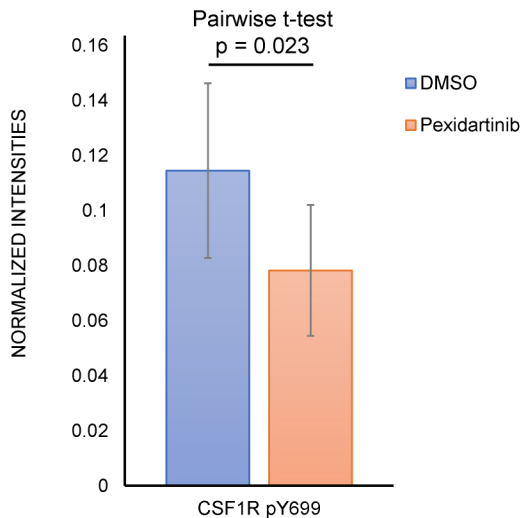
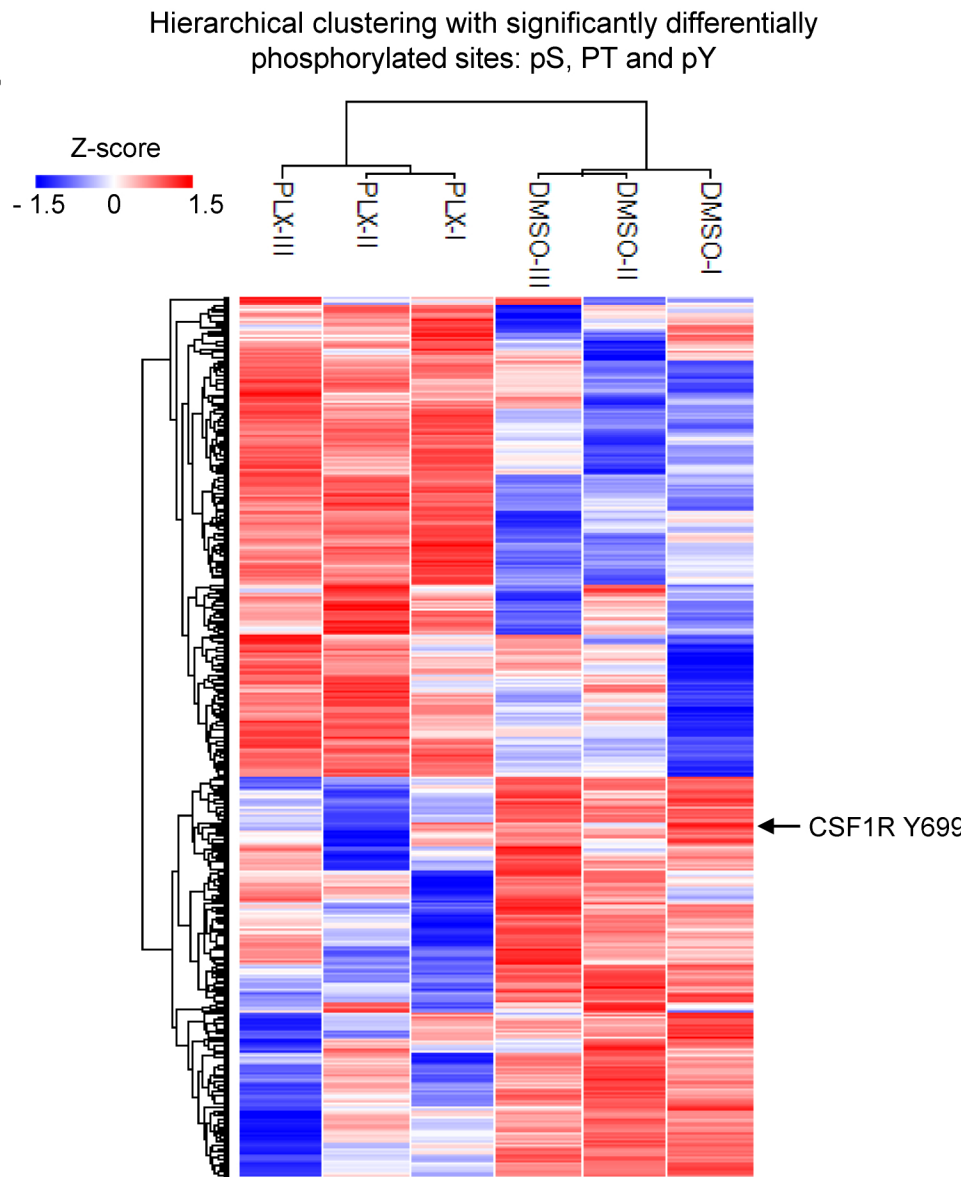


A.



B.



C.

DAVID analysis (KEGG pathways) based on the 551 proteins with at least one significantly (pairwise t-test) differentially phosphorylated sites

