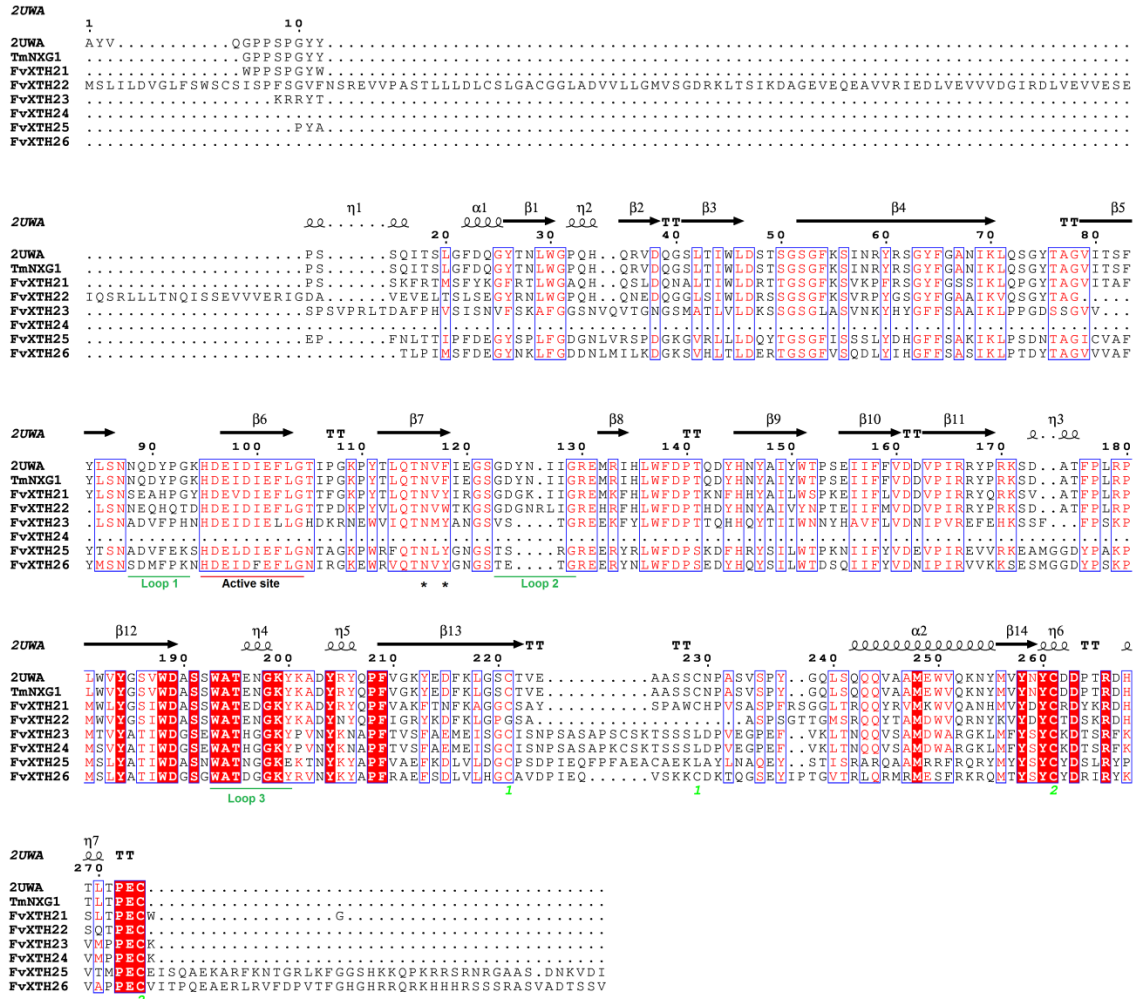




Supplementary Figure 2. The multiple alignment of deduced amino acid sequences of FvXTHs belonging to group IIIA and IIIB. The catalytic conserved domain (DEIDFEFLG), the secondary structures of β sheets (arrows) and α -helices (spiral), and loops 1, 2 and 3 (lines) are indicated.