

Table S3 Primers used in this study

Primer name	Sequence (5'-3')
TDH3-698	ataaaaaaacacgctttttcagttcg
15G-yGLuc	gggggggggggggggtagtcaccaccagcacc
15C-URA3-223	cccccccccccccaagctttcaattcatcttttttttttg
URA3-300c	tgttgtgaagtcattgacacag
KmARS7(201-260)-ADE2-797	caagacttctgaagtgaaaaccaactttcagcttcaaactaaaaatgaaaatc agtggcgctggcgcatctgttct
URA3+771c	ttcccagcctgcttttctgtaacgt
URA3+772	gcatatttgagaagatgcggccagc
TDH3-1c40	tttgtttgtttatgtgtgtttattcгааactaagttcttg
yGLuc+24c	agcgaacaagaccttgacacccat
yGLuc+27c	caaagcgaacaagaccttgacaccc
yGLuc+30c	gatcaaagcgaacaagaccttgaca
yGLuc+33c	acagatcaaagcgaacaagaccttg
yGLuc+36c	gatacagatcaaagcgaacaagacc
yGLuc+39c	agcgatacagatcaaagcgaacaag
yGLuc+42c	gacagcgatacagatcaaagcgaac
yGLuc+45c	agcgacagcgatacagatcaaagcg
yGLuc+48c	ttcagcgacagcgatacagatcaaa
yGLuc+51c	agcttcagcgacagcgatacagatc
yGLuc+54c	cttagcttcagcgacagcgatacag
yGLuc+57c	tggcttagcttcagcgacagcgata
yGLuc+60c	ggttggttagcttcagcgacagcg
yGLuc+63c	ttcggttggttagcttcagcgaca
yGLuc+66c	gttttcggttggttagcttcagcg
yGLuc+69c	ggtgttttcggttggttagcttca
yGLuc+72c	ttcgttgttttcggttggttagct
yGLuc+75c	gtcttcgttgttttcggttggttagct
yGLuc+78c	gaagtcttcgttgttttcggttggt
yGLuc+81c	ggtgaagtcttcgttgttttcggtt
yGLuc+90c	agcgacgatgttgaagtcttcgtt
yGLuc+108c	agcgaagtagaagcagcagcagc
yGLuc+12c-TDH3-1c	cttgacaccattttgtttgtttat
yGLuc+3-TDH3-1c	cattttgtttgtttatgtgtgtttattcga
yGLuc+4	ggtgtcaaggtcttgttcgctttga
yGLuc+7	gtcaaggtcttgttcgctttgatct
yGLuc+10	aaggtcttgttcgctttgatctgta
yGLuc+13	gtcttgttcgctttgatctgtatcg
yGLuc+16	ttgttcgctttgatctgtatcgctg
yGLuc+19	ttcgtttgatctgtatcgctgctg

yGLuc+22	gctttgatctgtatcgctgctgctg
yGLuc+25	ttgatctgtatcgctgctgctgaag
yGLuc+28	atctgtatcgctgctgctgaagcta
yGLuc+31	tgtatcgctgctgctgaagctaage
yGLuc+34	atcgtgctgctgaagctaagccaa
yGLuc+37	gctgctgctgaagctaagccaaccg
yGLuc+40	gctgctgaagctaagccaaccgaaa
yGLuc+43	gctgaagctaagccaaccgaaaaca
yGLuc+46	gaagctaagccaaccgaaaacaacg
yGLuc+49	gctaagccaaccgaaaacaacgaag
yGLuc+52	aagccaaccgaaaacaacgaagact
yGLuc+55	ccaaccgaaaacaacgaagacttca
yGLuc+58	accgaaaacaacgaagacttcaaca
yGLuc+61	gaaaacaacgaagacttcaacatcg
yGLuc+64	aacaacgaagacttcaacatcgctg
yGLuc+70	gaagacttcaacatcgctgctgctg
yGLuc+73	gacttcaacatcgctgctgctgctt
yGLuc+76	ttcaacatcgctgctgctgcttcta
yGLuc+79	aacatcgctgctgctgcttctaact
yGLuc+82	atcgtcgtgctgcttctaactteg
yGLuc+91	gctgcttctaacttegctaccaccg
yGLuc+100	aacttegctaccaccgacttgagcg
yGLuc+109	accaccgacttgagcgctgacagag
yGLuc+118	ttgagcgctgacagaggtaagtgc
Fc-yGLuc+45c	gaaagcgacagcgatacagatcaaagcg
Lc-yGLuc+45c	caaagcgacagcgatacagatcaaagcg
Ic-yGLuc+45c	gatagcgacagcgatacagatcaaagcg
Mc-yGLuc+45c	catagcgacagcgatacagatcaaagcg
Vc-yGLuc+45c	aacagcgacagcgatacagatcaaagcg
Sc-yGLuc+45c	ggaagcgacagcgatacagatcaaagcg
Pc-yGLuc+45c	tggagcgacagcgatacagatcaaagcg
Tc-yGLuc+45c	ggtagcgacagcgatacagatcaaagcg
Ac-yGLuc+45c	agcagcgacagcgatacagatcaaagcg
Yc-yGLuc+45c	gtaagcgacagcgatacagatcaaagcg
Hc-yGLuc+45c	gtgagcgacagcgatacagatcaaagcg
Qc-yGLuc+45c	ttgagcgacagcgatacagatcaaagcg
Nc-yGLuc+45c	gttagcgacagcgatacagatcaaagcg
Kc-yGLuc+45c	cttagcgacagcgatacagatcaaagcg
Dc-yGLuc+45c	gtcagcgacagcgatacagatcaaagcg
Cc-yGLuc+45c	acaagcgacagcgatacagatcaaagcg
Wc-yGLuc+45c	ccaagcgacagcgatacagatcaaagcg

TAA+(1-54)-hGLuc+49	atgatggcgcggtgggtgctctatttctgtacggccttcaggtcgcggcacct gccaagcccaccgagaacaacgaagacttcaacatcgt
SfGLU1+(1-54)-hGLuc+49	atgaaattcgggtgtttatccccgcttttctgtctattgttagtgctttacctgcca gcccaccgagaacaacgaagacttcaacatcgt
AmyLN5C+(1-27)c-TDH3-1 c	ggcgtaaagccgttttgtgtttcattttgtttgtttatgtgtgtttattcga
AmyLN5C+(28-69)-hGLuc+ 49	cgattgctgacgctgttatttgcgctcatcttctgtgctgccaagcccaccg agaacaacgaagacttcaacatcgt
KmPGU1+(1-54)-hGLuc+49	atgttattcagcaacacctattaatcgcagcagctagtgcattattagctgaag ccaagcccaccgagaacaacgaagacttcaacatcGT
hIL6+(1-33)c-TDH3-1c	tggaccgaaggcacttgtggagaaggagttcattttgtttgtttatgtgtgttta
hIL6+(34-69)-hGLuc+49	gttgcttctccctggggctgctcctgggtgtgctgtaagccaaccgaaaac aacgaag
hEPO+(1-21)c-TDH3-1c	aggacaaccgtgcacccccattttgtttgtttatgtgtgttta
hEPO+(22-60)-yGLuc+49	gcctggctgtggttctctctgctcctgctgctgctcctgctaagccaaccgaa aacaacgaag
hLIF+(1-30)c-TDH3-1c	gggcacaactcctgccgccaagaccttattttgtttgtttatgtgtgtttattcga
hLIF+(31-78)-yGLuc+49	ctgctgttggttctgactggaaacatggggcgaggagccccctccccgcta agccaaccgaaaacaacgaag
hZA2G+(1-18)c-TDH3-1c*	aggcaccattctaccattttgtttgtttatgtgtgttta
hZA2G+(19-51)-yGLuc+49*	gtcctgctgtctctgctgctgcttctgggtcctgctaagccaaccgaaaacaac gaag
3CG9-yGLuc+558c	cccggggcccttagtcaccaccagcacccttgatc
KmACT1+16	gcagaggctgctgcttttagttattg
KmACT+1111c	atggaccagattcgtcgtattcttg
hLIF+1	atgaaggtcttggcggcaggag
hLIF+694c	gcacctgaggagtgaattcc
3CG9-FLAGc-hLIF+609c	cccggggcccttacttgcgtcgtcgtcctttagtcgaaggcctgggccaaca cggcgtatg
URA3+772term3CG9	gcataattgagaagataaaaaactgtattataagtaaatgcatgtataactaaactc acaaattagacttcaatttaattatcagttaccgggccc
MKM(16)Ec-TDH3p-1c	ctccatcatcatcatcatcatcatcatcatcatcatcatcatcatcatcattttcattt tgtttgtttatgtgtgtttattcga
hLIF+4	aaggtcttggcggcaggagt

*hZA2G is an alternative name for hAZGP1.