Supplementary

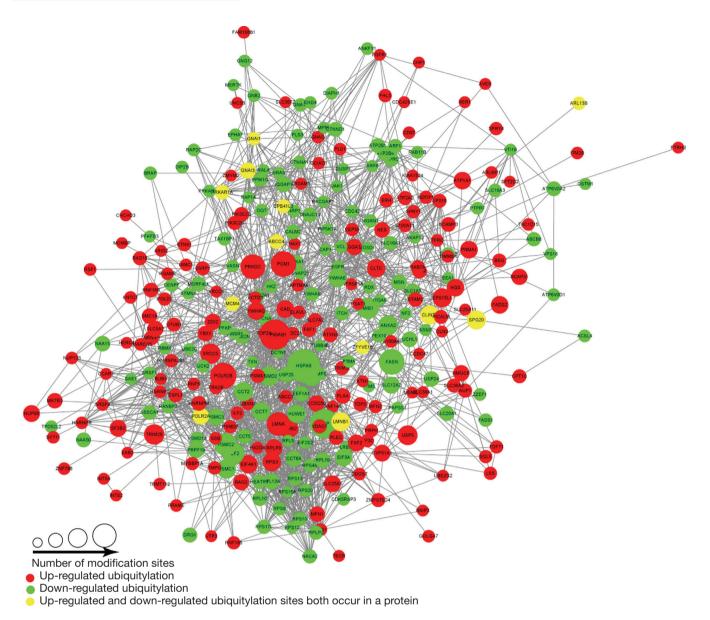


Figure S1 Protein-protein interaction network of differentially quantified Kub (lysine ubiquitination) proteins in perifosine-treated AS cells.

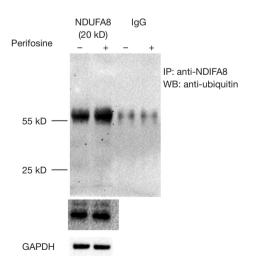


Figure S2 Detection of ubiquitylation of NDUFA8. Co-IP and western blotting were used to validate ubiquitylome, resulting increase of ubiquitylated NDUFA8 after 10 μ M perifosine treatment.

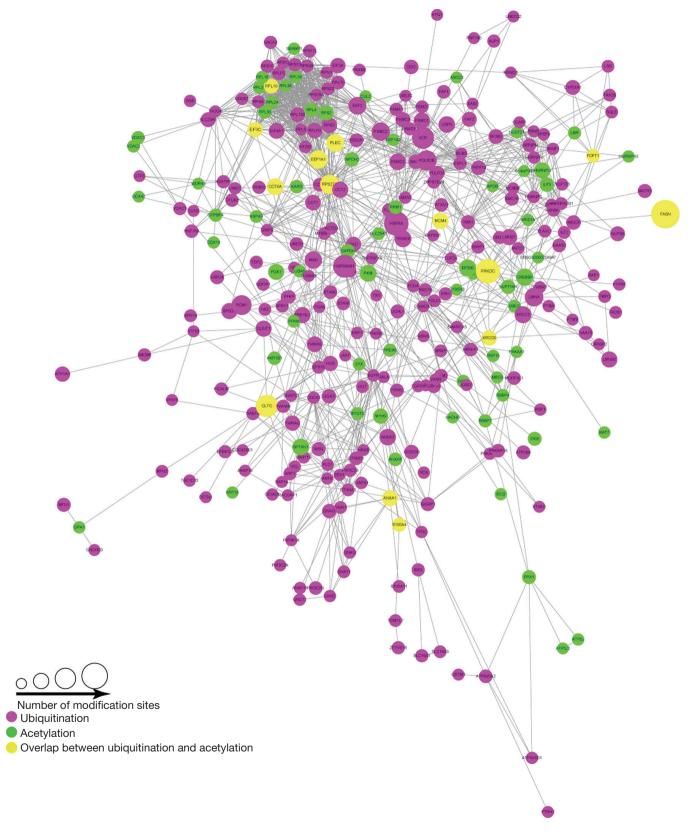


Figure S3 Protein-protein interaction network of differentially quantified Kub (lysine ubiquitination) and Kac (lysine acetylation) proteins in perifosine-treated AS cells.