

New Standard Primers

Details of Changes

Name	Sequence	Replaces former Primers from GATC / Eurofins
-96gIII	CCCTCATAGTTAGCGTAACG	
1392r	ACGGGCGGTGTGTGTAC	16S-1392r (ACGGGCGGTGTGTTRC)
1492r	TACGGTTACCTTGTACGACTT	1492r (TACGGYTACCTTGTACGACTT)
27f	AGA GTT TGA TCA TGG CTC A	16S-27f (AGAGTTTGATCMTGGCTCAG)
5AOX	GACTGGTTCCAATTGACAAGC	
ACYCDuetUP1	GGATCTCGACGCTCTCCCT	
BAC1	GCTCGTATGTTGTGTGGAATTG	
CMVfor	CGCAAATGGGCGGTAGGCGTG	CMV-F (CGCAAATGGGCGGTAGGCGTG)
CMVmin	CGCCATCCACGCTGTTTTG	
DuetDOWN1	GATTATGCGGCCGTGTACAA	
DuetUP2	TTGTACACGGCCGATAATC	
EGFP-Cfor	GATCACATGGTCCTGCTG	pEGFP_C2-FP (GATCACATGGTCCTGCTG)
EGFP-Nrev	CCGTCCAGCTCGACCAG	pEGFP-N (CCGTCCAGCTCGACCAG)
EuVH	GGCAGCAGCCACAGGTAAGA	
EuVL	TTGCTGTTGCACAGTGATTC	
Gal4AD	TACCACTACAATGGATG	
Gal4BD	TCATCGGAAGAGAGTAG	
HuCAL_VH_for	GATAAGCATGCGTAGGAGAAA	
IgG_const_for	AGCCAGCAACACCAAGG	
IntXL39	ATTAGGACAAGGCTGGTGG	
M13rev-29	CAGGAAACAGCTATGACC	M13-RP and M13 rev (-29) (same sequence)
M13rev-49	GAGCGGATAACAATTTACACAGG	M13 rev (-49) (same sequence)
M13uni-21	TGTAAAACGACGGCCAGT	M13-FP and M13 uni (-21) (same sequence)
M13uni-43	AGGGTTTTCCAGTCACGACGTT	M13 uni (-43) (same sequence)
pBABE3	ACCCTAACTGACACACATTCC	
pBABE5	CTTTATCCAGCCCTCAC	
pBAD-FP	ATGCCATAGCATTTTTATCC	
pBakPAC-FP	TAAAATGATAACCATCTCGC	
pBR1	CGAAAAGTGCCACCTGAC	
pBR3	TCCCCATCGGTGATGTC	
pcDNA3.1-RP_1	CAAACAACAGATGGCTGGC	
pcDNA3_for	GGCTAACTAGAGAACCCACTG	pcDNA3.1-FP (CTCTGGCTAACTAGAGAA)
pcDNA3_rev	GGCAACTAGAAGGCACAGTC	
pCEP-Forward	AGAGCTCGTTTAGTGAACCG	
pCEP-Reverse	GTGGTTTGTCCAACTCATC	EBV-RP (same sequence)
PCR2	TTAGCTCACTCATTAGG	
pCR3.1-BGHrev	TAGAAGGCACAGTCGAGG	BGH-Reverse (same sequence)
pDONOR-FP	TAACGCTAGCATGGATCTC	
pDONOR-RP	GCAATGTAACATCAGAGAT	
pEGFP_for	TTTAGTGAACCGTCAGATC	pEGFP-FP (TTTAGTGAACCGTCAGATC)
pEGFP_rev	TTTAAAGCAAGTAAACCTC	pEGFP_C2-RP (same sequence)
pENTattL1for	TCGCGTTAACGCTAGCATGGATCTC	
pENTattL2rev	ACATCAGAGATTTTGAGACACGGGC	
pET-24a	GGGTTATGCTAGTTATTGCTCAG	

Name	Sequence	Replaces former Primers from GATC / Eurofins
petup	ATGCGTCCGGCGTAGA	
pEX-For	GGAGCAGACAAGCCCGTCAGG	
pEX-Rev	CAGGCTTTACACTTTATGCTTCCGGC	
pFASTBAC-F	TCCGGATTATTCATACCGTCCC	
pFASTBAC-R	CCTCTACAAATGTGGTATGGCTG	
pGexfor	ATAGCATGGCCTTTGCAGG	
pGEX-3	GGAGCTGCATGTGTCAGAGG	pGex rev (GAGCTGCATGTGTCAGAGG)
pGEX-5	CTGGCAAGCCACGTTTGG	
pGEX5-FP	AACGTATTGAAGCTATCCC	
pGLrev	CTTTATGTTTTTGGCGTCTTCC	
pGL3for	CTAGCAAAATAGGCTGTCCC	
pIRES-RP	TATAGACAAACGCACACCG	
pJET1.2for	CRACTCACTATAGGGAGAGCGGC	
pJET1.2rev	AAGAACATCGATTTTCCATGGCAG	
pJET1_fwd	GCCTGAACACCATATCCATCC	
pJet1-FP	ACTACTCGATGAGTTTTTCGG	
pJet1-RP	TGAGGTGGTTAGCATAGTTC	
pLKO1	GACTATCATATGCTTACCGT	
pMalE	TCAGACTGTGCGATGAAGC	malE (GGTCGTCAGACTGTGCGATGAAGCC)
Polyhed_fwd	AAAATGATAACCATCTCG	
pQEfor	GTATCACGAGGCCCTTTTCGTCT	
pQErev	CATTACTGGATCTATCAACAGGAG	
pQE-FP	CGGATAACAATTTACACAG	
pQE-RP	GTTCTGAGGTCATTACTGG	
pRSET-RP	ATGCTAGTTATTGCTCAGC	
pTeSp-1	CCTCCATAGAAGACACC	
pTrcHis-RP	CTGATTTAATCTGTATCAGG	pTrcHis rev (CTTCTGCGTTCTGATTTAATCTG)
SP6	CATTTAGGTGACACTATAG	SP6 (ATTTAGGTGACACTATAGAA)
T3	AATTAACCCTCACTAAAGGG	T3 (ATTAACCCTCACTAAAGGGA)
T7	TAATACGACTCACTATAGGG	T7-981079 and T7minus1
T7term	CTAGTTATTGCTCAGCGGT	pET-RP (CTAGTTATTGCTCAGCGG)
T7-pET-mod	CCCGCGAAATTAATACGACTCAC	
Topo-1	TCGGATCCACTAGTAACG	
U6_fwd	GAGGGCCTATTTCCCATGATTCC	
v5epitoperev	CGTAGAATCGAGACCGAGGAGAGG	