SEQUENCE LISTING

<110> SANGAMO THERAPEUTICS, INC.

<120> GENETIC MODIFICATION OF CYTOKINE INDUCIBLE SH2-CONTAINING PROTEIN (CISH) GENE

<130> 8325-0167.40

<140> PCT/US2018/060038
<141> 2018-11-09

<150> 62/716,002
<151> 2018-08-08

<150> 62/583,724
<151> 2017-11-09

<160> 50

<170> PatentIn version 3.5

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<223> Description of Unknown:
FokI cleavage half domain sequence

<400> 1

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Lys Leu Lys Tyr Val Pro His Glu Tyr Ile Glu Leu Ile Glu Ile Ala
            20                  25                  30

Arg Asn Ser Thr Gln Asp Arg Ile Leu Glu Met Lys Val Met Glu Phe
        35                  40                  45

Phe Met Lys Val Tyr Gly Tyr Arg Gly Lys His Leu Gly Gly Ser Arg
    50                  55                  60

Lys Pro Asp Gly Ala Ile Tyr Thr Val Gly Ser Pro Ile Asp Tyr Gly
65                  70                  75                  80
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Val Ile Val Asp Thr Lys Ala Tyr Ser Gly Gly Tyr Asn Leu Pro Ile
85 90 95

Gly Gln Ala Asp Glu Met Gln Arg Tyr Val Glu Glu Asn Gln Thr Arg
100 105 110

Asn Lys His Ile Asn Pro Asn Glu Trp Trp Lys Val Tyr Pro Ser Ser
115 120 125

Val Thr Glu Phe Lys Phe Leu Phe Val Ser Gly His Phe Lys Gly Asn
130 135 140

Tyr Lys Ala Gln Leu Thr Arg Leu Asn His Ile Thr Asn Cys Asn Gly
145 150 155 160

Ala Val Leu Ser Val Glu Glu Leu Leu Ile Gly Gly Glu Met Ile Lys
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Ala Gly Thr Leu Thr Leu Glu Glu Val Arg Arg Lys Phe Asn Asn Gly
180 185 190

Glu Ile Asn Phe
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<400> 3
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<210> 14
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peptide

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<210> 30
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peptide

<400> 30
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<210> 31
<211> 7
<212> PRT
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<220>
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peptide

<400> 31
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<210> 32
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<220>
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peptide

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<210> 33
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<210> 34
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peptide

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<210> 35
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peptide

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<210> 37
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<220>
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<210> 39
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<220>
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<210> 40
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<213> Unknown

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<210> 42
<211> 28
<212> DNA
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<220>
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CISH target site sequence

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<210> 43
<211> 28
<212> DNA
<213> Unknown

<220>
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CISH target site sequence

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<210> 44
<211> 28
<212> DNA
<213> Unknown

<220>
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<210> 45
<211> 28
<212> DNA
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<220>
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<210> 46
<211> 28
<212> DNA
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CISH target site sequence

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<210> 47
<211> 28
<212> DNA
<213> Unknown

<220>
<223> Description of Unknown:
CISH target site sequence

<400> 47
ccggaaaggc caggatgcgt ggcctgga 28

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<212> DNA
<213> Unknown

<220>
<223> Description of Unknown:
CISH sequence

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cagcccttgc ctgctggggc cttcctcgag gaggtggcag agggtacccc agcccagaca 180

gagagtgagc caaaggtgct ggacccagag gaggatctgc tgtgcatagc caagaccttc 240

tcctaccttc gggaatctgg tgagtctgag gggggaggca ggccttttcc tgagtagtct 300

ggtgggagaa gcttagttct gcctttgagt ctgatttggg aggcatagcc cggcccagga 360

gagtctgatg ggagaggcac agccctctct agggaagcgt gaagctggag gcactgcctt 420

gacttgcact ggattggtta ccccagctaa tttccactat gtcccttggc cccctctgca 480

cttgcctagg ctggtattgg ggttccatta cggccagcga ggcccgacaa cacctgcaga 540

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tgtcagtgaa aaccactcgt ggccccacca atgtacgcat tgagtatgcc gactccagct 660

tccgtctgga ctccaactgc ttgtccaggc cacgcatcct ggcctttccg gatgtggtca 720

gccttgtgca gcactatgtg gcctcctgca ctgctgatac ccgaagcgac agccccgatc 780

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ccccagctgg acttgggccc 1100

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<213> Unknown

<220>
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'LAGLIDADG' family peptide motif sequence

<400> 49
Leu Ala Gly Leu Ile Asp Ala Asp Gly

1 5

<210> 50
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<212> DNA
<213> Unknown

<220>
<223> Description of Unknown:
CISH sequence

<400> 50
ggaaaggcca ggatgcgtgg cctggacaag ca 32