Gene lists for Table 2, promoter motifs

Genes that are require Sty1p, but not Atf1p, for stress induction:

pka1/tpk/git6	CAMP-dependent protein kinase catalytic subunit
SPBC56F2.06	hypothetical protein; sequence orphan
SPCC191.01	hypothetical protein; sequence orphan
SPAC11E3.14	conserved hypothetical protein; similar to S. cerevisiae YKR018C
SPCC1322.08	putative serine/threonine protein kinase
mpr1/spy1	stress response regulator phosphotransmitter; respone regulator phospotransferase; similar to S. cerevisiae YPD1
SPBP4H10.10	membrane protein of unknown function, possibly involved in respiration by similarity to YGR101W; rhomboid family
SPBC20F10.10	similarity to yeast Pho85p-associated PCL6 cyclin
SPCC417.05C	putative involvement in chitin biosynthesis; by similarity
tps1	alpha,alpha-trehalose-phosphate synthase
SPCC61.03	conserved hypothetical protein
SPAC607.08C	conserved hypothetical protein; similar to YFL034W
atf1/mts1/sss1/gad7.	transcription factor atf1; heterodimeric transcriptional activator; binds M26 recombination hotspot; stimulated by the stress-activated Sty1p MAP kinase
SPAC1E11.03	serine/threonine protein kinase; yeast yak1 homolog
ntp1	neutral trehalase; subcellular localization of GFP fusion- Cytoplasm
SPAC4H3.03C	putative family 15 glycosyl hydrolase
SPAC31A2.12	hypothetical protein; similar to S. cerevisiae ROD; similar to S. pombe SPCC584.15C
ptc1	protein phosphatase 2c homolog 1
SPBC20F10.03	hypothetical protein; contains Pfam-B_11425; similarity to mouse and rat interferon-related developmental regulator 1- nerve growth factor; also conserved in C. elegans
isp6	sexual differentiation process protein; putative subtilase-type proteinase

Genes repressed by Sty1p during stress:

SPBC32H8.02C	located in cell periphery and cytoplasm; sequence orphan
SPAC3H1.06C	MFS drug efflux transporter of unknown specificity
SPBC1683.03C	MFS transporter of unknown specificity
SPAC1142.05	possible copper transporter
SPAC17A2.12	helicase; putative DNA repair protein
SPCC1223.09	uricase
SPAC24H6.10C	putative phospho-2-dehydro-3-deoxyheptonate aldolase
SPAC6C3.02C	hypothetical protein; similar to S. cerevisiae YMR002W
SPCC285.05	purine transporter
SPCC1672.03C	putative guanine deaminase
SPAC22A12.06C	similar to S. pombe dihydrofolate reductase Dfr1p
SPAC750.05C	hypothetical protein; low similarity to S. cerevisiae YHL017W is probably spurious; highly similar to S. pombe SPAC1348.02; highly similar to S. pombe SPBPB2B2.19c; possibly S. pombe specific
SPAC5H10.10	putative NADPH dehydrogenase
SPAC1002.18	hypothetical protein, conserved in other yeasts (Gaillardin et al)
SPAC1399.01C	putative purine permease
SPAC11D3.02C	ELLA family protein; putative acetyl transferase
SPAC8C9.10C	hypothetical protein; sequence orphan
SPCC330.04C	unknown function, Protein with Pfam-B_13958
SPCC4B3.06C	hypothetical protein; similar to S. cerevisiae YLR011W
SPBC776.16	hypothetical protein; sequence orphan
SPBC359.03C	putative amino acid permease
aru1	arginase
ipk1	Inositol 1,3,4,5,6-pentakisphosphate (IP5) kinase
ats1	putative N-acetyltransferase; closest S. cerevisiae homologs HPA2/ HPA3 are histone acetyltransferases; S. pombe sequence more similar to bacterial acetyltransferases; human homolog SAT is a spermidine/spermine N
ght5	MFS glucose transporter.
ura4	orotidine 5'-phosphate decarboxylase; similar to S. cerevisiae URA3
SPCC18B5.02C	cinnamoyl-coa reductase pseudogene
SPAC1F8.06	hypothetical serine-rich protein; similar to S. pombe SPAC8A4.02C and SPAC977.07C; putative cell surface protein; possibly S. pombe specific
SPBC1685.12C	very hypothetical protein
SPAC458.06	conserved hypothetical protein
SPCC965.14C	putative Cytidine and deoxycytidylate deaminase zinc-binding protein
SPAC212.06C	pseudogene; truncated C terminal of DNA helicase in rearranged telomeric region
SPBP8B7.10C	similar to S. cerevisiae YOR078W; may be involved in protein synthesis
SPBPB2B2.19C	duplicated at telomere; almost identical to SPAC1348.02 and SPAC756.05; contains 5 predicted transmembrane helices; possibly S. pombe specific
SPCC1827.06C	aspartate semialdehyde dehydrogenase
SPBC3H7.05C	hypothetical protein; sequence orphan; contains 7 predicted transmembrane helices contains predicted N-term signal sequence
gpd2	glycerol-3-phosphate dehydrogenase [nad+]
liz1	MFS transporter of unknown specificity
SPAC1399.02	MFS drug transporter of unknown specificity
SPBC19C2.04C	putative ubiquitin carboxyl-terminal hydrolase

Genes repressed by Sty1p, but not by Atf1p, during stress:

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SPAC1142.05	possible copper transporter
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SPBC776.16	hypothetical protein; sequence orphan
aru1	arginase
ats1	putative N-acetyltransferase; GNAT superfamily; closest S. cerevisiae HPA2 and HPA3 (histone acetyltransferases); S. pombe sequence more similar to bacterial acetyltransferases; human homolog SAT is spermidine/spermine N
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