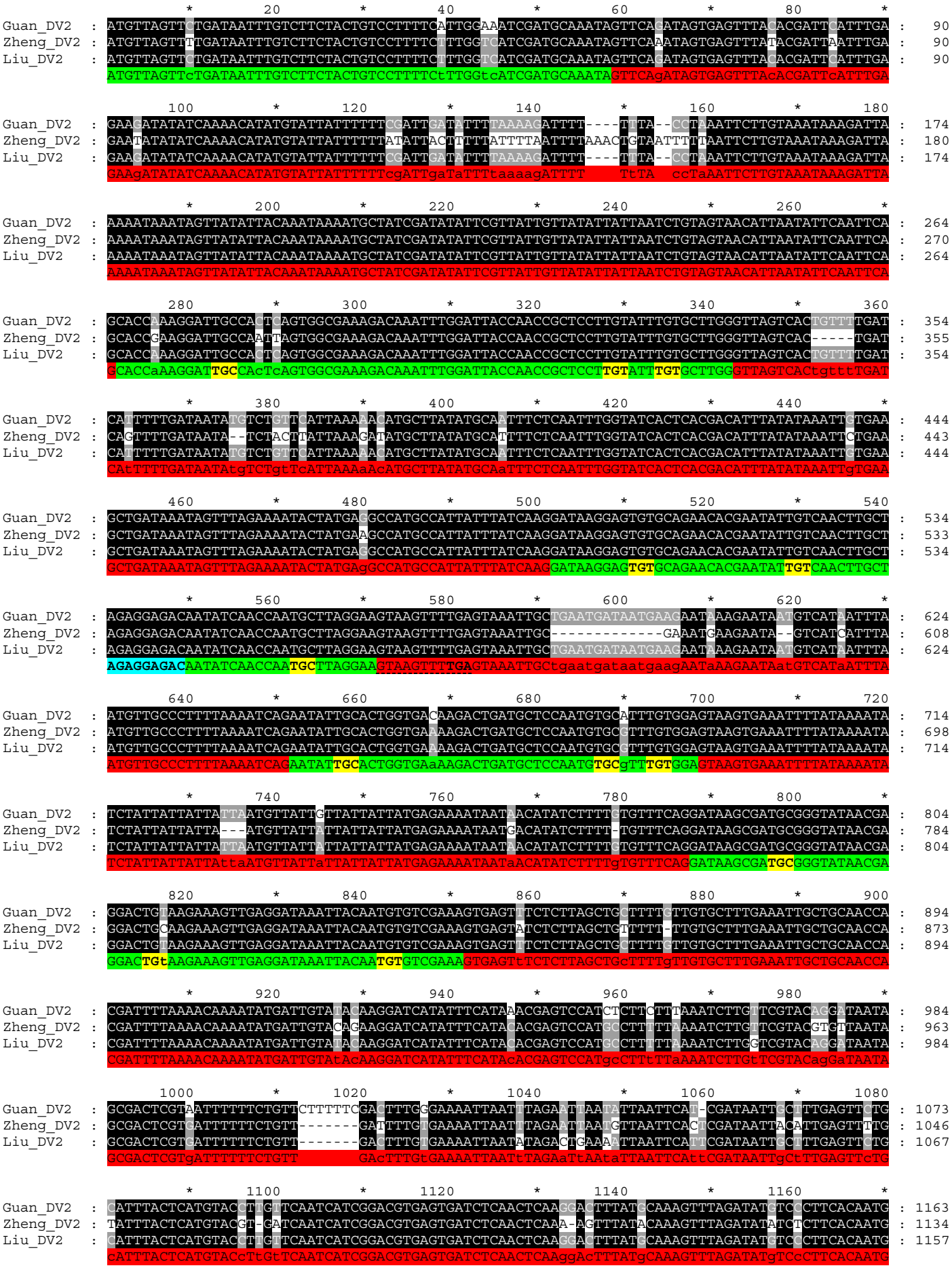
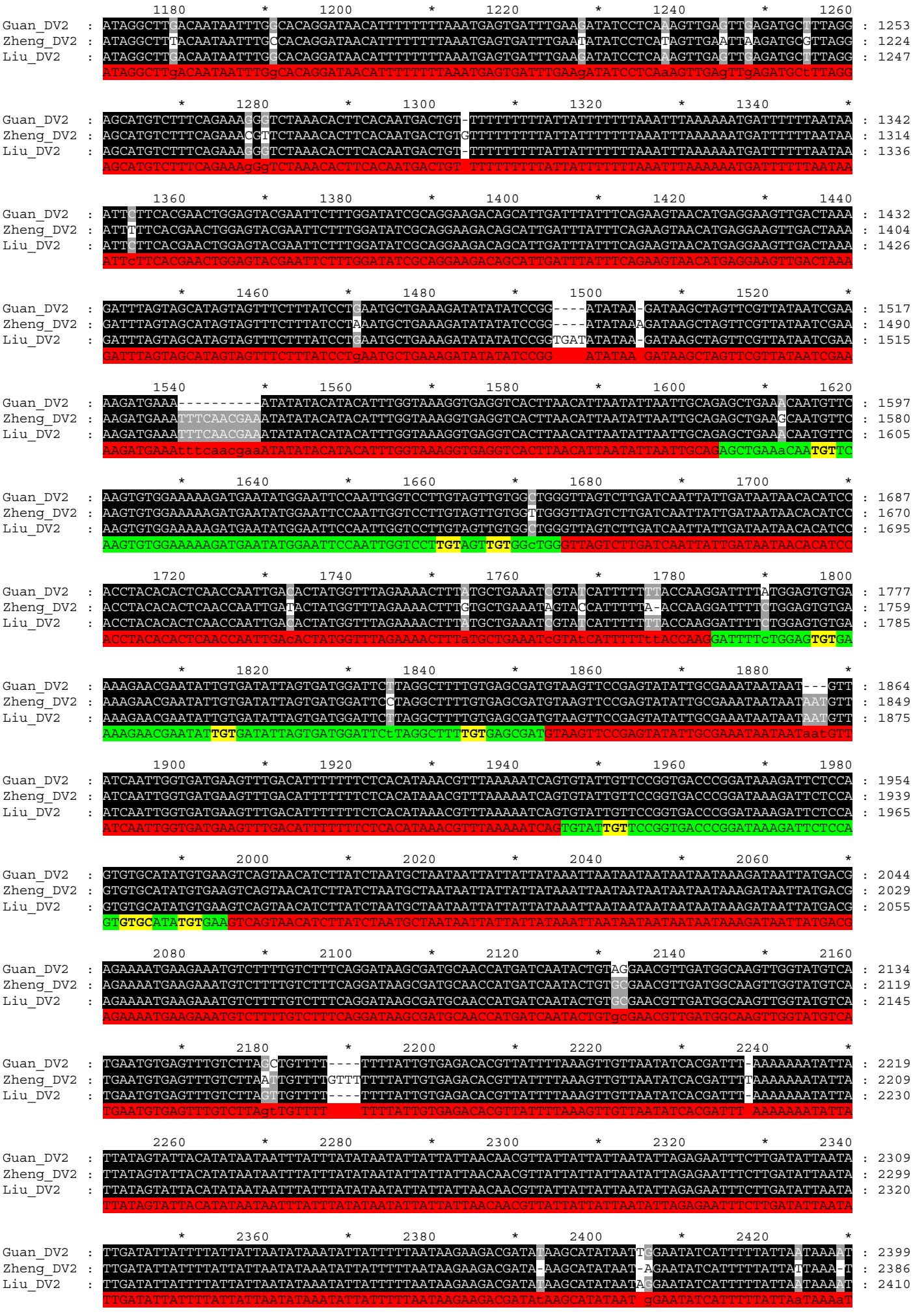
Figure S3A. Multiple sequence alignments of putative decorsin Hman_DV2 genes derived from the genome data of *H. manillensis* provided by Guan et al. (2020), Zheng et al. (2023) and Liu et al. (2023), respectively. The exons are labeled in green and the introns are labeled in red. Start and stop codons are marked in bold, the cysteine codons are marked in bold and yellow and the RGD/KGD motif encoding codons are marked in cyan and bold.





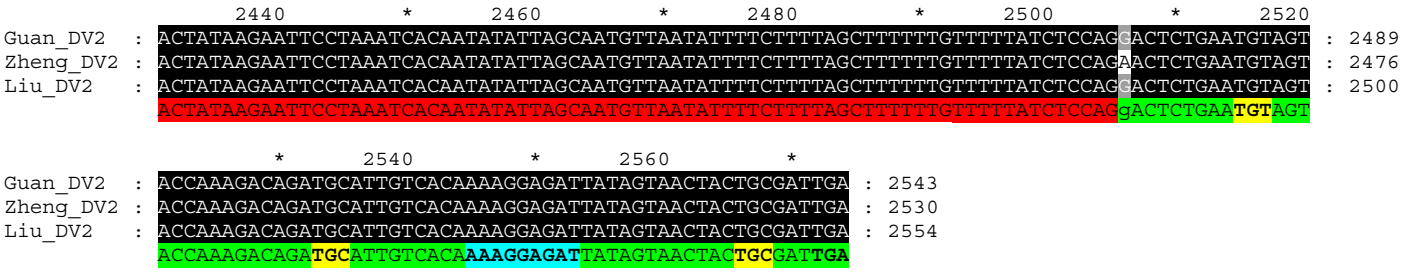


Figure S3B. Multiple sequence alignments of putative decorsin Hman_DV2 proteins derived from the genome data of *H. manillesis* provided by Guan et al. (2020), Zheng et al. (2023) and Liu et al. (2023), respectively. The cysteine residues are marked in bold and yellow and the RGD/KGD motifs are marked in cyan and bold. The signal peptide is underlined.

