

Primer Set	Pool	Concentration (uM)	FWD sequence	REV sequence	Forward Primer Reference Position	Reverse Primer Reference Position
1	1	40	TCCTGAGTACGGCTGAGTGAAG	CCTTCTTTGTTCTCCAACAGGCT	1115-1137	1475-1498
2	2	30	ATTAGGCTACGACCCAACGGAA	AATTTCTCTCTGCGGCTAGA	1363-1385	1711-1733
3	1	20	ACCTAGTGGTGGAAACAGGAACA	GTCTTGTTGTGGAGCTGGTTG	1628-1651	1987-2009
4	2	20	CCATCAAGCGGCTATGCAGATT	GCTGCATCTGTCTGTTCTGCTC	1893-1915	2246-2268
5	1	40	TGGGGTTGCAAAATGTGTCAGA	GCTTTCTTGGTCCCCTCTGTTG	2114-2137	2476-2498
6	2	20	GCCGGGACAGAAGGCTAGATTA	TTTAGCAGATCCACAGCTGGGT	2376-2398	2720-2742
7	1	30	AATTTCCCATGGCTCAAGTGC	TCCTATTCCTCCTACTATTTTTGGGGT	2644-2666	2963-2990
8	2	10	TGGATACAGGGGCTGATGATTCT	CCAAGTACCATCCTTTCCATCTTT	2881-2904	3249-3275
9	1	20	GTCGCCTTAAAGCCAGGAAAGG	GAGGTATGGAGAAATATGCATCACCT	3135-3157	3465-3491
10	2	10	TAGGAATACCACCCCTGCAGG	CCCTGTCATGTTCCAGGTCTGT	3382-3404	3708-3730
11	1	40	ACATGTGCTAGAACCCTTCAGGA	CAGTCCACTGAACTTCCTCTGTT	3602-3625	3996-4019
12	2	10	AGAGAGACCTGGACAGTGAATGA	TGTGCTAATAGTCTCACTCCATTGGT	3852-3875	4213-4239
13	1	40	GAGTCAGGACAATCAGTGGTCTTAT	ATCCTGCTTCCCTCTTTTGACTG	4100-4125	4473-4498
14	2	10	TCTCAACACCACCGCTAGTAAGA	TACCTTTGTGTGCTGGTACCCA	4354-4377	4728-4750
15	1	15	CCTACAGAATCAGAGAGCAGGCT	ATTTGCCTGCCATGTATAGCC	4632-4655	4949-4971
16	2	15	CCCAGAATAGTGCCAGACAGA	CAAAGGTGTGCTCTATCCCTGC	4869-4891	5202-5224
17	1	30	ATGGTGCTAACTTTGCTTCGCA	GGTCCCTTCCACAGTTGATCTCT	5134-5156	5497-5520
18	2	30	AGGGGATATGACTCCAGCAGAA	CCATCCGACCTTAAAATGGGGC	5357-5379	5742-5764
19	1	25	AGAGGTGGATAGCAGTCCAC	CTTCTCCGCTGTAAAGCAAGG	5609-5631	5959-5981
20	2	50	GGACAGATGTAACACCAAATATGCA	TTCCAAGACCTTTGCCAAACC	5885-5911	6229-6251
21	1	70	GGAGAAGAGACAATAGGAGAGGCC	CATCCCATGGTTCCCTTTGTGG	6107-6131	6458-6480
22	2	10	TCCCCCTCAGGACTAGCATAA	TGCTTCTAGAGGGCGGTATAGC	6385-6407	6704-6726
23	1	30	TCATGCATTTTCAAGGCGGATG	GCATTCCTCCAAGCTGGTACAC	6615-6637	6972-6994
24	2	10	TTGGGAATCAGCTGCTTATCGC	TGTCAATCCCATCTATCTGTCTCA	6870-6892	7231-7256
25	1	25	AGTCACAGAACAGGCAATAGAGGA	TCCCTTGTTCACATACCAAATCTGC	7102-7126	7451-7476
26	2	40	AGCTGTAAATTCAACATGACAGGGTT	GTCTATTATCCCTACCATGCCAGT	7358-7384	7749-7774
27	1	20	GGTGGTCTTTCATGCACAAGG	TGGGATGTTTGACAATGGTCTGC	7633-7655	7955-7978
28	2	20	AAAGCAGGCATGGTGTGGTTT	TGGAGTTACCGTGAGGTCTCC	7876-7898	8243-8265
29	1	20	CCAGAAGCCAAAGGAACAGCAT	TTGCGAGAAAACCCAAGAACC	8113-8135	8471-8493
30	2	20	AGATGTGAAGAGGTACTACTGGT	CTAAACGCACATCCCAAGCAT	8389-8414	8700-8722
31	1	15	TAGGGTCACTGCCATCGAGAAG	TCCAAGAAGCAAGGTCAAACCA	8632-8654	8921-8943
32	2	40	AGCTGGGATGTGTTGGCAATT	GCCAAGTCAAGAGGCGTATCAG	8879-8901	9206-9228
33	1	100	GGCAAAGAAAGAGACGGTGGAG	AAGAGAGTGAGCTCAAGCCCTT	9101-9123	9495-9517
34	2	40	CTGGAGATCTGCGACAGAGACT	GTAATAAATCCCTTCCAGTCCCC	9370-9392	9734-9758
35	1	40	TTGGTAGGGGTATCAGTGAGGC	CCATGCTAGAACCTCTCCCCAA	9621-9643	9971-9993
36	2	40	CAGATGAGGCACAGGAGGATGA	ACATCAAGAAAGTGGGCGTTCC	9880-9902	10205-10227
37	1	40	AGGGACTTCCACAAGGGGATG	TCGGTTTCCCAAGCAGAAAGG	10131-10153	10516-10538