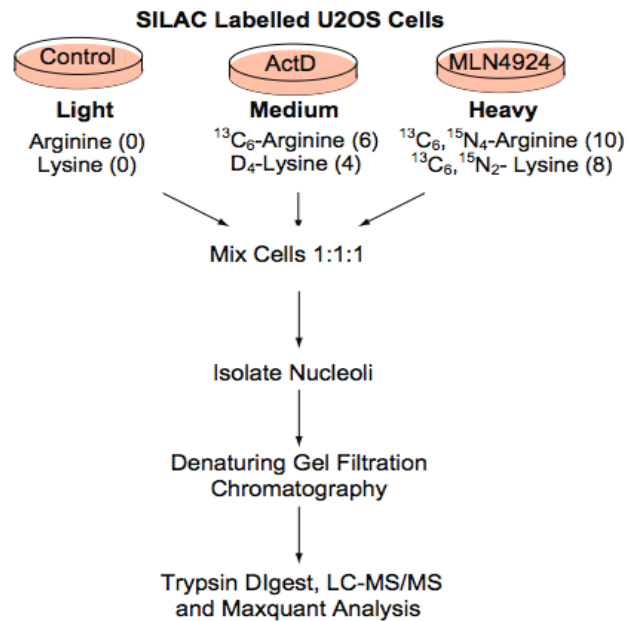


Quantitative proteomics to determine the effect of NEDD8 inhibition on the nucleolar proteome



1300 nucleolar proteins quantified
MLN4924 alters the nucleolar proteome on a similar scale to
the well-established nucleolar stressor Actinomycin D

