

NAMES	PRIMERS
GATA6 F	ATGGACCTGGGTGATAACAGCTG
GATA6 R	TTAGGCCAAGGCCATGGGGCACC
foxa3 F	ATGTTGAGCTCCGTGAAGATGG
foxa3 R	CCTTAAAGCATTCTTTTTCTCAG
prss1 F	ATGAAGGCTTTCATTCTTCTG
prss1 R	CCATGATAACGACCTCAAAAC
FOXA1 F	CTCACACACACGCACGCTGTGAG
FOXA1 R	GGGGATCACAACGAGTATTTTAC
lfabp F	TCTCCAGAAAGCATGGCCTTCAG
lfabp R	GTCAGTCAGAGCTTTATTTAAATAG
ifabp F	GTCATCATCATGACCTTCAACG
ifabp R	CTCACAGGTGCAAATGACACG
MYCN KI F	GGGTCATCTCTTTACAGGGTG
MYCN KI R	CTGACAAAAGCAGTTCTACTGTC
Grna1 (MYCN KI)	TAATACGACTCACTATAGGCTCTCGGGCGTCACACCGGTTTTAGAGCTAGAAATAGC
Grna2 (MYCN KI)	TAATACGACTCACTATAGGAGACTAAAGTGCCCCGTTGTTTTAGAGCTAGAAATAGC
Grna3 (MYCN KI)	TAATACGACTCACTATAGGCGTGGTCCAATTATTCGCGTTTTAGAGCTAGAAATAGC
PUC19 F	GACGAAAGGGCCTCGTGATACG
PUC19 R	GCCTGGGGTGCCCTAATGAGTGAG
Left Arm fusion F	TTAGGCACCCCAGGCGAGAATAAGGGCTGGCACTGGGC
Left Arm R	CGAACCACTCCCAGCGAGTCTGGCCTGCTCAAGTC
LINKER F	CAGGCCAGGACTCGCTCGGGAGGTGGTTCGGGTGGC
LINKER R	AATCAGACTCACCATAGGACCAGGATTTTCTTCAAC
EGFP fusion F	GAAAATCCTGGTCTATGGTGAGCAAGGGCGAGGAG
EGFP fusion R	GAGCGTAAAGGGTTACTTGTACAGCTCGTCCATGCC
RIGHT ARM F	GACAACGTCAAGTAACCCTTACGCTCATTGGCTGAATG
RIGHT Arm fusion R	CGAGGCCCTTCGTCGCTCTCTCTCTCTAGCTGTGTG
Mycn KO F	CTCTGTCCCCGAGCCGGGCA
Mycn KO R	GCTGCTGCTGCTCCCGTTTC
RPL3 F	CGAACGTGCTTCCAGCTTTATAAG
T7-RPL3 R	GTAATACGACTCACTATAGGGTTAGGCCATCTTCTCCTGGCCAG
RPL12 F	GGTGAAGGTGCAACAGTTTTTCG
T7-RPL12 R	GTAATACGACTCACTATAGGGTTACTCAGTTGGGCACTCAACTG
RPL15 F	ATGGGAGCGTACAAGTATATGC
T7-RPL15 R	GTAATACGACTCACTATAGGGATGTACAGTTAAGCACACTTTG
RPL19 F	GCTTTCGCTGACCACCAGCCATG
T7-RPL19 R	GTAATACGACTCACTATAGGGTTATTCTTGGTCTCATCCTCTTTG
RPL5 F	GCAGCGGAGTGAAACTCCCAG
T7-RPL5 R	GTAATACGACTCACTATAGGGGCGGGTCGAGCTCTAGTCCCTCC
RPL18 F	GTCCAAGATGGGAGTTGACATC
T7-RPL18 R	GTAATACGACTCACTATAGGGCCTTAGTGTCTGCGACCACGG
RPL30 F	ACGAGGCCATCTTGGTGTGATCAG
T7-RPL30 R	GTAATACGACTCACTATAGGGGTGGTGGAAAAGGCTCTACTTCTC
RPL24D1 F	ATGCGCATTGAGAAGTGTATTTTC
T7-RPL24D1 R	GTAATACGACTCACTATAGGGGTGATCAGCTGAATGCAGAGATG
RPL4 F	ATCGAGTTAGGAGCGTGGTTCCTG
T7-RPL4 R	GTAATACGACTCACTATAGGGCTAAGCTTGAGCTGGAGTAGCTTC
RPL32 F	GCGGCCGCGGATCCGACGTCTC
T7-RPL32 R	GTAATACGACTCACTATAGGGGATTATTCGTTCTCCTCGCTGC
RPL17 F	ATGGTCCGCTACTCTCGACCC

T7-RPL17 R	GTAATACGACTCACTATAGGGTTACTCCCGAGCCATAAGCTTCTG
RPL6 F	TCCACACACAATCTGCATTAATC
T7-RPL6 R	GTAATACGACTCACTATAGGGGCATTTTAGAAAACCAGTTTGTG
RPL36 F	GGACGCCATTAGCAAAGAGCAG
T7-RPL36 R	GTAATACGACTCACTATAGGGCACAAACAGTTTTATTTTTGATG
Rps9 F	TCTCTTTAGTGTTCGAGCCTGAC
Rps9 R	GTAATACGACTCACTATAGGGTTAATCTTCTCTTCATCATCACC
Rps4x F	ATGGCCCCGAGGACCGAAGAAGCACC
Rps4x R	GTAATACGACTCACTATAGGGGAAACCAAGGTGTGCTAGTCCTG
Rps6 F	CTCCAAGCGAGAAAGTCCTCCATC
Rps6 R	GTAATACGACTCACTATAGGGTCACTTCTGGCTTGACTCTGAC
Rps3A F	ATGGCAGTCGGCAAAAATAAGAGG
Rps3A R	GTAATACGACTCACTATAGGGCAGCACACAAAGGCTCAAATTTAC
Rps20 F	CACCGCTGTTCGCCTTTCAGCTG
Rps20 R	GTAATACGACTCACTATAGGGTTAAGCATCTGCAATTGTGACC
Rps8A F	CTCTTTCTAGCCGGCGCTGAG
Rps8A R	GTAATACGACTCACTATAGGGCTATTGCTTTCTTGGCTTTAATC
Rps271 F	ATGCCTTTGTCCAGAGATTTGATC
Rps271 R	GTAATACGACTCACTATAGGGGTGAAGGTCATTCATTGTAATC
Rps17 F	ATGGGACCGGTGAGGACCAAGACGG
Rps17 R	GTAATACGACTCACTATAGGGTTTTGCAGTCTTTATTGACATCTG
Mycn qF	TACCAAGTCAGCGCTTGAGG
Mycn qR	TTTGCTGCCTTGTGCTTGTG
RPL3 qF	AAGGGACATGGATGCAAGGG
RPL3 qR	AGGCTCCAATACAAGCCACC
RPL12 qF	GATGACATCGCAAAGGCCAC
RPL12 qR	CAGAGGCTGATGGCACTACC
RPL15 qF	CTGTGCATCATGGCGTCAAC
RPL15 qR	TAGGAGTTCAGGACCCGGAG
RPL19 qF	TCTTGCACGAAGGAAGGGTC
RPL19 qR	ACGTTACCCTTTGCCCTCAG
RPL5b qF	ATTTTGCCAGAAAGCGCCTG
RPL5b qR	CCCACACTGATGCCGTACTT
RPL18 qF	CAACAAGGACCGCAAGGTTTC
RPL18 qR	GGGAGCATCAGAACGACGAG
RPL30 qF	GTTGGCAAAAACCGGTGTCC
RPL30 qR	CTGCTGGTCTGGCATACTCC
RPL24d1 qF	CGCAACCCAAGAAAGACCAG
RPL24d1 qR	ATGATGAAGCGAGCCTGACG
RPL4 qF	CGTATCCAGCGTAAGGGACC
RPL4 qR	AGAGGTCATCCAGTTTGCGG
RPL32 qF	AGGGTCAGATGTTGATGCC
RPL32 qR	TTGAGGACACGTTGTGAGCG
RPL17 qF	CGGCCTGAGAACATCTCTGAA
RPL17 qR	AACACACTGGTGTCTTGACGA
RPL6 qF	TCATCATCGCGTTATGGCTGAG
RPL6 qR	CACCTTCGTCACTGGAGCTTT
RPL36 qF	ACGCCATTAGCAAAGAGCAG
RPL36 qR	TCTCTCGTAGGGTGCAGAAAC
RPs9 qF	GCTCATTCGTCAGAGGCACA
RPs9 qR	CCACCATATGGAGAGCGGAG

RP _s 4x qF	GGACCGAAGAAGCACCTGAA
RP _s 4x qR	CTCCCTCAGTTTGTGGGGAC
RP _s 6 qF	TTCAGCGTCTGGTTACACCC
RP _s 6 qR	TGACGTTTCTCCTTGGCCTC
RP _s 3A qF	AAGAGGCTGACCAAAGGTGG
RP _s 3A qR	ACTGACCTCGAACACACGTC
RP _s 20 qF	AGGTGGCATTACCGTATC
RP _s 20 qR	TTGGTAGGCATACGCACAGG
RP _s 8A qF	CGCCGCATCCACACAATAAG
RP _s 8A qR	CCAGGGTCTTGGTTCTACC
RP _s 271 qF	TGTTCAACAGTGCTCTGCCA
RP _s 271 qR	GGCATCAACAAGTCACGACAG
RP _s 17 qF	CCACATTAACAAGCGGGTGTG
RP _s 17 qR	TGCAGTTTGATGGAGATGCCT
Hsp70 P F	CCAAGTTTGTACAAAAAAGCAGGC
Hsp70 P R	AGCAACTTTTCTATACAAAGTTG
Sox17 promoter F	CTTTGTATAGAAAAGTTGCTTACACTCTCTGACAAAAGGTACAC
Sox17 promoter R	CTTTTTGTACAAACTTGGTCCACAGTGAAAAGTTTCAGGCAG
ZMYCN F	GTACAAAAAAGCAGGCTTCGGATCCATGCCAGCTAAAACCATGAGC
ZMYCN R	CAGAGAGAAGTTTGTAGCGGATCCGCGAGTCCTGGCCTGCTCAAGTC
Grna1 (mycn)	TAATACGACTCACTATAGGCTGGGGATTGCTCGCTGAGTTTTAGAGCTAGAAATAGC
Grna2 (mycn)	TAATACGACTCACTATAGGAAGCCAGTCCAACGGGTGTTTTAGAGCTAGAAATAGC
Grna3 (mycn)	TAATACGACTCACTATAGGAGTGTGTGGACCCTACCGTTTTAGAGCTAGAAATAGC
Constant GRNA	AAAAGCACCGACTCGGTGCCACTTTTTCAAGTTGATAACGGACTAGCCTTATTTTAACCTTGC TATTTCTAGCTCTAAAAC
T7 PROMOTER F	TAATACGACTCACTATAGG
Grna scaffold R	AAAAGCACCGACTCGGTGC