

**SUPPLEMENTAL TABLE 2 – ChIP primers used**

<b>Primer #</b>	<b>Location</b>	<b>Primer Sequence</b>
ADH1 #1	Prom, - 235 Prom, -18	TTCCTTCCTTCATTCACGCACACT GTTGATTGTATGCTTGGTATAGCTTG
ADH1 #2	ORF, +146 ORF, +372	ACGCTTGGCACGGTGACTG ACCGTCGTGGGTGTAACCAGA
ADH1 #3	ORF, +844 ORF, +1013	TTCAACCAAGTCGTCAAGTCCATCTCTA ATTTGACCCTTTTCCATCTTTTCGTAA
ADH1 #4	3' UTR, +1231 3' UTR, +1400	ACCGGCATGCCGAGCAAATGCCTG CCCAACTGAAGGCTAGGCTGTGG
GAL1 #1	3' UTR, +1764 3' UTR, +2079	CCACAACTTTAAAACACAGGGAC CCTCCTCGCGCTTGTCTACTAAAATC
GAL1 #2	ORF, +1039 ORF, +1331	GAAGAGTCTCTCGCCAATAAGAAACAGG GAACATTCGTAAAGTTTATCGCAAG
GAL1 #3	ORF, +427 ORF, +726	CCGGAAAGGTTTGCCAGTGCTC CGGAGTAGCCTTCAACTGCGGTTTG
GAL1 #4	Prom, -190 Prom, +54	GGTAATTAATCAGCGAAGCGATG TGCGCTAGAATTGAACTCAGGTAC
GAL10 #5	Prom, -58 ORF, +182	TTAAACTTCTTTGCGTCCATCC TGCTTGGTCAAGACCTCTAACC
GAL10 #6	ORF, +823 ORF, +1108	TGTCGTGAGTGGAACCTGGGTT GCATATCTTCAGCGGAAAATCTGGC
GAL10 #7	ORF, +1869 3' UTR, +2078	TGTCAAGGCTTTTCATCCCGATTC TTGGACCCGTAAGTTTCACCGT
GAL10 #8	3' UTR, +2102 3' UTR, +2356	TTGCCAGCTTACTATCCTTCTT CGGCTCGTGCTATATTCTTGT
GAL10 #9	3' UTR, +2323 3' UTR, +2604	ATGGACGGTAGCAACAAGAATA TGGGTTAAGGAAAATGACAGAA
GAL7 #10	Prom, -203 ORF, +38	CAAAAAGCGCTCGGACAACT TGGGAATGGCTAGAAAAATCAA
GAL7 #11	ORF, +361 ORF, +655	AGCCCAATCATAATCTAACCAT TGACGTAATCGGCAAACAAAT
GAL7 #12	ORF, +813 ChrIX, +1019	CGCCTCGATTTTAAAGCAACTA CCTAACAATTCAAACCAACCA
PYK1 #1	Prom, -288 Prom, +19	CATATGATGCTAGGTACCTTTAGTGTCTTC CAATCTTTCTAATCTAGACATTGTGATGATG
PYK1 #2	ORF, +508 ORF, +799	GGTAAGATCTGTTCCACAAGGGTG CAAGTCACCTCTGGCAACCATAACAC

PYK1 #3	ORF, +1104 ORF, +1372	GAAACTGTACTCCAAAGCCAACCTCC GGGCTTCAACATCATCAGTCCAGTC
PYK1 #4	3' UTR, +1470 3' UTR, +1720	CTCCAACACTTTGCAAGTCTCTACCG GTTTCCAAAAATGCAACACCTCATCG
PYK1 #5	3' UTR, +1695 3' UTR, +1995	CGATGAGGTGTTGCATTTTTGGAAAC GACACTAATTGAATCTGCCCGCGTC
PMA1 #1	ORF, +168 ORF, +376	CGACGACGAAGACAGTGATAACG ATTGAATTGGACCGACGAAAAACATAAC
PMA1 #2	ORF, +1010 ORF, +1235	GTTTGCCAGCTGTCGTTACCACCAC GCAGCCAAACAAGCAGTCAACATCAAG
PMA1 #3	ORF, +2018 ORF, +2290	CTATTATTGATGCTTTGAAGACCTCCAG TGCCCCAAAATAATAGACATACCCCATAA
PMA1 #4	3' UTR, +2886 3' UTR, +3098	GATACACTAAAAAGAATTAGGAGCCAAC CAAGAAAGAAAAAGTACCATCCAGAG
PMA1 #5	3' UTR, +3287 3' UTR, +3500	GAAAATATTTGGTATCTTTGCAAGATG GTAAATTTGTATACGTTTCATGTAAGTG
PMA1 #6	3' UTR, +3448 3' UTR, +3662	GCGCCATACAGACACTCAAGATAC GGCCTGGCGATTTGTTTGCTTTCTTG
PMA1 #7	3' UTR, +3619 3' UTR, +3905	CTTCATCACAAGAAAGCAAACAAATCG CGTGATGAGTGAGTTAAGTTCTGCTG
Chr V	no ORF, Top no ORF, Bottom	GGCTGTCAGAATATGGGGCCGTAGTA CACCCCGAAGCTGCTTTCACAATAC