

Additional file 5: Figure S2. Quantification of total and intracellular TEX101 in pooled SP samples by SRM assays. The heavy isotope-labeled peptides AGTETAILATK* (total TEX101) and QIQTSSSQTSPEEAMGTPR* (intracellular TEX101) with a trypsin-cleavable JPT tag (serine-alanine-[3-nitro]tyrosine-glycine) were used as internal standards for the absolute quantification of endogenous TEX101 proteoforms. Calibration curves were generated by spiking increasing amounts of the internal standards into 1 μ L of SP before proteomic sample preparation and trypsin digestion.

