

S2 Table: Sequence conservation of prefoldin subunits in model eukaryotes.

Gene name	gene ID	alias	subunit type	product size (aa)	Human homologue	% amino acid identity				
						<i>Homo sapiens</i>	<i>Mus musculus</i>	<i>Drosophila melanogaster</i>	<i>Caenorhabditis elegans</i>	<i>Saccharomyces cerevisiae</i>
<i>pdf1</i>	SPBC1D7.01	none	beta	112	<i>PFDN1</i>	21.4	21.4	19.6	25.9	28.4
<i>pdf2</i>	SPAC227.10	none	beta	114	<i>PFDN2</i>	31.6	31.6	29.0	21.1	41.4
<i>pdf3</i>	SPAC3H8.07c	<i>pac10</i>	alpha	169	<i>PFDN3</i>	43.2	43.2	39.3	34.3	44.9
<i>pdf4</i>	SPAC227.05	none	beta	123	<i>PFDN4</i>	34.2	32.5	26.7	34.7	33.6
<i>pdf5</i>	SPBC215.02	<i>bob1</i>	alpha	154	<i>PFDN5</i>	45.2	45.2	32.9	32.2	35.8
<i>pdf6</i>	SPAC3A11.13	none	beta	114	<i>PFDN6</i>	36.8	36.8	32.5	29.5	42.1