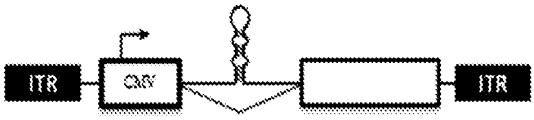
**Figure 1**

**scAAVrh.74.CMV.miR29c**

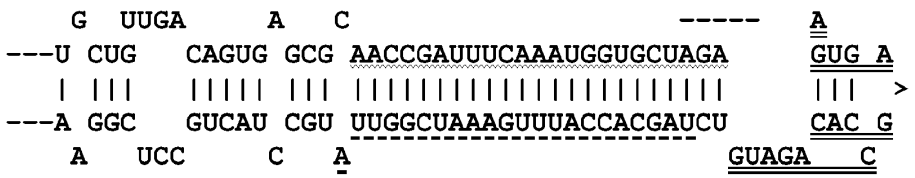



**SEQ ID NO: 2: miR-29C IN A miR-30 BACKBONE**

GGCCGGCC*tgtttgaatgaggcttcagtactttacagaat*CGTTGCCTGCACATCTTGGAAACACTTGCTGGGATTACT
TCTTCAGGTTAACCCAACAGAAGGCTCGAGAAGGTATATTGCTGTTGACAGTGAGCGCAACCGA
TTTCAAATGGTGCTAGAGTGAAGCCACAGATGTCTAGCACCATTTGAAATCGGTTATGCCTACTG
CCTCGGAATTCAAGGGGCTACTTTAGGAGCAATTATCTTGTTTACTAAAACTGAATACCTTGCTA
TCTCTTTGATACATTGGCCGGCC

**FSE-I cut site (restriction site)**
**miR-30 backbone**
**miR-30 stem loop (GTGAAGCCACAGATG; SEQ ID NO: 5)**
**miR-29c target (sense) strand (ACCGATTTCAAATGGTGCTAGA; SEQ ID NO:3))**
**miR-29c guide (antisense) strand (TCTAGCACCATTTGAAATCGGTTA; SEQ ID NO: 4)**

**Predicted hairpin structure (SEQ ID NO:6)**



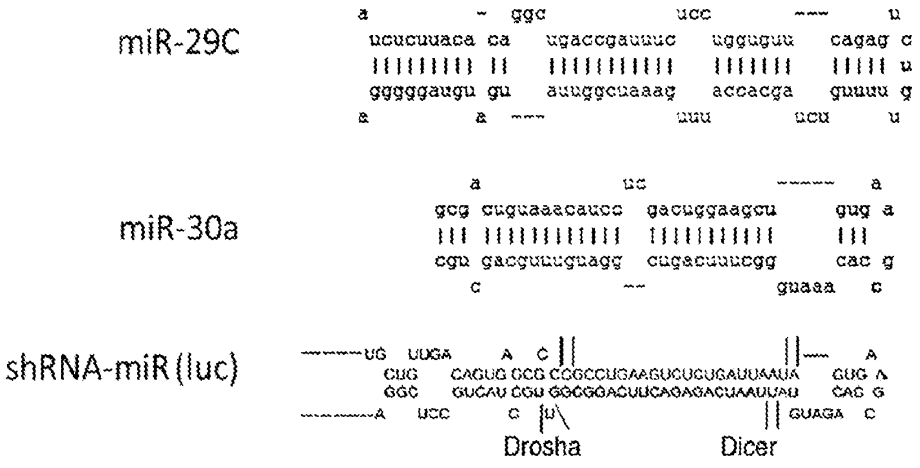



Figure 2A

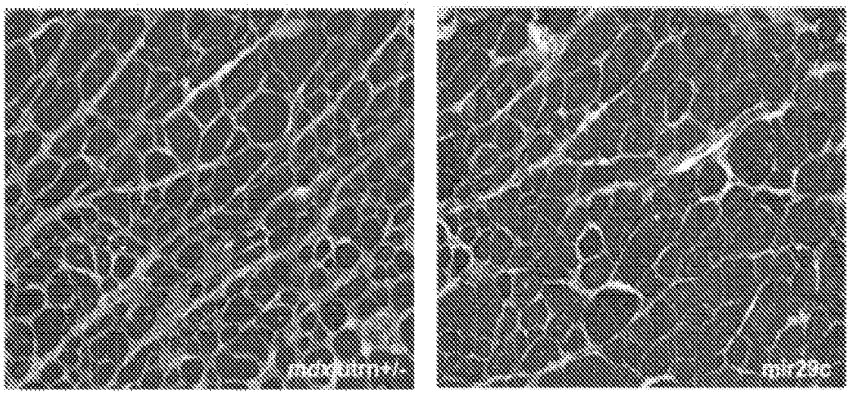



Figure 2B

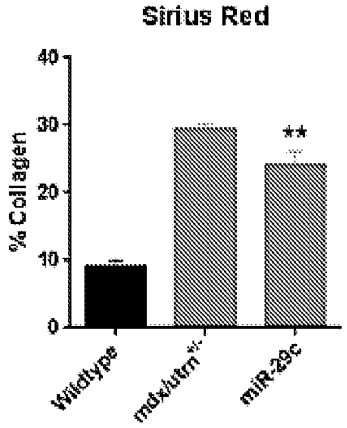



Figure 2C

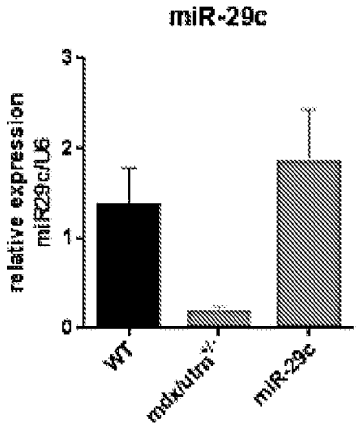


Figure 3A

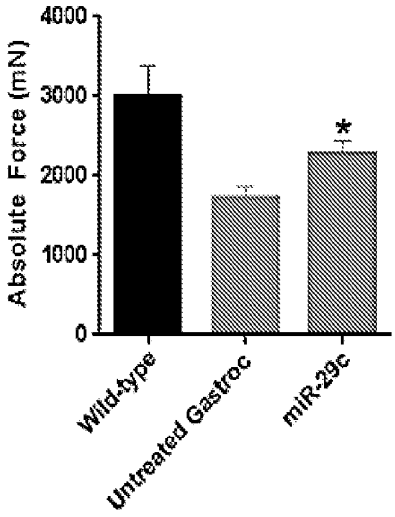



Figure 3B

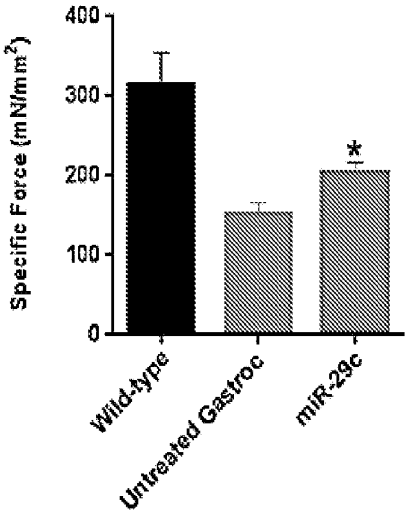



Figure 3C

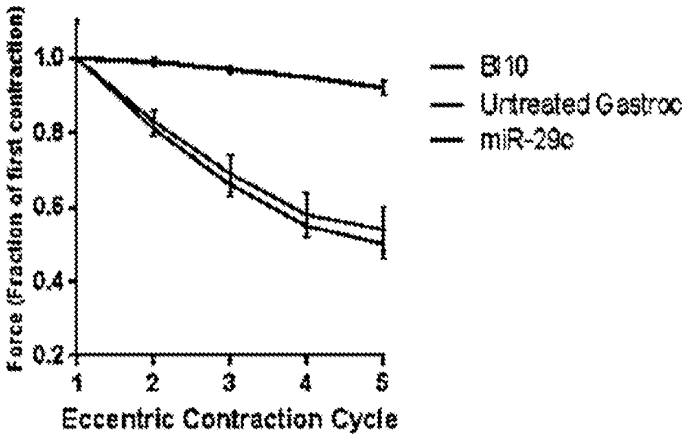


Figure 4C

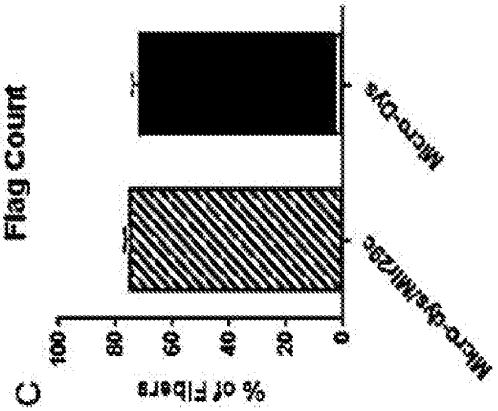



Figure 4B

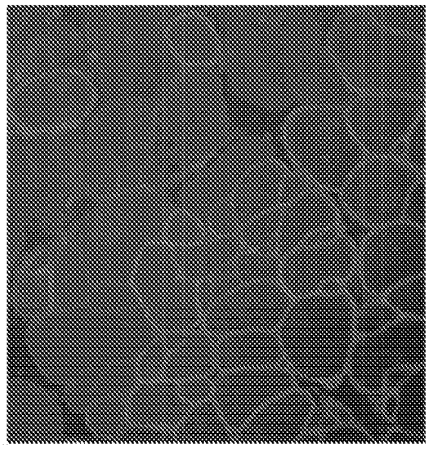
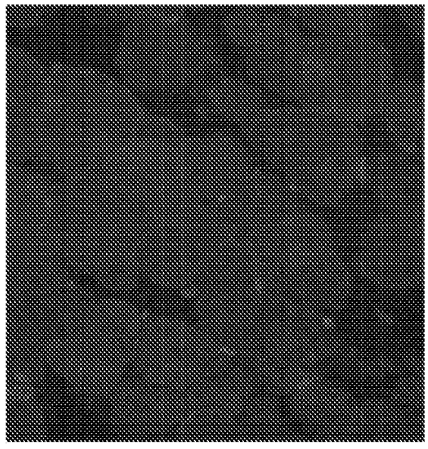


Figure 4A



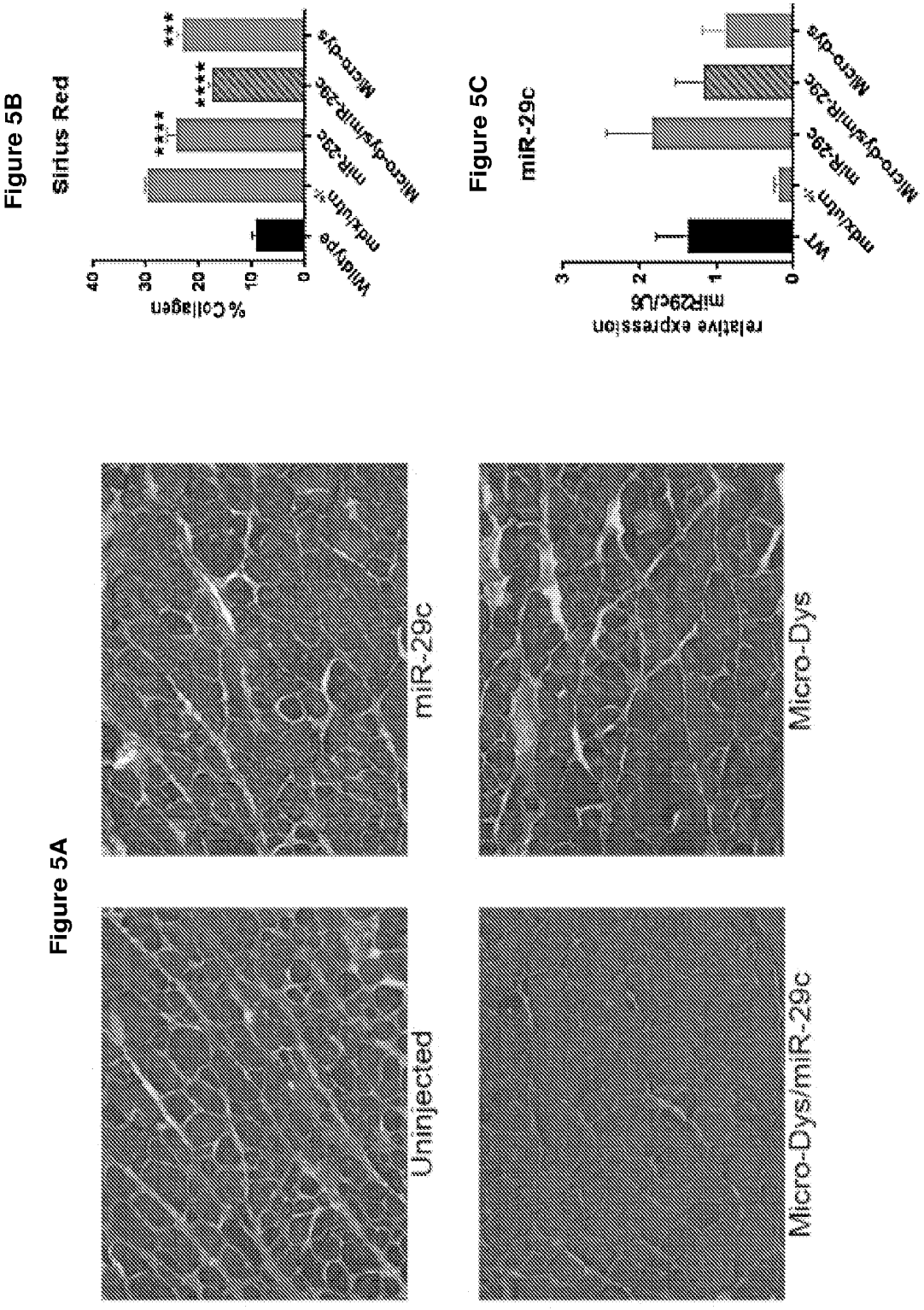

Figure 5A
Uninjected
miR-29c
Micro-Dys/miR-29c
Micro-Dys
Figure 5B
Sirius Red
% Collagen
0
10
20
30
40
****
****
***
Wild-type
mdx/utn-/-
mdx/utn-/- miR-29c
Micro-dys/miR-29c
Micro-dys
Figure 5C
miR-29c
relative expression
miR29c/U6
0
1
2
3
WT
mdx/utn-/-
mdx/utn-/- miR-29c
Micro-dys/miR-29c
Micro-dys

Figure 6A

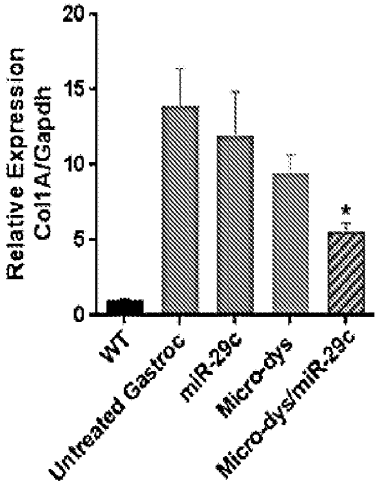



Figure 6B

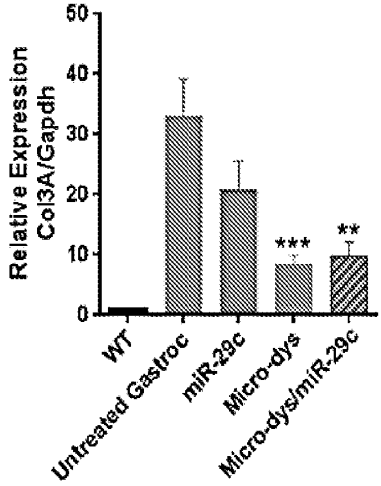



Figure 6C

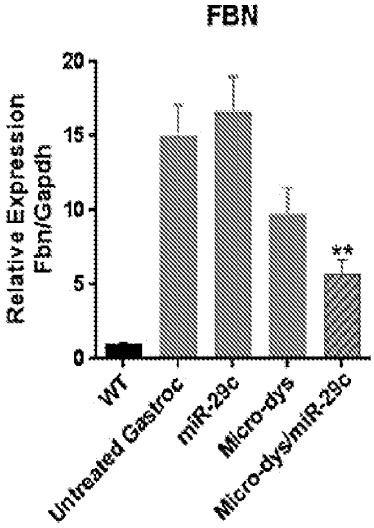



Figure 6D

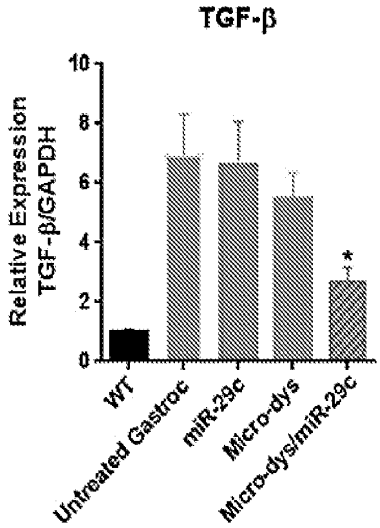


Figure 7A

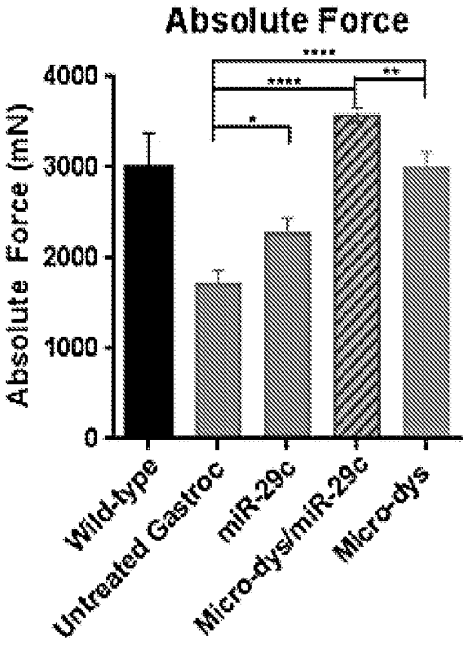



Figure 7B

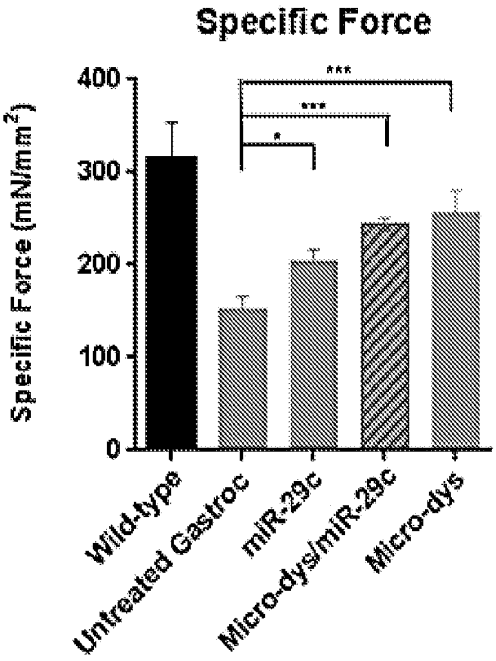



Figure 7C

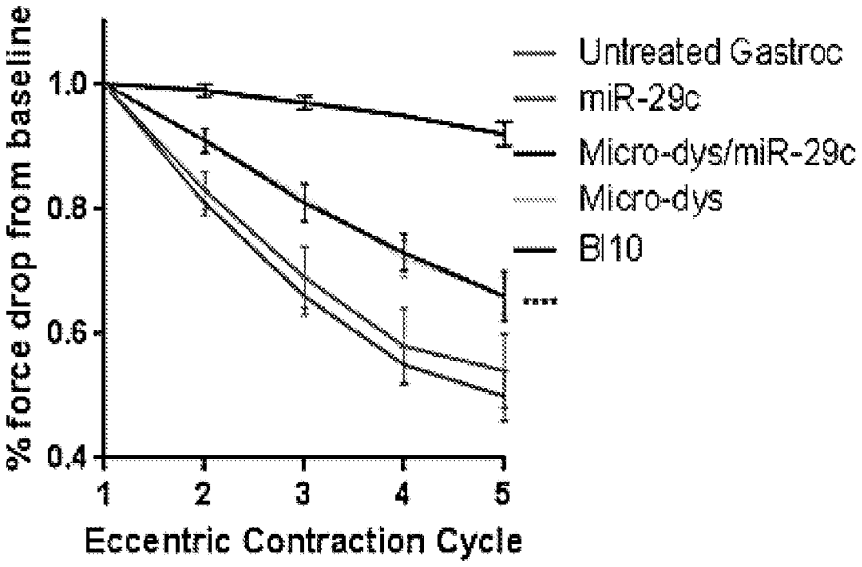
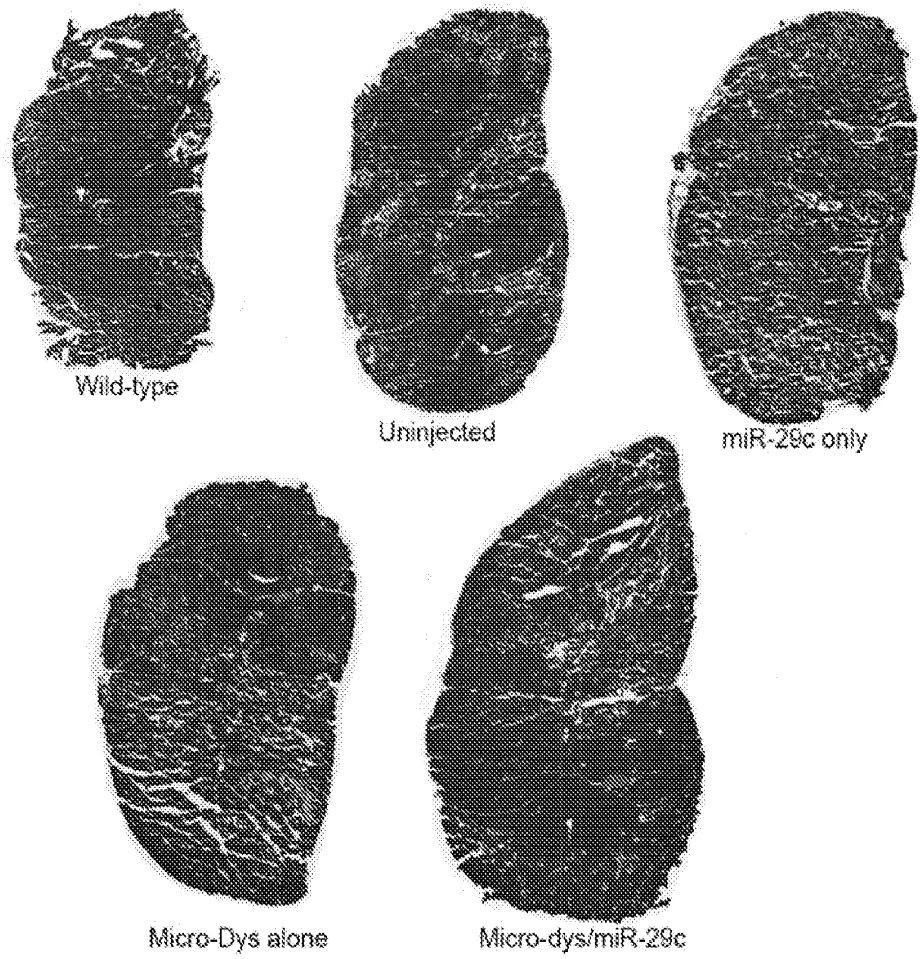
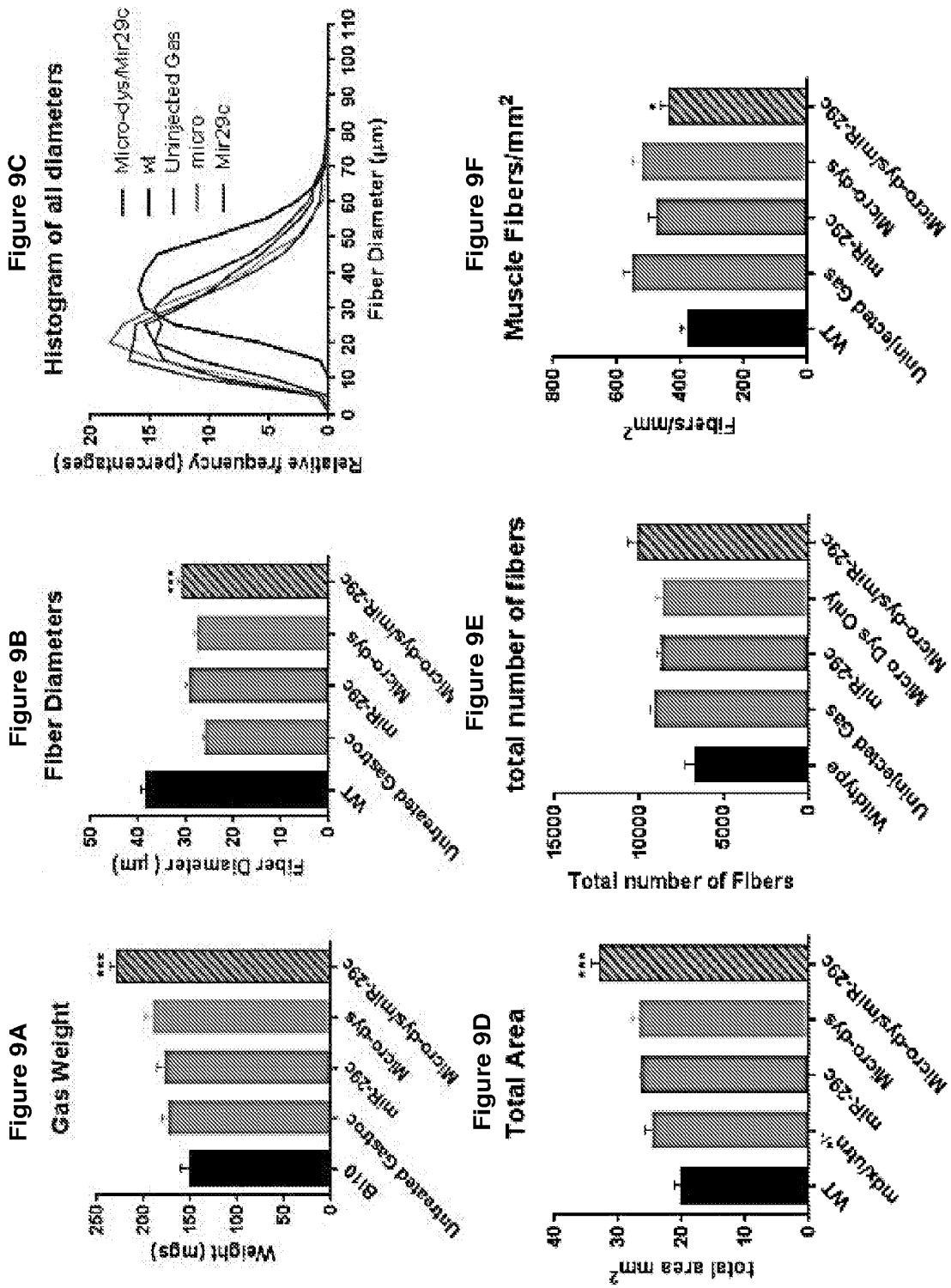


**Figure 8**


Wild-type
Uninjected
miR-29c only
Micro-Dys alone
Micro-dys/miR-29c


Figure 9A
Gas Weight
Weight (mgs)
Untreated Gastroc
mdx/utrn+/-
mir-29c
Micro-dys
Micro-dysfmir-29c
Figure 9B
Fiber Diameters
Fiber Diameter ( µm)
Untreated Gastroc
WT
mir-29c
Micro-dys
Micro-dysfmir-29c
Figure 9C
Histogram of all diameters
Micro-dys/Mir29c
wt
Uninjected Gas
micro
Mir29c
Relative frequency (percentages)
Fiber Diameter (µm)
Figure 9D
Total Area
total area mm2
WT
mdx/utrn+/-
mir-29c
Micro-dys
Micro-dysfmir-29c
Figure 9E
total number of fibers
Total number of Fibers
Wildtype
Uninjected Gas
mir-29c
Micro Dys Only
Micro-dysfmir-29c
Figure 9F
Muscle Fibers/mm2
Fibers/mm2
WT
Uninjected Gas
mir-29c
Micro-dys
Micro-dysfmir-29c

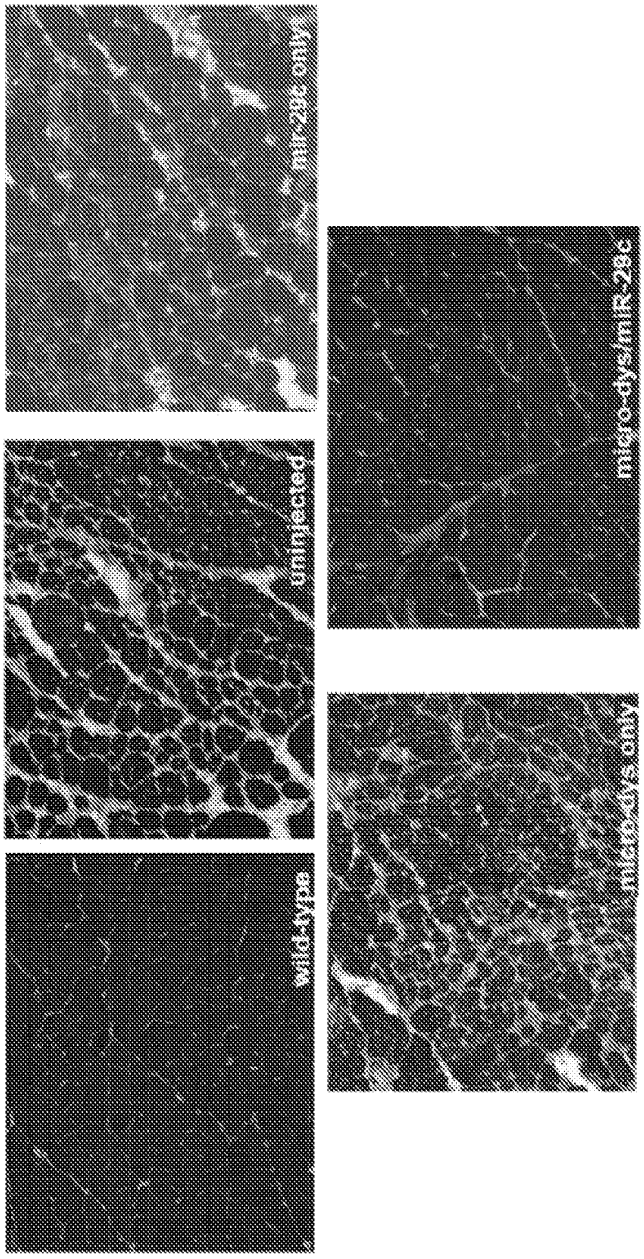

mdx-23c only
uninjected
wild-type
micro-dys/miR-23c
micro-dys only

Figure 10A

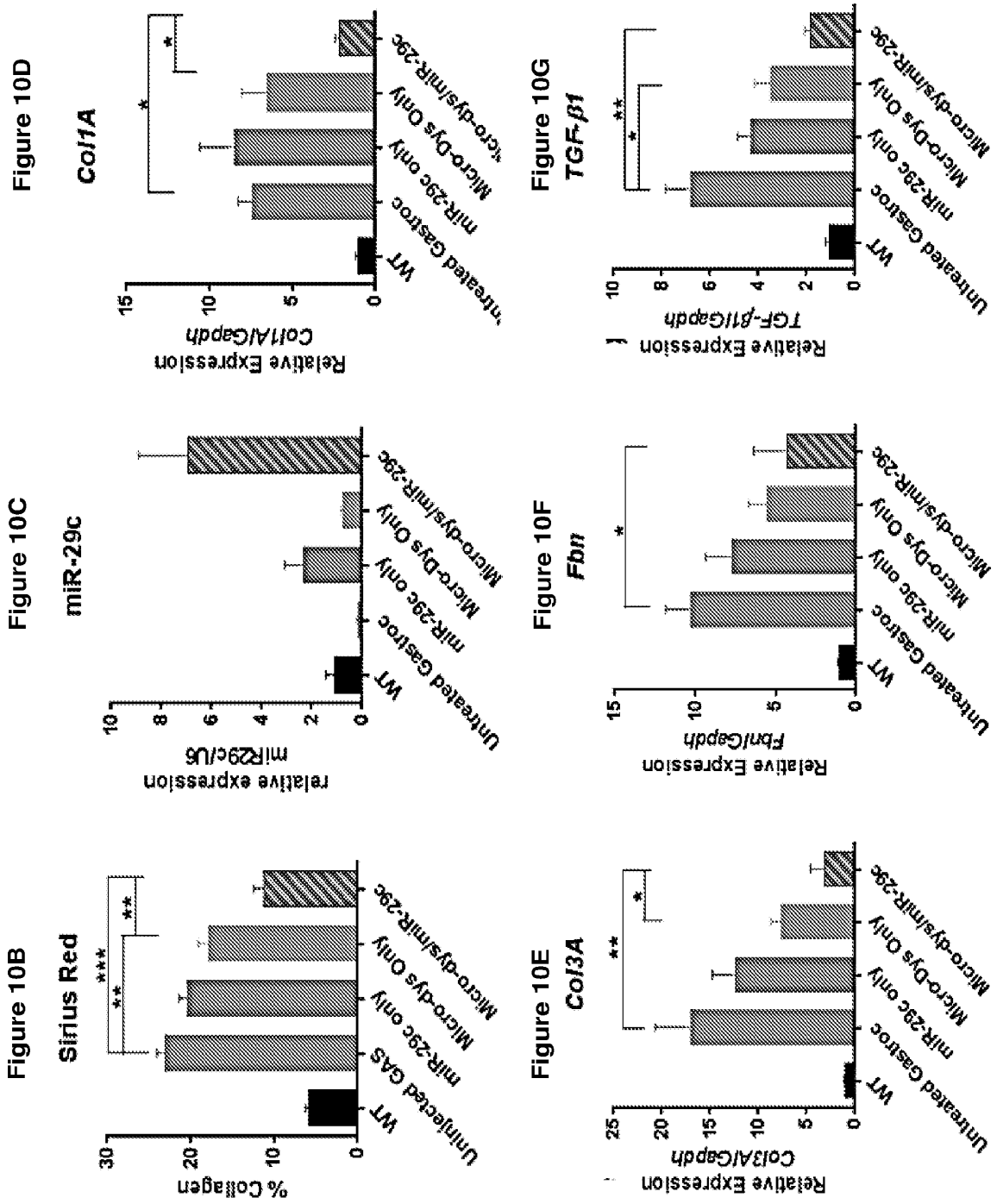

Figure 10B
Sirius Red
% Collagen
Figure 10C
miR-29c
relative expression
miR29c/U6
Figure 10D
Col1A
Relative Expression
Col1A/Gapdh
Figure 10E
Col3A
Relative Expression
Col3A/Gapdh
Figure 10F
Fbn
Relative Expression
Fbn/Gapdh
Figure 10G
TGF-β1
Relative Expression
TGF-β1/Gapdh
WT
Untreated Gastroc
miR-29c only
Micro-Dys Only
Micro-dys/miR-29c
Unifjected GAS

Figure 11

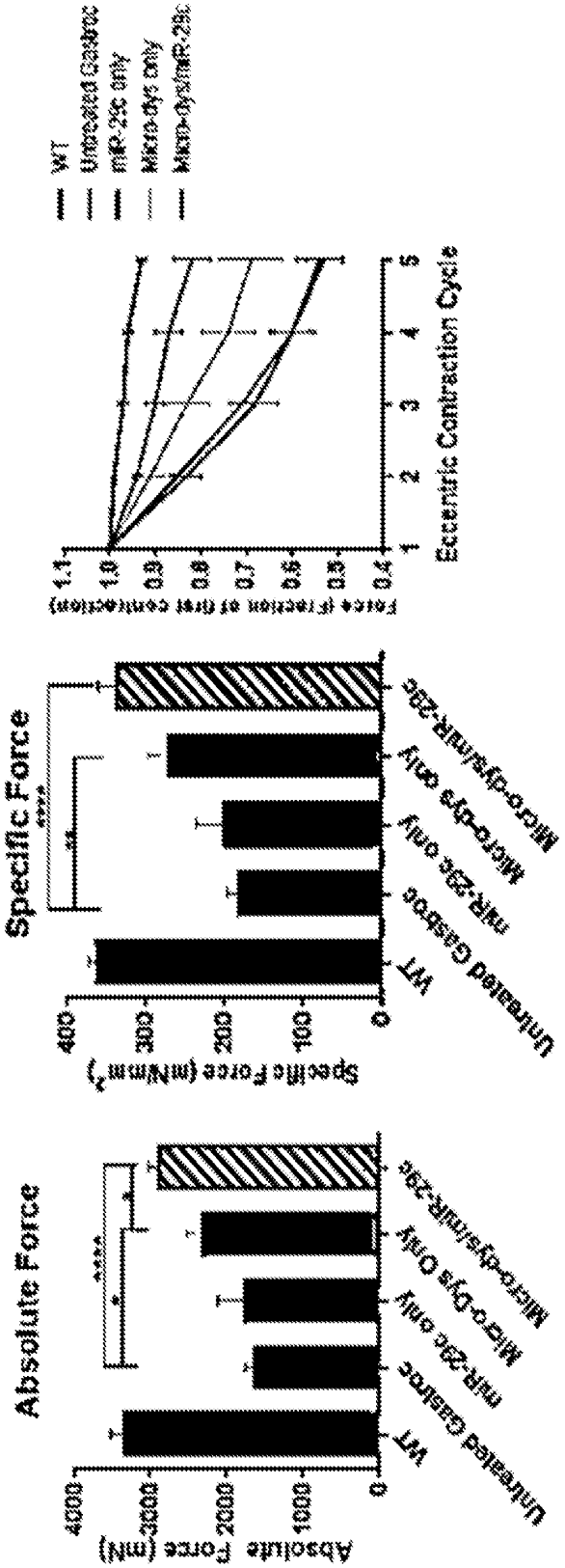
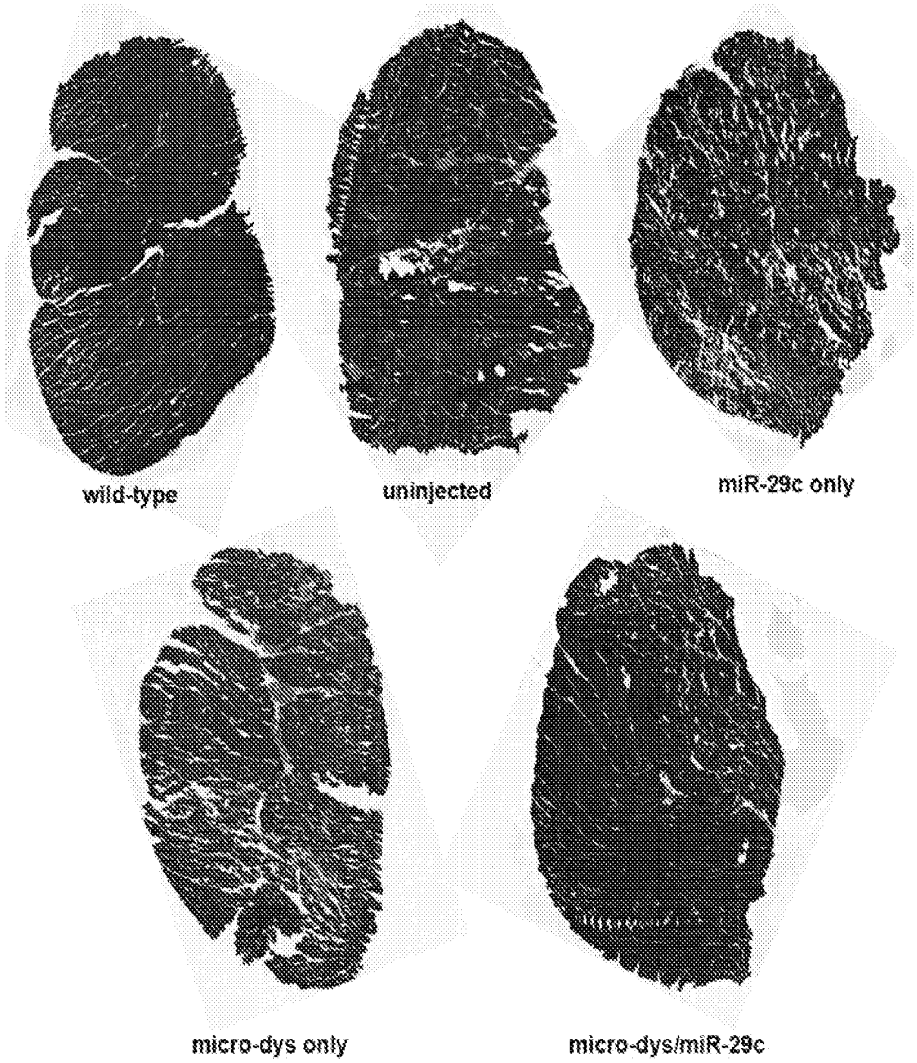

WT
Untreated Controls
mR-23c only
Micro-dys only
Micro-dys/mR-23c
Force (Fraction of first contraction)
1.1
1.0
0.9
0.8
0.7
0.6
0.5
0.4
1
2
3
4
5
Eccentric Contraction Cycle
Specific Force
****
**
Specific Force (mN/mm²)
400
300
200
100
0
WT
Untreated Gastroc
mR-23c only
Micro-dys only
Micro-dys/mR-23c
Absolute Force
****
*
Absolute Force (mN)
4000
3000
2000
1000
0
WT
Untreated Gastroc
mR-23c only
Micro-Dys Only
Micro-dys/mR-23c

**Figure 12**


wild-type
uninjected
miR-29c only
micro-dys only
micro-dys/miR-29c

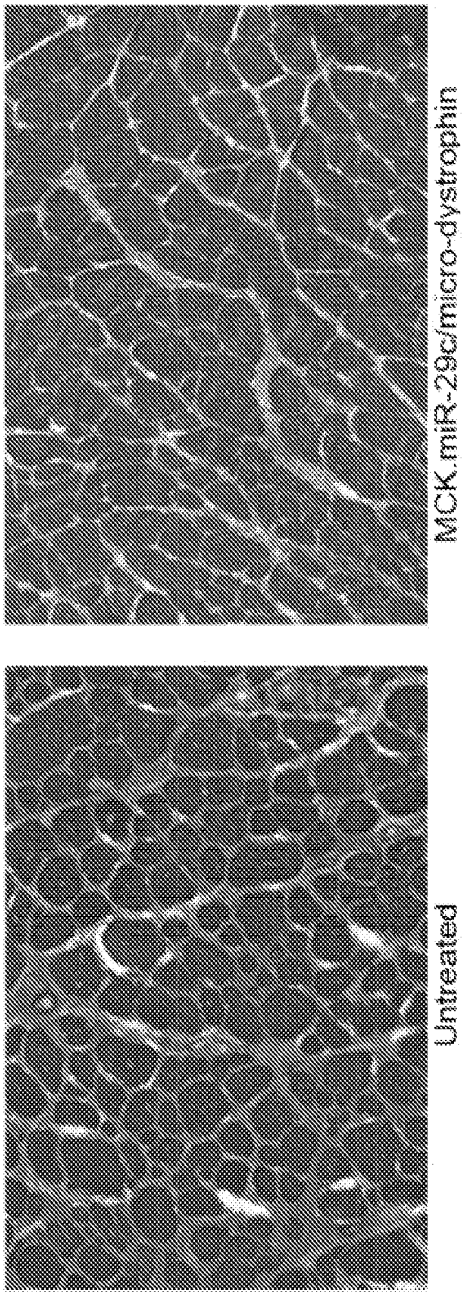
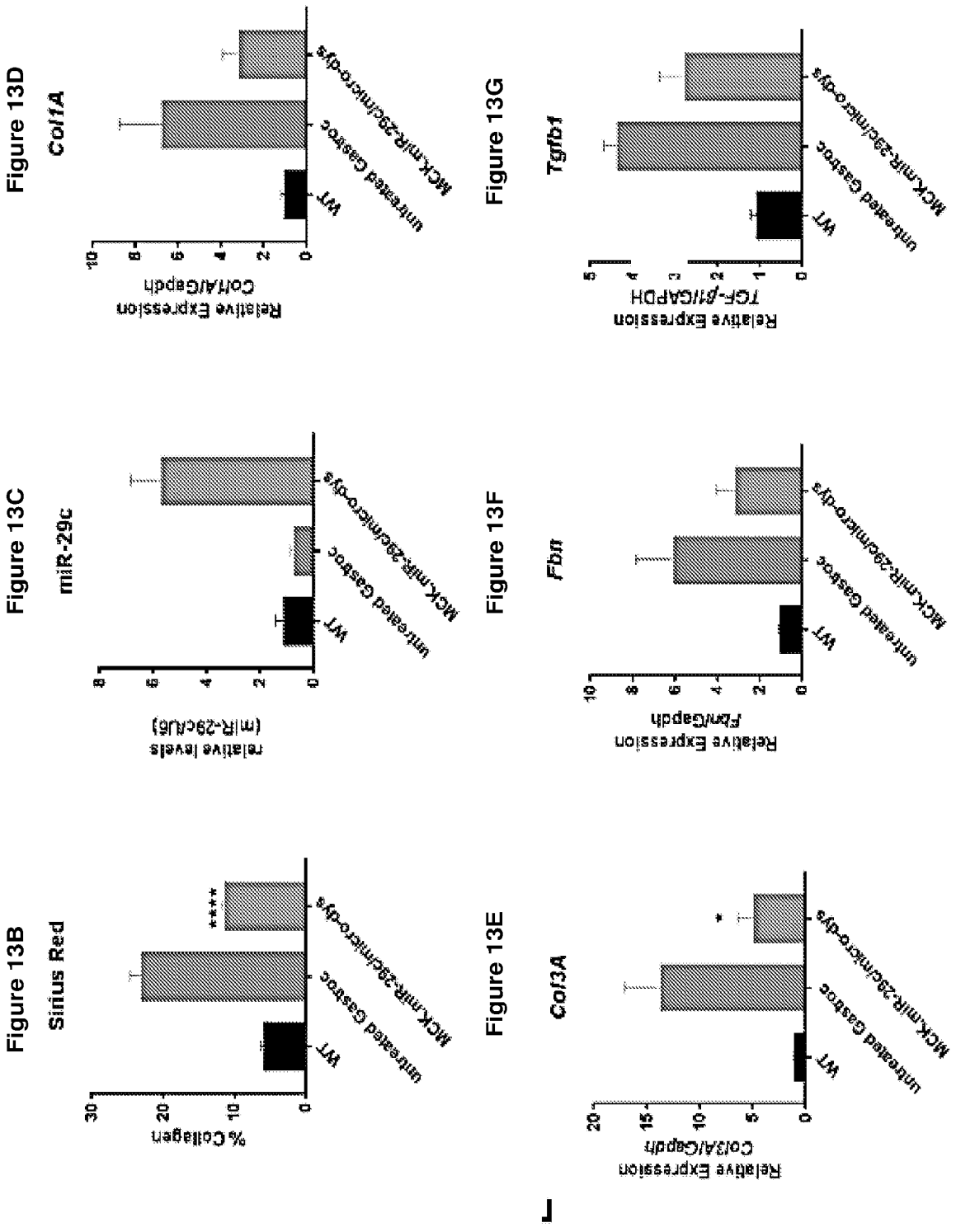



Figure 13A


Figure 13B
Sirius Red
% Collagen
WT
untreated Gastoc
MCK.miR-29cInducto-dys
****
Figure 13C
miR-29c
relative levels (miR-29cΔU6)
WT
untreated Gastoc
MCK.miR-29cInducto-dys
Figure 13D
Col1A
Relative Expression Col1A/Gapdh
WT
untreated Gastoc
MCK.miR-29cInducto-dys
Figure 13E
Col3A
Relative Expression Col3A/Gapdh
WT
untreated Gastoc
MCK.miR-29cInducto-dys
*
Figure 13F
Fbn
Relative Expression Fbn/Gapdh
WT
untreated Gastoc
MCK.miR-29cInducto-dys
Figure 13G
Tgfb1
Relative Expression TGF-β1/GAPDH
WT
untreated Gastoc
MCK.miR-29cInducto-dys

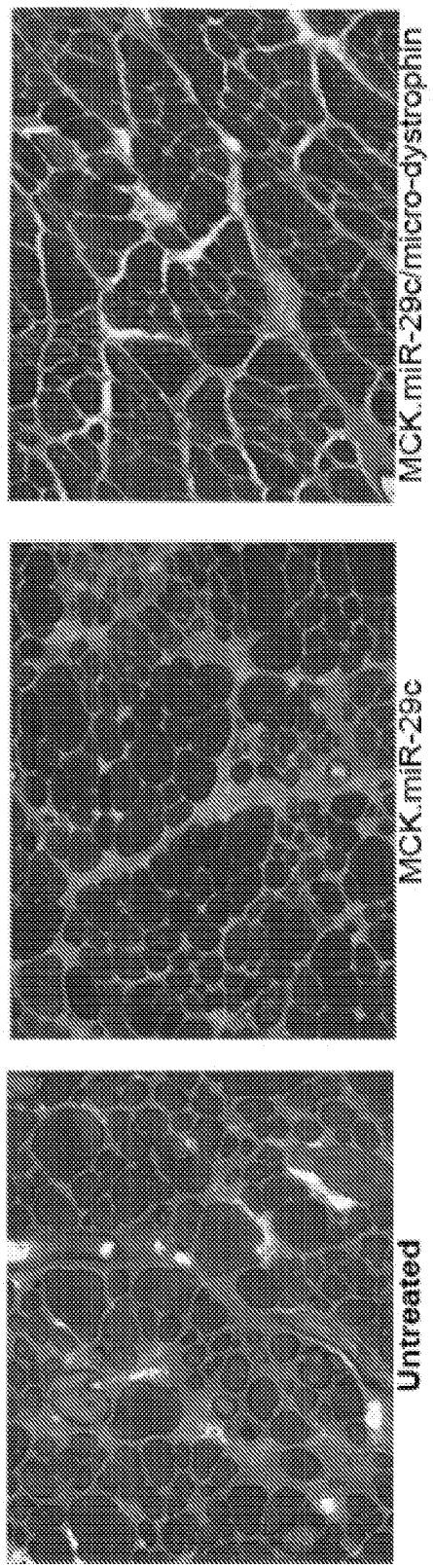



Figure 14A

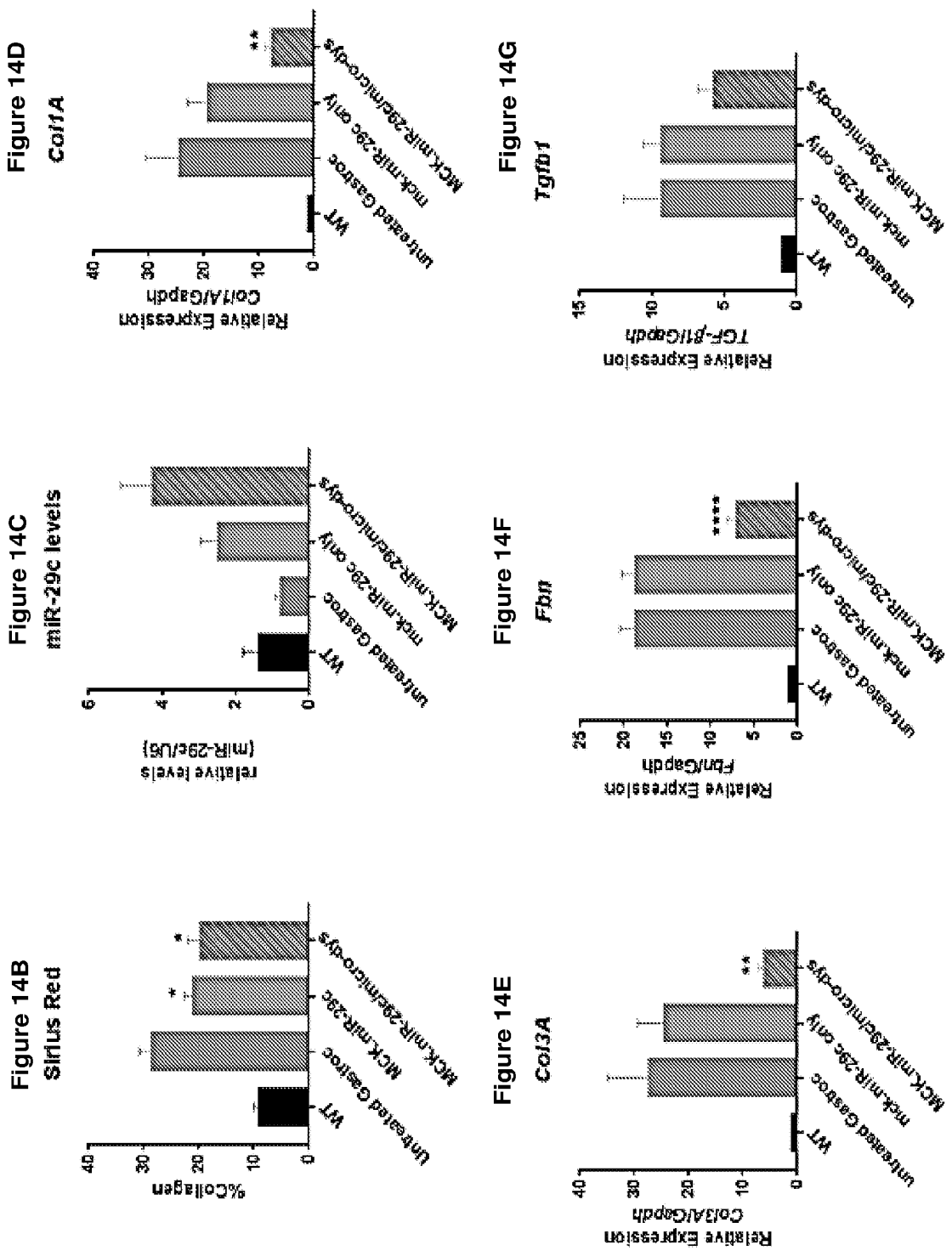

Figure 14B
Sirius Red
%Collagen
Figure 14C
miR-29c levels
relative levels
(miR-29c/U6)
Figure 14D
Col1A
Relative Expression
Col1A/Gapdh
Figure 14E
Col3A
Relative Expression
Col3A/Gapdh
Figure 14F
Fbn
Relative Expression
Fbn/Gapdh
Figure 14G
Tgfb1
Relative Expression
TGF-β1/Gapdh
WT
untreated
Gastroc
mck.miR-29c only
MCK.miR.29c/mice-dys

Figure 15A

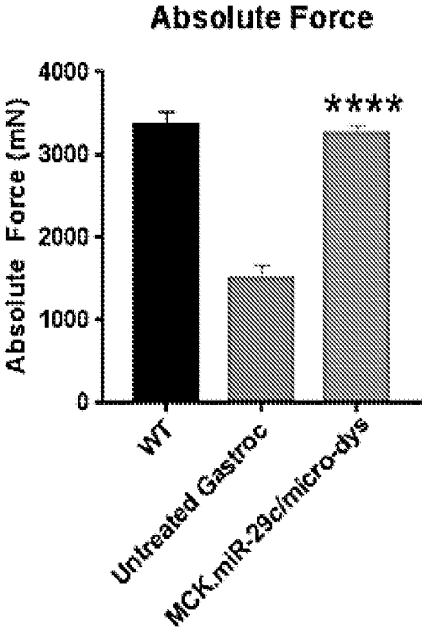



Figure 15B

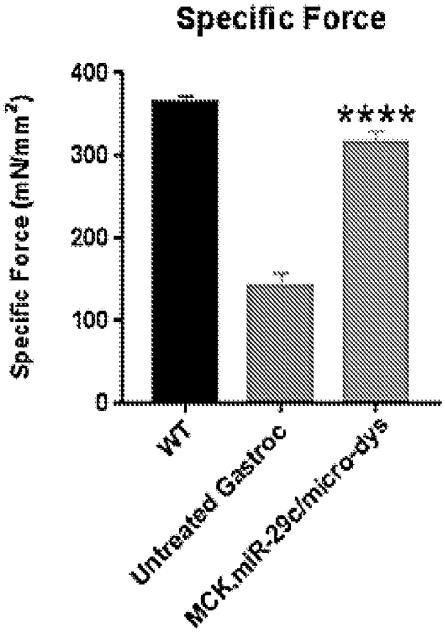



Figure 15C

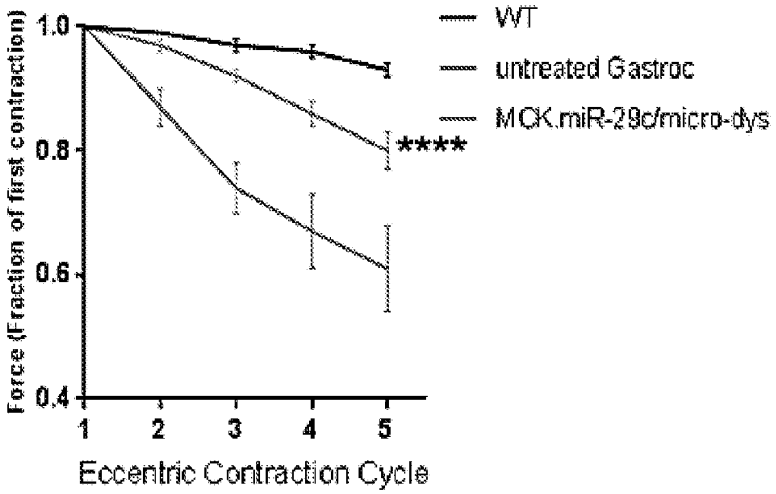


Figure 16A

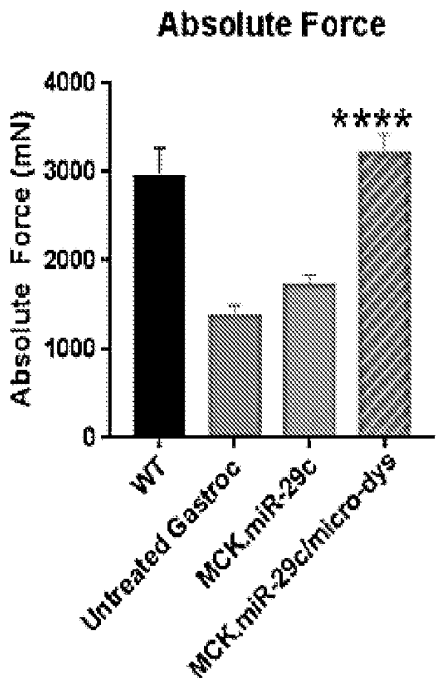



Figure 16B

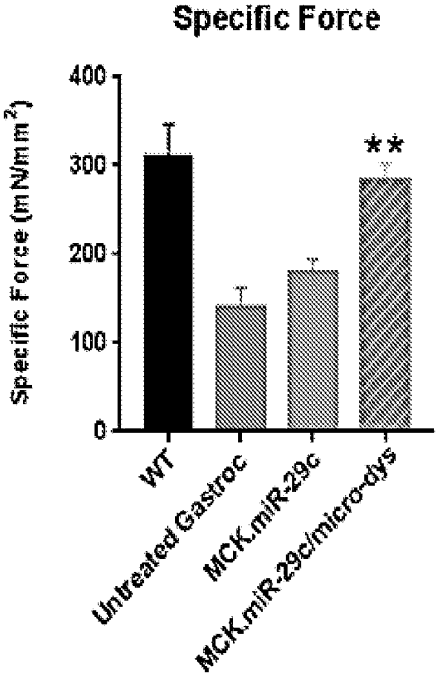



Figure 16C

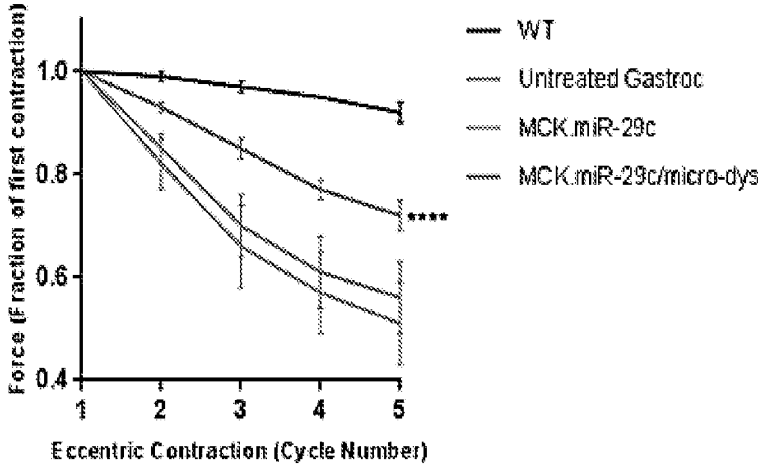


Figure 17A

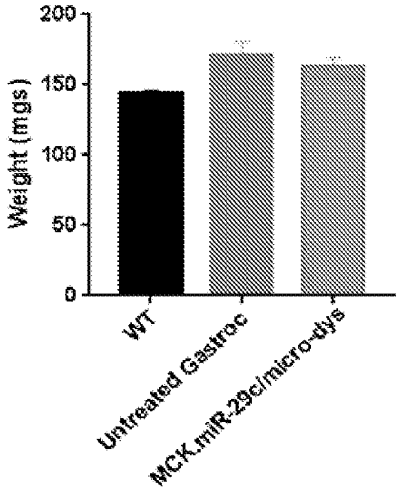



Figure 17B

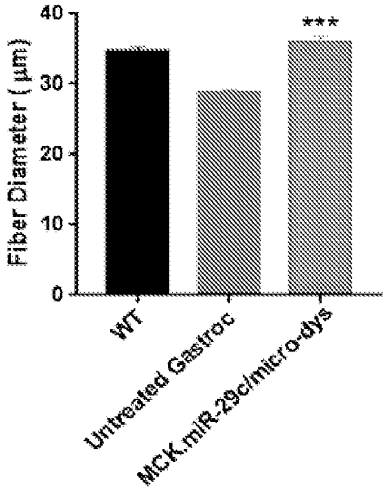



Figure 17C

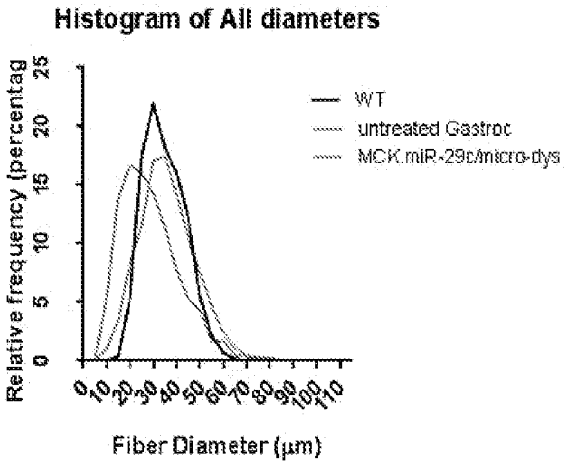
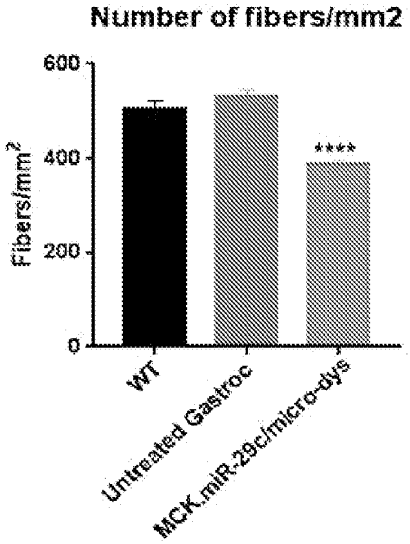



Figure 17D



## Figure 18A

**pAAV.CMV.miR29C Sequence (SEQ ID NO: 1)**

Main features:
CMV promoter- 120-526
EF1a Intron- 927-1087, 1380-1854
miR-29c- 1257-1284
shRNA-miR29-c with primary seed sequence- 1088-1375
PolyA- 1896-2091

CAGCAGCTGCGCGCTCGCTCGCTCACTGAGGCCGCCCGGGCAAAGCCCGGGCGTCGGGCGACCTTTGGTCGCCCG
GCCTCAGTGAGCGAGCGAGCGCGCAGAGAGGGAGTGGGGTTAAACTCGTTACATAACTTACGGTAAATGGCCCG
CCTGGCTGACCGCCCAACGACCCCCGCCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGA
CTTTCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATCAAGTGTATCATATGCC
AAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCGCCTGGCATTATGCCCAGTACATGACCTTATGGGAC
TTTCCTACTTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGGTGATGCGGTTTTGGCAGTACATCAATGG
GCGTGGATAGCGGTTTGACTCACGGGGATTTCCAAGTCTCCACCCCATTGACGTCAATGGGAGTTTGTTTTGGCAC
CAAAATCAACGGGACTTTCCAAAATGTCGTAACAACTCCGCCCCATTGACGCAAATGGGCGGTAGGCGTGTACGG
TGGGAGGTCTATATAAGCAGAGCTCGTTTAGTGAACCGTCAGATCGCCTGGAGACGCCATCCACGCTGTTTTGACC
TCCATAGAAGACACCGGGACCGATCCAGCCTCCGGACTCTAGAGGATCCGGTACTCGAGGAACTGAAAAACCAGA
AAGTTAACTGGTAAGTTTAGTCTTTTTGTCTTTTATTTCAGGTCCCGGATCCGGTGGTGGTGCAAATCAAAGAACTG
CTCCTCAGTGGATGTTGCCTTTACTTCTAGGCCTGTACGGAAGTGTTACTTCTGCTCTAAAAGCTGCGGAATTGTAC
CCGGGGCCGATCCACCGGTCTTTTTCGCAACGGGTTTGCCGCCAGAACACAGGTAAGTGCCGTGTGTGGTTCCCG
CGGGCGGCGACGGGGCCCGTGCGTCCCAGCGCACATGTTCGGCGAGGCGGGGCCTGCGAGCGCGGCCACCGAG
AATCGGACGGGGGTAGTCTCAAGCTGGCCGGCCTGTTTGAATGAGGCTTCAGTACTTTACAGAATCGTTGCCTGC
ACATCTTGGAAACACTTGCTGGGATTACTTCTTCAGGTTAACCCAACAGAAGGCTCGAGAAGGTATATTGCTGTTG
ACAGTGAGCGCAACCGATTTCAAATGGTGCTAGAGTGAAGCCACAGATGTCTAGCACCATTTGAAATCGGTTATG
CCTACTGCCTCGGAATTCAAGGGGCTACTTTAGGAGCAATTATCTTGTTTACTAAAACTGAATACCTTGCTATCTCTT
TGATACATTGGCCGGCCTGCTCTGGTGCCTGGCCTCGCGCCGCCGTGTATCGCCCCGCCCTGGGCGGCAAGGCTG
GCCCGGTCGGCACCAGTTGCGTGAGCGGAAAGATGGCCGCTTCCCGGCCCTGCTGCAGGGAGCTCAAAATGGAG
GACGCGGCGCTCGGGAGAGCGGGCGGGTGAGTCACCCACACAAAGGAAAAGGGCCTTTCCGTCCTCAGCCGTCG
CTTCATGTGACTCCACGGAGTACCGGGCGCCGTCCAGGCACCTCGATTAGTTCTCGAGCTTTTGGAGTACGTCGTC
TTTAGGTTGGGGGGAGGGGTTTTATGCGATGGAGTTTCCCCACACTGAGTGGGTGGAGACTGAAGTTAGGCCAG
CTTGGCACTTGATGTAATTCTCCTTGGAATTTGCCCTTTTTGAGTTTGGATCTTGGTTCATTCTCAAGCCTCAGACAG
TGGTTCAAAGTTTTTTTCTTCCATTTCAGGTGTCGTGAAAAGCTAGCGCTACCGGACTCAGATCTCGAGCTCAAGCT
GCGGGGATCCAGACATGATAAGATACATTGATGAGTTTGGACAAACCACAACTAGAATGCAGTGAAAAAAATGCT
TTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTAACAACAACAATT
GCATTCATTTTATGTTTCAGGTTCAGGGGGAGGTGTGGGAGGTTTTTTCACTAGTAGCATGGCTACGTAGATAAGT
AGCATGGCGGGTTAATCATTAACTACAAGGAACCCCTAGTGATGGAGTTGGCCACTCCCTCTCTGCGCGCTCGCTC
GCTCACTGAGGCCGGGCGACCAAAGGTCGCCCGACGCCCGGGCTTTGCCCGGGCGGCCTCAGTGAGCGAGCGAG
CGCGCCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCGCCCTTCCCAACAGTTGCGCAGCCTGAATGGCGAA
TGGAATTCCAGACGATTGAGCGTCAAAATGTAGGTATTTCCATGAGCGTTTTTCCTGTTGCAATGGCTGGCGGTAA
TATTGTTCTGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTCAGGCAAGTGATGTTATTACTAATCAAA
GAAGTATTGCGACAACGGTTAATTTGCGTGATGGACAGACTCTTTTACTCGGTGGCCTCACTGATTATAAAAACAC
TTCTCAGGATTCTGGCGTACCGTTCCTGTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTCTGATTCTAA
CGAGGAAAGCACGTTATACGTGCTCGTCAAAGCAAC

**Figure 18B**

CATAGTACGCGCCCTGTAGCGGCGCATTAAGCGCGGCGGGTGTGGTGGTTACGCGCAGCGTGACCGCTACACTTG
CCAGCGCCCTAGCGCCCGCTCCTTTCGCTTTCTTCCCTTCCTTTCTCGCCACGTTCGCCGGCTTTCCCCGTCAAGCTCT
AAATCGGGGGCTCCCTTTAGGGTTCCGATTTAGTGCTTTACGGCACCTCGACCCCAAAAAACTTGATTAGGGTGAT
GGTTCACGTAGTGGGCCATCGCCCTGATAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCTTTAATAGTG
GACTCTTGTTCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTCTTTTGATTTATAAGGGATTTTGCCGATTT
CGGCCTATTGGTTAAAAAATGAGCTGATTTAACAAAAATTTAACGCGAATTTTAACAAAATATTAACGTTTACAATT
TAAATATTTGCTTATACAATCTTCCTGTTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGATTGACATGCTA
GTTTTACGATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTCAGGCAATGACCTGATAGCCTTTGTAGAGACC
TCTCAAAAATAGCTACCCTCTCCGGCATGAATTTATCAGCTAGAACGGTTGAATATCATATTGATGGTGATTTGACT
GTCTCCGGCCTTTCTCACCCGTTTGAATCTTTACCTACACATTACTCAGGCATTGCATTTAAAATATATGAGGGTTCT
AAAAATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAAAAGTATTACAGGGTCATAATGTTTTTGGTACAAC
CGATTTAGCTTTATGCTCTGAGGCTTTATTGCTTAATTTTGCTAATTCTTTGCCTTGCCTGTATGATTTATTGGATGTT
GGAATTCCTGATGCGGTATTTTCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGGTGCACTCTCAGTACAA
TCTGCTCTGATGCCGCATAGTTAAGCCAGCCCCGACACCCGCCAACACCCGCTGACGCGCCCTGACGGGCTTGTCT
GCTCCCGGCATCCGCTTACAGACAAGCTGTGACCGTCTCCGGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCA
CCGAAACGCGCGAGACGAAAGGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATGATAATAATGGTTTCTT
AGACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACCCCTATTTGTTTATTTTTCTAAATACATTCAAATATG
TATCCGCTCATGAGACAATAACCCTGATAAATGCTTCAATAATATTGAAAAAGGAAGAGTATGAGTATTCAACATT
TCCGTGTCGCCCTTATTCCCTTTTTTGCGGCATTTTGCCTTCCTGTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAA
AAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTACATCGAACTGGATCTCAACAGCGGTAAGATCCTTGAGA
GTTTTCGCCCCGAAGAACGTTTTCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATT
GACGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGAGTACTCACCAGTCACA
GAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTATGCAGTGCTGCCATAACCATGAGTGATAACACTGCG
GCCAACTTACTTCTGACAACGATCGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTA
ACTCGCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGACACCACGATGCCTGTA
GCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACTACTTACTCTAGCTTCCCGGCAACAATTAATAGACT
GGATGGAGGCGGATAAAGTTGCAGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAAT
CTGGAGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGCCCTCCCGTATCGTAG
TTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGA
TTAAGCATTGGTAACTGTCAGACCAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAA
GGATCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTCGTTCCACTGAGCGTCA
GACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTTTTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAA
ACCACCGCTACCAGCGGTGGTTTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGC
AGAGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAAGAACTCTGTAGCACCGC
CTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGCTGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGA
CTCAAGACGATAGTTACCGGATAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGG
AGCGAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACGCTTCCCGAAGGGAGA
AAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAGGAGAGCGCACGAGGGAGCTTCCAGGGGGAAAC
GCCTGGTATCTTTATAGTCCTGTCGGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGG
GCGGAGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGGCCTTTTGCTCACATG
TTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTACCGCCTTTGAGTGAGCTGATACCGCTCGCCGCAG
CCGAACGACCGAGCGCAGCGAGTCAGTGAGCGAGGAAGCGGAAGAGCGCCCAATACGCAAACCGCCTCTCCCCG
CGCGTTGGCCGATTCATTAATG

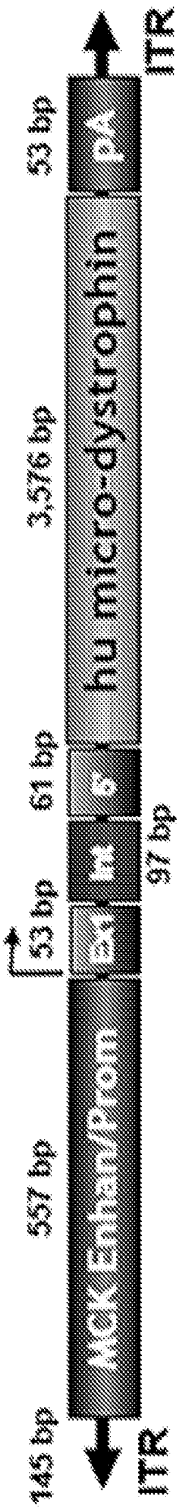

ITR
53 bp
PA
3,576 bp
hu micro-dystrophin
61 bp
5'
97 bp
I1
53 bp
E1
557 bp
MCK Enhan/Prom
145 bp
ITR

Figure 19

## Figure 20A

SEQ ID NO: 9
Main features:
MCK promoter
Chimeric intron sequence
Human micro-dystrophin sequence
Poly A tail
Ampicillin resistance
pGEX plasmid backbone with pBR322 origin or replication

GCCCAATACGCAAACCGCCTCTCCCCGCGCGTTGGCCGATTCATTAATGCAGCTGGC
GCGCTCGCTCGCTCACTGAGGCCGCCCGGGCAAAGCCCGGGCGTCGGGCGACCTTT
GGTCGCCCGGCCTCAGTGAGCGAGCGAGCGCGCAGAGAGGGAGTGGCCAACTCCAT
CACTAGGGGTTCCTTGTAGTTAATGATTAACCCGCCATGCTAATTATCTACGTAGCC
ATGTCTAGACAGCCACTATGGGTCTAGGCTGCCCATGTAAGGAGGCAAGGCCTGGG
GACACCCGAGATGCCTGGTTATAATTAACCCAGACATGTGGCTGCTCCCCCCCCCCA
ACACCTGCTGCCTGAGCCTCACCCCCACCCCGGTGCCTGGGTCTTAGGCTCTGTACA
CCATGGAGGAGAAGCTCGCTCTAAAAATAACCCTGTCCCTGGTGGGCTGTGGGGGA
CTGAGGGCAGGCTGTAACAGGCTTGGGGGCCAGGGCTTATACGTGCCTGGGACTCC
CAAAGTATTACTGTTCCATGTTCCCGGCGAAGGGCCAGCTGTCCCCCGCCAGCTAGA
CTCAGCACTTAGTTTAGGAACCAGTGAGCAAGTCAGCCCTTGGGGCAGCCCATACA
AGGCCATGGGGCTGGGCAAGCTGCACGCCTGGGTCCGGGGTGGGCACGGTGCCCGG
GCAACGAGCTGAAAGCTCATCTGCTCTCAGGGGCCCCTCCCTGGGGACAGCCCCTCC
TGGCTAGTCACACCCTGTAGGCTCCTCTATATAACCCAGGGGCACAGGGGCTGCCCC
CGGGTCACCACCACCTCCACAGCACAGACAGACACTCAGGAGCCAGCCAGCCAGGT
AAGTTTAGTCTTTTTGTCTTTTATTTCAGGTCCCGGATCCGGTGGTGGTGCAAATCAA
AGAACTGCTCCTCAGTGGATGTTGCCTTTACTTCTAGGCCTGTACGGAAGTGTTACTT
CTGCTCTAAAAGCTGCGGAATTGTACCCGCGGCCGCCACCATGCTGTGGTGGGAGG
AGGTGGAGGATTGTTATGAAAGGGAGGACGTGCAGAAGAAGACTTTTACCAAGTGG
GTGAACGCTCAGTTCAGCAAATTTGGGAAGCAGCACATCGAGAATCTGTTTTCCGAC
CTGCAGGATGGGAGACGGCTGCTGGATCTGCTGGAAGGACTGACTGGCCAGAAGCT
GCCCAAAGAGAAGGGGAGCACTAGGGTGCACGCCCTGAACAACGTGAACAAAGCT
CTGAGAGTGCTGCAGAACAACAACGTGGATCTGGTGAATATTGGCAGTACTGATAT
CGTGGACGGGAACCACAAACTGACACTGGGCCTGATCTGGAACATTATTCTGCACTG
GCAGGTGAAAAATGTGATGAAGAACATCATGGCCGGGCTGCAGCAGACCAATTCCG
AGAAGATCCTGCTGTCTTGGGTGCGGCAGAGCACCCGCAACTATCCCCAGGTGAAC
GTGATTAACTTCACTACATCCTGGAGCGACGGGCTGGCCCTGAATGCTCTGATTCAC
AGCCACAGGCCTGATCTGTTCGACTGGAATAGCGTGGTGTGCCAGCAGTCTGCCACA
CAGCGCCTGGAACATGCCTTCAATATCGCTCGGTACCAGCTGGGGATCGAAAAACT
GCTGGACCCAGAGGATGTGGACACTACATACCCAGATAAAAAGTCTATTCTGATGT
ACATTACTAGCCTGTTCCAGGTGCTGCCACAGCAGGTGTCTATTGAAGCCATTCAGG
AGGTGGAAATGCTGCCCCGCCCCCCCAAAGTGACTAAAGAGGAGCATTTTCAGCTG
CATCATCAGATGCATTACAGCCAGCAGATTACCGTGAGCCTGGCTCAGGGATATGA
GCGCACCAGTAGTCCAAAACCACGGTTCAAGTCCTACGCTTATAC

**Figure 20B**

CCAGGCTGCCTACGTGACAACTAGCGACCCTACTAGATCCCCCTTTCCATCCCAGCA
CCTGGAGGCCCCAGAGGACAAGAGCTTTGGGTCCAGCCTGATGG
AAAGCGAGGTGAATCTGGATCGGTACCAGACAGCCCTGGAGGAGGTGCTGAGCTGG
CTGCTGAGTGCTGAAGACACACTGCAGGCCCAGGGCGAAATTTCCAATGACGTGGA
AGTGGTGAAGGATCAGTTCCACACACACGAGGGCTATATGATGGACCTGACAGCTC
ACCAGGGGCGCGTGGGCAATATCCTGCAGCTGGGCTCTAAACTGATCGGCACCGGG
AAACTGAGTGAGGACGAGGAAACAGAAGTGCAGGAGCAGATGAACCTGCTGAACA
GCCGCTGGGAGTGTCTGAGAGTGGCTAGTATGGAGAAGCAGTCCAACCTGCACCGG
GTGCTGATGGACCTGCAGAACCAGAAACTGAAAGAGCTGAACGACTGGCTGACAAA
GACTGAGGAACGCACAAGGAAGATGGAGGAGGAGCCACTGGGACCCGACCTGGAG
GATCTGAAGAGACAGGTGCAGCAGCATAAGGTGCTGCAGGAGGATCTGGAACAGG
AGCAGGTGCGGGTGAACTCCCTGACACATATGGTGGTGGTGGTGGACGAATCTAGT
GGAGATCACGCCACCGCCGCCCTGGAGGAACAGCTGAAGGTGCTGGGGGACCGGTG
GGCCAACATTTGCCGGTGGACCGAGGACAGGTGGGTGCTGCTGCAGGACATCCTGC
TGAAATGGCAGAGGCTGACCGAGGAGCAGTGTCTGTTTAGTGCTTGGCTGAGCGAG
AAAGAGGACGCCGTGAACAAGATCCACACAACCGGCTTTAAGGATCAGAACGAAAT
GCTGTCTAGCCTGCAGAAACTGGCTGTGCTGAAGGCCGATCTGGAGAAAAAGAAGC
AGAGCATGGGCAAACTGTATAGCCTGAAACAGGACCTGCTGAGCACCCTGAAGAAC
AAGAGCGTGACCCAGAAGACAGAAGCCTGGCTGGATAACTTTGCCCGCTGCTGGGA
CAACCTGGTGCAGAAACTGGAGAAAAGTACAGCTCAGATCTCTCAGGCTGTGACCA
CAACCCAGCCTAGCCTGACCCAGACAACCGTGATGGAAACCGTGACCACCGTGACA
ACCCGCGAACAGATCCTGGTGAAACATGCCCAGGAAGAGCTGCCACCTCCACCTCC
CCAGAAGAAGAGAACCCTGGAGCGGCTGCAGGAGCTGCAGGAAGCCACTGACGAA
CTGGACCTGAAGCTGAGGCAGGCCGAAGTGATTAAGGGGTCTTGGCAGCCTGTGGG
CGATCTGCTGATTGATTCCCTGCAGGACCACCTGGAAAAGGTGAAGGCTCTGAGAG
GCGAAATTGCTCCACTGAAGGAGAACGTGAGTCATGTGAACGATCTGGCTAGACAG
CTGACAACACTGGGCATCCAGCTGAGCCCATACAATCTGAGCACACTGGAGGACCT
GAATACCAGGTGGAAGCTGCTGCAGGTGGCTGTGGAAGACCGGGTGCGGCAGCTGC
ATGAGGCCCATCGCGACTTCGGACCAGCCAGCCAGCACTTTCTGAGCACATCCGTGC
AGGGGCCCTGGGAGAGGGCCATTTCTCCCAACAAGGTGCCCTACTATATTAATCACG
AGACCCAGACCACTTGTTGGGACCATCCCAAGATGACAGAACTGTACCAGTCCCTG
GCCGATCTGAACAACGTGAGGTTTAGCGCTTACAGAACCGCTATGAAGCTGAGACG
GCTGCAGAAGGCCCTGTGCCTGGATCTGCTGTCCCTGTCCGCCGCCTGCGATGCCCT
GGATCAGCATAATCTGAAGCAGAACGATCAGCCAATGGATATCCTGCAGATCATCA
ACTGCCTGACCACTATCTACGACAGGCTGGAGCAGGAGCACAACAACCTGGTGAAC
GTGCCTCTGTGCGTGGATATGTGCCTGAACTGGCTGCTGAACGTGTATGACACTGGG
CGCACCGGCCGGATCAGAGTGCTGAGTTTTAAAACTGGGATTATCTCCCTGTGTAAG
GCCCACCTGGAGGACAAGTACAGGTACCTGTTCAAGCAGGTGGCTAGTAGCACTGG
ATTTTGTGACCAGCGCCGCCTGGGACTGCTGCTGCATGATAGTATCCAGATTCCTAG
ACAGCTGGGAGAGGTGGCTAGTTTCGGAGGATCTAACATCGAACCCAGCGTGCGCA
GCTGTTTCCAGTTTGCCAATAACAAACCTGAAATCGAGGCTGCTCTGTTCCTGGATT
GGATGCGCCTGGAACCACAGAGCATGGTGTGGCTGCCTGTGCTGCACAGAGTGG

**Figure 20C**

CTGCCGCCGAAACTGCCAAGCACCAGGCTAAATGCAACATCTGCAAGGAATGTCCC
ATTATCGGCTTTCGCTACAGGAGTCTGAAACATTTTAACTACGAT
ATTTGCCAGAGCTGCTTCTTTTCCGGAAGAGTGGCCAAAGGACACAAGATGCACTAC
CCTATGGTGGAATATTGCACCCCAACTACATCTGGCGAAGATGTGCGCGATTTTGCC
AAGGTGCTGAAGAATAAGTTTCGGACTAAGAGGTACTTCGCCAAGCACCCCCGCAT
GGGGTATCTGCCAGTGCAGACAGTGCTGGAAGGAGACAATATGGAGACCGATACAA
TGTGAGCGGCCGCAATAAAAGATCTTTATTTTCATTAGATCTGTGTGTTGGTTTTTTG
TGTGTCTAGAGCATGGCTACGTAGATAAGTAGCATGGCGGGTTAATCATTAACTACA
AGGAACCCCTAGTGATGGAGTTGGCCACTCCCTCTCTGCGCGCTCGCTCGCTCACTG
AGGCCGGGCGACCAAAGGTCGCCCGACGCCCGGGCTTTGCCCGGGCGGCCTCAGTG
AGCGAGCGAGCGCGCCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCGCCCTT
CCCAACAGTTGCGCAGCCTGAATGGCGAATGGAAGTTCCAGACGATTGAGCGTCAA
AATGTAGGTATTTCCATGAGCGTTTTTCCTGTTGCAATGGCTGGCGGTAATATTGTTC
TGGATATTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTCAGGCAAGTGATGTTA
TTACTAATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTGATGGACAGACTCTTT
TACTCGGTGGCCTCACTGATTATAAAAACACTTCTCAGGATTCTGGCGTACCGTTCCT
GTCTAAAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTCTGATTCTAACGAGGAA
AGCACGTTATACGTGCTCGTCAAAGCAACCATAGTACGCGCCCTGTAGCGGCGCATT
AAGCGCGGCGGGTGTGGTGGTTACGCGCAGCGTGACCGCTACACTTGCCAGCGCCC
TAGCGCCCGCTCCTTTCGCTTTCTTCCCTTCCTTTCTCGCCACGTTCGCCGGCTTTCCC
CGTCAAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGATTTAGTGATTTACGGCAC
CTCGACCCCAAAAAACTTGATTAGGGTGATGGTTCACGTAGTGGGCCATCGCCCTGA
TAGACGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCTTTAATAGTGGACTCTTGT
TCCAAACTGGAACAACACTCAACCCTATCTCGGTCTATTCTTTTGATTTATAAGGGAT
TTTGCCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATTTAACAAAAATTTAACGC
GAATTTTAACAAAATATTAACGTTTACAATTTAAATATTTGCTTATACAATCTTCCTG
TTTTTGGGGCTTTTCTGATTATCAACCGGGGTACATATGATTGACATGCTAGTTTTAC
GATTACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTCAGGCAATGACCTGATAGC
CTTTGTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGCATGAATTTATCAGCTAGA
ACGGTTGAATATCATATTGATGGTGATTTGACTGTCTCCGGCCTTTCTCACCCGTTTG
AATCTTTACCTACACATTACTCAGGCATTGCATTTAAAATATATGAGGGTTCTAAAA
ATTTTTATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAAAAGTATTACAGGGTCATA
ATGTTTTTGGTACAACCGATTTAGCTTTATGCTCTGAGGCTTTATTGCTTAATTTTGCT
AATTCTTTGCCTTGCCTGTATGATTTATTGGATGTTGGAAGTTCCTGATGCGGTATTT
TCTCCTTACGCATCTGTGCGGTATTTCACACCGCATATGGTGCACTCTCAGTACAATC
TGCTCTGATGCCGCATAGTTAAGCCAGCCCCGACACCCGCCAACACCCGCTGACGCG
CCCTGACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGACAAGCTGTGACCGTCTCC
GGGAGCTGCATGTGTCAGAGGTTTTCACCGTCATCACCGAAACGCGCGAGACGAAA
GGGCCTCGTGATACGCCTATTTTTATAGGTTAATGTCATGATAATAATGGTTTCTTAG
ACGTCAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACCCCTATTTGTTTATTTTTCT
AAATACATTCAAATATGTATCCGCTCATGAGACAATAACCCTGATAAATGCTTCAAT
AATATTGAAAAAGGAAGAGTATGAGTATTCAACATTTCCGTGTCGCCCTT

**Figure 20D**

ATTCCCTTTTTTGCGGCATTTTGCCTTCCTGTTTTTGCTCACCCAGAAACGCTGGTGA
AAGTAAAAGATGCTGAAGATCAGTTGGGTGCACGAGTGGGTTAC
ATCGAACTGGATCTCAACAGCGGTAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGT
TTTCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTG
ACGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTG
AGTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTA
TGCAGTGCTGCCATAACCATGAGTGATAACACTGCGGCCAACTTACTTCTGACAACG
ATCGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAAC
TCGCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTG
ACACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAA
CTACTTACTCTAGCTTCCCGGCAACAATTAATAGACTGGATGGAGGCGGATAAAGTT
GCAGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCT
GGAGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAA
GCCCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACG
AAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAG
ACCAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAG
GATCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTT
TTCGTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCC
TTTTTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAACCACCGCTACCAGCGGT
GGTTTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAG
CAGAGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTT
CAAGAACTCTGTAGCACCGCGTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGC
TGCTGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACC
GGATAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGG
AGCGAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCC
ACGCTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAA
CAGGAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCT
GTCGGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGG
CGGAGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGC
TGGCCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTA
TTACCGGGTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGC
GAGTCAGTGAGCGACCAAGCGGAAGAGC

**Figure 21A**

SEQ ID NO: 7 : human micro-dystrophin gene

ATGCTGTGGTGGGAGGAGGTGGAGGATTGTTATGAAAGGGAGGACGTGCAGAAGA
AGACTTTTACCAAGTGGGTGAACGCTCAGTTCAGCAAATTTGGGAAGCAGCACATC
GAGAATCTGTTTTCCGACCTGCAGGATGGGAGACGGCTGCTGGATCTGCTGGAAGG
ACTGACTGGCCAGAAGCTGCCCAAAGAGAAGGGGAGCACTAGGGTGCACGCCCTGA
ACAACGTGAACAAAGCTCTGAGAGTGCTGCAGAACAACAACGTGGATCTGGTGAAT
ATTGGCAGTACTGATATCGTGGACGGGAACCACAAACTGACACTGGGCCTGATCTG
GAACATTATTCTGCACTGGCAGGTGAAAAATGTGATGAAGAACATCATGGCCGGGC
TGCAGCAGACCAATTCCGAGAAGATCCTGCTGTCTTGGGTGCGGCAGAGCACCCGC
AACTATCCCCAGGTGAACGTGATTAACTTCACTACATCCTGGAGCGACGGGCTGGCC
CTGAATGCTCTGATTCACAGCCACAGGCCTGATCTGTTCGACTGGAATAGCGTGGTG
TGCCAGCAGTCTGCCACACAGCGCCTGGAACATGCCTTCAATATCGCTCGGTACCAG
CTGGGGATCGAAAAACTGCTGGACCCAGAGGATGTGGACACTACATACCCAGATAA
AAAGTCTATTCTGATGTACATTACTAGCCTGTTCCAGGTGCTGCCACAGCAGGTGTC
TATTGAAGCCATTCAGGAGGTGGAAATGCTGCCCCGCCCCCCCAAAGTGACTAAAG
AGGAGCATTTTCAGCTGCATCATCAGATGCATTACAGCCAGCAGATTACCGTGAGCC
TGGCTCAGGGATATGAGCGCACCAGTAGTCCAAAACCACGGTTCAAGTCCTACGCTT
ATACCCAGGCTGCCTACGTGACAACTAGCGACCCTACTAGATCCCCCTTTCCATCCC
AGCACCTGGAGGCCCCAGAGGACAAGAGCTTTGGGTCCAGCCTGATGGAAAGCGAG
GTGAATCTGGATCGGTACCAGACAGCCCTGGAGGAGGTGCTGAGCTGGCTGCTGAG
TGCTGAAGACACACTGCAGGCCCAGGGCGAAATTTCCAATGACGTGGAAGTGGTGA
AGGATCAGTTCCACACACACGAGGGCTATATGATGGACCTGACAGCTCACCAGGGG
CGCGTGGGCAATATCCTGCAGCTGGGCTCTAAACTGATCGGCACCGGGAAACTGAG
TGAGGACGAGGAAACAGAAGTGCAGGAGCAGATGAACCTGCTGAACAGCCGCTGG
GAGTGTCTGAGAGTGGCTAGTATGGAGAAGCAGTCCAACCTGCACCGGGTGCTGAT
GGACCTGCAGAACCAGAAACTGAAAGAGCTGAACGACTGGCTGACAAAGACTGAG
GAACGCACAAGGAAGATGGAGGAGGAGCCACTGGGACCCGACCTGGAGGATCTGA
AGAGACAGGTGCAGCAGCATAAGGTGCTGCAGGAGGATCTGGAACAGGAGCAGGT
GCGGGTGAACTCCCTGACACATATGGTGGT

**Figure 21B**

GGTGGTGGACGAATCTAGTGGAGATCACGCCACCGCCGCCCTGGAGGAACAGCTGA
AGGTGCTGGGGGACCGGTGGGCCAACATTTGCCGGTGGACCGAGGACAGGTGGGTG
CTGCTGCAGGACATCCTGCTGAAATGGCAGAGGCTGACCGAGGAGCAGTGTCTGTTT
AGTGCTTGGCTGAGCGAGAAAGAGGACGCCGTGAACAAGATCCACACAACCGGCTT
TAAGGATCAGAACGAAATGCTGTCTAGCCTGCAGAAACTGGCTGTGCTGAAGGCCG
ATCTGGAGAAAAAGAAGCAGAGCATGGGCAAACTGTATAGCCTGAAACAGGACCTG
CTGAGCACCCTGAAGAACAAGAGCGTGACCCAGAAGACAGAAGCCTGGCTGGATA
ACTTTGCCCGCTGCTGGGACAACCTGGTGCAGAAACTGGAGAAAAGTACAGCTCAG
ATCTCTCAGGCTGTGACCACAACCCAGCCTAGCCTGACCCAGACAACCGTGATGGA
AACCGTGACCACCGTGACAACCCGCGAACAGATCCTGGTGAAACATGCCCAGGAAG
AGCTGCCACCTCCACCTCCCCAGAAGAAGAGAACCCTGGAGCGGCTGCAGGAGCTG
CAGGAAGCCACTGACGAACTGGACCTGAAGCTGAGGCAGGCCGAAGTGATTAAGGG
GTCTTGGCAGCCTGTGGGCGATCTGCTGATTGATTCCCTGCAGGACCACCTGGAAAA
GGTGAAGGCTCTGAGAGGCGAAATTGCTCCACTGAAGGAGAACGTGAGTCATGTGA
ACGATCTGGCTAGACAGCTGACAACACTGGGCATCCAGCTGAGCCCATACAATCTG
AGCACACTGGAGGACCTGAATACCAGGTGGAAGCTGCTGCAGGTGGCTGTGGAAGA
CCGGGTGCGGCAGCTGCATGAGGCCCATCGCGACTTCGGACCAGCCAGCCAGCACT
TTCTGAGCACATCCGTGCAGGGGCCCTGGGAGAGGGCCATTTCTCCCAACAAGGTG
CCCTACTATATTAATCACGAGACCCAGACCACTTGTTGGGACCATCCCAAGATGACA
GAACTGTACCAGTCCCTGGCCGATCTGAACAACGTGAGGTTTAGCGCTTACAGAACC
GCTATGAAGCTGAGACGGCTGCAGAAGGCCCTGTGCCTGGATCTGCTGTCCCTGTCC
GCCGCCTGCGATGCCCTGGATCAGCATAATCTGAAGCAGAACGATCAGCCAATGGA
TATCCTGCAGATCATCAACTGCCTGACCACTATCTACGACAGGCTGGAGCAGGAGC
ACAACAACCTGGTGAACGTGCCTCTGTGCGTGGATATGTGCCTGAACTGGCTGCTGA
ACGTGTATGACACTGGGCGCACCGGCCGGATCAGAGTGCTGAGTTTTAAAACTGGG
ATTATCTCCCTGTGTAAGGCCCACCTGGAGGACAAGTACAGGTACCTGTTCAAGCAG
GTGGCTAGTAGCACTGGATTTTGTGACCAGCGCCGCCTGGGACTGCTGCTGCATGAT
AGTATCCAGATTCCTAGACAGCTGGGAGAGG

**Figure 21C**

TGGCTAGTTTCGGAGGATCTAACATCGAACCCAGCGTGCGCAGCTGTTTCCAGTTTG
CCAATAACAAACCTGAAATCGAGGCTGCTCTGTTCCTGGATTGGATGCGCCTGGAAC
CACAGAGCATGGTGTGGCTGCCTGTGCTGCACAGAGTGGCTGCCGCCGAAACTGCC
AAGCACCAGGCTAAATGCAACATCTGCAAGGAATGTCCCATTATCGGCTTTCGCTAC
AGGAGTCTGAAACATTTTAACTACGATATTTGCCAGAGCTGCTTCTTTTCCGGAAGA
GTGGCCAAAGGACACAAGATGCACTACCCTATGGTGGAATATTGCACCCCAACTAC
ATCTGGCGAAGATGTGCGCGATTTTGCCAAGGTGCTGAAGAATAAGTTTCGGACTAA
GAGGTACTTCGCCAAGCACCCCCGCATGGGGTATCTGCCAGTGCAGACAGTGCTGG
AAGGAGACAATATGGAGACCGATACAATGTGAGC

## Figure 22A

**pAAV.MCK.miR29C Sequence (SEQ ID NO: 12)**

Main features:
MCK enhancer- 190-395 (SEQ ID NO: 10)
MCK promoter- 396-753 (SEQ ID NO: 11)
EF1a Intron- 1155-1315, 1609-2083
miR-29c-1487-1512
shRNA-miR29-c with primary seed sequence- 1316-1608
SV40 PolyA- 2094-2146

CTGNNNNNNGCGCGCTCGCTCGCTCACTGAGGCCGCCCGGGCAAAGCCCGGGCGTC
GGGCGACCTTTGGTCGCCCGGCCTCAGTGAGCGAGCGAGCGCGCAGAGAGGGAGTG
GCCAACTCCATCACTAGGGGTTCCTTGTAGTTAATGATTAACCCGCCATGCTAATTA
TCTACGTAGCCATGTCTAGACAGCCACTATGGGTCTAGGCTGCCCATGTAAGGAGGC
AAGGCCTGGGGACACCCGAGATGCCTGGTTATAATTAACCCAGACATGTGGCTGCTC
CCCCCCCCCAACACCTGCTGCCTGAGCCTCACCCCCACCCCGGTGCCTGGGTCTTAG
GCTCTGTACACCATGGAGGAGAAGCTCGCTCTAAAAATAACCCTGTCCCTGGTGGGC
TGTGGGGGACTGAGGGCAGGCTGTAACAGGCTTGGGGGCCAGGGCTTATACGTGCC
TGGGACTCCCAAAGTATTACTGTTCCATGTTCCCGGCGAAGGGCCAGCTGTCCCCCG
CCAGCTAGACTCAGCACTTAGTTTAGGAACCAGTGAGCAAGTCAGCCCTTGGGGCA
GCCCATACAAGGCCATGGGGCTGGGCAAGCTGCACGCCTGGGTCCGGGGTGGGCAC
GGTGCCCGGGCAACGAGCTGAAAGCTCATCTGCTCTCAGGGGCCCCTCCCTGGGGA
CAGCCCCTCCTGGCTAGTCACACCCTGTAGGCTCCTCTATATAACCCAGGGGCACAG
GGGCTGCCCCCGGGTCACCACCACCTCCACAGCACAGACAGACACTCAGGAGCCAG
CCAGCCAGGTAAGTTTAGTCTTTTTGTCTTTTATTTCAGGTCCCGGATCCGGTGGTGG
TGCAAATCAAAGAACTGCTCCTCAGTGGATGTTGCCTTTACTTCTAGGCCTGTACGG
AAGTGTTACTTCTGCTCTAAAAGCTGCGGAATTGTACCCGCCTAGAGGATCCGGTAC
TCGAGGAACTGAAAAACCAGAAAGTTAACTGGTAAGTTTAGTCTTTTTGTCTTTTAT
TTCAGGTCCCGGATCCGGTGGTGGTGCAAATCAAAGAACTGCTCCTCAGTGGATGTT
GCCTTTACTTCTAGGCCTGTACGGAAGTGTTACTTCTGCTCTAAAAGCTGCGGAATT
GTACCCGGGGCCGATCCACCGGTCTTTTTCGCAACGGGTTTGCCGCCAGAACACAGG
TAAGTGCCGTGTGTGGTTCCCGCGGGCGGCGACGGGGCCCGTGCGTCCCAGCGCAC
ATGTTCGGCGAGGCGGGGCCTGCGAGCGCGGCCACCGAGAATCGGACGGGGGTAGT
CTCAAGCTGGCCGGCCTGTTTGAATGAGGCTTCAGTACTTTACAGAATCGTTGCCTG
CACATCTTGGAAACACTTGCTGGGATTACTTCTTCAGGTTAACCCAACAGAAGGCTC
GAGAAGGTATATTGCTGTTGACAGTGAGCGCAACCGATTTCAAATGGTGCTAGAGT
GAAGCCACAGATGTCTAGCACCATTTGAAATCGGTTATGCCTACTGCCTCGGAATTC
AAGGGGCTACTTTAGGAGCAATTATCTTGTTTACTAAAACTGAATACCTTGCTATCT
CTTTGATACATTGGCCGGCCTGCTCTGGTGCCTGGCCTCGCGCCGCCGTGTATCGCC
CCGCCCTGGGCGGCAAGGCTGGCCCGGTCGGCACCAGTTGCGTGAGCGGAAAGATG
GCCGCTTCCCGGCCCTGCTGCAGGGAGCTCAAAATGGAGGACGCGGCGCTCGGGAG
AGCGGGCGGGTGAGTCACCCACACAAAGGAAAAGGGCCTTTCCGTCCTCAGCCGTC
GCTTCATGTGACTCCACGGAGTACCGGGCGCCGTCCAGGCACCTCGATTAGTT

**Figure 22B**

CTCGAGCTTTTGGAGTACGTCGTCTTTAGGTTGGGGGGAGGGGTTTTATGCGATGGA
GTTTCCCCACACTGAGTGGGTGGAGACTGAAGTTAGGCCAGCTTGGCACTTGATGTA
ATTCTCCTTGGAATTTGCCCTTTTTGAGTTTGGATCTTGGTTCATTCTCAAGCCTCAG
ACAGTGGTTCAAAGTTTTTTTCTTCCATTTCAGGTGTCGTGAAAAGCTAGTGCGGCC
GCAATAAAAGATCTTTATTTTCATTAGATCTGTGTGTTGGTTTTTTGTGTGTCTAGAC
ATGGCTACGTAGATAATTAGCATGGCGGGTTAATCATTAACTACAAGGAACCCCTAG
TGATGGAGTTGGCCACTCCCTCTCTGCGCGCTCGCTCGCTCACTGAGGCCGGGCGAC
CAAAGGTCGCCCGACGCCCGGGCTTTGCCCGGGCGGCCTCAGTGAGCGAGCGAGCG
CGCNNNNNNCAGCTGGCGTAATAGCGAAGAGGCCCGCACCGATCGCCCTTCCCAAC
AGTTGCGCAGCCTGAATGGCGAATGGAAGTTCCAGACGATTGAGCGTCAAAATGTA
GGTATTTCCATGAGCGTTTTTCCTGTTGCAATGGCTGGCGGTAATATTGTTCTGGATA
TTACCAGCAAGGCCGATAGTTTGAGTTCTTCTACTCAGGCAAGTGATGTTATTACTA
ATCAAAGAAGTATTGCGACAACGGTTAATTTGCGTGATGGACAGACTCTTTTACTCG
GTGGCCTCACTGATTATAAAAACACTTCTCAGGATTCTGGCGTACCGTTCCTGTCTA
AAATCCCTTTAATCGGCCTCCTGTTTAGCTCCCGCTCTGATTCTAACGAGGAAAGCA
CGTTATACGTGCTCGTCAAAGCAACCATAGTACGCGCCCTGTAGCGGCGCATTAAGC
GCGGCGGGTGTGGTGGTTACGCGCAGCGTGACCGCTACACTTGCCAGCGCCCTAGC
GCCCGCTCCTTTCGCTTTCTTCCCTTCCTTTCTCGCCACGTTCGCCGGCTTTCCCCGTC
AAGCTCTAAATCGGGGGCTCCCTTTAGGGTTCCGATTTAGTGATTTACGGCACCTCG
ACCCCAAAAAACTTGATTAGGGTGATGGTTCACGTAGTGGGCCATCGCCCTGATAGA
CGGTTTTTCGCCCTTTGACGTTGGAGTCCACGTTCTTTAATAGTGGACTCTTGTTCCA
AACTGGAACAACACTCAACCCTATCTCGGTCTATTCTTTTGATTTATAAGGGATTTTG
CCGATTTCGGCCTATTGGTTAAAAAATGAGCTGATTTAACAAAAATTTAACGCGAAT
TTTAACAAAATATTAACGTTTACAATTTAAATATTTGCTTATACAATCTTCCTGTTTT
TGGGGCTTTTCTGATTATCAACCGGGGTACATATGATTGACATGCTAGTTTTACGATT
ACCGTTCATCGATTCTCTTGTTTGCTCCAGACTCTCAGGCAATGACCTGATAGCCTTT
GTAGAGACCTCTCAAAAATAGCTACCCTCTCCGGCATGAATTTATCAGCTAGAACGG
TTGAATATCATATTGATGGTGATTTGACTGTCTCCGGCCTTTCTCACCCGTTTGAATC
TTTACCTACACATTACTCAGGCATTGCATTTAAAATATATGAGGGTTCTAAAAATTTT
TATCCTTGCGTTGAAATAAAGGCTTCTCCCGCAAAAGTATTACAGGGTCATAATGTT
TTTGGTACAACCGATTTAGCTTTATGCTCTGAGGCTTTATTGCTTAATTTTGCTAATT
CTTTGCCTTGCCTGTATGATTTATTGGATGTTGGAAGTTCCTGATGCGGTATTTTCTC
CTTACGCATCTGTGCGGTATTTCACACCGCATATGGTGCACTCTCAGTACAATCTGCT
CTGATGCCGCATAGTTAAGCCAGCCCCGACACCCGCCAACACCCGCTGACGCGCCCT
GACGGGCTTGTCTGCTCCCGGCATCCGCTTACAGACAAGCTGTGACCGTCTCCGGGA
GCTGCATGTGTCAGAGGTTTTCACCGTCATCACCGAAACGCGCGAGACGAAAGGGC
CTCGTGATACGCCTATTTTTATAGGTTAATGTCATGATAATAATGGTTTCTTAGACGT
CAGGTGGCACTTTTCGGGGAAATGTGCGCGGAACCCCTATTTGTTTATTTTTCTAAAT
ACATTCAAATATGTATCCGCTCATGAGACAATAACCCTGATAAATGCTTCAATAATA
TTGAAAAAGGAAGAGTATGAGTATTCAACATTTCCGTGTCGCCCTTATTCCCTTTTTT
GCGGCATTTTGCCTTCCTGTTTTTGCTCACCCAGAAACGCTGGTGAAAGTAAAAGAT
GCTGAAGATCAGTTGGGTGCACGAGTGGGTTACATCGAACTGGATCTCAACAGCGG
TAAGATCCTTGAGAGTTTTCGCCCCGAAGAACGTT

**Figure 22C**

TTCCAATGATGAGCACTTTTAAAGTTCTGCTATGTGGCGCGGTATTATCCCGTATTGA
CGCCGGGCAAGAGCAACTCGGTCGCCGCATACACTATTCTCAGAATGACTTGGTTGA
GTACTCACCAGTCACAGAAAAGCATCTTACGGATGGCATGACAGTAAGAGAATTAT
GCAGTGCTGCCATAACCATGAGTGATAACACTGCGGCCAACTTACTTCTGACAACGA
TCGGAGGACCGAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACTC
GCCTTGATCGTTGGGAACCGGAGCTGAATGAAGCCATACCAAACGACGAGCGTGAC
ACCACGATGCCTGTAGCAATGGCAACAACGTTGCGCAAACTATTAACTGGCGAACT
ACTTACTCTAGCTTCCCGGCAACAATTAATAGACTGGATGGAGGCGGATAAAGTTGC
AGGACCACTTCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGG
AGCCGGTGAGCGTGGGTCTCGCGGTATCATTGCAGCACTGGGGCCAGATGGTAAGC
CCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTATGGATGAACGA
AATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAGAC
CAAGTTTACTCATATATACTTTAGATTGATTTAAAACTTCATTTTTAATTTAAAAGGA
TCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTC
GTTCCACTGAGCGTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTT
TTTTCTGCGCGTAATCTGCTGCTTGCAAACAAAAAAACCACCGCTACCAGCGGTGGT
TTGTTTGCCGGATCAAGAGCTACCAACTCTTTTTCCGAAGGTAACTGGCTTCAGCAG
AGCGCAGATACCAAATACTGTCCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAA
GAACTCTGTAGCACCGCGTACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGC
TGCCAGTGGCGATAAGTCGTGTCTTACCGGGTTGGACTCAAGACGATAGTTACCGGA
TAAGGCGCAGCGGTCGGGCTGAACGGGGGGTTCGTGCACACAGCCCAGCTTGGAGC
GAACGACCTACACCGAACTGAGATACCTACAGCGTGAGCTATGAGAAAGCGCCACG
CTTCCCGAAGGGAGAAAGGCGGACAGGTATCCGGTAAGCGGCAGGGTCGGAACAG
GAGAGCGCACGAGGGAGCTTCCAGGGGGAAACGCCTGGTATCTTTATAGTCCTGTC
GGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTTGTGATGCTCGTCAGGGGGGCGG
AGCCTATGGAAAAACGCCAGCAACGCGGCCTTTTTACGGTTCCTGGCCTTTTGCTGG
CCTTTTGCTCACATGTTCTTTCCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTAC
CGGGTTTGAGTGAGCTGATACCGCTCGCCGCAGCCGAACGACCGAGCGCAGCGAGT
CAGTGAGCGACCAAGCGGAAGAGC