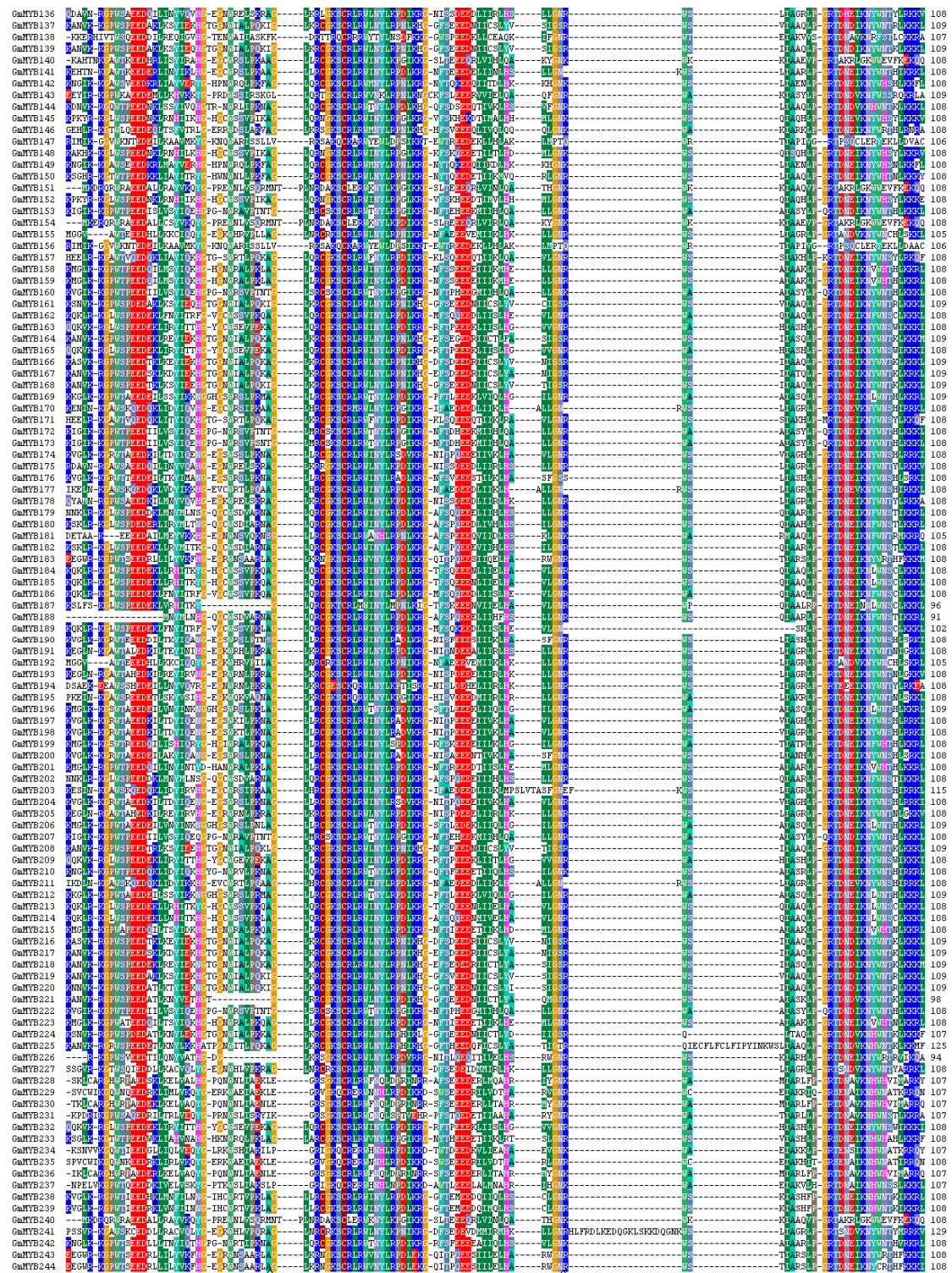


[illegible]



Additional file 2. Alignment of all MYB proteins used in this study.

CLUSTALW amino acid sequence alignment of the 244 soybean MYB domains. The shading of the alignment represents different degrees of conservation among sequences, respectively. The triangles indicate the most conserved intron insertion sites in the MYB domain.