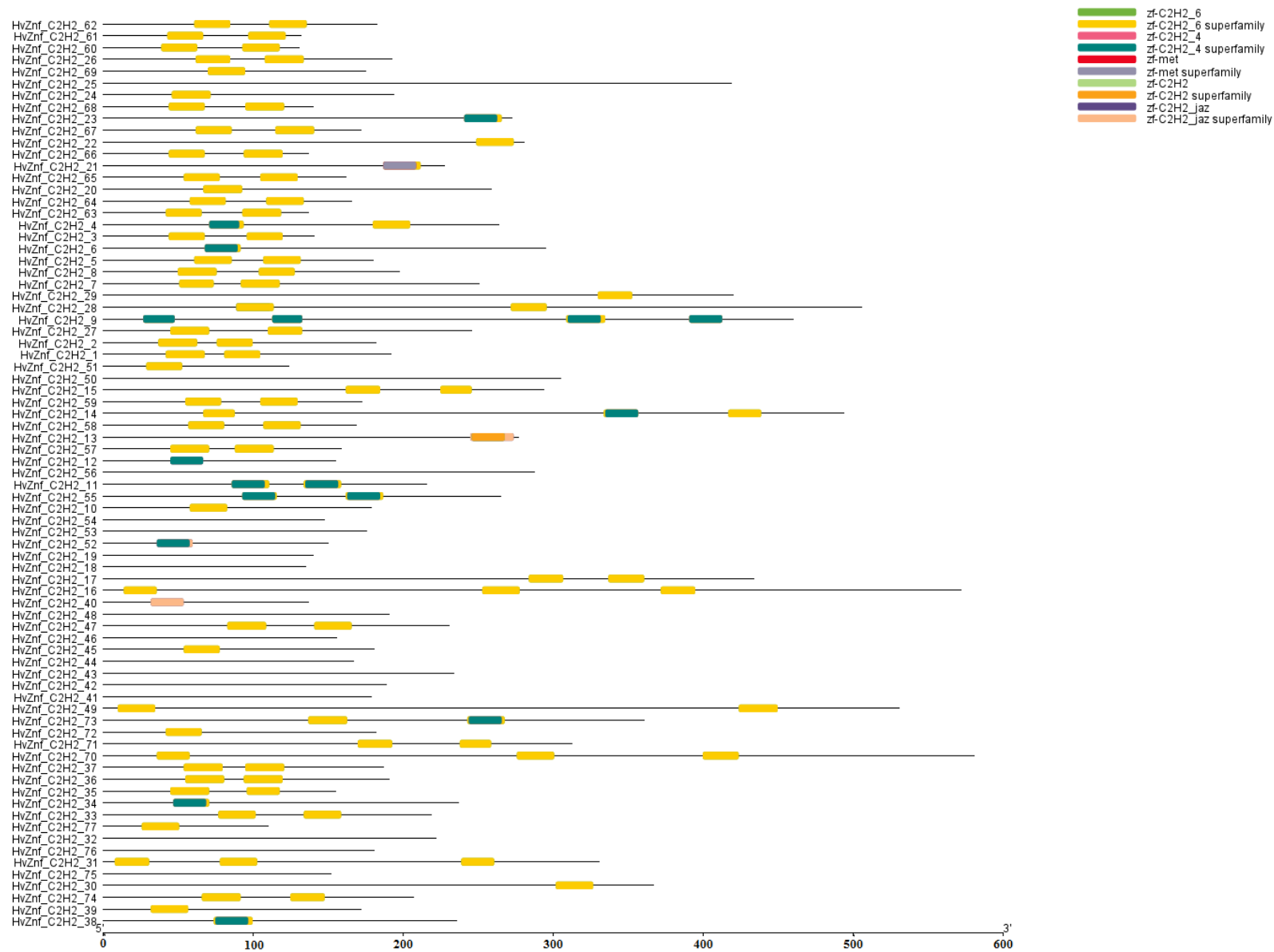
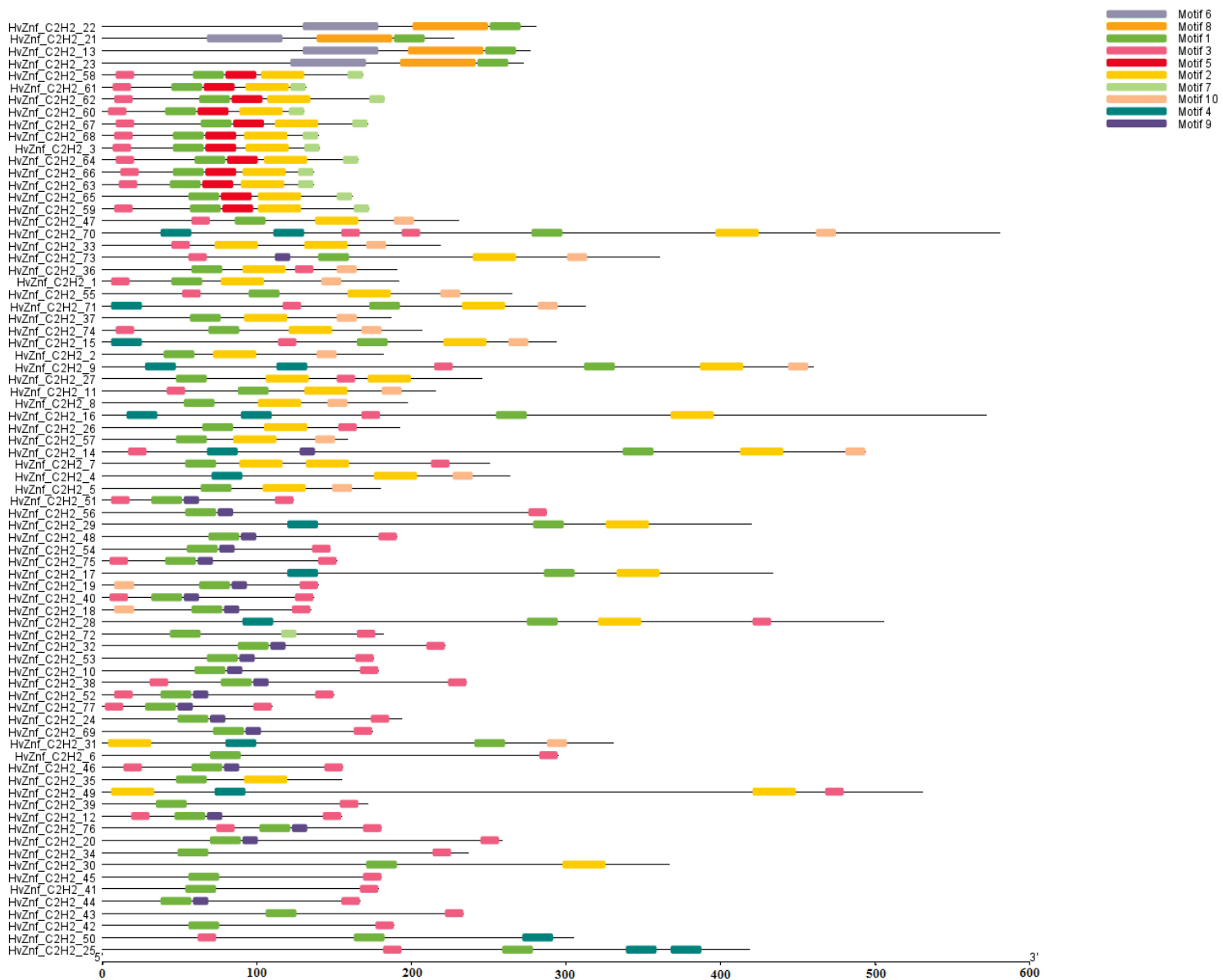
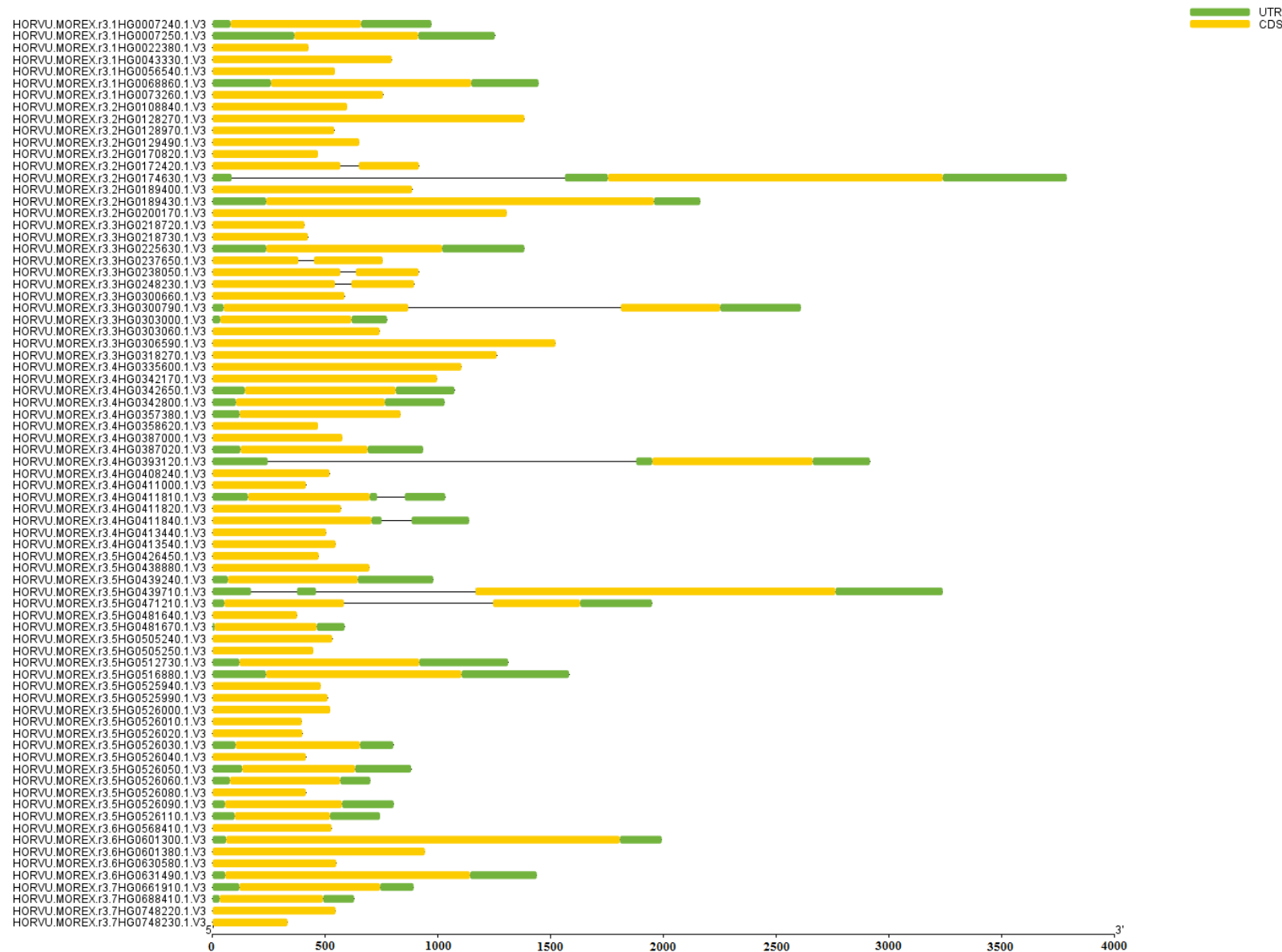




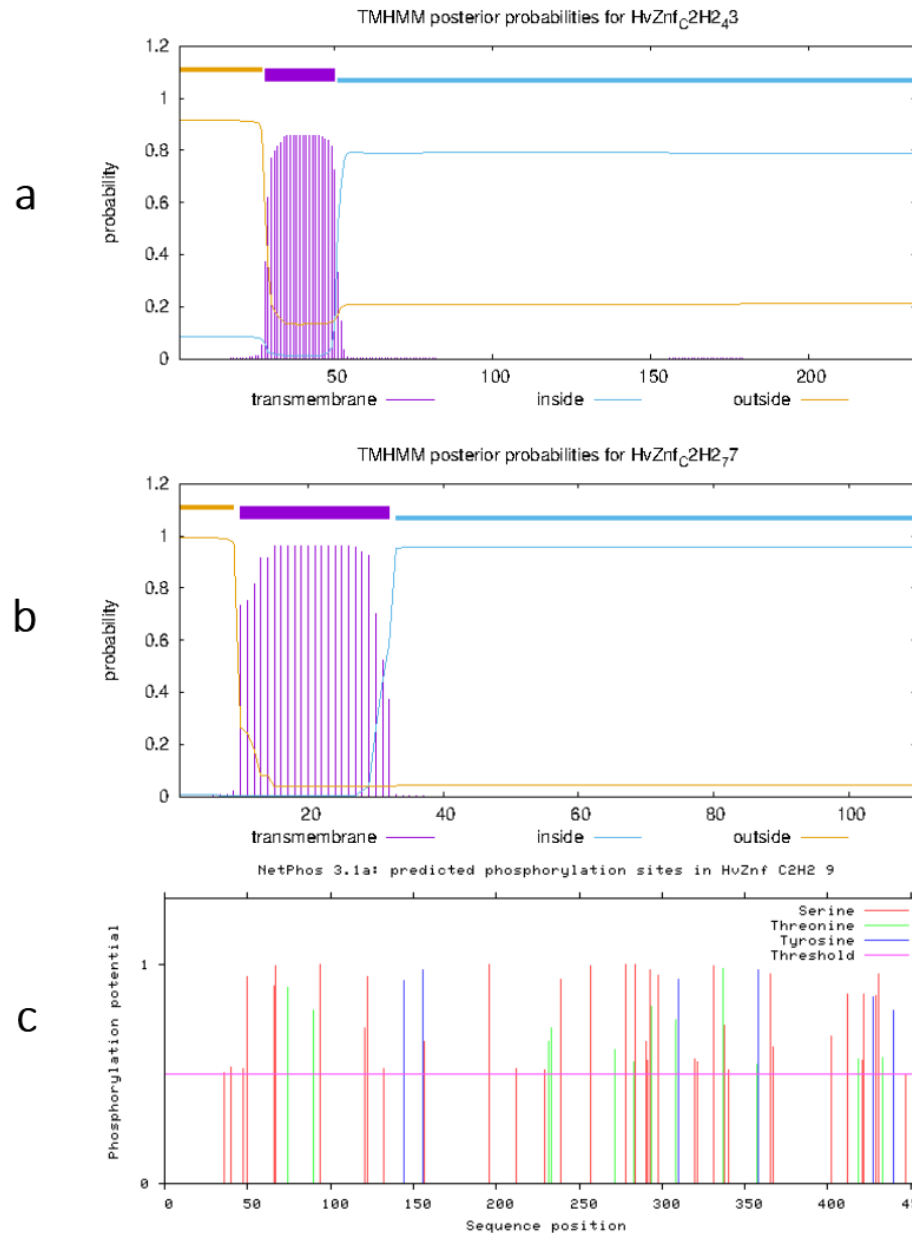
Domain of HvZnf_C2H2 proteins



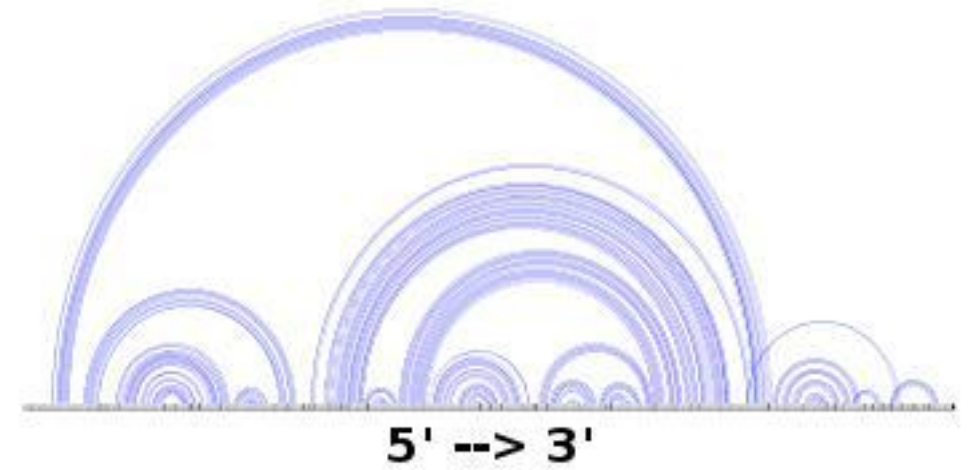
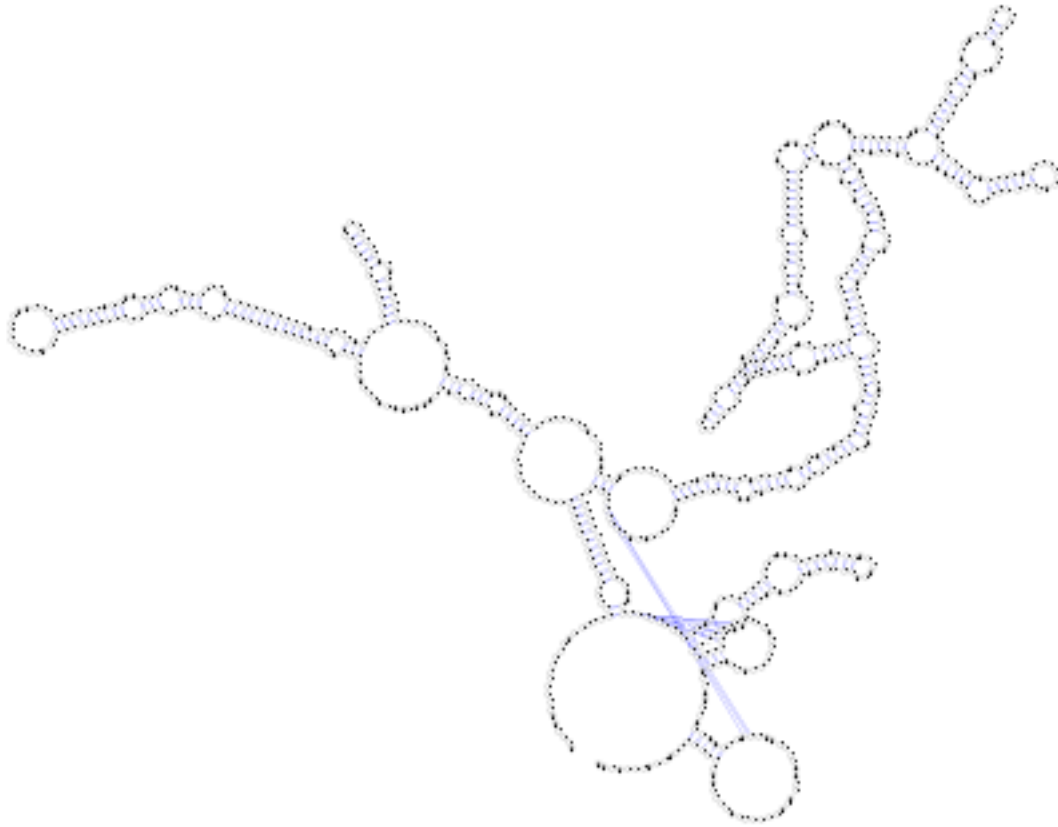
Motif of HvZnf_C2H2 proteins



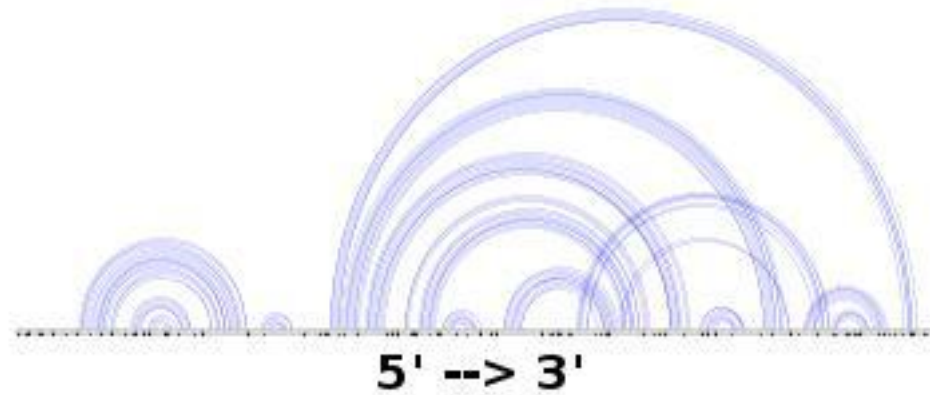
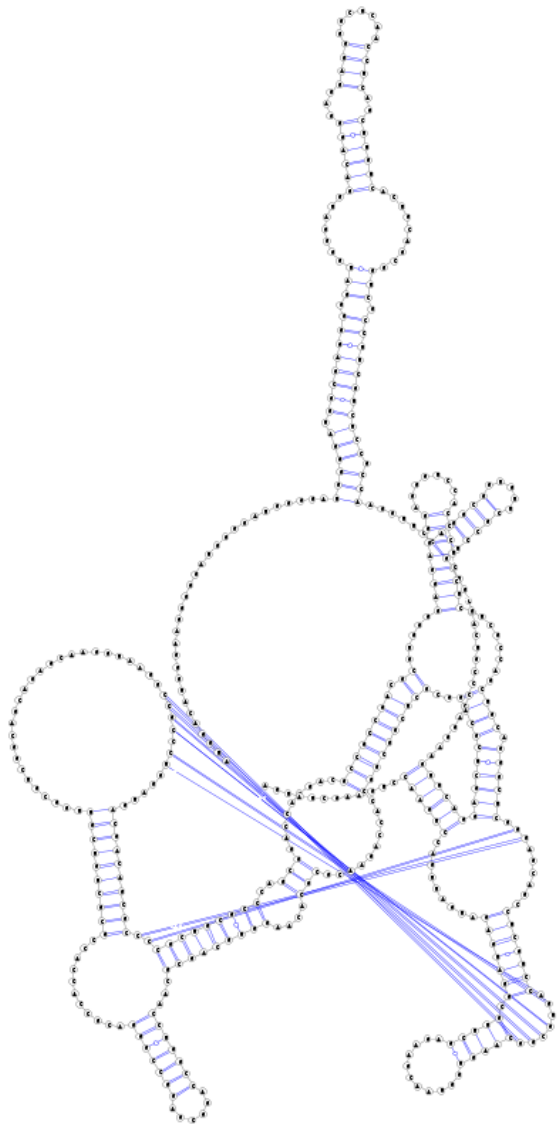
Gene structure of HvZnf_C2H2 proteins



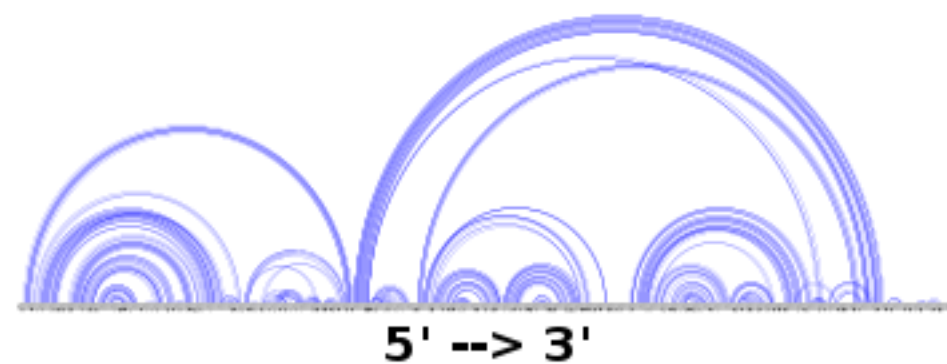
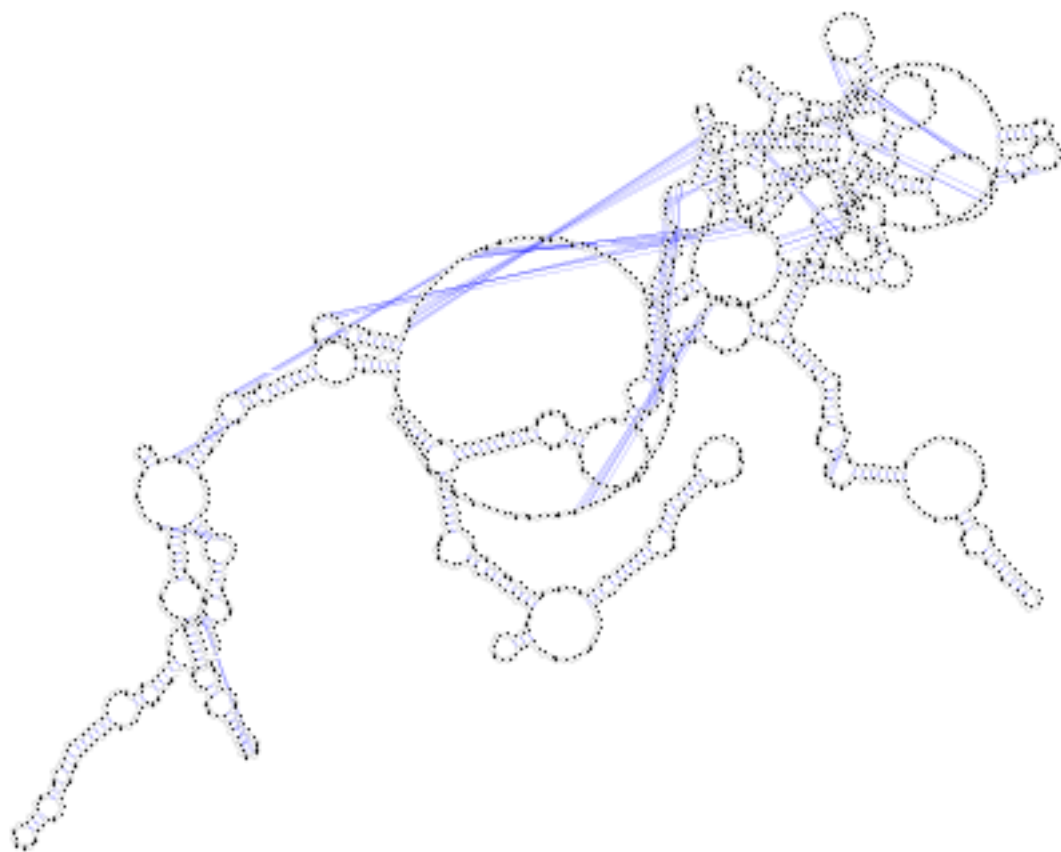
a. the transmembrane helical in HvZnf_C2H2_43 protein. b. the transmembrane helical in HvZnf_C2H2_77 proteins. c. HvZnf_C2H2_9 protein putative phosphorylation with serine, threonine and tyrosine.



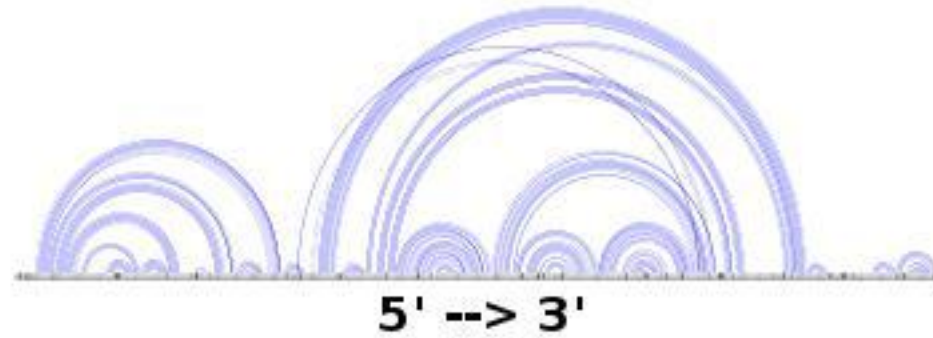
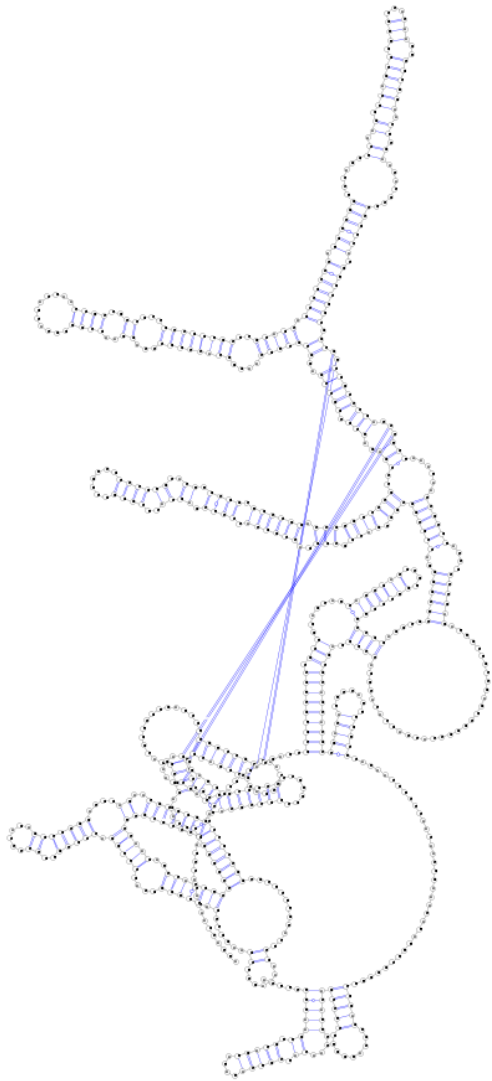
RNA secondary structures with pseudoknots for *HvZnf_C2H2_4*



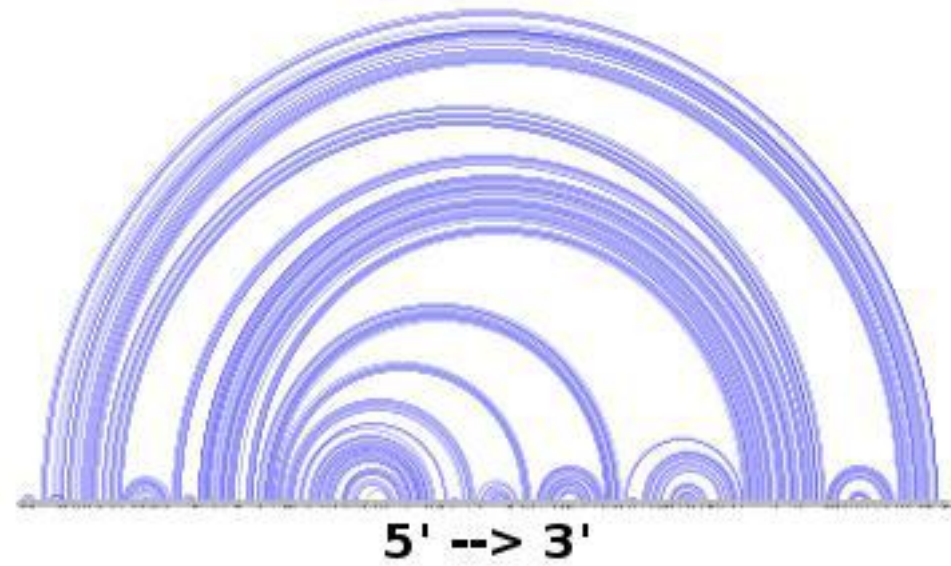
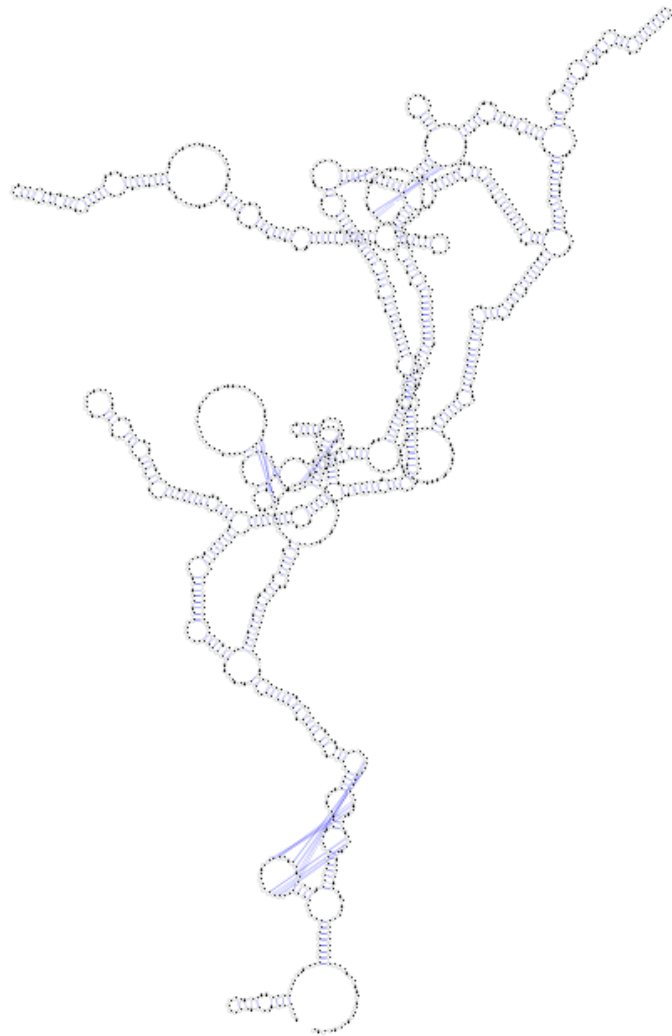
RNA secondary structures with pseudoknots for *HvZnf_C2H2_12*



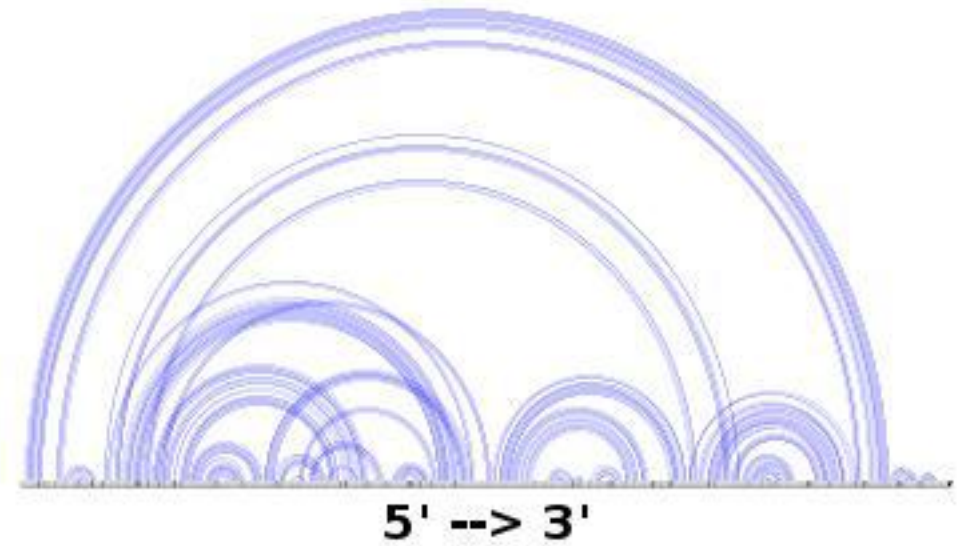
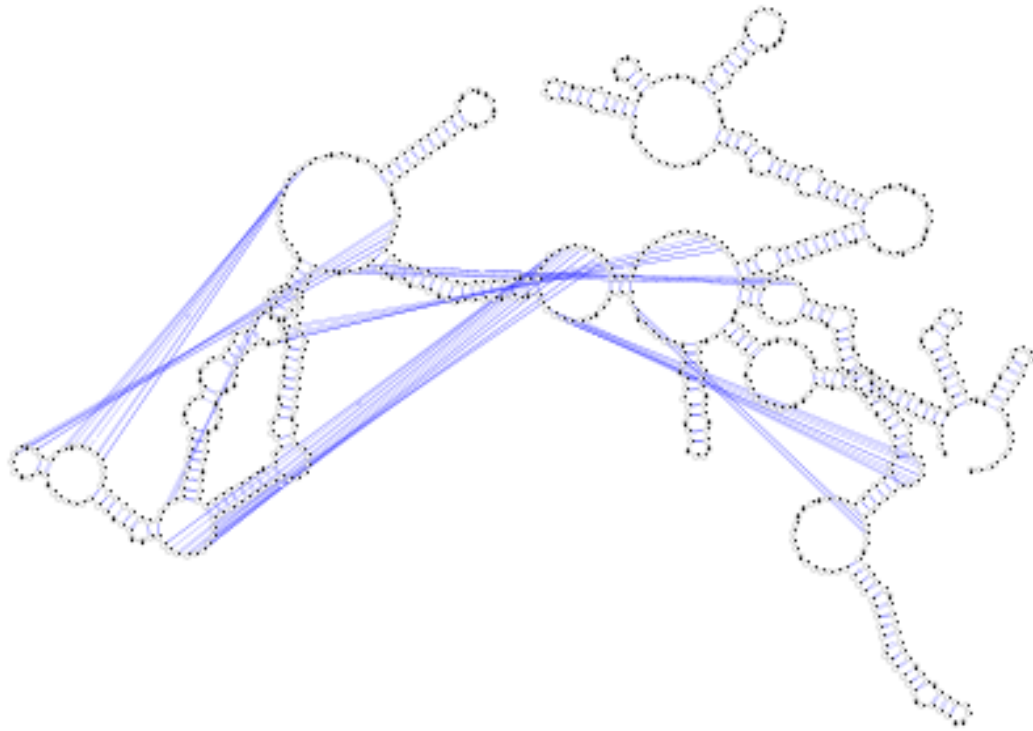
RNA secondary structures with pseudoknots for *HvZnf_C2H2_14*



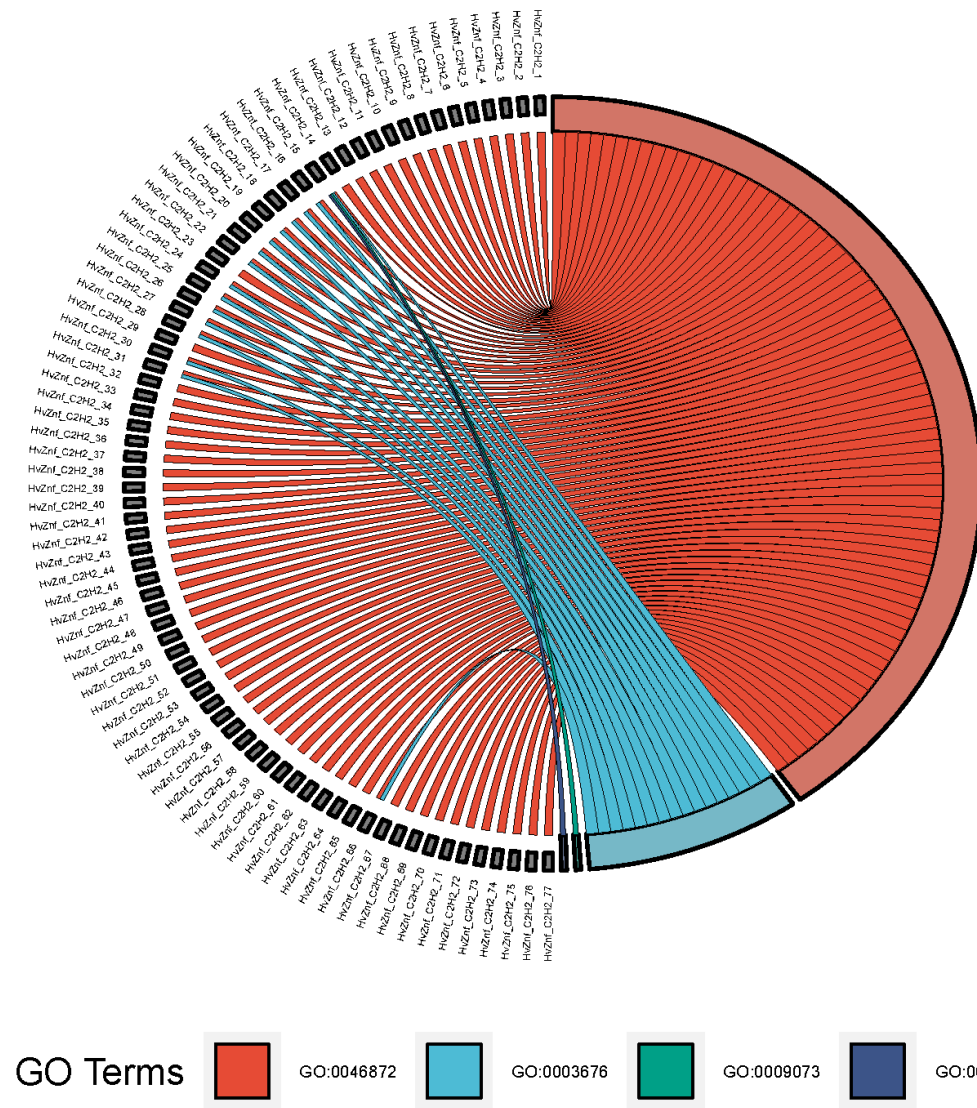
RNA secondary structures with pseudoknots for *HvZnf_C2H2_27*



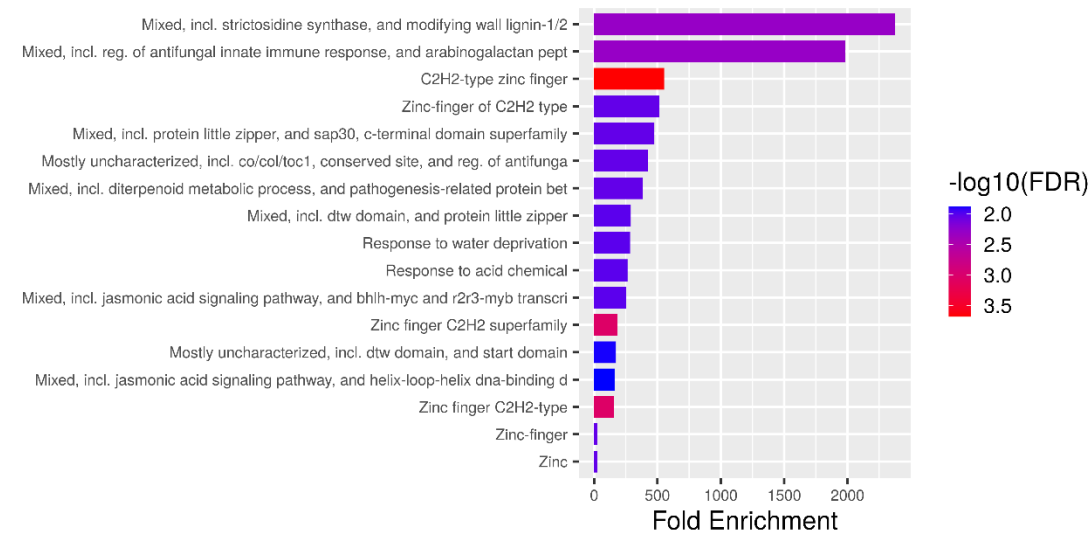
RNA secondary structures with pseudoknots for *HvZnf_C2H2_28*



RNA secondary structures with pseudoknots for *HvZnf_C2H2_55*



Gene ontology terms, GO:
 0003676 (nucleic acid binding),
 GO: 0004107 (chorismate
 synthase activity), GO: 0009073
 (aromatic amino acid family
 biosynthetic process) and GO:
 0046872 (metal ion binding).



Gene ontology enrichment analysis were confirmed the functional role of HvZnf_C2H2 as a stress responsive