

Přílohy

Primery použité k NGS

Gen	Sekvence 5'-3'	Sekvence 3'-5'	Délka amplikonu
BDNF	CTAAGCCAGCGCCCGAAAC	CAGCTCTTCTAGTGAGAAGGATTCA	238
BDNF	GGGTCCACACAAACCTCAC	GGCCGGACAGAGGAGC	190
BDNF	AGCCATGATTTACCCAAATGTTTC	CTGGTCTCCTCTTTGTTTTGTTT	218
BDNF	AACTGAATAATTTACCCTGTTATGT	CATAGACAAAAGGCATTGGAAC	235
BDNF	ACAATTGCTGGATGTGTCTCCTA	TGAAAGAAGGAAGACTTTACGTAGT	233
BDNF	GCTTTCTAGCCGACCAACAGAA	GCTGCAGAACAGAAGGAGTACAT	225
BDNF	TGGTCACAACGAAAATGTTTTGCT	GCCAAGGAGTGAAGGGTTTTAGA	230
BDNF	TCATGCTACTGGCAATGTGGATT	TGACTACTGATAAGCATGGAGA	235
BDNF	CCCTGGTTTCTCAAGTCTAGTT	AGTGCCTACATTATCTAACTGTGCTC	231
BDNF	TGTCATGAAAACAATGTGTCTGG	AGAAAAGTTGAGCTTTCTTAGC	187
BDNF	TGTTAAGCAACATGTCAAGATT	GCGGATATTGCAAAGGGTATT	231
BDNF	CTGTTTTCTGAAAGAGGACAGTT	ACAGTACAGTGGTTCTACAATCT	220
BDNF	AGTCTTAAAGCAAGGAACACACG	AAACTTAACCGCATCAGAAGCCTA	216
BDNF	ACATAGGTCCTTCGTCAAAAGC	GAGATGGGGAAAAGAATCCTACAA	226
BDNF	GACTAACCCGAGTCAAGAATCCC	CCAGGAATTGAGACTGAAGTGGAC	238
BDNF	ACTAAGCAACAACGCAAGTGTA	AGAGTATCTCTGTGGGAAGAGGA	222
BDNF	TTTTCAGGTGTGAAATGGGCTGA	GGTGCCTTGTCAATGGAAGAG	236
BDNF	TGGGCCTTTCTAGCTATGTGATG	GGGAGCCCTGAGTTCTAACATAAA	238
BDNF	TTCTGAATGTCCAGACACCTTCC	TGTGGCATTGTGATCATGAGATT	232
BDNF	GCTTCGAGGGGTGTTCCAG	GCGGGAAGGTGTGTCTGTG	227
BDNF	AGCAGATATTTCAAGCATTCTTAC	AGGGTGAGGGATAAGCAAAGAG	233
BDNF	CAGCCTTCATGCAACCAAAGTAT	GTCTGGTGCAGCTGGAGTTTAT	225
BDNF	AAGTTGTGCGCAAATGACTGTTT	ACCATTAAGGGGAAGATAGTGGA	226
BDNF	CCCAGAACCCCGAGATTTTAT	TTCACCATGTAACAAAACAAAACAA	222
BDNF	TTGCAAAATCTTTCACAACATACA	TTTACTGCTTTTAACTTCTGATAGCG	240
BDNF	GGAAAAGGAACGTGTTTTCTGC	TTCTAAGCATGTTTGTATTCTG	240
BDNF	CCCCTAAGCCAGTAAAGCAATGA	TTGCATGTGCTTTAATTGTGAA	233
BDNF	AATTCCACTGAAACGTGGAGGTA	GGATGCAAAAATAACCCGAAGGAC	221
BDNF	CCCCAAAACCCACACTCTATT	GCGAACTAGCATGAAATCTCCCT	224
BDNF	CAAGGCCACAGACATTTACTTAC	TCAGTGGCTCTTTTACCAATAA	239
BDNF	TGGCTTTTTCTGCTAATACACAC	GGGACTTCCAGGAAACTTCAA	216
BDNF	AAAGGCACCTGACTACTGAGCAT	AGAAGCAAACATCCGAGGACAAG	240
BDNF	GGCCTGCCGAGAGTGT	TAGGCTAGGAATCCTTCCAGGG	239
BDNF	CCCACTACGAGGTACAC	TCACGTAAACAGCGAGGTTAGTC	233
BDNF	CCATCCTGTTTTACAGGTAAGGAA	CATGACATTAGCCCCAGGCATAG	216
BDNF	AGAGGCCTTCGTTTTGGAATGTC	TGTAAGCAAAGGTAGAGCCCTG	216
BDNF	ACAGGTGATGGAAGAACTGGAA	CATGACTGTGCATCCAGAAGA	230
BDNF	CGGGCCTGTCTCAC	GAGTTGAAAGCTAGGGGAGC	240
BDNF	CTCTCCATGAACAGACAGGATGG	TTGGTAAGGGAAAAGGGAACAGG	234
BDNF	GTCCGAAAGGACCTTCTACTCC	gctGGATGCTTCTTTCTGGTTT	231
BDNF	TACAATAGGCTTCTGATGCGGTT	CTTTGTGGCAGAAGTGTTCAT	226
BDNF	CATGCAGTGTTCCTCCAAAGAT	TGCAAAGGACCATGTTGCTAACT	239
BDNF	GGATCCCCAGTCAACTCTCTC	GGGCTTTAATGAGACCCACC	217
BDNF	CCCAACTGCCTGCTATGTA	CAAGGCGAAGGTTTTCTTACCTG	240
BDNF	CTGCAGCTTCTTTTGTGTAACC	GATGCTGCAACATGTCCATGAG	239
BDNF	CGGTGGGTGTCTCATAAAGCC	TTTGACCAATCGAAGCTCAACCG	227
BDNF	CGAAAGTGTACGCAATGATGTC	CTTCTCCCTACAGTTCACCAG	225

BDNF	CTCAATGAGGGGACCAA	GATAGGAGTCCATTAGCACCTT	222
BDNF	ATTGATAGTGAAATTCATGGC	GCACTAGAGTGTCTATTTGAGGC	238
BDNF	CCATGTGGCCATCTGATTGTAA	AAGAAGGAGAACTGAAGCCACA	232
BDNF	ACAAATTGAAATCTTTGCTGATGGT	GGGAGCCTGAAATAGTGGTTCTT	236
BDNF	CTCTCATGCTCTCCCTCTACCT	CTAGCGGTGTAGGCTGGAATAG	234
BDNF	CGCTCCCCTAGCTTTCAACTC	TGCGCGAACTCTGTTTAAGCC	216
BDNF	TGCTGGAAGGTAATGTGTCTTGT	TCAAAGGAAACGTGTCTCTCGG	233
BDNF	ACTGTTCTCCATGCTTATACGAGT	CTCATTCTTTGGCGTTGGTCAG	216
BDNF	AAGACGCTTTTTAAGGGCGACAC	GTGACAAACCGTAAGGAAGTGGGA	232
BDNF	CGCCGTACCCTACTACTAATAC	GTTGCGCCAATGAAGAAAACAA	222
BDNF	gcgctaccgATACCCGTTC	CTGGAACACCCCTCGAAGC	208
BDNF	TGTAATCTCCCCTTCTTCTGCC	AACCCACATCTCTACCCATCTG	230
BDNF	ATTATCCAGAGGTGGGATGGTGG	TGCGGGAGGAATTTCTGAGTG	240
BDNF	GCATTTCCAAAGTTAACCCAGT	ACCCTGTGTGCGCCTTAAAAA	217
BDNF	CCATACAGAAGCGTGTGGGTAG	TCCTCTGCATTAAGCTACAAGT	239
BDNF	CAGAAGACAAAGCAACTGGCATC	ACTCCTCACTTTCTCTGGGAACT	229
BDNF	TCTTCCCCTTTAATGGTCAATGT	TCCCTGTATCAAAGGCCAACTG	233
BDNF	ACAAAGAGACCACAGCAAGACTTTA	AACAACCCCATCAAAACTTA	238
MC4R	AATTGAAAGCAGGCTGCAAATGG	TAGCCAAGAACAAGAATCTGCAT	224
MC4R	CATGCCTGCTGTGAGTAAATGTC	GCATGGCAGCTTCAAGGAAAAT	223
MC4R	TTATTAAGAACCAGCCAGTAGT	GCACCTTTTTCTAGCTATAACAC	205
MC4R	ATGGATATTCTCAACCAGTACCC	ATCGATCTCTGATTTATGCACT	234
MC4R	ACAGTGCCTACAACCTATAACAT	CACGCAATATAGGAACATGCATAA	233
MC4R	CTCACATCGCTGCCTCTC	GCTTCTGCATCTGAATCTGTGTC	235
MC4R	CTGCTCTTGCTTTCTTGTCTC	TCCACACATCATAAAAATCAGCA	221
MC4R	AAGGTAATCGCTCCCTTCATATT	ATAAGTTGATCTGGGCAGCTT	233
MC4R	AAGTCTGAAGTCGAGAAGCAAGC	AGTATCTAGCCAAAAGCAGAGT	229
MC4R	GGTGAAGAACATGGTGTGAGG	GTGATCTGTAGCTCCTTGCTT	228
MC4R	CTATGTTACGAAAGCACGCAAAGT	TGCCAGTCTCTGTATTATTTCCA	180
MC4R	GCTTCTTCTGACCAATCCAAT	GCTGCCTGAAGATAGCTTGTTTT	226
MC4R	CACAAACACCTCAGGAGAGACAA	AGGAAATAACTGAGACGACTCCC	231
MC4R	CATTTGAAACGCTCACCAGCATA	GCAGTTACAGACTGCACAGCAA	237
MC4R	AGACATGGTATGGCTGGAAGAAC	CAGAATGCAGCTTATTATTTCTGA	217
MC4R	TGCTCTGTCCCCATTTAATATCT	CCCCATTCTTCTCCACTTA	238
MC4R	CACGGTGGGTGGAGTTC	AGCCAAGATTTAAAGTGATGATG	240
MC4R	AGACATCATGTGTGTGTAAAATGA	ACATAGTTCTTCCAGCCATACCA	218
MC4R	CATGAAGCACACAAATATGGAT	TTCTCTATGTCCACATGTTCC	217
POMC	GAGACGTCTCGCGCTTC	CACGGGCTGCCCTCAT	227
POMC	CCAACTATCTGATCCCTCGAA	TAAAGCTGCCTGTAGTTAGGAAA	238
POMC	ATGGCAGGATTTGAAGAGGATGC	CCACTGGTTAAGGGAAATTTGG	240
POMC	CTTCTTGCCACCCGGCTT	AAGTACGTATGGGCCACTTC	234
POMC	GGACGGGGACAGGGGAT	CTGCCGGGAAGGTCAAAGTC	218
POMC	AGAGCAAGGGGCTTTGGG	GTGGCGGCCGAGAAGAAG	236
POMC	CAGCTCCAGTCCCATCTAATGTC	CCCACCAATCTTGTTGCTTCTG	232
POMC	CTTCTTCCCCTCCTTCGC	CGCCTTCGGCAGGACAG	232
POMC	TCAAAGTCAGAGGTGGATGTGAAA	GCTGTTCAAAAACGCCATCATCA	228
POMC	CGTCGGGGCCATCTCC	CCCCGCAGCGATGGTG	211
POMC	GGGAGACCGCTGGAAAG	TATTTACCGCAAATGCGAACC	240
POMC	CAGCGGAAGTGCTCCATCC	GGCGCCAGTGAAGGTG	223
POMC	AGGTTGCTTCCGTGGTGTGAG	AATGTTGGCTCAAGGTCCTTCT	234
SIM1	CTGCTCTTTCGAGGGATTTA	GCGCCTGCATTATCCT	240
SIM1	TTGAAACCACAAAGAAGTCAGTTT	ATTGCATTGTGACTATCCAGTGA	224

SIM1	TTTAAAGTAGGTATGGCAGGTGA	TGCCAAATGTTGAGAAACATGAA	217
SIM1	CCATCTCTTAACTGTTCTCATGC	GGAATAATAGTGAAGTGGCTTCG	227
SIM1	CCCAAAATGGTCAAATGCTCAAT	ATACAAGATTTCCGCCCTCAAGA	218
SIM1	GCTTTCCTTAGCTGTAATCTGTT	AGGGTGAATTCTGAGATCACTTA	237
SIM1	TGAGGATCACTGGATAGTCACA	ACATAGCACTAGCATTGAGGGTT	196
SIM1	GTCATCAAATGCTTAACTTACCT	AGATTGACCCAGTTTTAAGTACA	185
SIM1	CAATGCTACTAAAACTATACATTGC	AAGAAATGAAATTGGCCTGGAAA	240
SIM1	ATTGAGCCTAGTATCTGAAGTGT	GACATCAGTGACCAGAGAAATG	225
SIM1	ACTTTCAATTAGAGATCATATTTGGC	GATACCCTGAACGACCCCTC	226
SIM1	AGCAAAGTTGCAATAACAAAACT	CTCAGAGTTCCTTCTCTAGGAC	240
SIM1	AAACCACACATAAGTGAATCTGC	ATAGGTAGGTGACAATCTGAACG	230
SIM1	TCAAGTCTGCCAGTACATCCC	GCTCCTTAGGCAGGTTACAGTT	229
SIM1	AGCACTATCTGTACAACCCCTTAC	TTATGTAATGTCAAGTTTTGTGAA	231
SIM1	TTCAGTGTTCCTTCTCCCTCC	GCTGGTCTTTACTGCCATCAAT	229
SIM1	ATCTTCTGTGTGAAATCCCGAA	AAAGGATTTCTTTGGAAGCATCATA	240
SIM1	ATCATTATGAAAACAGGATGTATGT	TTCAGTGAATCTATCAGTCTCA	233
SIM1	ACCCAGTACTCATCTTTCATATT	GCCAAAACTCTTGCCATATTTA	186
SIM1	CTGCGGTCCGAAAACGTGC	GCTGTTTTTTGTTTGTGTTTTCTC	239
SIM1	TGGTCACCCCTGGTATAAACAG	CCAATCAAATGCAGATAAAAAACCCA	180
SIM1	TGAAGTGTCTACTATAATGTTTCC	TAAATCTATATGGGGTTGGGGTT	217
SIM1	TCCACAATTAGTTCACAGTAAG	ATAAAATGTTGAATGCCAGAT	222
SIM1	CTCCTGCTGTCTGATGAGGAGAT	CCCACCAACAGGTGAAGTCTG	233
SIM1	TTCTCAGGATCTATTTGCCAACC	TGAACAAGTATGCCTGAATTTACT	240
SIM1	TATCACAGAACATGAACGCTGGG	gctaagagcAGTACTCTGAAGGAA	237
SIM1	CGATGGAAACAATCTAGCCCTAT	TTAATGTCACAATCACTACAGGT	234
SIM1	CTGAGGAGCCAGCCTTTT	GTTTTTGGAAATGGGACGAGTG	237
SIM1	ACAGAGTCAAAGAAAATTACCCA	CCTCTAATTATATGGGAACCAGAA	238
SIM1	ACTCAAAGTGCATATTGGTAACT	TGCAATGTATAGTTTTTAGTGACAT	238
SIM1	TTCTACAAAACCTTAGCATTCTT	GGACTTGAGTATGTGAGAATTGA	229
SIM1	AGAGCTAAAATGGTTCCTGTTTCT	TGATATCTGCAAAGTGTCCCA	238
SIM1	GTGAAAAATTACCTTCTCTGATTGT	TCTTTGACTCTGTCACTAAACT	218
SIM1	TGGCAAAGTTATGGAAATCAAGAAT	TGACCCAAGGTAAGTTAAGACATTTG	239
SIM1	TCTATCCACACCTGGCAAAAAA	CCTCCCCATCAATACTTTCTCTT	216
SIM1	TTCAAACCATGTCGCCAAAAA	CCCTGCAGAGACACAGAATACAA	229
SIM1	TCTCTGCTGGATATGGTCACA	CCCATGAACCCAGCAAAATTGA	239
SIM1	AGCATTATTCCTTCTGATTACTCTCA	TCACAGATGGGTATTTCTGGTGG	220
SIM1	AAATGGGAACACTTTGCAAGATA	ATGAGTAGACACTTCAAAAAAGATAA	236
SIM1	AGTTACAGGTAGGGCCAAAATTA	CATATGTGCCAATGTGACAAC	231
SIM1	TATAATTGATCCTCATCCCCAT	TTACACAGAAAAACCACTTGCTC	216
SIM1	CCCCCTACCCTGCTTCC	CACTAACTTCTGTCTCTGCTC	236
SIM1	TTTCAGCAAAACATCAGCTTCCG	CTCTCCAAACTGCTTTGGCTCTC	233
SIM1	GCCATATTGCAAAAAATGTAAGC	TGCTCTTTAAATGACATGACCA	218
SIM1	CAGGCACTCAGACCTCAAAG	GGCACTCGGGTAATACAGTCG	227
SIM1	ATCATATTTTCTGTGTGCGTAT	AATGAATGACAACAGTTATGGAC	227
SIM1	CCTGTGGCGCTCACTTATACAAA	TGCTGTAATGCAAAAAAGTGCAA	227
SIM1	GTTTCAGAATCTGCGAATGTCTT	TTTTGTCCAGTAAAAACCAGGAG	238
SIM1	GTAGTCCCAGAGGTAGGGACATT	GTGTCACACGACCTGAGCTTC	222
SIM1	TTACCTGAGGGTATGGGGAAGTC	TTCTATGGTGGCTGATTAAAGGG	217
SIM1	TTGATGTGTGTCCTCAATATCTCA	AAACTGACTTCTTTGTGGTTTCA	233
SIM1	CTGGATCATGAATTGGAGGAGGG	GCTTGTGAGTGGCATTGAGGG	236
SIM1	TAAATCTTCTATTTTCTTCTGT	CTGTGATAGAGCAGAAGTGAACCTA	216
SIM1	TCAGAAGATTTCTTATCTGTGCT	TGGGCAGAGAATCCCACTTA	225

SIM1	GCAAAGTCACTTACCTTCTGGGA	GCGGAGCTTAAGTTGTGAGAAGA	230
SIM1	ACTTAAACTGGGTCAATCTACCA	AAGAACCAGGGAAAATGAACTGA	239
SIM1	GTCAAAATTTATCTATTCTGGTTCCC	AGTTTAATACCCATTACCAGCA	204
SIM1	GCCTCTCCACACCACC	TTCATGATCCAGAAAAGGGAGCC	221
SIM1	CTTTCTGGGGAGGCCTTTGT	CTGGCTCCAGCACGAC	238
SIM1	GTCATGTGGCACTTCACTGGT	CCTGGAGTGCTCAATTGGAAGTA	228
SIM1	GGACATAGTTGACGCTGACGATA	TCTTACCACCAGGGATTGCATT	223
SIM1	GCCACGACTAATTGGGGAAAAAC	GTCCCTTACCAGGGCATT	239
SIM1	CTCATTTTGAATGTCACCTCAGC	GAAGTTTCCCAATCTTTCTTGG	229
SIM1	AAACATTTGCCTATTCCTCAGAG	TCAAACCAATTCACCCAGAAAAAG	229
SIM1	GTGAGGACCTTAGTGAAGCAG	GGACAAAGCATCCATAATCAGAC	220
SIM1	CCAGAAAGATGAGCTTATGTCC	CTCTTCTACCTGTGGCTGAGT	240
SIM1	CAAGAACAGTGATTGATGGCAGT	TATGTCGATTTTCCGAGGACCAG	234
SIM1	TATCTGACCTGTAGTGATTGTGA	ACATAGCTCATGGTTTTGTAGAAT	229
SIM1	ATGCTGAGCCCTTAAATTGTGTT	GAGGTAATTTTACCTTGCTTTTCA	216
SIM1	GCCCCAGCCTGTCTAACC	GACCACCAAGTACTACAGTTCC	226
SIM1	CTGCTTAAGTCACAAAGCATCTA	TAATTTTGGCCCTACCTGTAAC	221
SIM1	TGGGCTCATAGGATAGACGAAA	CTTCGACGGCTGTACCAAAAC	233
SIM1	AACCTGCCAGGGCCTAAT	GTGAGGCAGGCAGGTAATTC	210
SIM1	AACCCTGAATGCTAGTGCTATGT	ACAAGGACACAAGGGAACATCTG	220
SIM1	AGCACATTTTATGTATCCCCAA	GTGCAGCAACATATTTCTCATA	225
SIM1	CCATTTCTTTAGCTTCTGTCCC	TGGAATACATAAAATGGAGTTTGA	216
SIM1	GTTACCGGGTGAAGTTAACAG	CAATTATAACACACACACACA	240
SIM1	AGTTTCTAATGGTTTCGCTGTCAT	GCACTATCTCGGATAAGTAGTCCC	229