

Figure S2

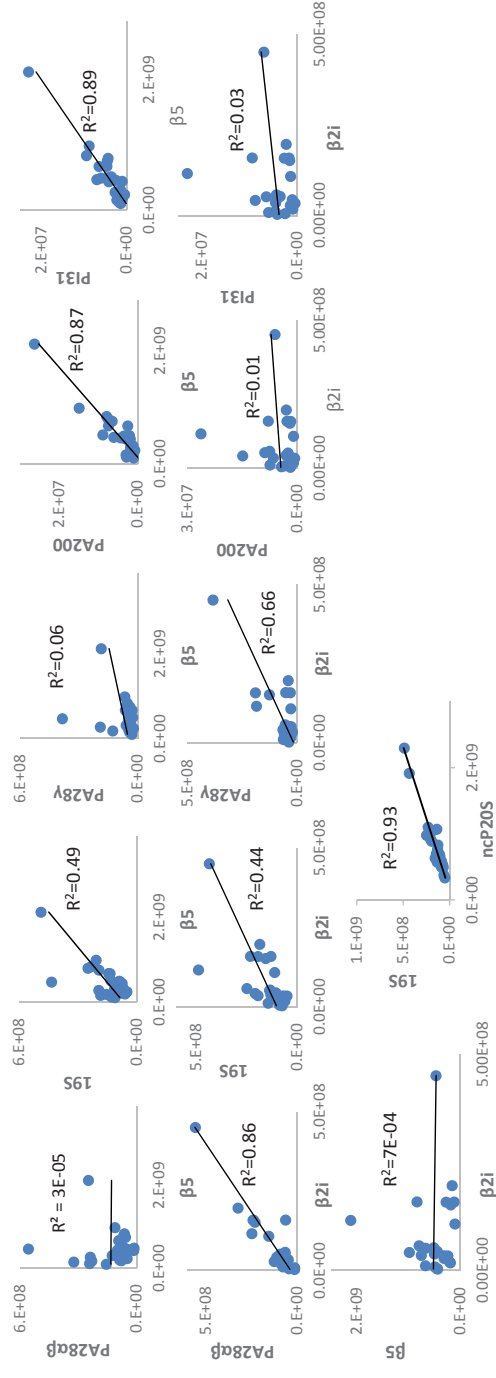
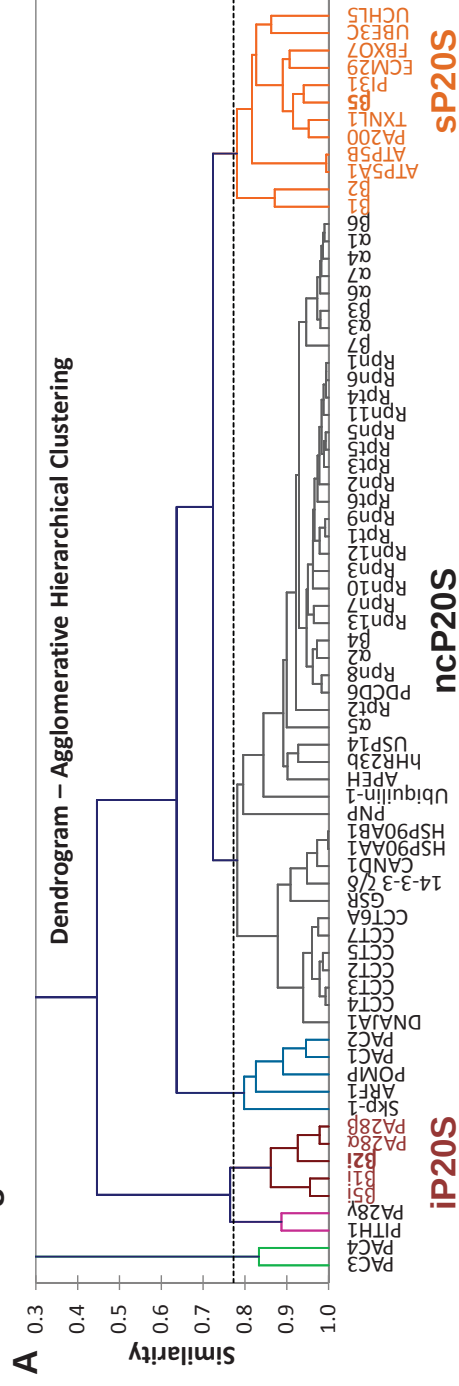


Figure S2: A- The abundances of 73 known PIPs were analyzed using an unsupervised Agglomerative Hierarchical Clustering approach. The clusters nicely grouping the subunits of the iP20S (in dark red), the ncP20S (in black), and the sP20S (in orange) were highlighted. B- Correlations between the abundances of the iP20S (represented by $\beta 2i$) or the sP20S (represented by $\beta 5$) and the 5 main proteasome regulators (PA28 $\alpha\beta$, 19S, PA28 γ , PA200, and P131) across the 24 AP-MS experiments.

PA28 $\alpha\beta$ profile corresponds to the median of PA28 α and PA28 β profiles and the 19S profile corresponds to the median of Rpt1-6, Rpn1-3, 5-14 profiles.