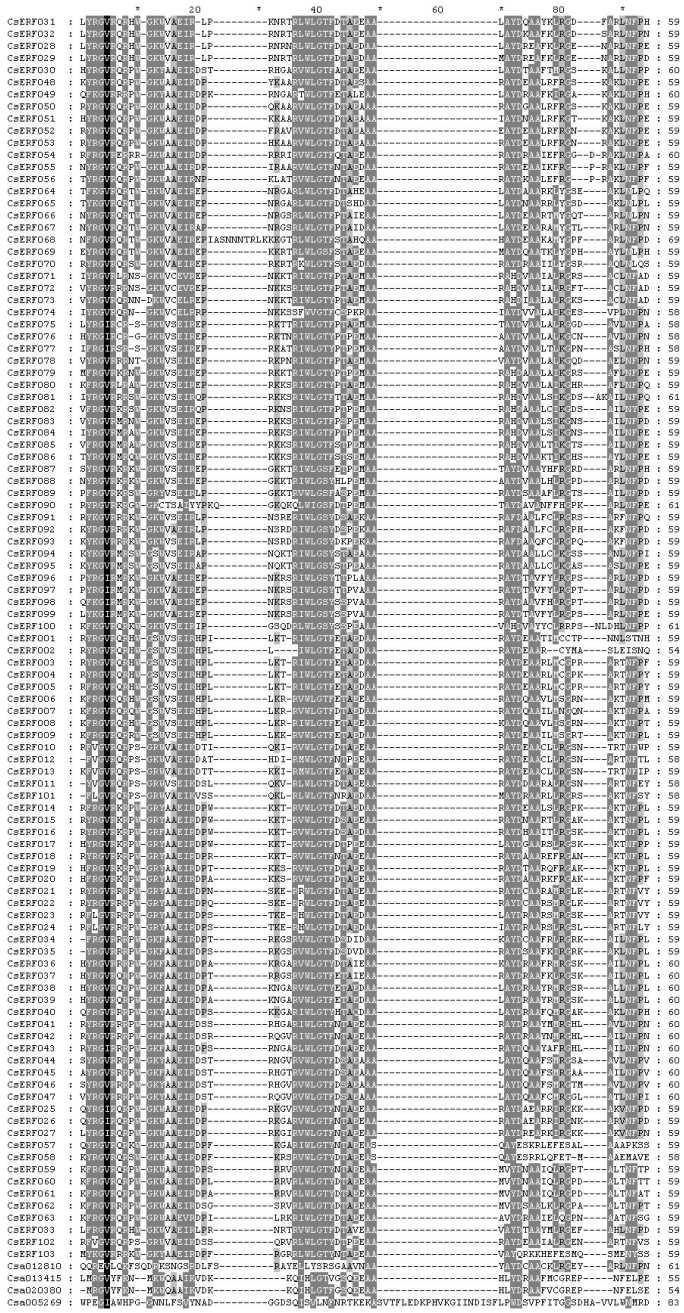

Figure S1 - Multiple sequence alignment of the AP2/ERF domains in 103 cucumber ERF, and 4 low homology proteins, Csa020380, Csa005269, Csa013415, and Csa012810**.** The protein sequences were aligned using Clustal X software.