Promoter-Construction	**Template**	**Primers (Forward and Reverse)**	**PCR fragment size**	**Cloning vector (Restriction Sites)**
T7-EGFP	pEGFP-N3	**Pr1_Fw**: aacagcccatggtgagcaagggcgaggagctg **Pr2_Rv**: cttcttctcgagttacttgtacagctcgtccatgccgag	740 bp	pET28a (NcoI/XhoI)
T7-EGFPΔG4	pEGFP-N3	**Pr3_Fw**: aacagcccatggtgagcaaggaggagctgttcaccggg **Pr4_Rv**: cttcttctcgagttacttgtacagctcgtccatgccgag	737 bp	pET28a (NcoI/XhoI)
Trc-EGFP	pEGFP-N3	**Pr5_Fw**: cacaggaaacagcatatggtgagcaagggcgaggagctg **Pr6_Rv**: cttcttctcgagttacttgtacagctcgtccatgccgag	747 bp	pJOQ (NdeI/XhoI)
Trc-EGFPΔG4	pEGFP-N3	**Pr7_Fw**: cacaggaaacagcatatggtgagcaaggaggagctgttcaccggg **Pr8_Rv**: cttcttctcgagttacttgtacagctcgtccatgccgag	744 bp	pJOQ (NdeI/XhoI)
Trc-rsGFP	pQBI25-f	**Pr9_Fw**: cacaggaaacagcatatggctagcaaaggagaagaactc **Pr10_Rv**: cgggctctcgagttatcagttgtacagttcatccatgccatg	750 bp	pJOQ (NdeI/XhoI)
Trc-rsGFPΔG4	pQBI25-f	**Pr11_Fw**: cacaggaaacagcatatggctagcaaagaagaactcttcactggagttgtc **Pr12_Rv**: cgggctctcgagttatcagttgtacagttcatccatgccatg	747 bp	pJOQ (NdeI/XhoI)
Trc-EGFPecol	pQBI25-f	**Pr13_Fw**: cctttccgcttgcatatggtgagcaaaggagaagaactcttc **Pr14_Rv:** gcattgaacaccataggtcagagtagtgactag **Pr15_Fw:** ctagtcactactctgacctatggtgttcaatgc **Pr16_Rv:** ttcaatgttgtggcgaatcttgaagttcacttt **Pr17_Fw:** aaagtgaacttcaagattcgccacaacattgaa **Pr18_Rv**: cgggctctcgagttatcatttgtacagttcatccatgcccagtgtaatcccagcagctgttac	228 bp 339 bp 249 bp (750 bp)	pJOQ (NdeI/XhoI)
Trc-EGFPecolΔG4	Trc-EGFPecol	**Pr19_Fw:** ctcaggacacagcatatggtgagcaaagaagaactcttcactggagttgtc **Pr20_Rv**: cgggctctcgagttatcatttgtacagttcatccatgcccagtgtaatcccagcagctgttac	747 bp	pJOQ (NdeI/XhoI)

Pr2_Rv = Pr4_Rv = Pr6_Rv =Pr8_Rv
Pr10_Rv = Pr12_Rv
Pr18_Rv = Pr20_Rv