

Primer	Sequence (5' to 3')
mChORFF-EagI	TCAC <u>CGGCCG</u> GATGGTTTCAAAAGGTGAAGAAG
mChORFR-MluI	TCA <u>ACGCGT</u> TTATTTATATAATTCATCCATACCACC
ACT1prF-BamI	TCATCAGGAT <u>CCCCAGCCT</u> CGTTTATAATAAACTTAGTC
ADH1-3'UTRR-ApaI	TCATCAGGG <u>CCCATCAAT</u> GCCAGAGATCAAACC
YPT72DISF	AACGATACAGAGTTTAATTTAATTTAATTCAATTCAATTTAATTA ATTAATCATATACACTTAATTTTCATGTGGAATTGTGAGCGGATA
YPT72DISR	AAACGAATTCGTGATATCTATTTGTTCTCTTTCTTCACCTGTGTA TTTGAAATTGATTGTTATATTATTATTTCCCAGTCACGACGTT
YPT72AMPR2	TCATCAGGATCCATTA ^{AAA} ACTGGTAATGATTTGG
YPT72DETF	TGTTGCATTTTATCGAGGGGC
YPT72DETR	CAGGCACTCGCCTCCGAGTCC
VPS11DISF	TTCTCCTTCGAGGACAATATCATCTCCTCTATTATCATTATCATCA TGGAGACAGTTCCAATTGTTTGATTGTGGAATTGTGAGCGGATA
VPS11DISR	CTATTGATCTTCCATTACACCTTTACCAATATAATCAGAAACAAA TTTAAATTTATCATTACTAGAATGCTTTCCCAGTCACGACGTT
VPS11AMPR2	TCAGGATCCAAATTACTGCATGAGAACACC
VPS11DETF3	GTAGTGGTACAACAGGAATCGG
VPS11DETR3	GATTTATCAATGCCATGATCGG
URA3INTF2	TTATACCATCCAAATCCCGCG
LUXINTDETF	CTGACCTTTAGTCTTTCTGCG

LUXINTDETR	CAGTAGTACTT <u>GTTGTT</u> GTATCG
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Engineered restriction sites are underlined.