

S3 Table. Identification of proteins in spots that exhibit significant Pro-Q Diamond-staining intensity changes in the absence of CaPpz1

Protein Name ¹	Protein ID ¹	Systemic gene name ²	Standard gene name ¹	Protein abbreviation ²	Spot ID	Fold change	p value
40S ribosomal protein S0	O42817	<i>C3_05370C</i>	<i>YST1</i>	Yst1	22	-2.79	0.041
40S ribosomal protein S7-A	Q5AJ93	<i>C3_01490W</i>	<i>RPS7A</i>	Rps7A	04	-2.58	0.047
60S acidic ribosomal protein P0*	A0A1D8PQS0	<i>C7_00990W</i>	<i>RPP0</i>	Rpp0	20	2.56	0.046
					21	2.63	0.005
60S ribosomal protein L20	A0A1D8PLC9	<i>C4_01520C</i>	<i>RPL20B</i>	Rpl20B	10	-4.81	0.019
ATP-dependent RNA helicase eIF4A	P87206	<i>C1_01350C</i>	<i>TIF1</i>	Tif1	23	-2.05	0.014
Cys-Gly metallodipeptidase DUG1	Q5AKA5	<i>C5_04300C</i>	<i>DUG1</i>	Dug1-like**	36	1.37	0.008
Deoxyhypusine hydroxylase*	Q59Z14	<i>C2_07290W</i>	<i>LIA1</i>	Lia1-like**	17	-3.09	0.019
					18	-2.32	0.008
D-lactate dehydrogenase	A0A1D8PJK5	<i>C3_03040W</i>	<i>AIP2</i>	Aip1	34	-2.05	0.026
Elongation factor 2	Q5A0M4	<i>C2_03100W</i>	<i>EFT2</i>	Eft2	28	2.79	0.008
Elongation factor 3	P25997	<i>C5_01580C</i>	<i>CEF3</i>	Cef3	27	-2.38	0.008
Guanylate kinase	A0A1D8PNS0	<i>C5_03790W</i>	<i>GUK1</i>	Guk1	03	-4.57	0.003
Heat shock protein SSA1	P41797	<i>C1_13480W</i>	<i>HSP70</i>	Hsp70/Ssa1	32	-3.32	0.024
Inosine triphosphate pyrophosphatase	Q59N80	<i>C5_03860W</i>	<i>HAM1</i>	Ham1	01	-2.79	0.034
Peptidyl-prolyl cis-trans isomerase	Q5ALM6	<i>C2_02320C</i>	<i>CPR3</i>	Cpr3	02	-2.59	0.018
Peptidyl-prolyl cis-trans isomerase	A0A1D8PKL0	<i>C3_06360C</i>	<i>CYP5</i>	Cyp5	07	-2.02	0.0003
Peroxisredoxin TSA1	Q9Y7F0	<i>C3_06180C</i>	<i>TSA1</i>	Tsa1	16	-2.18	0.028
Rad23p	A0A1D8PGE3	<i>C2_01850W</i>	<i>RAD23</i>	Rad23	39	1.95	0.006
Ribosomal 60S subunit protein L9B	Q5AEN2	<i>C3_02470C</i>	<i>RPL9B</i>	Rpl9B	13	-3.23	0.049
Transketolase 1	Q5A750	<i>C1_08320W</i>	<i>TKL1</i>	Tkl1	29	1.78	0.045
Ubiquitin-activating enzyme E1*	A0A1D8PKJ3	<i>C3_06500W</i>	<i>UBA1</i>	Uba1	30	1.90	0.006
					31	-2.31	0.037

Based on ¹<http://www.uniprot.org/> and ² <http://www.candidagenome.org/>.

*The same protein was identified in two separate spots.

** The abbreviation is based on the name of the *S. cerevisiae* ortholog.