

Description of Additional Supplementary Files

Title: Supplementary Data 1.

Description: LiP-SMap experiment of HeLa cell lysates treated with rapamycin at a concentration of 2 μ M.

Title: Supplementary Data 2.

Description: LiP-SMap experiment of HeLa live cell cultures treated with rapamycin at a concentration of 2 μ M.

Title: Supplementary Data 3.

Description: LiP-Scores obtained after analysis with the LiP-Quant machine learning-based classifier of HeLa cell lysates treated with the following compounds: rapamycin, FK506, calyculin A, fostriecin, selumetinib, staurosporine.

Title: Supplementary Data 4.

Description: Consistency of LiP-Quant scores and differential peptide abundance intensities with rapamycin and FK506.

Title: Supplementary Data 5.

Description: LiP scores obtained after analysis with the LiP-Quant machine learning-based classifier of *S.cerevisiae* cell lysates from *wild type* and $\Delta Frb1$ mutant strains.

Title: Supplementary Data 6.

Description: KinHub kinase reference database (<http://www.kinhub.org>).

Title: Supplementary Data 7.

Description: Benchmarking of LiP-Quant and Deep-LiP-Quant with TPP parametric (Savitski, 2014) and TPP non-parametric analysis (Childs, 2019).

Title: Supplementary Data 8.

Description: Deep-LiP-Quant experiment of HeLa cell lysates treated with staurosporine and analyzed with a 4-hours LC gradient.

Title: Supplementary Data 9.

Description: LiP-Scores obtained after analysis with the LiP-Quant machine learning-based classifier of HeLa membrane-enriched fractions.

Title: Supplementary Data 10.

Description: Single concentration LiP-Quant rapamycin control experiment of HeLa cell lysates treated with mild detergent during cell lysis.

Title: Supplementary Data 11.

Description: Nuanced specificity in target binding of fostriecin and calyculin A.

Title: Supplementary Data 12.

Description: Predictions of binding sites with center of mass of high confidence LiP peptides.

Title: Supplementary Data 13.

Description: LiP-Scores obtained after analysis with the LiP-Quant machine learning-based classifier of *Botrytis cinerea* cell lysates treated with the compound BAYE-004.

Title: Supplementary Data 14.

Description: Peptide Frequency Library (PFL) that identifies proteins that are common contaminants in LiP-Quant experiments.

Title: Supplementary Data 15.

Description: Peptide ranking for calculation of drug binding sites in the LiP-Quant experiments reported in this manuscript.

Title: Supplementary Data 16.

Description: List of reagents or resources used in this manuscript.