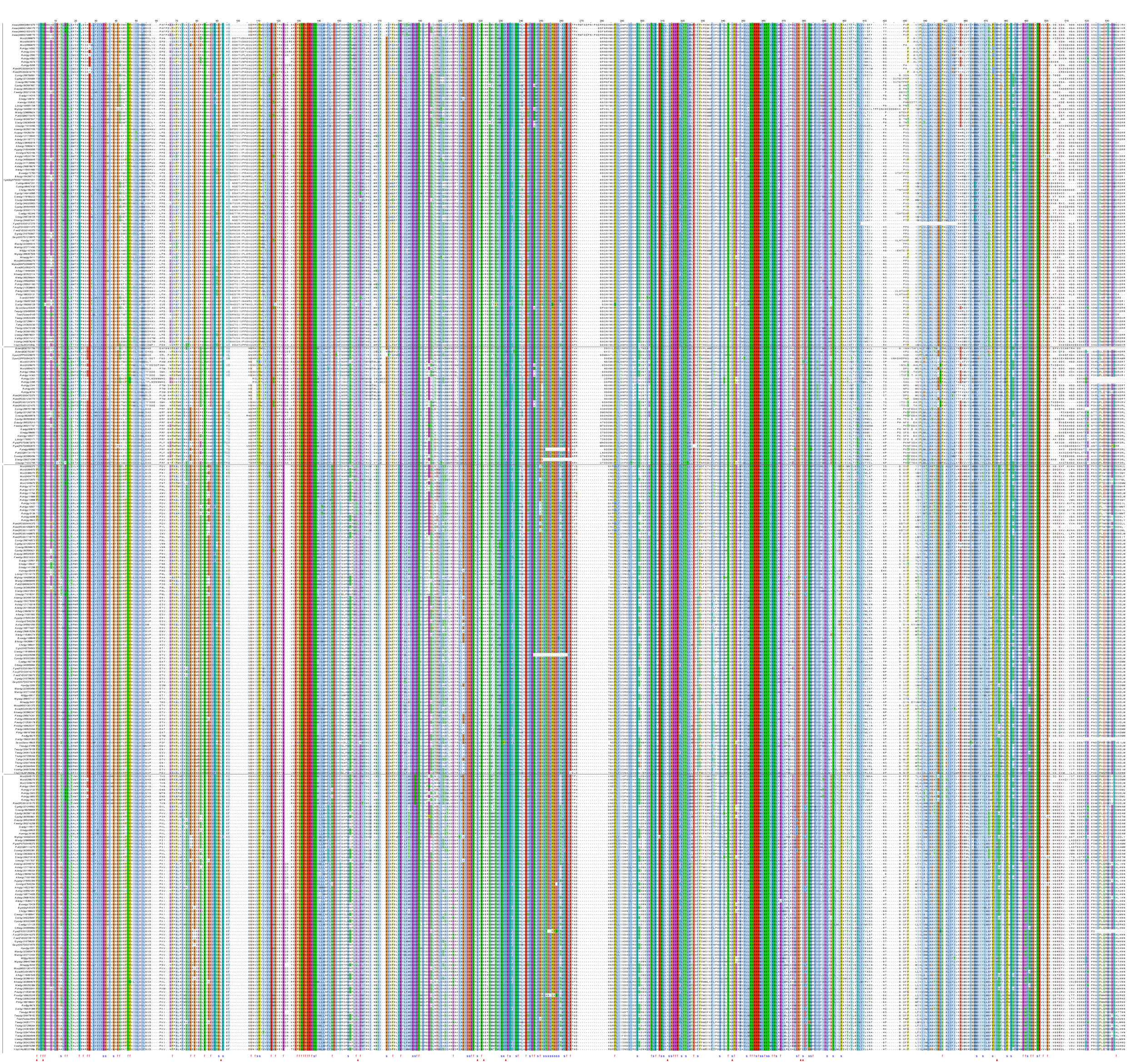

Figure S3. Multiple sequence alignment of CHS genes in division II. Red triangles indicate type II specificity-determining sites identified by SPEER-SERVER (Chakraborty et al., 2012).The letters f and s represent functional sites and structural sites identified by ConSurf (Ashkenazy et al., 2010), respectively.