

Supporting information for Dasgupta *et al.* (2002) *Proc. Natl. Acad. Sci. USA* **99** (5), 2666–2671. (10.1073/pnas.052397899)

Table 1. Quantitation of Northern (Fig. 1)

Gene name	Ratio of <i>mot1-14</i> to <i>MOT1+</i>	Ratio of <i>mot1-42</i> to <i>MOT1+</i>	Ratio of <i>taf145-869</i> to <i>TAF145+</i>	Ratio of <i>tsm1 ts</i> to <i>TSM1+</i>	Ratio of <i>toa2-3</i> to <i>TOA2+</i>
<i>MOT1</i> -regulated genes					
<i>INO1</i>	5.9	4.9	0.75	1.2	0.97
<i>THI5</i>	4.6	4.5	0.78	1.2	0.38
<i>YDR533C</i>	4.5	4.2	1.6	2.5	0.70
<i>YGR043C</i>	5.2	5.3	3.2	1.9	1.1
<i>AGA1</i>	4.5	3.0	0.35	1.1	0.36
<i>HSP26</i>	3.8	4.5	4.2	1.7	2.3
<i>TSL1</i>	3.2	1.8	2.1	1.4	1.4
<i>BNA1</i>	0.26	0.49	0.15	0.46	0.19
<i>URA1</i>	0.28	0.50	0.45	1.2	0.54
<i>YDR539W</i>	0.26	0.38	0.32	0.51	0.20
Genes not affected by <i>MOT1</i> mutation					
<i>RPL5</i>	1.0	1.2	0.44	1.6	0.80
<i>RPS5</i>	1.1	1.2	0.38	1.5	0.76
<i>ACT1</i>	1.3	1.6	0.78	2.0	1.1
<i>SAN1</i>	0.7	0.9	0.6	0.9	1.0
<i>RAD16</i>	0.8	1.2	1	0.9	0.7