

Table S1. Mitochondrial proteins differentially co-immunoprecipitated with 2B

Symbol	Entrez Gene Name	Protein Accession Number	Heavy/Light Ratio	Heavy/Light Count
SDHB	succinate dehydrogenase complex iron sulfur subunit B	P21912	1183.487	6
UQCRB	ubiquinol-cytochrome c reductase binding protein	P14927	966.47	3
ATP5MG	ATP Synthase Membrane Subunit G	O75964	848.136	2
UQCRC1	ubiquinol-cytochrome c reductase core protein 1	P31930	597.584	9
VDAC3	voltage-dependent anion channel 3	Q9Y277	548.569	2
NDUFS1	NADH:ubiquinone oxidoreductase core subunit S1	P28331	478.214	4
ATP5PF	ATP synthase peripheral stalk subunit F6	P18859	331.494	4
NDUFV2	NADH:ubiquinone oxidoreductase core subunit V2	P19404	320.749	2
NDUFA7	NADH:ubiquinone oxidoreductase subunit A7	O95182	220.892	3
ATP5PD	ATP Synthase Peripheral Stalk Subunit D	O75947	206.964	6

Symbol refers to the official symbol of the protein-coding gene; Entrez gene name refers to the full gene name provided by Entrez database; protein accession number is an unique identifier assigned to the protein by the UniProtKB/Swiss-Prot database; the heavy/light ratio refers to the ratio of the quantification values of the heavy (i.e. deuterated) and light (i.e. non-deuterated) quantification channels; heavy/light count refers to the number of peptide ratios used for calculation of a particular protein ratio.