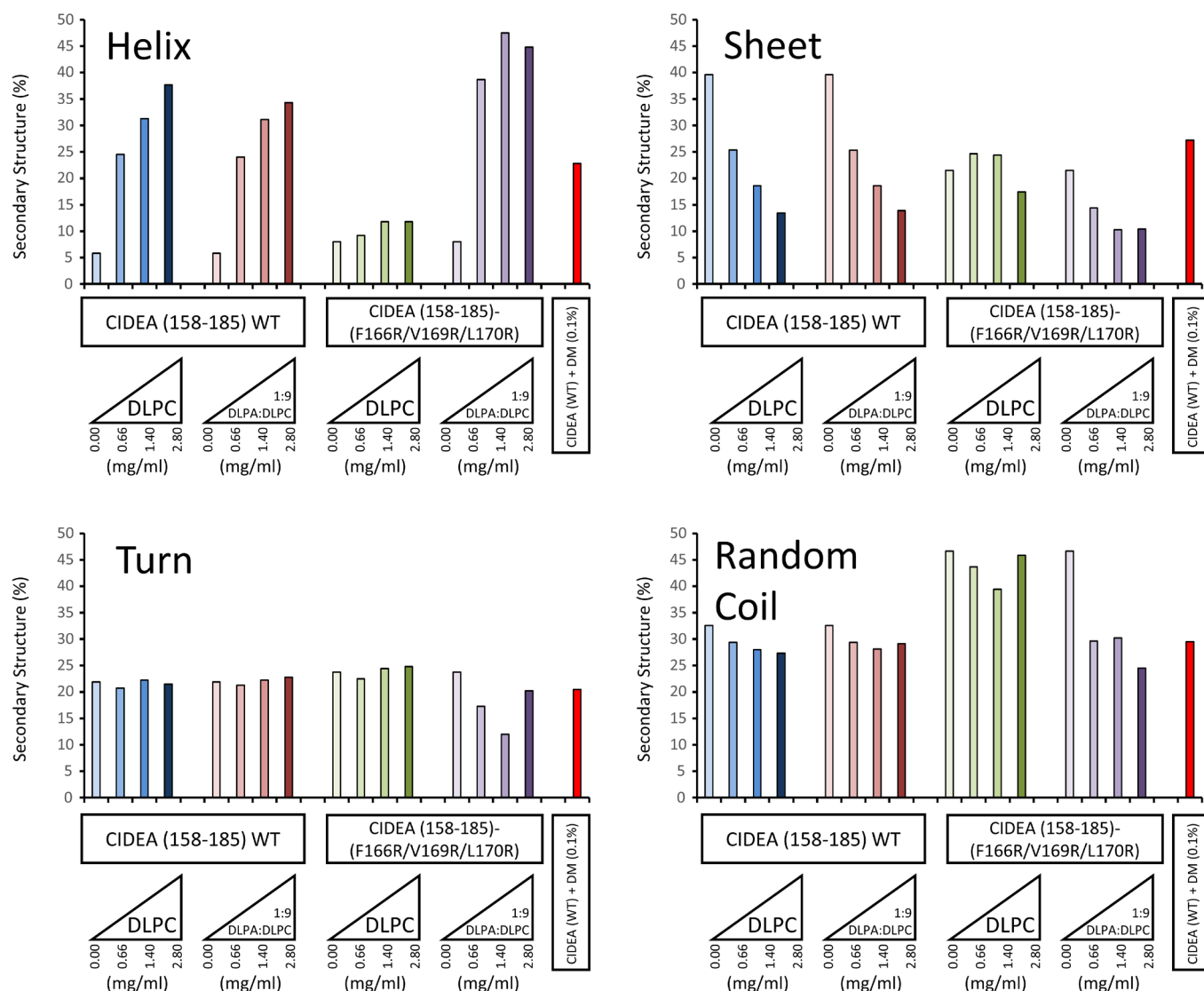


Figure 5-Figure Supplement 1

Secondary structure estimation of CIDEA amino acids 158-185 (wild type and F166R/V169R/L170R) based on Circular Dichroism data by using CDPro software with base set 4 (43 soluble proteins) using the CONTINLL program.



DM = 0.1% n-dodecyl- β -D-maltopyranoside