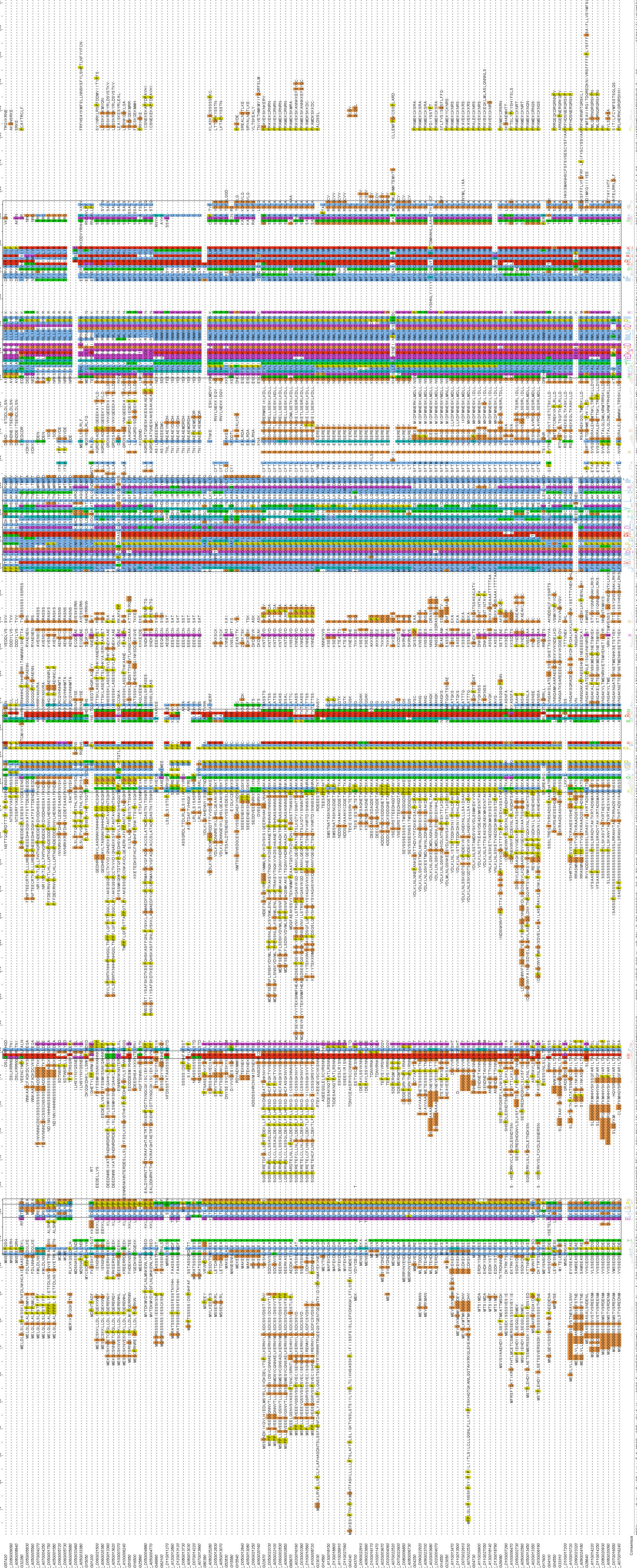


Supplementary Figure 2. Multiple sequence alignment of Aux/IAA proteins from *Arabidopsis* and upland cotton showed conserved residues or motifs.



Consensus

Domain I

Domain II

Domain III

Domain IV

GDP

KR rate motif