

Figure S6

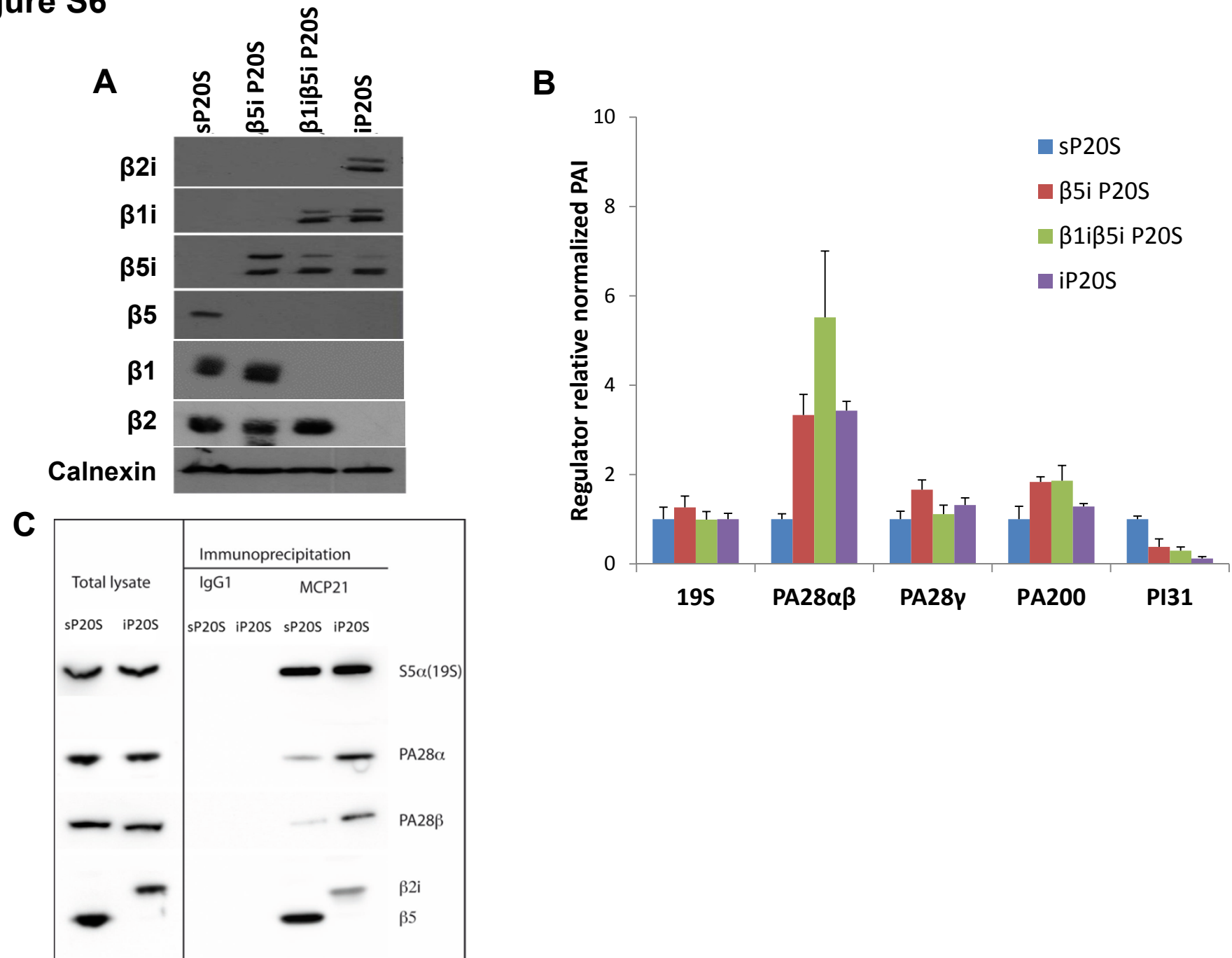


Figure S6: Interaction of the four different 20S proteasome subtypes (sP20S, β 5i P20S, β 1i β 5i P20S or iP20S) with the main 20S proteasome regulators in HEK EBNA cell lines transfected with β 5i (β 5i intermediate P20S), β 1i and β 5i (β 1i β 5i intermediate P20S) or β 1i, β 2i, β 5i (iP20S).

A) Immunodetection of the catalytic subunits in the total cell lysates from the four HEK EBNA cell lines expressing only standard catalytic subunits (sP20S), a mixture of standard and catalytic subunits (β 5i intermediate P20S and β 1i β 5i intermediate P20S), or only immuno catalytic subunits (iP20S). Calnexin is used as a loading control.

B) Relative normalized abundance indexes of proteasome regulators in HEK EBNA cells containing only sP20S, iP20S, β 5i intermediate P20S or β 1i β 5i intermediate P20S. The normalized abundance indexes for each regulator was set to 1 for standard proteasome conditions (n=4).

C) Comparison of the quantity of 19S and PA28 $\alpha\beta$ regulators associated with the 20S proteasome in HEK T-Rex cells containing only iP20S or sP20S. Western blot analyses were performed on total cell lysates or proteasome MCP21 immunoprecipitates with antibodies against the S5a (Rpt1 – 19S subunit), PA28 α , PA28 β , β 2i and β 5 subunits.