Supplementary Materials: Transcriptomic Analysis of the Spider Venom Gland Reveals Venom Diversity and Species Consanguinity

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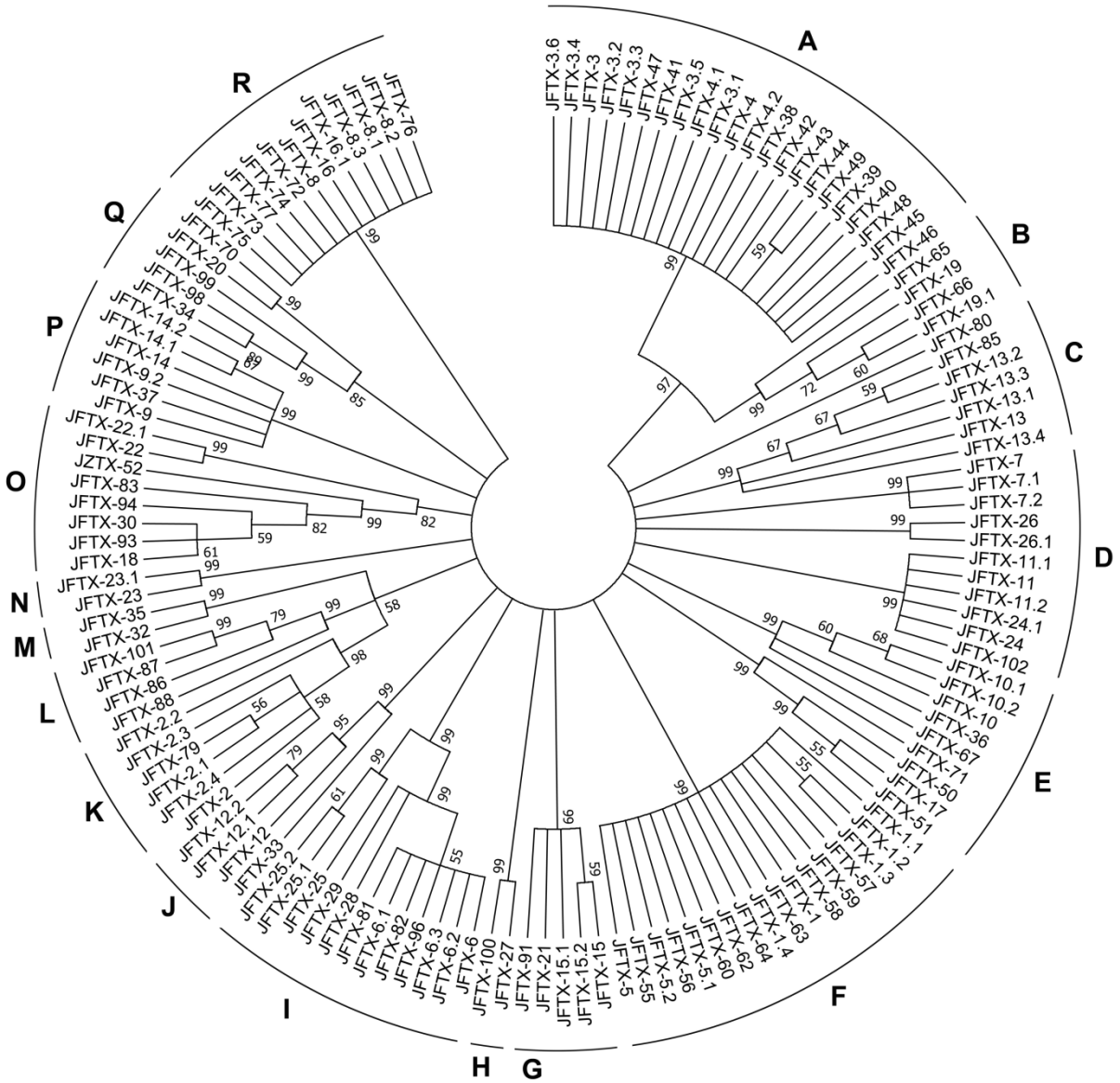

Figure 1. Phylogenetic tree of putative toxin precursors from *S. jiafu* venom glands. The phylogenetic analysis was conducted by using the neighbor-joining method of the MEGA 7 software package.