



Figure S2. Graph showing the change in NSR1/PEX6 expression caused by deletion of Sds3 in the W303 background (ACY605 strain used in microarray analysis), the BY4741 background used in the YKO collection (ACY978), and the *sds3Δ* strain from the YKO collection. Each bar shows the expression level compared to that found in the wild-type strain (both 20min after treatment with 0.4M KCl), as measured by qPCR (Methods). In this experiment cells were grown and harvested as described for the DNA microarray experiments but the mRNA was purified using a RiboPure RNA purification kit (Ambion).