



Legend: **Peptide Sequence**      **Ion**   **XC**      **Delta Cn**

## Blk-Interacting Proteins

A) gi|5901922|ref|NP\_008996.1| (NM\_007065) CDC37 cell division cycle 37 homolog (S. cerevisiae)

Shown:	<b>CIDSGLWVPNSK</b>	2	3.308	0.3
Others:	<b>LQAEAQLR</b>	2	3.199	0.196

B) gi|4502203|ref|NP\_001650.1| (NM\_001659) ADP-ribosylation factor 3

NOTE: Could be ARF 2,3,4...

Shown:	<b>DAVLLVFANK</b>	2	3.842	0.353
Others:	<b>ILMVGLDAAGK</b>	2	2.916	0.272

C) gi|4557797|ref|NP\_000260.1| (NM\_000269) non-metastatic cells 1, protein (NM23A)

Shown:	<b>GDFCIQVGR</b>	2	3.086	0.292
Others:	<b>DRPFFAGLVK</b>	2	2.494	0.233

## Tere1- Interacting Proteins

D) gi|23308577|ref|NP\_006614.2| phosphoglycerate dehydrogenase

Shown:	<b>GGIVDEGALLR</b>	2	4.15	0.202
Others:	<b>GTIQVITQGTSLK</b>	2	4.233	0.404
	<b>ILQDGGLQVVEK</b>	2	4.027	0.279
	<b>TLGIIGLGR</b>	2	2.705	0.135
	<b>DLPLLLFR</b>	2	2.473	0.085

E) gi|4757810|ref|NP\_004037.1| ATP synthase, H<sup>+</sup> transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle

Shown:	<b>TSIAIDIINQK</b>	2	4.757	0.28
Others:	<b>VVDALGNAIDGK</b>	2	3.906	0.375
	<b>VLSIGDGIAR</b>	2	2.673	0.209
	<b>ILGADTSVDLEETGR</b>	2	5.069	0.386
	<b>HALIYDDLSK</b>	2	3.736	0.333