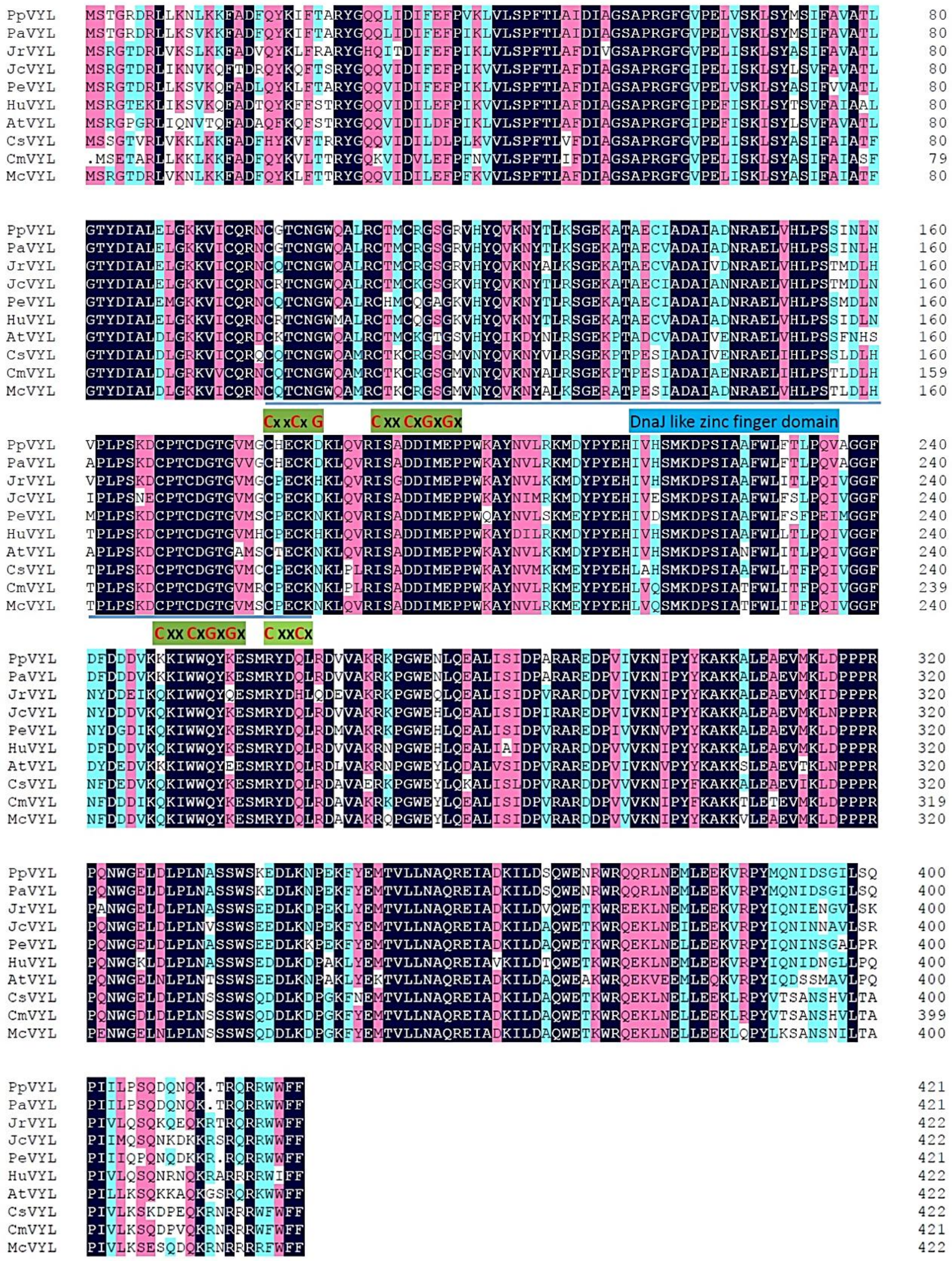Supplementary Figure 3 Sequence alignment of the DnaJ-like zinc finger protein from 10 CsVYL's homologous. The DnaJ-like CR domain are highlighted in blue underline. The four zinc binding motifs are shaded green. Plant species used for the multiple sequence alignment: Cs, *Cucumis sativus* (XP_004141219.1); Cm, *Cucumis melo* (XP_008452442.1); Pp, *Prunus persica* (ONI12706.1); Pa, *Prunus avium* (XP_021801691.1); Jr, *Juglans regia* (XP_018850702.1); Jc, *Jatropha curcas* (XP_012076623.1); Pe, *Populus euphratica* (XP_011029493.1); Hu, *Herrania umbratica* (XP_021288076.1); At, *Arabidopsis thaliana* (OAO89767.1); Mc, *Momordica charantia* (XP_022139600.1).