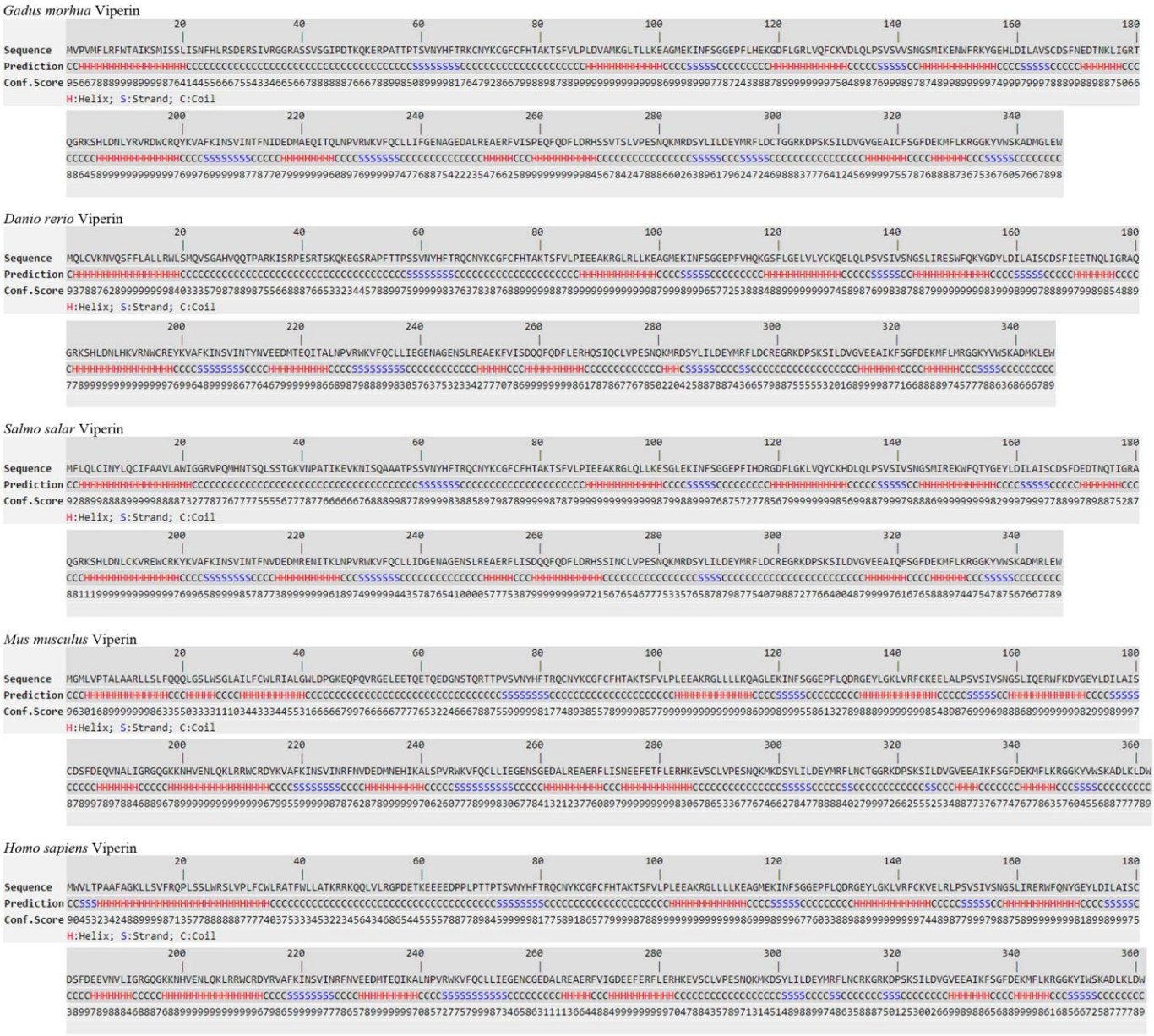

Supplementary Figure S1: Predicted secondary structure of orthologous Viperins from several species using I-TASSER website. Confidence score values range from 0 (the lowest confidence) to 9 (the highest confidence).