

**Supplemental Figure 1.** Hierarchical clustering of kinase log2 LFQ z-scores of WT, KITmutant, PDGFRA-mutant tumors and normal gastric tissues, as determined by MIB-MS.



**Supplemental Figure 2. Defining kinome signatures of GIST tumor subtypes using MIB-MS profiling** (A, C and E) Scatterplot depicts the overlap in kinases elevated or reduced determined by LFQ or s-SILAC comparing (A) *KIT*-mutant vs WT, (C) *PDGFRA*-mutant vs WT and (E) *KIT*-mutant vs. *PDGFRA*-mutant. Regression analysis (R<sup>2</sup>) among quantitative methods was performed in Perseus Software. Differential expressed kinases commonly identified by LFQ and s-SILAC quantitation (FDR <0.05) are labeled. (**B**, **D** and **F**) Bar graph depicts high-confident kinases log2 LFQ values overexpressed in (B) *KIT*-mutant vs. WT, (D) *PDGFRA*-mutant vs. WT or (F) *KIT*-mutant vs. *PDGFRA*-mutant GIST primary tumors determined by LFQ and/or s-SILAC quantitation (FDR <0.05).



**Supplemental Figure 3.** MK-1775 and avapritinib have enhanced combination on in vitro GIST882 cell growth.

(A) Panels 1 & 2, Dose response curves for single agents (avapritinib, MK-1775) in GIST882 cell line. Red box indicates estimation of LD20 concentration for each single drug. Panel 3, Dose response curve representing increasing series of combinations in GIST882 cell line. Red box indicates estimation of LD20 concentration for combination of drugs. Panel 4, Single point (blue) on isobole curve for 20% kill. Red line indicates 50% isobole for strictly additive effect. CI<sub>LD20</sub> in GIST882 is 0.0237 and is found within the synergistic triangle (region below the red line). (B) Representative images of GIST882 spheroids after 120-hour treatment at indicated concentrations. (C, left panel) Bars represent average viability  $\pm$  SEM after 120-hour treatment at indicated drug concentrations for GIST882 spheroids as a percentage of vehicle-treated spheroids. (C, right panel) Bars represent the average spheroid data were analyzed using GraphPad Prism, with comparisons of treatment groups performed in one-way ANOVA and post hoc comparisons made using Bonferroni multiple comparisons method \* p = 0.0249, \*\* p = 0.0042, \*\*\* p = 0.0006, \*\*\*\* p = < 0.0001.