

Name	Sequence	Length	Tm [°C]	GC [%]
-96gIII	CCCTCATAGTTAGCGTAACG	20	57.3	50
1392r	ACGGGCGGTGTGTGTAC	17	57.6	65
1492r	TACGGTTACCTTGTACGACTT	22	56.5	41
27f	AGA GTT TGA TCA TGG CTC A	19	52.4	42
5AOX	GACTGGTTCCAATTGACAAGC	21	57.9	48
ACYCDuetUP1	GGATCTCGACGCTCTCCCT	19	61,0	63
BAC1	GCTCGTATGTTGTGTGGAATTG	22	58.4	45.5
CMVfor	CGCAAATGGGCGGTAGGCGTG	21	65.7	67
CMVmin	CGCCATCCACGCTGTTTTG	19	58.8	58
DuetDOWN1	GATTATGCGGCCGTGTACAA	20	57,3	50
DuetUP2	TTGTACACGGCCGCATAATC	20	57,3	50
EGFP-Cfor	GATCACATGGTCCTGCTG	18	56	55.6
EGFP-Nrev	CCGTCCAGCTCGACCAG	17	60	70.6
EuVH	GGCAGCAGCCACAGGTAAGA	20	61.4	60
EuVL	TTGCTGTTGCACAGTGATTC	20	55.2	45
Gal4AD	TACCACTACAATGGATG	17	47.9	41
Gal4BD	TCATCGGAAGAGAGTAG	17	50.4	47
HuCAL_VH_for	GATAAGCATGCGTAGGAGAAA	21	55.9	43
IgG_const_for	AGCCCAGCAACACCAAGG	18	58.2	61
IntXL39	ATTAGGACAAGGCTGGTGG	19	56.7	53
M13rev-29	CAGGAAACAGCTATGACC	18	53.7	50
M13rev-49	GAGCGGATAACAATTTACACAGG	24	61.0	46
M13uni-21	TGTAAAACGACGGCCAGT	18	53.7	50
M13uni-43	AGGGTTTTCCCAGTCACGACGTT	23	62.4	52
pBABE3	ACCCTAACTGACACACATTCC	21	57.9	48
pBABE5	CTTTATCCAGCCCTCAC	17	52.8	53
pBAD-FP	ATGCCATAGCATTTTTATCC	20	51.1	35
pBakPAC-FP	TAAAATGATAACCATCTCGC	20	51.1	35
pBR1	CGAAAAGTGCCACCTGAC	18	56	55.6
pBR3	TCCCATCGGTGATGTC	17	55.2	58.8
pcDNA3.1-RP_1	CAAACAACAGATGGCTGGC	19	56.7	52.6
pcDNA3_for	GGCTAACTAGAGAACCCACTG	21	59.8	52
pcDNA3_rev	GGCAACTAGAAGGCACAGTC	20	59.3	55
pCEP-Forward	AGAGCTCGTTTGTAGTGAACCG	20	57.3	50
pCEP-Reverse	GTGGTTTGTCCAACTCATC	20	55.2	45
PCR2	TTAGCTCACTCATTAGG	17	47.9	41.2
pCR3.1-BGHrev	TAGAAGGCACAGTCGAGG	18	56.0	56
pDONOR-FP	TAACGCTAGCATGGATCTC	19	54.5	47.4
pDONOR-RP	GCAATGTAACATCAGAGAT	19	50.2	36.8
pEGFP_for	TTTAGTGAACCGTCAGATC	19	52.4	42.1
pEGFP_rev	TTTAAAGCAAGTAAACCTC	20	49.1	30
pENTattL1for	TCGCGTTAACGCTAGCATGGATCTC	25	64.6	52
pENTattL2rev	ACATCAGAGATTTTGAGACACGGGC	25	63.0	48
pET-24a	GGGTTATGCTAGTTATTGCTCAG	23	58.9	43.5

Name	Sequence	Length	T _m [°C]	GC [%]
petup	ATGCGTCCGGCGTAGA	16	54.3	62
pEX-For	GGAGCAGACAAGCCCGTCAGG	21	65.7	67
pEX-Rev	CAGGCTTTACACTTTATGCTTCCGGC	26	64.8	50
pFASTBAC-F	TCCGGATTATTCATACCGTCCC	22	60.3	50
pFASTBAC-R	CCTCTACAAATGTGGTATGGCTG	23	60.6	48
pGexfor	ATAGCATGGCCTTTGCAGG	19	56.7	53
pGEX-3	GGAGCTGCATGTGTCAGAGG	19	58.8	57.9
pGEX-5	CTGGCAAGCCACGTTTGG	18	58.2	61.1
pGEX5-FP	AACGTATTGAAGCTATCCC	19	52.4	42.1
pGLrev	CTTTATGTTTTTGGCGTCTTCC	22	56.5	41
pGL3for	CTAGCAAATAGGCTGTCCC	20	57.3	50
pIRES-RP	TATAGACAAACGCACACCG	19	54.5	47.4
pJET1.2for	CGACTCACTATAGGGAGAGCGGC	23	66.0	61
pJET1.2rev	AAGAACATCGATTTTCCATGGCAG	24	59.3	42
pJET1_fwd	GCCTGAACACCATATCCATCC	21	59.8	52
pJet1-FP	ACTACTCGATGAGTTTTCCGG	20	55.3	45
pJet1-RP	TGAGGTGGTTAGCATAGTTC	20	55.3	45
pLKO1	GACTATCATATGCTTACCGT	20	53.2	40
pMalE	TCAGACTGTCGATGAAGC	18	53.7	50
Polyhed_fwd	AAAATGATAACCATCTCG	18	46.9	33
pQEfor	GTATCACGAGGCCCTTTCGTCT	22	62.1	55
pQErev	CATTACTGGATCTATCAACAGGAG	24	59.3	42
pQE-FP	CGGATAACAATTTACACAG	20	53.2	40
pQE-RP	GTTCTGAGGTCATTACTIONG	19	54.5	47.4
pRSET-RP	ATGCTAGTTATTGCTCAGC	19	52.4	42.1
pTeSp-1	CCTCCATAGAAGACACC	17	52.8	52.9
pTrcHis-RP	CTGATTTAATCTGTATCAGG	20	51.1	35
SP6	CATTTAGGTGACACTATAG	19	50.2	37
T3	AATTAACCCTCACTAAAGGG	20	53.2	40
T7	TAATACGACTCACTATAGGG	20	53.2	40
T7term	CTAGTTATTGCTCAGCGGT	19	54.5	47
T7-pET-mod	CCCGCGAAATTAATACGACTCAC	23	60.6	48
Topo-1	TCGGATCCACTAGTAACG	18	53.7	50
U6_fwd	GAGGGCCTATTTCCCATGATTCC	23	62,4	52
v5epitoperev	CGTAGAATCGAGACCGAGGAGAGG	24	66.1	58