

Universal Primer List

Primer Name	Sequence(5 - 3)	Primer Name	Sequence(5 - 3)
1100R	GGGTTGCGCTCGTTG	1492R	TACGGYTACCTTGTACGACTT
27F	AGAGTTTGATCMTGGCTCAG	337F	GACTCCTACGGGAGGCWGCAG
35S-A	AAGGGTCTTGCGAAGGATAG	35S-B	AGTGGAAGGAAGGTGGCT
518F	CCAGCAGCCGCGTAATACG	785F	GGATTAGATACCCTGGTA
800R	TACCAGGGTATCTAATCC	907R	CCGTCAATTCMTTTRAGTTT
a-Factor	TACTATTGCCAGCATTGCTGC	AD_Reverse	AGATGGTGCACGATGCACAG
Amp-R	ATAATACCGCGCCACATAGC	AOX1_Forward	GACTGGTTCCAATTGACAAGC
AOX1_Reverse	GCAAATGGCATTCTGACATCC	BEX2F	GGGCGCAGGACCYGRGGA
BEX2R	GGTCACTACCGKCCTCG	BEX3F	GGGCCAGGGTCTCACA
BEX3R	CCCACTGCCCTGGTACC	BGH-R	TAGAAGGCACAGTCGAGG
BINT1	GGAGGCCATCCCCGGCGACCTAT	Bluescript_KS	TCGAGGTCGACGGTATC
Bluescript_SK	CGCTCTAGAACTAGTGGATC	Bx1	GGGAGGAGCGAGGGGACCSCAG
CMV-F	CGCAAATGGGCGGTAGGCGTG	CYC1_Reverse	GCGTGAATGTAAGCGTGAC
DsRed1-C	AGCTGGACATCACCTCCCACAACG	DsRed1-N	GTACTGGAAGTGGGGGACAG
EBV-RP	GTGGTTTGTCCAAACTCATC	EGFP-C	CATGGTCTGCTGGAGTTCGTG
EGFP-CF	AGCACCCAGTCCGCCCTGAGC	EGFP-CR	CGTCCATGCCGAGAGTG
EGFP-N	CGTCGCCGTCCAGCTCGACCAG	EGFP-NR	CGTCGCCGTCCAGCTC
GAL1_Forward	AATATACCTCTATACTTTAACGTC	Gal4AD	TACCACTACAATGGATG
GLprimer1	TGTATCTTATGGTACTGTAAGTG	GLprimer2	CTTTATGTTTTTGGCGTCTTCCA
HCO2198	TAAACTTCAGGGTGACCAAAAAATCA	ITS1	TCCGTAGGTGAACCTGCGG
ITS2	GCTGCGTTCTTCATCGATGC	ITS3	GCATCGATGAAGAACGCAGC
ITS4	TCCTCCGCTTATTGATATGC	ITS5	GGAAGTAAAAGTCGTAACAAGG
Kan-R	ATCGCGAGCCCATTTATACC	KAN2-FP	ACCTACAACAAAGCTCTCATCAACC
KAN2-RP	GCAATGTAACATCAGAGATTTTGAG	LCO1490	GGTCAACAAATCATAAAGATATTGG

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LR0R	ACCCGCTGAACTTAAGC	LR16	TTCCACCCAAACACTCG
LR3	CCGTGTTTCAAGACGGG	LR3R	GTCTTGAAACACGGACC
LR7	TACTACCACCAAGATCT	M13-FP	TGTAAAACGACGGCCAGT
M13F	GTA AACGACGGCCAGT	M13F-pUC	GTTTTCCAGTCACGAC
M13FA	GTA AACGACGGCCAG	M13R	GCGGATAACAATTCACACAGG
M13R-pUC	CAGGAAACAGCTATGAC	mCherry-F	CCCCGTAATGCAGAAGAAGA
mCherry-R	TTGGTCACCTTCAGCTTGG	MT_Forward	CATCTCAGTGCAACTAAA
NS1	GTAGTCATATGCTTGTCTC	NS2	GGCTGCTGGCACCAGACTTGC
NS24	AAACCTTGTTACGACTTTTA	NS3	GCAAGTCTGGTGCCAGCAGCC
NS4	CTTCGTC AATTCCTTTAAG	NS5	AACTTAAAGGAATTGACGGAAG
NS6	GCATCAGACCTGTTATTGCCTC	NS7	GAGGCAATAACAGGTCTGTGATGC
NS8	TCCG CAGGTTACCTACGGA	pBABE-3	ACCCTAACTGACACACATTCC
pBABE-5	CTTTATCCAGCCCTCAC	pBacPAC-RP	GTCTGTAATCAACAACGC
pBAD-F	ATGCCATAGCATT TTTATCCA	pBAD-FP	ATGCCATAGCATT TTTATCC
pBAD-R	GATTTAATCTGTATCAGG	pDONOR-FP	TAACGCTAGCATGGATCTC
pEGFP_N	CCGTCCAGCTCGACCAG	pEGFP-FP	TTTAGTGAACCGTCAGATC
pEGFP-RP	AACAGCTCCTCGCCCTTG	pESP-RP	TCCAAAAGAAGTCGAGTGG
pET-24a	GGGTTATGCTAGTTATTGCTCAG	pET-RP	CTAGTTATTGCTCAGCGG
pFastBac_Forward	GGATTATTCATACCGTCCCA	pFastBac_Reverse	CAATGTGGTATGGCTGATT
pGEX3	GGAGCTGCATGTGTCAGAGG	pGEX5	GGCAAGCCACGTTTGGTG
pJET1_2F	CGACTCACTATAGGGAGAGCGGC	pJET1_2R	AAGAACATCGATTTTCCATGGCAG
pMalE	TCAGACTGTCGATGAAGC	pQE-F	CCCGAAAAGTGCCACCTG
pQE-R	GTTCTGAGGTCATTACTGG	pREP-fwd_primer	GCTCGATACAATAAACGCC
pRH_Forward	CTGTCTCTATACTCCCTATAG	pRH_Reverse	CAAAATTC AATAGTTACTATCGC
pTrcHis_Forward	GAGGTATATATTAATGTATCG	Puro-F	GCAACCTCCCCTTCTACGAGC

Primer Name	Sequence(5 - 3)	Primer Name	Sequence(5 - 3)
QE_Promoter	CCGAAAAGTGCCACCTG	RVprimer3	CTAGCAAATAGGCTGTCCC
RVprimer4	GACGATAGTCATGCCCCGCG	SP6	ATTTAGGTGACTATAG
STag_18mer_Primer	GAACGCCAGCACATGGAC	SV40-pArev	CCTCTACAAATGTGGTATGG
SV40-Promoter	GCCCCTAACTCCGCCATCC	T3	ATTAACCCTCACTAAAG
T7_17mer	AATACGACTCACTATAG	T7_EEV	ATGTCGTAATAACCCCGCCCCG
T7promoter	TAATACGACTCACTATAGGG	T7terminator	GCTAGTTATTGCTCAGCGG
U-19mer_Primer	GTTTTCCCAGTCACGACGT	WPRE-R	CATAGCGTAAAAGGAGCAACA