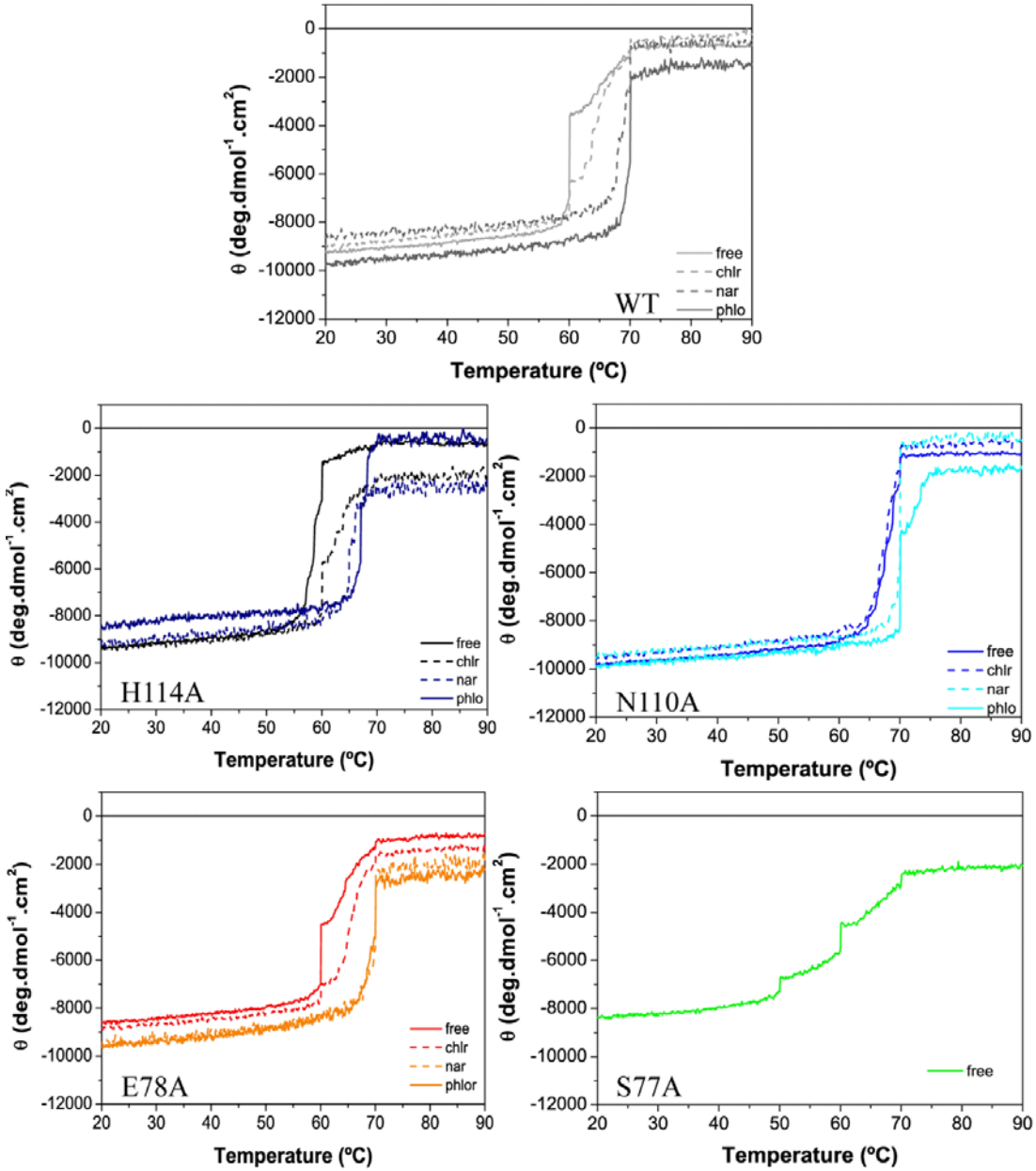


***SI_Figure: Protein unfolding of TtgR WT and mutants in the presence of ligands monitored by CD ellipticity at 222 nm***