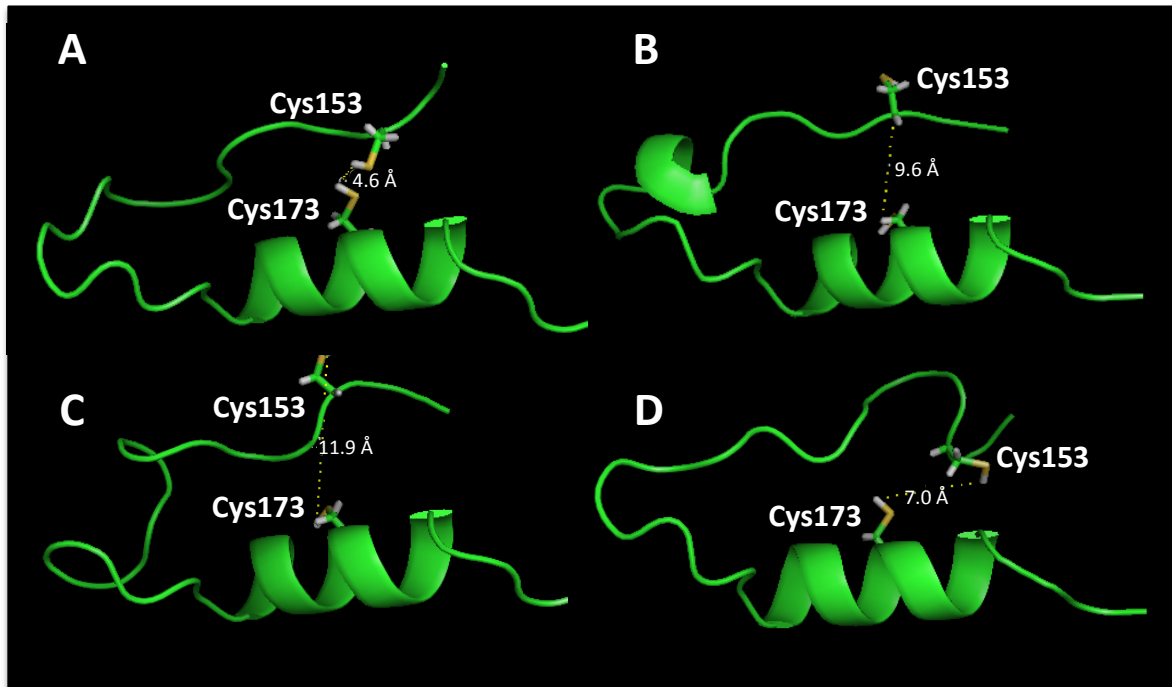


Supplemental figure 3



Model of the tertiary structure of the 170's loop showing the distance between Cys153 and Cys173 in the different subunits. The predicted structure was obtained with SWISS-MODEL (<https://swissmodel.expasy.org/>) by using as a template the modelled crystal structure 1DBZ of the C153S mutant of pea chloroplast FBPase. Images were generated with PyMOL.