

Contig	Annotation	E-value	log2FoldChange	
CGI_10004018	O-methyltransferase mdmC	4E-23	↑	661.416.733.671.093
CGI_10002823	Heat shock protein 68	0	↑	603.574.568.795.695
CGI_10024833	N-acetyltransferase ESCO1	1E-63	↑	501.546.305.790.145
CGI_10020140	Polyglutamine-binding protein 1	9E-25	↑	500.045.316.862.761
CGI_10000185	Relaxin receptor 2	6E-68	↑	480.606.825.033.121
CGI_10017043	Mitotic apparatus protein p62	1E-20	↑	461.347.300.084.844
CGI_10002384	Plasminogen	2E-46	↑	424.764.556.373.692
CGI_10000014	Elongation factor Tu, mitochondrial	2E-32	↑	411.456.864.340.155
CGI_10017684	Probable ATP-dependent DNA helicase	8E-22	↑	399.710.178.590.712
CGI_10010647	Heat shock protein 68	0	↑	364.038.502.681.902
CGI_10012787	NmrA-like family domain-containing pr	2E-38	↓	-389.963.545.870.178
CGI_10016364	Baculoviral IAP repeat-containing prot	2E-30	↓	-403.605.204.044.111
CGI_10019361	Baculoviral IAP repeat-containing prot	3E-48	↓	-405.844.884.917.493
CGI_10016363	Baculoviral IAP repeat-containing prot	5E-37	↓	-411.726.973.782.534
CGI_10014170	Nacrein-like protein	9E-79	↓	-412.080.938.949.555
CGI_10022071	Cysteine sulfinic acid decarboxylase	0	↓	-412.724.673.515.028
CGI_10009981	Sodium- and chloride-dependent glyci	0	↓	-425.976.108.157.904
CGI_10002314	Short-chain dehydrogenase/reductase	9E-57	↓	-429.268.398.900.263
CGI_10023094	Mitotic checkpoint serine/threonine-p	7E-65	↓	-436.786.643.610.721
CGI_10005650	Ras-related protein Rab-11A	4E-90	↓	-487.338.881.451.019