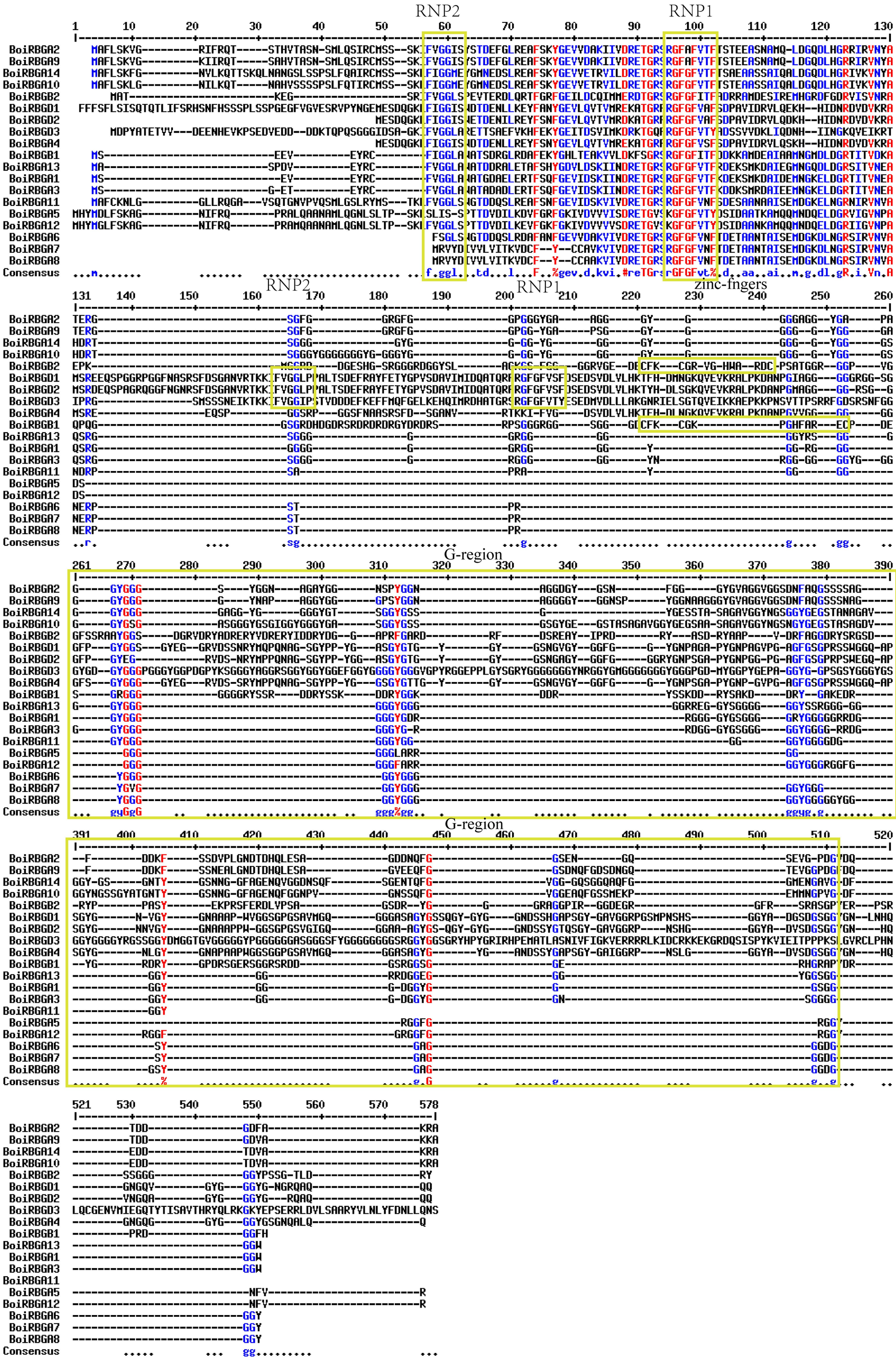

Figure S2. Amino acid sequence alignment of BoiRBGs. The yellow boxes around sequences show conserved motifs, including RRM and glycine-rich domains.