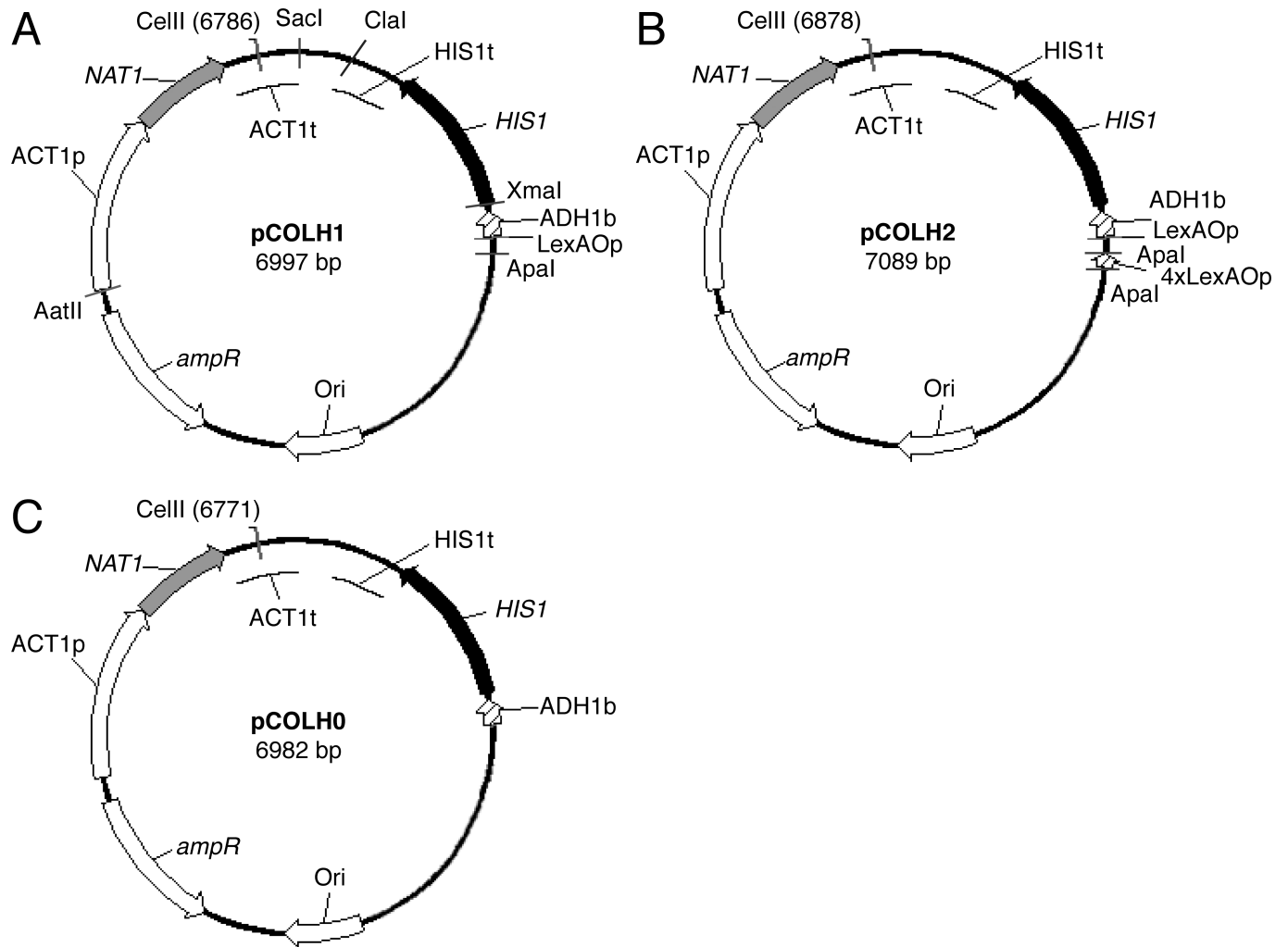


Supplementary Table 1. Primers

Primer name	Restriction site	Sequence
a) Construction of pCOLH0, pCOLH1, pCOLH2 and pC2H-HIS1		
HIS1-F	XmaI	<u>CCCGGGAACGTCATGGATTTAGTCAATCAT</u>
HIS1-R	Clal	AGTCATCGATCTAGAAAACCGTACCAGGTG
NAT1-F	AatII	ATCGGACGTCCAGAGCTATTAAGATCACCAG
NAT1-R	NarI	ATCGGGCGCCCTTCTAGAATTTTATGATG
HIS1IS-F	SacI	<u>AAGAGCTCATAGTCAGTAGCTAAACTG</u>
HIS1IS-R	SacI	<u>AAGAGCTCCTAGCCTATAGGTACCCTG</u>
b) Construction of pC2H-LACZ		
MPA-F	AatII	ATCGGACGTCTAGATGTTTATGATACTAAGG
MPA-R	NarI	ATCGGGCGCCTCTAGAACTCAGTATATCTTC
RPS1-F	PacI	ATCGTTAATTAACATGGCTGTCGGTAAAAAC
RPS1-R	MluNI	ATCGTGGCCATTAACAGATTCTAAAAACAAC
HIS1TLac-F	SacI	ATCGGAGCTCAAGAAGTGATAGTTTCTCATAAAT
HIS1TLac-R	SacI	ATCGGAGCTCCTAGAAAACCGTACCAGGTG
4xLexA-F	Apal	<u>GGGCCCTTGCGAATATTTGTTTCGTATTTGCGAATATTTGT</u> <u>TCGTATTTGCGAATATTTGTTTCGTATTTGCGAATATTTGTT</u> <u>CGTATGGGCC</u>
4xLexA-R	Apal	<u>GGGCCCATACGAACAAATATTCGCAAATACGAACAAATA</u> <u>TTCGCAAATACGAACAAATATTCGCAAATACGAACAAAT</u> <u>ATTCGCAAGGGCCC</u>
c) Construction of pC2HB		
MCSB-F	BspEI, SalI	<u>CTCCGGAAGATCTGGTGCTGGTGGTGGTGGTATCCGTATG</u> <u>ATGTGCCTGACTACGCAGGTGCTGGTGGTGGTGGTGCTA</u> <u>GCAGGCCTTTCGAAAGGCGCGCCGTCGACC</u>
MCSB-R	BspEI, SalI	<u>CCGGGTCGACGGCGGCCTTTCGAAAGGCCTGCTAGCA</u> <u>CCAACACCACCAGCACCTGCGTAGTCAGGCACATCATA</u> <u>GGATAACCAACACCACCAGCACCATCTTCCGGAGACG</u> <u>T</u>
LexA-F	AccIII	ATCGTCCGGAATGGGTGCTCCTCCAAAAAAGAAGAGAAA GGTAGCTAGAGAATTAACAAAACGACAAAGCG
LexA-R	BglII	ATCGAGATCTCATTCGCGGTACAAACC
MET3P-F	AatII	<u>GACGCTTCCAATTAGAGTTGTGGA</u>
MET3P-R	BspEI	<u>TCCGGAGTTTTCTGGGAGGGTATTTAC</u>
ACT1TB-F	SalI	ATCGGTCGACGGATCCGAGTGAAATTCTGG
ACT1TB-R	DraIII	ATCGCACGAAGTGTGAAGGGGGAGATTTTCAC
BIS-F	PvuI	ACTGTACGTAATCACCTTGTCTTGCCCGTG
BIS-R	SnaBI	ACTGCGATCGACCAGATCAAAATGCTAGAAC
NotIB	AseI	<u>TTCCATTAATGCGGCCGATTAATGGAA</u>
NewMCSB-F	AseI, BsaJI	<u>AAGGCGCGCCACGCGTAGTACTTAACCGCGGAAAA</u>
NewMCSB-R	AseI, BsaJI	<u>TTTTCCGCGGTTAAGTACTACGCGTGGCGGCCTT</u>
d) Construction of pC2HP		
MCSP-F1	AatII	<u>CTCCGGAATGGGTGCTCCTCCAAAAAAGAAGAGAAAGGT</u> <u>AGCTCCGCGGGCTAGCGGTGCTGGTGGTGGTGGTACTA</u> <u>CAAAGACCATGACGGTG</u>
MCSP-F2	SgrAI	ATTATAAAGATCATGACATCGATTACAAGGATGACGATG ACAAGGGAGCAGGAGGCGTCCGGAAGGCCTGAAGACGGG ATCGTTCGAACG
MCSP-R1	SgrAI	<u>CCGGCGTTCGAACGATCCCGTCTTCAGGCCTTCCGACGCC</u> <u>TCCTGCTCCCTTGTTCATCGTCATCCTTGTAAATCGATGTCA</u> <u>TGATCTTTATAATCACCGTCATGGTCTTTGTAGTC</u>

MCSP-R2	AatII	ACCAACACCACCAGCACCGCTAGCCCGCGGAGCTACCTT TCTCTTCTTTTTTGGAGGAGCACCCATTCCGGAGACGT
MET3P-F	AatII	<u>GACGTC</u> TCCAATTAGAGGTTGTGGA
MET3P-R	BspEI	<u>TCCGGAGT</u> TTTTCTGGGGAGGGTATTTAC
ACT1TP-F	SgrAI	ATCGCGCCGGTGTGAAGGGGGAGATTTTCAC
ACT1TP-R	BstBI	ATCGTTCGAAGAGTGAAATTCTGGAAATCTGG
PIS-F	PflMI	CCAAATATTGGTAACAGAGGTAACAACACTCTGG
PIS-R	ApaI	<u>GGGCCCTT</u> GAGAATTGGAAGAAGTTGGC
NotIP		GCGGCCGC
VP16P-F	SacII	ATTACCGCGGTCTCCAGGGCCCCG
VP16P-R	NheI	ATTAGCTAGCACCACCGTATTCATC
NewMCSP-F	StuI, EcoRI	AAAGGCCTCCC GGGAGGCGGCCACGCGTAGTACTTAAG ATATCCC
NewMCSP-R	StuI, EcoRI	GGGATATCTTAAAGTACTACGCGTGGCGGCCTCCCGGA GGCCTT
e) Cloning of bait and prey genes of interest		
INO4-F	NheI	<u>GCTAGCATGT</u> CGTCCGATAGTAGTTCACC
INO4-R	StuI	<u>AGGCCTT</u> TATTTTGC GATACACTACCATCG
INO2-F	NheI	<u>GCTAGCATGT</u> TCTTCCACTGTTACCC
INO2-R	StuI	<u>AGGCCT</u> CTACCTATTTTCTATTAATTTT
KIS1-F	NheI	ACTGGCTAGCCTCAGTCCAAAATCTAC
KIS1-R	SalI	ACTGGTCGACTCACAGTTGCTGCTCTTG
SNF4-F	StuI	ACTGAGGCCTACTGACATAGCACCACC
SNF4-R	BstBI	ACTGTTCGAACTAATCTTCTCCAAATAATATG
CEK1-F	NheI	<u>GCTAGCATGAT</u> GAATATAGATCAACATC
CEK1-R	StuI	<u>AGGCCTT</u> TATAATGGCTTCATAATCTC
CEK2-F	StuI	AAGGAGGCCTATGAAGAAATCTACTGGCCCCAC
CEK2-R	SalI	AAGGGTCGACTTACGACATGACTATTTTCG
CPH1-F	StuI	<u>AGGCCTATGT</u> CAATTACTAAAACATACAATGG
CPH1-R	StuI	<u>AGGCCT</u> CTATGTTTGTGACTGTTTACTTC
VP16Cterm-F	StuI	GGAGGCCTTCTCCAGGGCCCCGTTTCAC
VP16Cterm-R	AscI	AGGCGCGCCATTAACCACCGTATTCATCAATACC
CPH1Nterm-F	XmaI	TCCCCCGGGATGTCAATTACTAAAACATACAATGG
CPH1Nterm-R	NheI	AAGGAGCTAGCTGTTTGTGACTGTTTACTTCTTC
HST7-F	StuI	AAGGAGGCCTATGACAAGAACAACCTCGTATAG
HST7-R	BstBI	AAGGTTCGAATTACACTTTGCATTTTCTGATC
HST7-Y2H-F	XmaI	<u>CCCGGC</u> CATGACAAGAACAACCTCGTATAG
HST7-Y2H-R	XmaI	<u>CCCGGC</u> TTACACTTTGCATTTTCTGATC
CEK1-Y2H-F	XmaI	<u>CCCGGC</u> CATGATGAATATAGATCAACATC
CEK1-Y2H-R	XmaI	<u>CCCGGC</u> TTATAATGGCTTCATAATCTC



Supplementary Figure 1. (A) Plasmid pCOLH1 contains reporter gene *HIS1* with upstream a basal *ADH1* promoter and one LexAOp sequence. (B) Plasmid pCOLH2 contains reporter gene *HIS1* with upstream a basal *ADH1* promoter and five LexAOp sequences. (C) Plasmid pCOLH0 contains reporter gene *HIS1* with upstream a basal *ADH1* promoter and no LexAOp sequences. All three plasmids integrate at the *ACT1* terminator and *NAT1* is used for selective growth on nourseothricin. ADH1b: *ADH1* basal promoter; p: promoter; t: terminator; Ori: origin of replication.