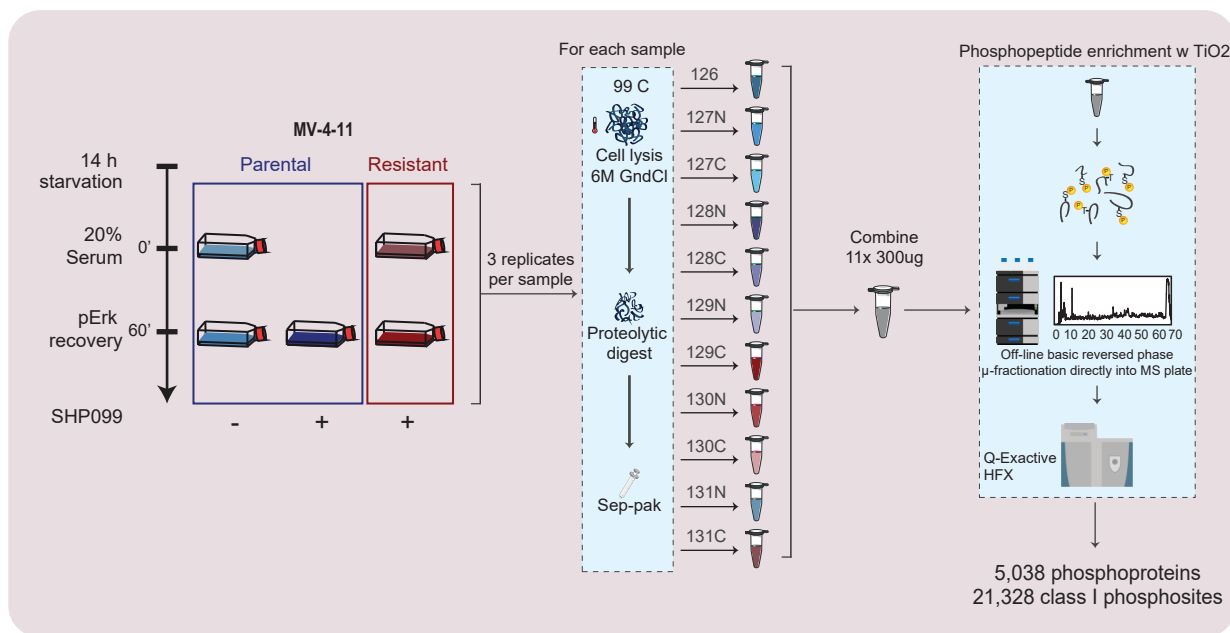
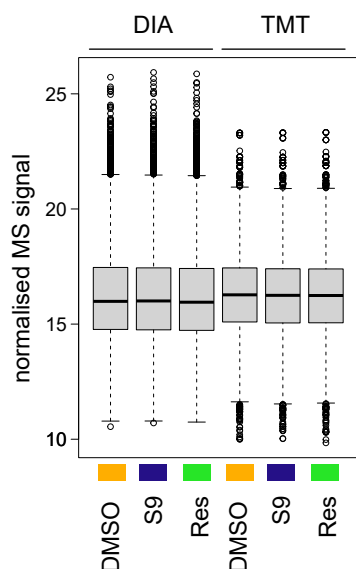


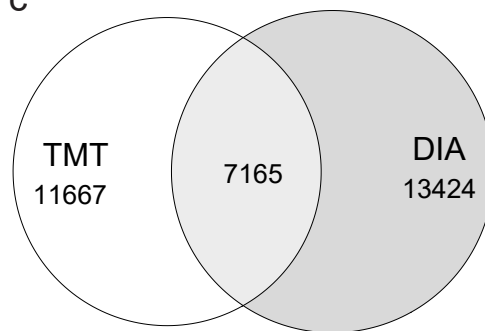
a



b



c



Supplementary Figure S2.

a) Experimental pipeline utilized for TMT analysis of the phosphoproteome of MV-4-11 and MV-4-11/R 1h after SHP099 treatment.

b) Boxplot of the normalized MS signal in the three conditions measured both with a DIA and TMT-based approach: 60 min. of SHP099 treatment in MV-4-11 cells. “Res” = cells with acquired resistance to SHP099; “S9” = SHP099-treated parental cells.

c) Proportional euler diagram of the phosphorylation sites quantified in the TMT and corresponding conditions of the DIA data sets.