

US 20190256858A1

(19) United States (12) Patent Application Publication (10) Pub. No.: US 2019/0256858 A1 Zajac-Kaye et al.

Aug. 22, 2019 (43) **Pub. Date:**

(54) AAV DELIVERY OF SHRNA FOR TREATMENT OF PANCREATIC CANCER

- (71) Applicant: University of Florida Research Foundation, Incorporated, Cainesville, FL (US)
- (72) Inventors: Maria Zajac-Kaye, Gainesville, FL (US); Kyungah Maeng, Cambridge, MA (US)
- (73) Assignee: University of Florida Research Foundation, Incorporated, Cainesville, FL (US)
- (21) Appl. No.: 16/347,461
- (22) PCT Filed: Nov. 3, 2017
- (86) PCT No.: PCT/US2017/059843 § 371 (c)(1),
 - (2) Date: May 3, 2019

Related U.S. Application Data

(60) Provisional application No. 62/416,893, filed on Nov. 3, 2016.

Publication Classification

(51)	Int. Cl.	
, í	C12N 15/113	(2006.01)
	C12N 7/00	(2006.01)
	A61P 35/00	(2006.01)

(52) U.S. Cl. C12N 15/1137 (2013.01); C12N 7/00 CPC (2013.01); A61P 35/00 (2018.01); C12N 2330/51 (2013.01); C12N 2320/32 (2013.01); C12N 2310/335 (2013.01); C12N 2310/14 (2013.01); C12N 2310/122 (2013.01); C12N 2750/14142 (2013.01); C12N 2750/14143 (2013.01); C12N 2310/11 (2013.01); C12N 2310/531 (2013.01)

(57)ABSTRACT

Aspects of the disclosure relate to methods and compositions for treating pancreatic cancer (e.g., islet cell tumors). In some aspects, adeno-associated virus (AAV) may be used to deliver an interfering RNA that targets thymidylate synthase (TS).

Specification includes a Sequence Listing.









Figure 2D





Figure 3C



Figure 3D









MEF-Men1-/-









scAAV-mIP-GFP-shRNA constructs:



Figure 7D



Figure 7E



Figure 7F

AAV DELIVERY OF SHRNA FOR TREATMENT OF PANCREATIC CANCER

RELATED APPLICATIONS

[0001] This application claims priority under 35 U.S.C. § 119(e) to U.S. provisional application No. 62/416,893, filed Nov. 3, 2016, entitled "AAV DELIVERY OF SHRNA FOR TREATMENT OF PANCREATIC CANCER," the content of which is incorporated herein by reference in its entirety.

BACKGROUND

[0002] The incidence of pancreatic neuroendocrine tumors (PanNET) is increasing and few therapeutic options are available. There remains a need for additional therapeutic treatments for PanNET and other pancreatic cancers.

SUMMARY

[0003] Aspects of the application relate to methods and compositions for treating pancreatic cancer (e.g., islet cell tumor).

[0004] Aspects of the disclosure also include a recombinant adeno-associated viral (rAAV) particle comprising a nucleic acid vector that comprises (a) a heterologous nucleic acid region comprising a sequence that encodes an interfering RNA that comprises a region of complementarity with a thymidylate synthase mRNA and (b) inverted terminal repeat (ITR) sequences flanking the heterologous nucleic acid region. In some embodiments, the interfering RNA is a small hairpin RNA (shRNA) or a microRNA. In some embodiments, the region of complementarity is 100% complementary to at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides of any of SEQ ID NOs .: 6-11. In some embodiments, the interfering RNA comprises at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides of SEQ ID NOs: 1, 2, 3 and/or 4, optionally where one or more Ts or Us in the sequence(s) may be substituted by one or more Us or Ts, respectively. [0005] In some embodiments, the particle is an AAV8 particle or a modified AAV8 particle. In some embodiments, the modified AAV8 particle comprises an AAV8 capsid protein comprising a Y275F, Y447F, or Y733F mutation, or any combination thereof. In some embodiments, the AAV8 capsid protein comprises both a Y447F and Y733F mutation. In some embodiments, the interfering RNA is under expression control of a promoter sequence as described herein. In some embodiments, the interfering RNA is under expression control of an RNA polymerase III promoter. In some embodiments, the interfering RNA is under expression control of a pancreas-specific promoter, optionally wherein the pancreas-specific promoter is an islet-specific promoter, further optionally wherein the islet-specific promoter is an insulin promoter (e.g., a human insulin promoter).

[0006] In some embodiments, the disclosure includes a composition comprising any recombinant adeno-associated viral (rAAV) particle described herein, for example according to any one of the above-mentioned embodiments or another embodiment described herein. In some embodi-

ments, the composition comprises one or more physiologically acceptable carriers and/or one or more physiologically acceptable adjuvants.

[0007] In some embodiments, the disclosure includes a method of decreasing thymidylate synthase expression in a cell (e.g., a human cell), the method comprising administering to the cell of an rAAV particle described herein or a composition described herein. In some embodiments, the cell is a cell of the pancreas.

[0008] In some embodiments, the disclosure includes a method of decreasing thymidylate synthase expression in a subject (e.g., a human subject), the method comprising administering to the subject an rAAV particle as described herein or a composition as described herein. In some embodiments, the administration results in delivery of the rAAV particle to the pancreas (e.g., to tumor cells in the pancreas, such as islet tumor cells). In some embodiments, the subject is a subject having one or more symptoms of a pancreatic condition, disease, or disorder (e.g., pancreatic cancer) or a subject that has been diagnosed with a pancreatic condition, disease, or disorder (e.g., pancreatic cancer). [0009] In some embodiments, the disclosure includes a method of treating pancreatic cancer in a subject, the method comprising administering to the subject an rAAV particle as described herein or a composition as described herein. In some embodiments, the pancreatic cancer is islet cell carcinoma. In some embodiments, the subject is a human subject. In some embodiments, the subject is a subject having one or more symptoms of pancreatic cancer or a subject that has been diagnosed with pancreatic cancer.

[0010] In some embodiments, aspects of the application include a synthetic ribonucleic acid (RNA) molecule comprising a sense strand of sequence AACCUUUGGGA-GAUGCACAUAUUUGUGAAGCCACA-

GAUGAAAUAUGUGCAUCUC CCAAAGUUUUUUGUU (SEQ ID NO: 1) and an antisense strand of sequence AACAAAAACUUUGGGAGAUGCACAUAUUU-

CAUCUGUGGCUUCACAAAUAUGUGC AUCUC-CCAAAGGUU (SEQ ID NO: 2). In some embodiments, the RNA is a small hairpin RNA (shRNA).

[0011] Other aspects include a shRNA having a targeted sequence that comprises RNA of sequence AAAUAUGUG-CAUCUCCCAAAG (SEQ ID NO: 3) or RNA of sequence CUUUGGGAGAUGCACAUAUUU (SEQ ID NO: 4).

[0012] In some embodiments, the synthetic RNA further comprises an unpaired overhang sequence at the 5' and/or 3' end. In some embodiments, the unpaired overhang sequence comprises a sequence of repeating bases. In some embodiments, the sequence of repeating bases comprises repeating uracil (U) bases. In some embodiments, the unpaired overhang sequence is UU.

[0013] In some embodiments of any of the sequences herein, one or more Ts or Us in the sequence(s) may be substituted by one or more Us or Ts, respectively.

[0014] In some embodiments, a composition comprises a synthetic RNA described herein. In some embodiments, the composition further comprises one or more physiologically acceptable carriers and/or one or more physiologically acceptable adjuvants.

[0015] In some embodiments, the shRNA is encoded by a vector (e.g., a DNA vector). In some embodiments, the shRNA is selected from SEQ ID NOs: 1-4. In some embodiments, the vector is an expression plasmid. In some embodiments, the vector is a viral vector. In some embodiments, the

2

viral vector is a recombinant adeno-associated viral (rAAV) vector. In some embodiments, a rAAV nucleic acid encodes an shRNA and is packaged in an rAAV particle.

[0016] In some embodiments, an rAAV particle or vector described herein may encode two or more shRNAs as described herein.

[0017] In some embodiments, a method of decreasing thymidylate synthase expression in a subject in provided. In some embodiments, the method comprises administering to the subject the composition or vector described herein. In some embodiments, the invention includes a method of treating pancreatic cancer in a subject, the method comprising administering to the subject a composition or a vector described herein. In some embodiments, the composition or vector is delivered using an rAAV. In some embodiments, the pancreatic cancer is islet cell carcinoma. In some embodiments, the rAAV is AAV8. In some embodiments, the interfering RNA is under expression control of promoter sequences. In some embodiments, the interfering RNA is shRNA, and wherein the shRNA is under expression control of an RNA polymerase III promoter. In some embodiments, the interfering RNA is shRNA, and wherein the shRNA is under expression control of a pancreas-specific promoter.

[0018] In some embodiments, the subject is a mammal. In some embodiments, the mammal is a rodent or a dog. In some embodiments, the mammal is a human.

[0019] These and other aspects are described in the following drawings, examples, and claims.

BRIEF DESCRIPTION OF DRAWINGS

[0020] The drawings and following brief descriptions provide non-limiting examples of aspects of the compositions and methods described herein.

[0021] FIG. **1** shows the conversion of deoxyuridine monophosphate (dUMP) to deoxythymidine monophosphate (dTMP) with thymidylate synthetase.

[0022] FIGS. **2**A-2D show generation of novel hTS/ Men1^{-/-} genetically engineered mouse model. (FIG. **2**A) Schematic representation of breeding strategy. Arrows indicate primer location for genotyping analysis. (FIG. **2**B) Genotyping results of hTS/Men1^{-/-} and Men1^{-/-} mice. (FIG. **2**C) Immunoblot anlaysis for TS expression in the pancreases of hTS/Men1^{-/-} and Men1^{-/-} mice. (FIG. **2**D) PanNET progression of hTS/Men1^{-/-} GEMMs. Representative H&E images and pathological photographs of pancreatic islet lesions in hTS/fMen1^{-/-} mice (Scale bar, 100 µm).

[0023] FIGS. 3A-3E show hTS overexpression significantly reduced survival of Men1-null mice. (FIG. 3A) Overall survival analysis of total Men1^{-/-} vs. hTS/Men1^{-/-} mice. (FIG. 3B) Survival analysis of Men1^{-/-} vs. hTS/ Men1^{-/-} mice that developed only PanNETs. (FIG. 3C) mice. (FIG. 3D) Overall survival analysis of total Men1^{+/-} vs. hTS/Men1^{+/-}mice. (FIG. 3E) Separate survival analysis of male and female.

[0024] FIGS. **4**A-**4**B show TS overexpression induces PanNET progression. (FIG. **4**A) Pancreatic islet lesions in Men1^{-/-} vs. hTS/Men1^{-/-} mice. Animals were euthanized at 5, 6.5 and 8 months and pancreas were isolated for histopathologic analysis (n=16 per group at each time point). (FIG. **4**B) Pancreatic islet carcinoma incidence. The percentage of mice with islet tumor lesions is shown (*p<0.01, **p<0.001). **[0025]** FIGS. **5A-5**C show TS overexpression induces somatic mutations. (FIG. **5**A) Overview of the λ Select-CII Mutation Detection System. (FIG. **5**B) Mutation frequencies in Men1^{-/-}/BB and hTS/fMen1^{-/-}/BB mice (n=3 per group at each time point). (FIG. **5**C) The type of mutations in the pancreas and tumors of Men1^{-/-}/BB and hTS/Men1^{-/-}/BB mice at 5 and 10 months of age.

[0026] FIGS. **6**A-**6**B show overexpression induces DNA double strand breaks. (FIG. **6**A) Immunofluorescence image of γ H2AX foci in MEF-Men1^{*WT*}-vector, MEF-Men1^{*WT*}-hTS cells, MEF Men1^{-/-}-vector and MEF-Men1^{-/-}-hTS cells. Representative nuclei are shown (Scale bars, 10 µm). (FIG. **6**B) Quantification of γ H2AX foci (*p<0.01).

[0027] FIGS. 7A-7F show AAV-TS shRNA inhibits Pan-NET progression. (FIG. 7A) Vector map of scAAV-mIP-GFP-NSshRNA construct. (FIG. 7B) Schematics of scAAVmIP-GFP-NSshRNA (AAV-shNS) or scAAV-mIP-GFP-TSshRNA (AAV-shTS) treatment in hTS/Men1^{-/-} mice. (FIG. 7C) Survival analysis of pancreas tissues from hTS/ Men1 mice after TS shRNA injection (n=24 per group). (FIG. 7D) TS mRNA expression levels in tumors. (FIG. 7E) TS protein expression levels in tumors. (FIG. 7F) The percentage of islet tumor lesion (n=9 per group).

DETAILED DESCRIPTION

[0028] Aspects of the application provide methods and compositions that are useful for treating pancreatic cancer (e.g., islet cell tumors). In some embodiments, a method of treating pancreatic cancer (e.g., PanNET) in a subject includes administering to the subject a composition or vector described herein. In some embodiments, the composition or vector for treating pancreatic cancer is a TS inhibitor. In some embodiments, the TS inhibitor is a TS shRNA (e.g., SEQ ID NOs: 1-4). In some embodiments, the TS shRNA can decrease cancer progression and increase survival of the subject.

[0029] The pancreas secretes enzymes that help digestion and hormones that assist with the regulation of the metabolism of sugars. The exocrine cells and endocrine cells of the pancreas can form different types of tumors in a subject. Pancreatic cancers can include exocrine pancreatic cancers (e.g., pancreatic adenocarcinoma) and pancreatic neuroendocrine tumors (e.g., PanNETs) or islet cell tumors (e.g., insulinomas). Pancreatic NETs may be functional or nonfunctional. Functional tumors make extra amounts of hormones, such as gastrin, insulin, and glucagon, that cause disease symptoms. Nonfunctional tumors do not make extra amounts of hormones. Signs and symptoms are caused by the tumor as it spreads and grows. Most nonfunctional tumors are malignant. Multiple endocrine neoplasia type 1 (MEN1) syndrome is a risk factor for PanNETs.

[0030] PanNETs can be detected or diagnosed via lab tests and/or imaging tests. Physical exam, blood chemistry studies, chromogranin A tests, CAT scans, MRI, somatostatin receptor scintigraphy, endoscopic ultrasound, endoscopic retrograde cholangiopancreatography (ERCP), angiogram, laparotomy, intraoperative ultrasound, biopsy, bone scans, specific PanNET lab tests, and combinations of these methods can be used for detection/diagnosis.

[0031] Thymidylate synthetase (EC 2.1.1.45) catalyzes the conversion of deoxyuridine monophosphate (dUMP) to deoxythymidine monophosphate (dTMP). Thymidine is one

of the nucleotides in DNA. With inhibition of TS, an imbalance of deoxynucleotides and increased levels of dUTP arise. TS plays a crucial role in the early stages of DNA biosynthesis; DNA damage or deletion occur on a daily basis as a result of both endogenous and environmental factors. Therefore, synthesis and insertion of undamaged DNA is vital for normal body functions and avoidance of cancerous activity. In addition, inhibition in synthesis of important nucleotides necessary for cell growth is important. TS has become an important target for cancer treatment.

[0032] Human thymidylate synthetase (TS) is encoded by the gene TYMS (GenBank Ref. No. AAH83512). Nonlimiting examples of cDNA sequences (SEQ ID NOs: 6-11) of TS are provided below, with the corresponding mRNA sequences having the Ts substituted with Us.

Homo No .	<i>sapiens</i> thym BC002567);	idylate syn	thetase, mR	NA (cDNA cl	one MGC:159	0; GenBank R	ef.	_
1	cgcgccactt	ggcctgcctc	cgtcccgccg	cgccacttcg	cctgcctccg	SEQ tecccegece	ID NO:	6
61	gccgcgccat	gcctgtggcc	ggctcggagc	tgccgcgccg	gcccttgccc	cccgccgcac		
121	aggagcggga	cgccgagccg	cgtccgccgc	acgggggagct	gcagtacctg	gggcagatcc		
181	aacacatcct	ccgctgcggc	gtcaggaagg	acgaccgcac	gggcaccggc	accctgtcgg		
241	tattcggcat	gcaggcgcgc	tacagcctga	gagatgaatt	ccctctgctg	acaaccaaac		
301	gtgtgttctg	gaagggtgtt	ttggaggagt	tgctgtggtt	tatcaaggga	tccacaaatg		
361	ctaaagagct	gtcttccaag	ggagtgaaaa	tctgggatgc	caatggatcc	cgagactttt		
421	tggacagcct	gggattctcc	accagagaag	aaggggactt	gggcccagtt	tatggcttcc		
481	agtggaggca	ttttgggggca	gaatacagag	atatggaatc	agattattca	ggacagggag		
541	ttgaccaact	gcaaagagtg	attgacacca	tcaaaaccaa	ccctgacgac	agaagaatca		
601	tcatgtgcgc	ttggaatcca	agagatette	ctctgatggc	gctgcctcca	tgccatgccc		
661	tctgccagtt	ctatgtggtg	aacagtgagc	tgtcctgcca	gctgtaccag	agatcgggag		
721	acatgggcct	cggtgtgcct	ttcaacatcg	ccagctacgc	cctgctcacg	tacatgattg		
781	cgcacatcac	gggcctgaag	ccaggtgact	ttatacacac	tttgggagat	gcacatattt		
841	acctgaatca	catcgagcca	ctgaaaattc	agetteageg	agaacccaga	cctttcccaa		
901	agctcaggat	tcttcgaaaa	gttgagaaaa	ttgatgactt	caaagctgaa	gactttcaga		
961	ttgaagggta	caatccgcat	ccaactatta	aaatggaaat	ggctgtttag	ggtgctttca		
1021	aaggagcttg	aaggatattg	tcagtcttta	ggggttgggc	tggatgccga	ggtaaaagtt		
1081	ctttttgctc	taaaagaaga	aggaactagg	tcaaaaatct	gtccgtgacc	tatcagttat		
1141	taatttttaa	ggatgttgcc	actggcaaat	gtaactgtgc	cagttettte	cataataaaa		
1201	ggctttgagt	taactcactg	agggtatctg	acaatgctga	ggttatgaac	aaagtgagga		
1261	gaatgaaatg	tatgtgctct	tagcaaaaac	atgtatgtgc	atttcaatcc	cacgtactta		
1321	taaagaaggt	tggtgaattt	cacaagctat	ttttggaata	tttttagaat	attttaagaa		
1381	tttcacaagc	tattccctca	aatctgaggg	agctgagtaa	caccatcgat	catgatgtag		
1441	agtgtggtta	tgaactttat	agttgtttta	tatgttgcta	taataaagaa	gtgttctgca		
1501	ttcgtaaaaa	aaaaaaaaaa	aaaa					
Homo comp 1	<i>sapiens</i> thym lete cds (Gen ggggggcgcgc	idylate syn bank Ref. N ggaaggggtc	thetase, mR o. BC013919 ctgccaccgc	NA (cDNA cl .1, SEQ ID : gccacttggc	one MGC:228 NO: 7) ctgcctccgt	84 IMAGE:404 cccgccgcgc	8625),	
61	cacttcgcct	gcctccgtcc	cccgcccgcc	gcgccatgcc	tgtggccggc	teggagetge		
121	cacaccaacc	cttqcccccc	qccqcacaqq	aqcqqqacqc	cqaqccqcqt	ccqccqcacq		
1.81	aggaggtgga	atacctocco	cadat ccaac	acatecteed	ctacaacata	aggaaggagg		
101	gggageegea	geacergygy	atata	tagaat	aaaaa	aggaaggacg		
∠41	accycacygg	caccygeade	cigloggtat	loggoatgda	ggegegetad	ayeergagag		

4

			-con	ntinued		
301	atgaattccc	tctgctgaca	accaaacgtg	tgttctggaa	gggtgttttg	gaggagttgc
361	tgtggtttat	caagggatcc	acaaatgcta	aagagctgtc	ttccaaggga	gtgaaaatct
421	gggatgccaa	tggatcccga	gactttttgg	acagcctggg	attetecace	agagaagaag
481	gggacttggg	cccagtttat	ggcttccagt	ggaggcattt	tggggcagaa	tacagagata
541	tggaatcaga	ttattcagga	cagggagttg	accaactgca	aagagtgatt	gacaccatca
601	aaaccaaccc	tgacgacaga	agaatcatca	tgtgcgcttg	gaatccaaga	gatetteete
661	tgatggcgct	gcctccatgc	catgccctct	gccagttcta	tgtggtgaac	agtgagctgt
721	cctgccagct	gtaccagaga	tcgggagaca	tgggcctcgg	tgtgcctttc	aacatcgcca
781	gctacgccct	gctcacgtac	atgattgcgc	acatcacggg	cctgaagcca	ggtgacttta
841	tacacacttt	gggagatgca	catatttacc	tgaatcacat	cgagccactg	aaaattcagc
901	ttcagcgaga	acccagacct	ttcccaaagc	tcaggattct	tcgaaaagtt	gagaaaattg
961	atgacttcaa	agctgaagac	tttcagattg	aagggtacaa	tccgcatcca	actattaaaa
1021	tggaaatggc	tgtttagggt	gctttcaaag	gagctcgaag	gatattgtca	gtctttaggg
1081	gttgggctgg	atgccgaggt	aaaagttctt	tttgctctaa	aagaaaaagg	aactaggtca
1141	aaaatctgtc	cgtgacctat	cagttattaa	tttttaagga	tgttgccact	ggcaaatgta
1201	actgtgccag	ttctttccat	aataaaaggc	tttgagttaa	ctcactgagg	gtatctgaca
1261	atgctgaggt	tatgaacaaa	gtgaggagaa	tgaaatgtat	gtgctcttag	caaaaacatg
1321	tatgtgcatt	tcaatcccac	gtacttataa	agaaggttgg	tgaatttcac	aagctatttt
1381	tggaatattt	ttagaatatt	ttaagaattt	cacaagctat	tccctcaaat	ctgagggagc
1441	tgagtaacac	catcgatcat	gatgtagagt	gtggttatga	actttaaagt	tatagttgtt
1501	ttatatgttg	ctataataaa	gaagtgttct	gcattcgcca	aaaaaaaaaa	aaaaaaaaa
1561	aaaaaaaaa					
Synthe	tic constru mRNA part	ct <i>Homo sap</i> ial cds (Ge	<i>iens</i> clone nbank Ref	FLH058707.0 No AY89370	1L thymidyl 6 1 SEO ID	ate synthetase
1	atgcctgtgg	ccggctcgga	gctgccgcgc	cggcccttgc	cccccgccgc	acaggagcgg
61	gacgccgagc	cgcgtccgcc	gcacgggggag	ctgcagtacc	tggggcagat	ccaacacatc

121 ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc 181 atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc 241 tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag 301 ctgtcttcca agggagtgaa aatctgggat gccaatggat cccgagactt tttggacagc 361 ctgggattct ccaccagaga agaaggggac ttgggcccag tttatggctt ccagtggagg 421 cattttgggg cagaatacag agatatggaa tcagattatt caggacaggg agttgaccaa 481 ctgcaaagag tgattgacac catcaaaacc aaccctgacg acagaagaat catcatgtgc 541 gettggaate caagagatet teetetgatg gegetgeete catgeeatge eetetgeeag 601 ttctatgtgg tgaacagtga gctgtcctgc cagctgtacc agagatcggg agacatgggc 661 ctcggtgtgc ctttcaacat cgccagctac gccctgctca cgtacatgat tgcgcacatc 721 acgggcctga agccaggtga ctttatacac actttgggag atgcacatat ttacctgaat

cacategage caetgaaaat teagetteag egagaaeeea gaeetteee aaageteagg

attcttcgaa aagttgagaa aattgatgac ttcaaagctg aagactttca gattgaaggg

901 tacaatccgc atccaactat taaaatggaa atggctgttt tg

781

841

301 ctqtcttcca aqqqaqtqaa aatctqqqat qccaatqqat cccqaqactt tttqqacaqc 361 ctgggattet ccaccagaga agaaggggae ttgggeecag tttatggett ccagtggagg 421 cattttgggg cagaatacag agatatggaa tcagattatt caggacaggg agttgaccaa 481 ctqcaaaqaq tqattqacac catcaaaacc aaccctqacq acaqaaqaat catcatqtqc 541 gettggaate caagagatet teetetgatg gegetgeete catgecatge cetetgecag 601 ttctatgtgg tgaacagtga gctgtcctgc cagctgtacc agagatcggg agacatgggc 661 ctcqqtqtqc ctttcaacat cqccaqctac qccctqctca cqtacatqat tqcqcacatc 721 acgggcctga agccaggtga ctttatacac actttgggag atgcacatat ttacctgaat 781 cacategage caetgaaaat teagetteag egagaaceea gaeetteee aaageteagg 841 attettegaa aagttgagaa aattgatgae tteaaagetg aagaetttea gattgaaggg 901 tacaatccgc atccaactat taaaatggaa atggctgttt ag Synthetic construct *Homo sapiens* clone FLH117831.01L thymidylate synthetase (TYMS) mRNA, partial cds (Genbank Ref. No. AY890751.1, SEQ ID NO: 10) atgeetgtgg eeggetegga getgeegege eggeeettge eeeegeege acaggagegg 1 61 gacgccgagc cgcgtccgcc gcacgggggag ctgcagtacc tggggcagat ccaacacatc 121 ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc 181 atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc 241 tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag 301 ctgtcttcca agggagtgaa aatctgggat gccaatggat cccgagactt tttggacagc 361 ctgggattct ccaccagaga agaaggggac ttgggcccag tttatggctt ccagtggagg 421 cattttqqqq caqaatacaq aqatatqqaa tcaqattatt caqqacaqqq aqttqaccaa 481 ctgcaaagag tgattgacac catcaaaacc aaccctgacg acagaagaat catcatgtgc 541 gettggaate caagagatet teetetgatg gegetgeete catgecatge cetetgecag 601 ttctatgtgg tgaacagtga gctgtcctgc cagctgtacc agagatcggg agacatgggc 661 ctcggtgtgc ctttcaacat cgccagctac gccctgctca cgtacatgat tgcgcacatc 721 acgggcctga agccaggtga ctttatacac actttgggag atgcacatat ttacctgaat 781 cacategage caetgaaaat teagetteag egagaaceea gaeetteee aaageteagg 841 attettegaa aagttgagaa aattgatgae tteaaagetg aagaetttea gattgaaggg 901 tacaatccgc atccaactat taaaatggaa atggctgttt tg Synthetic construct Homo sapiens clone FLH117835.01X thymidylate synthetase (TYMS) mRNA, complete cds (Genbank Ref. No. AY888178.1, SEQ ID NO: 11) 1 atgeetgtgg eeggetegga getgeegege eggeeettge eeeegeege acaggagegg

gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc

ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc

atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc

-COntinued Synthetic construct Homo sapiens clone FLH058712.01X thymidylate synthetase (TYMS) mRNA, complete cds (Genbank Ref. No. AY890751.1, SEQ ID NO: 9) 1 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg

gacgccgagc cgcgtccgcc gcacgggggag ctgcagtacc tgggggcagat ccaacacatc

ctccqctqcq qcqtcaqqaa qqacqaccqc acqqqcaccq qcaccctqtc qqtattcqqc

atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc

tqqaaqqqtq ttttqqaqqa qttqctqtqq tttatcaaqq qatccacaaa tqctaaaqaq

61

121

181

241

61

121

181

5

4

			-con	ltinued		
241	tggaagggtg	ttttggagga	gttgctgtgg	tttatcaagg	gatccacaaa	tgctaaagag
301	ctgtcttcca	agggagtgaa	aatctgggat	gccaatggat	cccgagactt	tttggacagc
361	ctgggattct	ccaccagaga	agaagggggac	ttgggcccag	tttatggctt	ccagtggagg
421	cattttgggg	cagaatacag	agatatggaa	tcagattatt	caggacaggg	agttgaccaa
481	ctgcaaagag	tgattgacac	catcaaaacc	aaccctgacg	acagaagaat	catcatgtgc
541	gcttggaatc	caagagatct	tcctctgatg	gcgctgcctc	catgccatgc	cctctgccag
601	ttctatgtgg	tgaacagtga	gctgtcctgc	cagctgtacc	agagatcggg	agacatgggc
661	ctcggtgtgc	ctttcaacat	cgccagctac	gccctgctca	cgtacatgat	tgcgcacatc
721	acgggcctga	agccaggtga	ctttatacac	actttgggag	atgcacatat	ttacctgaat
781	cacatcgagc	cactgaaaat	tcagcttcag	cgagaaccca	gacctttccc	aaagctcagg
841	attcttcgaa	aagttgagaa	aattgatgac	ttcaaagctg	aagactttca	gattgaaggg
901	tacaatccgc	atccaactat	taaaatggaa	atggctgttt	ag	

[0033] In some aspects, the disclosure provides interfering RNAs (e.g., shRNAs or microRNAs) that contain a region of complementarity with a region of a TS mRNA (e.g., one of SEQ ID Nos: 6-11). Interfering RNAs, such as shRNAs or microRNAs, are further described herein. An interfering RNA may comprise a region of complementarity that is at least 80% complementary to (optionally one of at least 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% complementary to) the consecutive nucleotides of a TS mRNA (e.g., one of SEQ ID Nos: 6-11). In some embodiments, the region of complementarity is at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides long. In some embodiments, the region of the TS transcript is at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides long. In some embodiments, the interfering RNA is expressed by a nucleic acid as described herein (e.g. contained with an rAAV particle or vector). Methods for producing interfering RNAs are known in the art (see, e.g., Rao et al. Tissue-specific and cell type-specific RNA interference in vivo, Nature Protocols 1, 1494-1501 (2006); Moore et al. Short Hairpin RNA (shRNA): Design, Delivery, and Assessment of Gene Knockdown. Methods Mol Biol. 2010; 629: 141-158 and Wahid t al. MicroRNAs: Synthesis, mechanism, function, and recent clinical trials. Biochimica et Biophysica Acta. Volume 1803, Issue 11, November 2010, Pages 1231-1243) and are commercially available (see, e.g., services available from Dharmacon, Sigma-Aldrich, Origene, Thermofisher, System Biosciences, etc.).

[0034] In some embodiments, the interfering RNA is an shRNA. In some embodiments, an shRNA comprises an antisense sequence that is complementary to a target RNA and a sense sequence that is the reverse complement of the antisense sequence, typically separated by a spacer or loop sequence. A spacer or loop can be of a sufficient length to permit the antisense and sense sequences to anneal and form a double-stranded structure (or stem). The spacer can then be

cleaved away to form a double-stranded RNA (and, optionally, subsequent processing steps that may result in addition or removal of one, two, three, four, or more nucleotides from the 3' end and/or the 5' end of either or both strands). In some embodiments, the stem of the shRNA comprises 19-29 basepairs and the loop comprises 4-8 nucleotides, optionally with a dinucleotide overhang at the 3' end of the shRNA. In some embodiments, the stem of the shRNA comprises a region of complementarity with a region of at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides of a TS transcript (e.g., one of SEQ ID Nos: 6-11) as described herein and/or comprises at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides of SEQ ID NOs: 1, 2, 3 and/or 4.

[0035] In some embodiments, the interfering RNA is a microRNA (miRNA). MiRNAs are small non-coding RNAs, belonging to a class of regulatory molecules that control gene expression by binding to complementary sites on a target RNA transcript. Typically, miRNAs are generated from large RNA precursors (termed pri-miRNAs) that are processed in the nucleus into approximately 70 nucleotide pre-miRNAs, which fold into imperfect stem-loop structures. These pre-miRNAs typically undergo an additional processing step within the cytoplasm where mature miRNAs of 18-25 nucleotides in length are excised from one side of the pre-miRNA hairpin by an RNase III enzyme, Dicer. As used herein, miRNAs including pri-miRNA, pre-miRNA, mature miRNA or fragments of variants thereof that retain the biological activity of mature miRNA. In one embodiment, the size range of the miRNA can be from 21 nucleotides to 170 nucleotides, although miRNAs of up to 2000 nucleotides can be utilized. In some embodiments, the miRNA comprises a region of complementarity with a region of at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides of a TS transcript (e.g., one of SEQ ID Nos: 6-11) as described herein and/or comprises at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides of SEQ ID NOs: 1, 2, 3 and/or 4.

[0036] In some embodiments, the interfering RNA is under expression control of a promoter sequence as described herein. In some embodiments, the interfering RNA is under expression control of an RNA polymerase III promoter. In some embodiments, the interfering RNA is under expression control of a pancreas-specific promoter, optionally wherein the pancreas-specific promoter is an islet-specific promoter, further optionally wherein the isletspecific promoter is an insulin promoter (e.g., a human insulin promoter). Promoters are described in more detail herein.

[0037] Aspects of the disclosure also include a recombinant adeno-associated viral (rAAV) particle comprising a nucleic acid vector that comprises (a) a heterologous nucleic acid region comprising a sequence that encodes an interfering RNA that comprises a region of complementarity with a thymidylate synthase mRNA and (b) inverted terminal repeat (ITR) sequences flanking the heterologous nucleic acid region. In some embodiments, the interfering RNA is a small hairpin RNA (shRNA) or a microRNA. In some embodiments, the region of complementarity is complementary to at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides of any of SEQ ID NOs.: 6-11. In some embodiments, the interfering RNA comprises at least 8 (e.g., at least 8, at least 9, at least 10, at least 11, at least 12, at least 13, at least 14, at least 15, at least 16, at least 17, at least 18, at least 19, at least 20, or at least 21, e.g., 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, or 21) contiguous nucleotides of SEQ ID NOs: 1, 2, 3 and/or 4.

[0038] In some embodiments, the particle is an AAV8 particle or a modified AAV8 particle. AAV8 particles and capsid proteins (including modified capsid proteins) are further described herein. In some embodiments, the modified AAV8 particle comprises an AAV8 capsid protein comprising at least one phenylalanine-to-tyrosine mutation, e.g., a Y275F, Y447F, or Y733F mutation, or any combination thereof (amino acid positions relative to the AAV8 VP1 protein, SEQ ID NO: 5). In some embodiments, the AAV8 capsid protein comprises both a Y447F and Y733F mutation (amino acid positions relative to the AAV8 VP1 protein, SEQ ID NO: 5). In some embodiments, the disclosure includes a method of decreasing thymidylate synthase expression in a cell, the method comprising administering to the cell of an rAAV particle described herein or a composition described herein. In some embodiments, the cell is a cell of the pancreas (e.g., a human pancreas cell).

[0039] In some embodiments, the disclosure includes a method of decreasing thymidylate synthase expression in a subject (e.g., a human subject), the method comprising administering to the subject an rAAV particle described herein or a composition described herein. In some embodiments, the administration results in delivery of the rAAV particle to the pancreas.

[0040] In some embodiments, the disclosure includes a method of treating pancreatic cancer in a subject, the method comprising administering to the subject an rAAV particle described herein or a composition described herein. In some embodiments, the pancreatic cancer is islet cell carcinoma. In some embodiments, the pancreatic cancer is a pancreatic cancer associated with an increase in TS expression compared to a baseline level of TS expression (e.g., in pancreatic cells that are not cancerous). In some embodiments, the subject is a human subject.

[0041] In some aspects, this application describes a synthetic ribonucleic acid (RNA) molecule comprising a sense strand of sequence AACCUUUGGGAGAUGCA-CAUAUUUGUGAAGCCACAGAUGAAAUAUGUG-

CAUCUC CCAAAGUUUUUGUU (SEQ ID NO: 1) and an antisense strand of sequence AACAAAAACUUUGGGA-GAUGCACAUAUUUCAUCUGUGGCUUCA-

CAAAUAUGUGC AUCUCCCAAAGGUU (SEQ ID NO: 2).

[0042] In some embodiments, the RNA is a small interfering RNA (siRNA). In some embodiments, the RNA is a small hairpin RNA (shRNA). In some embodiments, the shRNA has a targeted sequence that comprises RNA of sequence AAAUAUGUGCAUCUCCCAAAG (SEQ ID NO: 3) or RNA of sequence CUUUGGGAGAUGCA-CAUAUUU (SEQ ID NO: 4). In some embodiments, the RNA is an artificial micro RNA (miRNA). In some embodiments, the synthetic RNA comprises an unpaired overhang sequence at the 5' and/or 3' end. In some embodiments, the unpaired overhang sequence comprises a sequence of repeating bases. In some embodiments, the sequence of repeating bases comprises repeating uracil (U) bases. In some embodiments, the unpaired overhang sequence is UU. [0043] In some embodiments, a composition comprises any synthetic RNA variation described herein. In some embodiments, the composition further comprises one or more physiologically acceptable carriers and/or one or more physiologically acceptable adjuvants.

[0044] In some embodiments, the invention includes a vector encoding a shRNA described herein (e.g., SEQ ID NOs: 1-4). In some embodiments, the vector is an expression plasmid. In some embodiments, the vector comprises an adeno-associated viral (AAV) vector. In some embodiments, the AAV vector is AAV8. The AAV vector can be any AAV vector. In some embodiments, the invention includes a method of decreasing thymidylate synthase expression in a subject (e.g., rodent, dog, or human), the method comprising administering to the subject compositions or vectors described herein. In some embodiments, a recombinant AAV is used that includes a nucleic acid encoding the shRNA packaged in a particle comprising AAV capsid proteins. The rAAV can be of any serotype.

[0045] In some embodiments, a method of treating pancreatic cancer (e.g., PanNET) in a subject includes administering to the subject an AAV particle, a composition or vector described herein. In some embodiments, the method further comprises delivering at least one additional therapeutic agent for treatment of pancreatic cancer. In some embodiments, the at least one additional treatment comprises one or more of surgery, octreotide, doxorubicin, streptozocin, fluorouracil, capecitabine, temozolomide, cisplatin, etoposide, an mTOR inhibitor (e.g., everolimus), sunitinib, and combinations thereof.

[0046] In some embodiments, the interfering RNA is under expression control of a single promoter sequence (e.g., insulin promoter). In some embodiments, the promoter is a pancreas-specific promoter. In some embodiments, the promoter is a islet-specific promoter. In some embodiments, the islet specific promoter is an insulin promoter (e.g., a rat insulin promoter (Chai et al., Gene Therapy (2009) 16, RNA polymerase III promoter or other suitable constitutive or inducible promoter) or as artificial microRNAs (miRNAs) driven by a promoter (e.g., using an RNA polymerase II promoter or other suitable constitutive or inducible promoter). In some embodiments, the pancreas-specific promoter is an islet-specific promoter. In some embodiments, the islet-specific promoter is an insulin promoter.

TABLE 1

		Non-limiting sequences
SEQ NO :	ID Name	RNA sequence
1	TSshRNA64- sense	AACCTTTGGGAGATGCACATATTTGTGAAGCCACAGATGAAATATG TGCATCTCCCAAAGTTTTTGTT
2	TS shRNA-64- antisense	AACAAAAACTTTGGGAGATGCACATATTTCATCTGTGGCTTCACAA ATATGTGCATCTCCCAAAGGTT
3	TS targeted sequence	AAAUAUGUGCAUCUCCCAAAG
4	TS targeted sequence	CUUUGGGAGAUGCACAUAUUU
5	AAV8 Capsid Sequence	Exemplary AAV8 capsid protein1MAADGYLPDWLEDNLSEGIREWWALKPGAPKPKANQQKQDDGRGLVLPGY51KYLGPFNGLDKGEPVNAADAAALEHDKAYDQQLQAGDNPYLRYNHADAEF101QERLQEDTSFGGNLGRAVFQAKKRVLEPLGLVEEGAKTAPGKKRPVEPSP151QRSPDSSTGIGKKGQQPARKRLNFGQTGDSESVPDPQPLGEPPAAPSGVG201PNTMAAGGAPMADNNEGADGVGSSSGNWHCDSTWLGDRVITTSTRTWAL251PTYNNHLYKQISNGTSGGATNDNYFGYSTPMGYFDPNRFHCHFSPRDWQ301RLINNNWGFRHQCLPPFPADVFMIPQYGYLTLNNGSQAVGRSSFYCLEY451YQLPYVLGSAHQGCLPPFPADVFMIPQYGYLTLNNGSQAVGRSSFYCLEY451TQTTGGTANTQTLGFSQGGPNTMANQAKNWLPGPCYRQRVSTTTGQINN501SNFAWTAGTKYHLNGRNSLANFGIAMATHKDEERFFPSNGLLFGQNA551ARDNADYSDVMLTSEEEIKTTNPVATEEYGIVADNLQQQNTAPQIGTNS561QGALPGMWQNRDVYLQGFIWAKIPHTDGNFHPSPLMGGFGLKHPPQLL651IKNTPVPADPPTTFNQSKLNSFITQYSTGQVSVEIEWELQKENSKRMPE701IQYTSNYYKSTSVDFAVNTEGVSEPRPIGTRYLTRNL*

1202-1209; doi:10.1038/gt.2009.114; published online 3 September 2009) or a human insulin promoter (Melloul et al., Proc Natl Acad Sci USA. 1993 May 1; 90(9): 3865-3869). In some embodiments, the interfering RNA is shRNA, and wherein the shRNA is under expression control of an RNA polymerase III promoter. In some embodiments, the interfering RNA is an artificial miRNA, and wherein the artificial miRNA is under expression control of an RNA polymerase II promoter. In some embodiments, the shRNA is under expression control of a constitutive or inducible promoter.

[0047] In some embodiments, a single adeno-associated virus (AAV) vector is used to deliver the RNA agent (e.g., small hairpin RNA or artificial microRNA).

[0048] In some embodiments, small hairpin RNAs, artificial microRNAs (a-miRs) and/or RNA enzymes (ribozymes) can be designed to degrade thymidylate mRNA by targeting sequences that are common in mouse, human, and dog. Such molecules can be useful to test inhibition in cell culture, in mice, and/or in dogs, to develop inhibitors that can work in human patients.

[0049] In some embodiments, one or more of the interfering RNAs can be delivered using an adeno-associated virus (AAV) vector either as short hairpin RNAs (shRNAs) driven by a promoter (e.g., an pancreas-specific promoter, **[0050]** Accordingly, compositions herein can be administered to a subject in need of treatment of pancreatic cancer. In some embodiments, the subject has or is suspected of having one or more of the conditions, diseases, and disorders disclosed herein. In some embodiments, the subject is a human. In some embodiments, the subject is a human. In some embodiments, the subject is a non-human primate. Non-limiting examples of non-human primate subjects include macaques (e.g., cynomolgus or rhesus macaques), marmosets, tamarins, spider monkeys, owl monkeys, vervet monkeys, squirrel monkeys, baboons, gorillas, chimpanzees, and orangutans. Other exemplary subjects include domesticated animals such as dogs and cats; livestock such as horses, cattle, pigs, sheep, goats, and chickens; and other animals such as mice, rats, guinea pigs, and hamsters.

[0051] In some embodiments, the disclosure includes a composition comprising any recombinant adeno-associated viral (rAAV) particle or interfering RNA described herein. In some embodiments, the composition comprises one or more physiologically acceptable carriers and/or one or more physiologically acceptable adjuvants. In some embodiments, the composition comprising any recombinant adeno-associated viral (rAAV) particle or interfering RNA described herein is administered to a subject having one or more symptoms of a pancreatic condition, disease, or disorder (e.g., pancreatic cancer) or is administered after the

subject has been diagnosed with a pancreatic condition, disease, or disorder (e.g., pancreatic cancer).

[0052] In some embodiments, the number of rAAV particles contained in a composition or administered to a cell or a subject may be on the order ranging from 10^6 to 10^{14} particles/mL or 10³ to 10¹⁵ particles/mL, or any values therebetween for either range, such as for example, about $10^6, 10^7, 10^8, 10^9, 10^{10}, 10^{11}, 10^{12}, 10^{13}, \text{ or } 10^{14}, \text{ particles}/$ mL. In one embodiment, rAAV particles of higher than 10¹³ particles/mL are be administered. In some embodiments, the number of rAAV particles administered to a subject may be on the order ranging from 10^6 to 10^{14} vector genomes(vgs)/ mL or 10^3 to 10^{15} vgs/mL, or any values therebetween for either range, such as for example, about 10⁶, 10⁷, 10⁸, 10⁹, 10^{10} , 10^{11} , 10^{12} , 10^{13} , or 10^{14} vgs/mL. In some embodiments, the number of rAAV particles administered to a subject may be on the order ranging from 10^6 to 10^{14} vector genomes(vgs) or 10^3 to 10^{15} vgs, or any values therebetween for either range, such as for example, about 10^3 , 10^4 , 10^5 , 10^6 , 10^7 , 10^8 , 10^9 , 10^{10} , 10^{11} , 10^{12} , 10^{13} , 10^{14} or 10^{15} vgs. In one embodiment, rAAV particles of higher than 10^{13} vgs/mL are administered. The rAAV particles can be administered as a single dose, or divided into two or more administrations as may be required to achieve therapy of the particular disease or disorder being treated. In some embodiments, 0.0001 mL to 10 mLs of the rAAV particles at any of the concentrations (vgs/mL) described herein are delivered to a subject.

[0053] In some embodiments, rAAV viral titers range from $1 \times 10^{10} - 5 \times 10^{13}$ vg/ml. In some embodiments, rAAV viral titers can be 1×10^{10} , 2.5×10^{10} , 5×10^{10} , 1×10^{11} , 2.5×10^{11} , 5×10^{11} , 1×10^{12} , 2.5×10^{12} , 5×10^{12} , 1×10^{13} , 2.5×10^{13} , or 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are less than 1×10^{10} vg/mL. In some embodiments, rAAV viral titers are greater than 1×10^{15} vg/mL. In one embodiment, rAAV particles are greater than 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are state than 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are greater than 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are state than 1×10^{15} vg/mL. In some embodiments, rAAV viral titers are state than 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are state than 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are state than 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are state than 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are state than 5×10^{13} vg/mL. In some embodiments, rAAV viral titers are administered via methods further described herein.

[0054] The rAAV particles can be administered as a single dose, or divided into two or more administrations as may be required to achieve therapy of the particular disease or disorder being treated. In some embodiments, from 1 to 500 microliters of a composition described in this application is administered to a subject. For example, in some embodiments, about 1, about 10, about 50, about 100, about 200, about 300, about 400, or about 500 microliters can be administered to a subject. However, it should be appreciated that smaller or larger volumes could be administered in some embodiments.

[0055] In some embodiments, an rAAV particle, rAAV vector or interfering RNA as described herein is administered to a subject once. In some embodiments, an rAAV particle, rAAV vector or interfering RNA as described herein is administered to a subject more than once. In some embodiments, an rAAV particle, rAAV vector or interfering RNA as described herein is administered to a subject more than once. In some embodiments, an rAAV particle, rAAV vector or interfering RNA as described herein is administered to a subject as a single dose or as multiple doses.

[0056] In some embodiments, the disclosure provides formulations of one or more rAAV-based compositions disclosed herein in pharmaceutically acceptable solutions for administration to a cell or an animal, either alone or in combination with one or more other modalities of therapy, and in particular, for therapy of human cells, tissues, and diseases affecting man. **[0057]** If desired, rAAV particle or nucleic acid vectors may be administered in combination with other agents as well, such as, e.g., proteins or polypeptides or various pharmaceutically-active agents, including one or more systemic or topical administrations of therapeutic polypeptides, biologically active fragments, or variants thereof. In fact, there is virtually no limit to other components that may also be included, given that the additional agents do not cause a significant adverse effect upon contact with the target cells or host tissues. The rAAV particles may thus be delivered along with various other agents as required in the particular instance. Such compositions may be purified from host cells or other biological sources, or alternatively may be chemically synthesized as described herein.

[0058] Formulation of pharmaceutically-acceptable excipients and carrier solutions is well-known to those of skill in the art, as is the development of suitable dosing and treatment regimens for using the particular compositions described herein in a variety of treatment regimens, including e.g., oral, parenteral, intravenous, intranasal, intra-articular, and intramuscular administration and formulation.

[0059] Typically, these formulations may contain at least about 0.1% of the therapeutic agent (e.g., rAAV particle or host cell) or more, although the percentage of the active ingredient(s) may, of course, be varied and may conveniently be between about 1 or 2% and about 70% or 80% or more of the weight or volume of the total formulation. Naturally, the amount of therapeutic agent(s) (e.g., rAAV particle) in each therapeutically-useful composition may be prepared in such a way that a suitable dosage will be obtained in any given unit dose of the compound. Factors such as solubility, bioavailability, biological half-life, route of administration, product shelf life, as well as other pharmacological considerations will be contemplated by one skilled in the art of preparing such pharmaceutical formulations, and as such, a variety of dosages and treatment regimens may be desirable.

[0060] In certain circumstances it will be desirable to deliver an rAAV particle or host cell in suitably formulated pharmaceutical compositions disclosed herein either subcutaneously, intraocularly, intravitreally, parenterally, subcutaneously, intravenously, intracerebro-ventricularly, intramuscularly, intrathecally, orally, intraperitoneally, by oral or nasal inhalation, or by direct injection to one or more cells, tissues, or organs by direct injection.

[0061] The pharmaceutical forms of the rAAV particle or host cell compositions suitable for injectable use include sterile aqueous solutions or dispersions. In some embodiments, the form is sterile and fluid to the extent that easy syringability exists. In some embodiments, the form is stable under the conditions of manufacture and storage and is preserved against the contaminating action of microorganisms, such as bacteria and fungi. The carrier can be a solvent or dispersion medium containing, for example, water, saline, ethanol, polyol (e.g., glycerol, propylene glycol, and liquid polyethylene glycol, and the like), suitable mixtures thereof, and/or vegetable oils. Proper fluidity may be maintained, for example, by the use of a coating, such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants.

[0062] The term "carrier" refers to a diluent, adjuvant, excipient, or vehicle with which the rAAV particle or host cell is administered. Such pharmaceutical carriers can be sterile liquids, such as water and oils, including those of

petroleum oil such as mineral oil, vegetable oil such as peanut oil, soybean oil, and sesame oil, animal oil, or oil of synthetic origin. Saline solutions and aqueous dextrose and glycerol solutions can also be employed as liquid carriers. **[0063]** The compositions of the present disclosure can be administered to the subject being treated by standard routes including, but not limited to, pulmonary, intranasal, oral, inhalation, parenteral such as intravenous, topical, transdermal, intradermal, transmucosal, intraperitoneal, intramuscular, intracapsular, intraorbital, intravitreal, intraacriac, transtracheal, subcutaneous, subcuticular, intraarticular, subcapsular, subarachnoid, intraspinal, epidural and intrasternal injection.

[0064] The compositions of the present disclosure can be delivered to the pancreas through a variety of routes. They may be delivered systemically through an oral route or by subcutaneous, intravenous or intramuscular injection. Alternatively, they may be delivered by means of a catheter or by means of an implant, wherein such an implant is made of a porous, non-porous or gelatinous material, including membranes such as silastic membranes or fibers, biodegradable polymers, or proteinaceous material. They can be administered prior to the onset of the condition, to prevent its occurrence, or immediately after the onset of the pathological condition or during the occurrence of an acute or protracted condition.

[0065] For administration of an injectable aqueous solution, for example, the solution may be suitably buffered, if necessary, and the liquid diluent first rendered isotonic with sufficient saline or glucose. These particular aqueous solutions are especially suitable for intravenous, intramuscular, intravitreal, subcutaneous and intraperitoneal administration. In this connection, a sterile aqueous medium that can be employed will be known to those of skill in the art in light of the present disclosure. For example, one dosage may be dissolved in 1 ml of isotonic NaCl solution and either added to 1000 ml of hypodermoclysis fluid or injected at the proposed site of infusion, (see for example, "Remington's Pharmaceutical Sciences" 15th Edition, pages 1035-1038 and 1570-1580). Some variation in dosage will necessarily occur depending on the condition of the subject being treated. The person responsible for administration will, in any event, determine the appropriate dose for the individual subject. Moreover, for human administration, preparations should meet sterility, pyrogenicity, and the general safety and purity standards as required by, e.g., FDA Office of Biologics standards.

[0066] Sterile injectable solutions are prepared by incorporating the rAAV particles or host cells in the required amount in the appropriate solvent with several of the other ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the various sterilized active ingredients into a sterile vehicle which contains the basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, the preferred methods of preparation are vacuum-drying and freeze-drying techniques which yield a powder of the active ingredient plus any additional desired ingredient from a previously sterile-filtered solution thereof.

[0067] The amount of rAAV particle, nucleic acid vector, or host cell compositions and time of administration of such compositions will be within the purview of the skilled

artisan having benefit of the present teachings. It is likely, however, that the administration of therapeutically-effective amounts of the disclosed compositions may be achieved by a single administration, such as for example, a single injection of sufficient numbers of infectious particles to provide therapeutic benefit to the patient undergoing such treatment. Alternatively, in some circumstances, it may be desirable to provide multiple, or successive administrations of the rAAV particle or host cell compositions, either over a relatively short, or a relatively prolonged period of time, as may be determined by the medical practitioner overseeing the administration of such compositions.

[0068] The composition may include rAAV particles or host cells, either alone, or in combination with one or more additional active ingredients, which may be obtained from natural or recombinant sources or chemically synthesized. In some embodiments, rAAV particles are administered in combination, either in the same composition or administered as part of the same treatment regimen, with a proteasome inhibitor, such as Bortezomib, or hydroxyurea.

[0069] To "treat" a disease as the term is used herein, means to reduce the frequency or severity of at least one sign or symptom of a disease or disorder experienced by a subject, e.g., cancer such as pancreatic cancer. In some embodiments, with respect to cancer, "treat" means causing remission of the cancer, slowing the course of cancer progression, slowing or inhibiting tumor growth, and/or slowing or inhibiting tumor metastasis. In some embodiments, "treat" means slowing progression of symptoms (e.g., symptoms of pancreatic cancer) or reversing the course of the disease (e.g., the course of pancreatic cancer). The compositions described above are typically administered to a subject in an effective amount, that is, an amount capable of producing a desirable result. The desirable result will depend upon the active agent being administered. For example, an effective amount of a rAAV particle may be an amount of the particle that is capable of transferring a heterologous nucleic acid to a host organ, tissue, or cell.

[0070] Pancreatic cancers include symptoms such as pain, gastrointestinal issues, whole body fatigue or loss of appetite, dark urine, weight loss, or yellow skin and eyes. Non-functional pancreatic NET may grow for a long time without causing signs or symptoms. It may grow large or spread to other parts of the body before it causes signs or symptoms including, but not limited to, diarrhea, indigestion, a lump in the abdomen, pain in the abdomen or back, and/or yellowing of the skin and whites of the eyes.

[0071] Functional PanNETs have various signs and symptoms depending on the type of hormone being made (e.g., gastrin, insulin, glucagon, vasoactive intestinal peptide (VIP), and somatostatin). Symptoms can include stomach ulcers that keep coming back, pain in the abdomen, which may spread to the back, the pain may come and go and it may go away after taking an antacid, the flow of stomach contents back into the esophagus (gastroesophageal reflux), diarrhea, low blood sugar, this can cause blurred vision, headache, and feeling lightheaded, tired, weak, shaky, nervous, irritable, sweaty, confused, or hungry, fast heartbeat, skin rash on the face, stomach, or legs, high blood sugar, this can cause headaches, frequent urination, dry skin and mouth, or feeling hungry, thirsty, tired, or weak, blood clots, blood clots in the lung can cause shortness of breath, cough, or pain in the chest, blood clots in the arm or leg can cause pain, swelling, warmth, or redness of the arm or leg, sore tongue or sores at the corners of the mouth, very large amounts of watery diarrhea, dehydration, this can cause feeling thirsty, making less urine, dry skin and mouth, headaches, dizziness, or feeling tired, low potassium level in the blood, this can cause muscle weakness, aching, or cramps, numbness and tingling, frequent urination, fast heartbeat, and feeling confused or thirsty, cramps or pain in the abdomen, weight loss for no known reason, steatorrhea, gallstones, and/or yellowing of the skin and whites of the eyes.

[0072] Toxicity and efficacy of the compositions utilized in methods of the disclosure can be determined by standard pharmaceutical procedures, using either cells in culture or experimental animals to determine the LD50 (the dose lethal to 50% of the population). The dose ratio between toxicity and efficacy the therapeutic index and it can be expressed as the ratio LD50/ED50. Those compositions that exhibit large therapeutic indices are preferred. While those that exhibit toxic side effects may be used, care should be taken to design a delivery system that minimizes the potential damage of such side effects. The dosage of compositions as described herein lies generally within a range that includes an ED50 with little or no toxicity. The dosage may vary within this range depending upon the dosage form employed and the route of administration utilized.

[0073] Aspects of the disclosure relate to recombinant adeno-associated virus (rAAV) particles for delivery of one or more nucleic acid vectors comprising a gene of interest into various tissues, organs, and/or cells. In some embodiments, the rAAV particles comprise an rAAV capsid protein as described herein, e.g., comprising one or more amino acid substitutions. In some embodiments, the gene of interest encodes a polypeptide or protein of interest (e.g., a therapeutic polypeptide or protein). In some embodiments, the gene of interest encodes an RNA of interest (e.g., a therapeutic mRNA, siRNA, shRNA, microRNA, antisense RNA, tRNA, rRNA, or a ribozyme).

[0074] Recombinant AAV (rAAV) particles may comprise at a minimum one or more heterologous nucleic acid regions comprising a sequence encoding a gene of interest (e.g., an RNA of interest such as a siRNA or shRNA) and (b) one or more regions comprising inverted terminal repeat (ITR) sequences (e.g., wild-type ITR sequences or engineered ITR sequences) flanking the one or more heterologous nucleic acid regions. In some embodiments, the nucleic acid vector is between 4 kb and 5 kb in size (e.g., 4.2 to 4.7 kb in size). This nucleic acid vector may be encapsidated by a viral capsid, such as an AAV1, AAV2, AAV3, AAV4, AAV5, or AAV8 capsid, which may comprise a modified capsid protein as described herein. In some embodiments, the nucleic acid vector is circular. In some embodiments, the nucleic acid vector is single-stranded. In some embodiments, the nucleic acid vector is double-stranded (including partially double-stranded). In some embodiments, a double-stranded nucleic acid vector may be, for example, a self-complementary vector that contains a region of the nucleic acid vector that is complementary to another region of the nucleic acid vector, initiating the formation of the double-strandedness of the nucleic acid vector. In some embodiments, a subject is treated for pancreatic cancer with the TS shRNA described herein after diagnosis of pancreatic cancer (e.g., PanNET). [0075] The rAAV particle may be of any AAV serotype, including any derivative or pseudotype (e.g., 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 2/1, 2/5, 2/8, or 2/9). As used herein, the serotype of an rAAV viral vector (e.g., an rAAV particle) refers to the serotype of the capsid proteins of the recombinant virus. In some embodiments, the rAAV has 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more mutations. In some embodiments, the rAAV capsid sequence includes tyrosine to phenylalanine mutations. In some embodiments, the rAAV has 1, 2, 3 or more $Y \rightarrow F$ mutations. In some embodiments, the rAAV particle is not AAV8. In some embodiments, the rAAV particle is AAV8. In some embodiments, the rAAV particle is an AAV8 serotype comprising an rAAV capsid protein as described herein. In some embodiments, the rAAV particle is a modified AAV8. In some embodiments, the rAAV8 has 1, 2, 3 or more $Y \rightarrow F$ (tyrosine to phenylalanine) mutations. In some embodiments, the recombinant AAV8 has a Y275F, Y447F, or Y733F mutation, or any combination thereof (amino acid positions relative to the AAV8 VP1 protein, SEQ ID NO: 5)). In some embodiments, the AAV8 has both a Y447F and Y733F mutation (amino acid positions relative to the AAV8 VP1 protein, SEQ ID NO: 5).

[0076] Non-limiting examples of derivatives and pseudotypes include rAAV2/1, rAAV2/5, rAAV2/8, rAAV2/ 9, AAV2-AAV3 hybrid, AAVrh.10, AAVhu.14, AAV3a/3b, AAVrh32.33, AAV-HSC15, AAV-HSC17, AAVhu.37, AAVrh.8, CHt-P6, AAV2.5, AAV6.2, AAV2i8, AAV-HSC15/17, AAVM41, AAV9.45, AAV6(Y445F/Y731F), AAV2.5T, AAV-HAE1/2, AAV clone 32/83, AAVShH10, AAV2 $(Y \rightarrow F)$, AAV8 (Y733F), AAV2.15, AAV2.4, AAVM41, and AAVr3.45. Such AAV serotypes and derivatives/pseudotypes, and methods of producing such derivatives/pseudotypes are known in the art (see, e.g., Mol Ther. 2012 April; 20(4):699-708. doi: 10.1038/mt.2011.287. Epub 2012 Jan. 24. The AAV vector toolkit: poised at the clinical crossroads. Asokan Al, Schaffer D V, Samulski R J.). In some embodiments, the rAAV particle is a pseudotyped rAAV particle, which comprises (a) a nucleic acid vector comprising ITRs from one serotype (e.g., AAV2) and (b) a capsid comprised of capsid proteins derived from another serotype (e.g., AAV1, AAV3, AAV4, AAV5, AAV6, AAV7, AAV8, AAV9, or AAV10). Methods for producing and using pseudotyped rAAV vectors are known in the art (see, e.g., Duan et al., J. Virol., 75:7662-7671, 2001; Halbert et al., J. Virol., 74:1524-1532, 2000; Zolotukhin et al., Methods, 28:158-167, 2002; and Auricchio et al., Hum. Molec. Genet., 10:3075-3081, 2001).

[0077] Methods of producing rAAV particles and nucleic acid vectors are also known in the art and commercially available (see, e.g., Zolotukhin et al. Production and purification of serotype 1, 2, and 5 recombinant adeno-associated viral vectors. Methods 28 (2002) 158-167; and U.S. Patent Publication Numbers US20070015238 and US20120322861, which are incorporated herein by reference; and plasmids and kits available from ATCC and Cell Biolabs, Inc.). For example, a plasmid containing the nucleic acid vector may be combined with one or more helper plasmids, e.g., that contain a rep gene (e.g., encoding Rep78, Rep68, Rep52 and Rep40) and a cap gene (e.g., encoding VP1, VP2, and VP3, including a modified VP3 region as described herein), and transfected into a producer cell line such that the rAAV particle can be packaged and subsequently purified.

[0078] In some embodiments, the one or more helper plasmids include a first helper plasmid comprising a rep gene and a cap gene (e.g., encoding a rAAV capsid protein as described herein) and a second helper plasmid comprising

a E1a gene, a E1b gene, a E4 gene, a E2a gene, and a VA gene. In some embodiments, the rep gene is a rep gene derived from AAV8 and the cap gene is derived from AAV8 and may include modifications to the gene in order to produce the modified capsid protein described herein. Helper plasmids, and methods of making such plasmids, are known in the art and commercially available (see, e.g., pDM, pDG, pDP1rs, pDP2rs, pDP3rs, pDP4rs, pDP5rs, pDP6rs, pDG(R484E/R585E), and pDP8.ape plasmids from PlasmidFactory, Bielefeld, Germany; other products and services available from Vector Biolabs, Philadelphia, Pa.; Cellbiolabs, San Diego, Calif.; Agilent Technologies, Santa Clara, Calif.; and Addgene, Cambridge, Mass.; pxx6; Grimm et al. (1998), Novel Tools for Production and Purification of Recombinant Adenoassociated Virus Vectors, Human Gene Therapy, Vol. 9, 2745-2760; Kern, A. et al. (2003), Identification of a Heparin-Binding Motif on Adeno-Associated Virus Type 2 Capsids, Journal of Virology, Vol. 77, 11072-11081.; Grimm et al. (2003), Helper Virus-Free, Optically Controllable, and Two-Plasmid-Based Production of Adeno-associated Virus Vectors of Serotypes 1 to 6, Molecular Therapy, Vol. 7, 839-850; Kronenberg et al. (2005), A Conformational Change in the Adeno-Associated Virus Type 2 Capsid Leads to the Exposure of Hidden VP1 N Termini, Journal of Virology, Vol. 79, 5296-5303; and Moullier, P. and Snyder, R. O. (2008), International efforts for recombinant adeno-associated viral vector reference standards, Molecular Therapy, Vol. 16, 1185-1188).

[0079] An exemplary, non-limiting, rAAV particle production method is described next. One or more helper plasmids are produced or obtained, which comprise rep and cap ORFs for the desired AAV serotype and the adenoviral VA, E2A (DBP), and E4 genes under the transcriptional control of their native promoters. The cap ORF may also comprise one or more modifications to produce a modified capsid protein as described herein. HEK293 cells (available from ATCC®) are transfected via CaPO4-mediated transfection, lipids or polymeric molecules such as Polyethylenimine (PEI) with the helper plasmid(s) and a plasmid containing a nucleic acid vector described herein. The HEK293 cells are then incubated for at least 60 hours to allow for rAAV particle production. Alternatively, in another example Sf9-based producer stable cell lines are infected with a single recombinant baculovirus containing the nucleic acid vector. As a further alternative, in another example HEK293 or BHK cell lines are infected with a HSV containing the nucleic acid vector and optionally one or more helper HSVs containing rep and cap ORFs as described herein and the adenoviral VA, E2A (DBP), and E4 genes under the transcriptional control of their native promoters. The HEK293, BHK, or Sf9 cells are then incubated for at least 60 hours to allow for rAAV particle production. The rAAV particles can then be purified using any method known the art or described herein, e.g., by iodixanol step gradient, CsCl gradient, chromatography, or polyethylene glycol (PEG) precipitation.

[0080] The disclosure also contemplates host cells that comprise at least one of the disclosed rAAV particles or nucleic acid vectors. Such host cells include mammalian host cells, with human host cells being preferred, and may be either isolated, in cell or tissue culture. In the case of genetically modified animal models (e.g., a mouse), the transformed host cells may be comprised within the body of a non-human animal itself. In some embodiments, the host cell is a cell of erythroid lineage, such as a CD36+ burst-

forming units-erythroid (BFU-E) cell or a colony-forming unit-erythroid (CFUE-E) progenitor cell.

[0081] In some embodiments, the vector is a PEMBOLds-INS-GFP. In some embodiments, the vector used is a vector well known in the art (e.g., Wang et al., Diabetes 2006 April; 55(4): 875-884. dx.doi.org/10.2337/diabetes.55.04. 06.db05-0927; Wang et al., Gene Ther10 :2105-2111, 2003; Wang et al., Nat Biotechno123 :321-328, 2005).

[0082] In some embodiments, compositions described herein (e.g., siRNA, shRNA) are formulated in a nanoparticle. In some embodiments, compositions described herein (e.g., siRNA, shRNA) are formulated in a lipid nanoparticle. In some embodiments, compositions described herein (e.g., siRNA, shRNA) are formulated in a lipid-polycation complex, referred to as a cationic lipid nanoparticle. The formation of the lipid nanoparticle may be accomplished by methods known in the art and/or as described in U.S. Pub. No. 20120178702, herein incorporated by reference in its entirety. As a non-limiting example, the polycation may include a cationic peptide or a polypeptide such as, but not limited to, polylysine, polyornithine and/or polyarginine and the cationic peptides described in International Pub. No. WO2012013326 or US Patent Pub. No. US20130142818; each of which is herein incorporated by reference in its entirety. In some embodiments, compositions described herein (e.g., siRNA, shRNA) are formulated in a lipid nanoparticle that includes a non-cationic lipid such as, but not limited to, cholesterol or dioleoyl phosphatidylethanolamine (DOPE).

Other Embodiments

[0083] Other Non-Limiting Embodiments of the Disclosure Include Those Below.

[0084] 1. A synthetic ribonucleic acid (RNA) molecule comprising:

[0085] a sense strand of sequence AACCUUUGGGA-GAUGCACAUAUUUGUGAAGCCACA-

GAUGAAAUAUGUGCAUCUC CCAAAGUUUUUGUU (SEQ ID NO: 1) and an antisense strand of sequence AACAAAAACUUUUGGGAGAUGCACAUAUUU-

CAUCUGUGGCUUCACAAAUAUGUGC AUCUC-CCAAAGGUU (SEQ ID NO: 2).

[0086] 2. The synthetic RNA molecule of clause 1, wherein the RNA is a small hairpin RNA (shRNA).

[0087] 3. A shRNA of having a targeted sequence that comprises RNA of sequence AAAUAUGUGCAUCUC-CCAAAG (SEQ ID NO: 3) or RNA of sequence CUUUGGGAGAUGCACAUAUUU (SEQ ID NO: 4).

[0088] 4. The synthetic RNA of any one of clauses 1 to 3, further comprising an unpaired overhang sequence at the 5' and/or 3' end.

[0089] 5. The synthetic RNA of clause 4, wherein the unpaired overhang sequence comprises a sequence of repeating bases.

[0090] 6. The synthetic RNA of clause 4 or 5, wherein the sequence of repeating bases comprises repeating uracil (U) bases.

[0091] 7. The synthetic RNA of clause 4 or 5, wherein the unpaired overhang sequence is UU.

[0092] 8. A composition comprising the synthetic RNA of any one of clauses 1 to 7.

[0093] 9. The composition of clause 8, further comprising one or more physiologically acceptable carriers and/or one or more physiologically acceptable adjuvants.

[0096] 12. The vector of any one of clauses 9 or 10, wherein the vector is an expression plasmid.

[0097] 13. The vector of any one of clauses 9-11, wherein the vector is a viral vector.

[0098] 14. The viral vector of clause 12, wherein the viral vector comprises an adeno-associated viral vector.

[0099] 15. A method of decreasing thymidylate synthase expression in a subject, the method comprising administering to the subject the composition of clauses 8-9 or the vector of clauses 10-14.

[0100] 16. A method of treating pancreatic cancer in a subject, the method comprising administering to the subject the composition of clauses 8-9 or the vector of clauses 10-14.

[0101] 17. The method of clause 16, wherein the composition or vector is delivered using an rAAV.

[0102] 18. The method of clause 16, wherein the pancreatic cancer is islet cell carcinoma.

[0103] 19. The method of clause 17, wherein the rAAV is AAV8.

[0104] 20. The method of clause 16, wherein the interfering RNA is under expression control of promoter sequences. **[0105]** 21. The method of clause 20, wherein the interfering RNA is shRNA, and wherein the shRNA is under expression control of an RNA polymerase III promoter.

[0106] 22. The method of clause 20, wherein the interfering RNA is shRNA, and wherein the shRNA is under expression control of a pancreas-specific promoter.

[0107] 23. The method of any one of clauses 16-22, wherein the subject is a mammal.

[0108] 24. The method of clause 23, wherein the mammal is a rodent or a dog.

[0109] 25. The method of clause 23, wherein the mammal is a human.

[0110] 26. A transgenic mouse comprising pancreatic islet cells that express human thymidylate synthase (TS) and further comprising a conditional Men1 null allele.

EXAMPLES

Example 1

Thymidylate Synthase (TS) Overexpression

[0111] It is hypothesized that human thymidylate synthase accelerates the development of MEN1-driven PanNET and that new strategies to incorporate TS inhibition within current cancer treatment will prevent development and progression of PanNET.

[0112] Thymidylate synthase, essential for cell proliferation, DNA biosynthesis and repair, exhibits oncogene-like activity. High levels of TS correlate with poor prognosis and overall survival in cancer patients. Rahman et at. (Cancer Cell. 2004 April; 5(4):341-51) describes a link between TS-regulated DNA synthesis and the induction of a neoplastic phenotype. Transgenic mice have also been shown to overexpress human TS and then subsequently develop islet hyperplasia or islet cell tumor (Chen et al. Oncogene. 2007 Jul. 19; 26(33):4817-24). Analysis of 320 gastroenteropancreatic neuroendocrine tumors identifies TS expression as independent biomarker for survival (Lee et al. Int J Cancer. 2013 Dec. 18). **[0113]** The lack of suitable animal models that recapitulate human disease has limited development and testing of new treatments for PanNET. A mouse model designated hTS/ Men1^{-/-}, where hTS is overexpressed in pancreatic islet cells carrying a conditional Men1 null allele, was established. Since hTS overexpression accelerates PanNET development in Men1 null mice the research goals focus on defining the mechanism underlying the ability of high levels of hTS to accelerate tumor growth and to develop new treatment strategies.

[0114] Elevated level of thymidylate synthase (TS) plays a direct causal role in tumorigenesis in vitro and overexpression of human TS (hTS) in transgenic mice promotes development of adenomas in the endocrine pancreas in vivo. Pancreatic islet tumor formation in hTS transgenic mice occurred with a long latency period, suggesting that additional somatic events are required to promote PanNET formation and progression. MEN1 was recently shown to be the most commonly mutated tumor suppressor gene in sporadic PanNETs (44% of PanNET patients have MEN1 mutations). Mice with conditional knockouts of Men1 gene in the pancreatic islets develop pancreatic islet tumors with long latency after homozygous inactivation of Men1 gene, suggesting that further sequential somatic events are required for tumor formation.

[0115] hTS transgenic mice were crossed with conditional Men1 null mice that normally develop pancreatic islet carcinoma with long latency. The effect of hTS overexpression on the lifespan of Men1^{-/-} mice vs. hTS/Men1^{-/-} mice was compared. To test whether high levels of hTS results in earlier PanNET development, hTS/Men1^{-/-} and control Men1^{-/-} mice were sacrificed at 5, 6.5 and 8 month of age and tumor development was compared. To determine whether hTS increase mutation frequency, hTS/Men1^{-/-} mice were crossed with Big Blue® transgenic mouse that serves as a mutation detection system. Mutation frequency was analyzed in tumors isolated from hTS/Men1^{-/-} and Men1^{-/-} mice. AAV vectors were used (e.g., AAV8) for delivery of TS shRNA and measured TS levels, tumor progression and survival of hTS/Men1^{-/-} mice.

[0116] The newly established $hTS/Men1^{-/-}$ mice model developed aggressive PanNET with 100% penetrance associated with overexpressed TS in Men1 null mice. TS expression induced islet carcinoma with shortened latency as compared to Men1^{-/-} mice. The hTS/Men1^{-/-} mice develop islet carcinoma as early as 6 month of age whereas Men1⁻ mice develop islet carcinoma at 8 months of age. A significant decrease of overall survival was observed in hTS/ Men1^{-/-} mice as compared to Men1^{-/-} mice (p<0.001). In addition, it was observed that overexpression of TS results in the increase of mutational frequency in tumors derived from the hTS/Men1^{-/-} as compared to control Men1^{-/-} mice. Mutations such as transitions, transversions, insertions and deletions were 3.2 fold higher in tumors isolated from hTS/Men1^{-/-} as compared to control Men1^{-/-} mice. It was shown that high levels of TS increase mutational frequency that may accelerate PanNET progression in hTS/Men1mice. To evaluate the effect of TS inhibition in PanNET progression, AAV-TS shRNA was delivered to the endocrine pancreas of hTS/Men1^{-/-} mice and it was shown that AAV-TS shRNA, specifically targeted pancreatic islet cells, decreased TS expression, significantly decreased PanNET progression and increased survival of hTS/Men1^{-/-} mice.

[0117] The conversion of deoxyuridine monophosphate (dUMP) to deoxythymidine monophosphate (dTMP) with thymidylate synthetase is shown in FIG. 1.

[0118] FIGS. **2**A-2D show generation of novel hTS/ Men1^{-/-} genetically engineered mouse model. FIG. **2**A shows schematic representation of breeding strategy. Arrows indicate primer location for genotyping analysis. FIG. **2**B shows genotyping results of hTS/Men1^{-/-} and Men1^{-/-} mice. FIG. **2**C shows immunoblot anlaysis for TS expression in the pancreases of hTS/Men1^{-/-} and Men1^{-/-} mice. FIG. **2**D shows PanNET progression of hTS/Men1^{-/-} GEMMs. Representative H&E images and pathological photographs of pancreatic islet lesions in hTS/fMen1^{-/-} mice (Scale bar, 100 µm).

[0119] FIGS. **3**A-**3**E show hTS overexpression significantly reduced survival of Men1-null mice. FIG. **3**A shows overall survival analysis of total Men1^{-/-} vs. hTS/Men1^{-/-} mice. FIG. **3**B shows survival analysis of Men1^{-/-} vs. hTS/Men1^{-/-} mice that developed only PanNETs. (FIG. **3**C) mice. FIG. **3**D shows overall survival analysis of total Men1^{+/-} vs. hTS/Men1^{+/-} mice. FIG. **3**E shows separate survival analysis of male and female.

[0120] FIGS. **4**A-**4**B show TS overexpression induces PanNET progression. FIG. **4**A shows Pancreatic islet lesions in Men1^{-/-} vs. hTS/Men1^{-/-} mice. Animals were euthanized at 5, 6.5 and 8 months and pancreas were isolated for histopathologic analysis (n=16 per group at each time point). FIG. **4**B shows pancreatic islet carcinoma incidence. The percentage of mice with islet tumor lesions is shown (*p<0. 01, **p<0.001).

[0121] FIGS. **5**A-**5**C show TS overexpression induces somatic mutations. FIG. **5**A shows an overview of the λ Select-cII Mutation Detection System. FIG. **5**B shows mutation frequencies in Men1^{-/-}/BB and hTS/fMen1^{-/-}/BB mice (n=3 per group at each time point). FIG. **5**C shows the type of mutations in the pancreas and tumors of Men1^{-/-}/BB and hTS/Men1^{-/-}/BB mice at 5 and 10 months of age.

[0122] FIGS. **6A-6**B show overexpression induces DNA double strand breaks. FIG. **6**A shows an immunofluorescence image of λ H2AX foci in MEF-Men1^{*WT*}-vector, MEF-Men1^{*WT*}-hTS cells, MEF Men1^{-/-}-vector and MEF-Men1^{-/-}-hTS cells. Representative nuclei are shown (Scale bars, 10 µm). FIG. **6**B shows quantification of γ H2AX foci (*p<0.01).

[0123] FIGS. 7A-7F show AAV-TS shRNA inhibits Pan-NET progression. FIG. 7A shows a vector map of scAAVmIP-GFP-NSshRNA and scAAV-mIP-GFP-TSshRNA (containing SEQ ID NO:1 and 2) construct. FIG. 7B shows schematics of scAAV-mIP-GFP-NSshRNA (AAV-shNS) or scAAV-mIP-GFP-TSshRNA (AAV-shTS) treatment in hTS/ Men1^{-/-} mice. FIG. 7C shows survival analysis of pancreas tissues from hTS/Men1 mice after TS shRNA injection (n=24 per group). FIG. 7D shows TS mRNA expression levels in tumors. FIG. 7E shows TS protein expression levels in tumors. FIG. 7F shows the percentage of islet tumor lesion (n=9 per group).

[0124] The novel animal model described herein will allow development of new strategies for targeting TS in combination with other FDA approved drugs for the treatment of PanNET. The data provided herein also show that an interfering RNA (e.g., shRNA) that targets TS can be delivered using AAV and is effective for treating pancreatic cancer.

Equivalents

[0125] While several inventive embodiments have been described and illustrated herein, those of ordinary skill in the art will readily envision a variety of other means and/or structures for performing the function and/or obtaining the results and/or one or more of the advantages described herein, and each of such variations and/or modifications is deemed to be within the scope of the inventive embodiments described herein. More generally, those skilled in the art will readily appreciate that all parameters, dimensions, materials, and configurations described herein are meant to be exemplary and that the actual parameters, dimensions, materials, and/or configurations will depend upon the specific application or applications for which the inventive teachings is/are used. Those skilled in the art will recognize, or be able to ascertain using no more than routine experimentation, many equivalents to the specific inventive embodiments described herein. It is, therefore, to be understood that the foregoing embodiments are presented by way of example only and that, within the scope of the appended claims and equivalents thereto, inventive embodiments may be practiced otherwise than as specifically described and claimed. Inventive embodiments of the present disclosure are directed to each individual feature, system, article, material, kit, and/or method described herein. In addition, any combination of two or more such features, systems, articles, materials, kits, and/or methods, if such features, systems, articles, materials, kits, and/or methods are not mutually inconsistent, is included within the inventive scope of the present disclosure.

[0126] All definitions, as defined and used herein, should be understood to control over dictionary definitions, definitions in documents incorporated by reference, and/or ordinary meanings of the defined terms.

[0127] All references, patents and patent applications disclosed herein are incorporated by reference with respect to the subject matter for which each is cited, which in some cases may encompass the entirety of the document.

[0128] The indefinite articles "a" and "an," as used herein in the specification and in the claims, unless clearly indicated to the contrary, should be understood to mean "at least one."

[0129] The phrase "and/or," as used herein in the specification and in the claims, should be understood to mean "either or both" of the elements so conjoined, i.e., elements that are conjunctively present in some cases and disjunctively present in other cases. Multiple elements listed with "and/or" should be construed in the same fashion, i.e., "one or more" of the elements so conjoined. Other elements may optionally be present other than the elements specifically identified by the "and/or" clause, whether related or unrelated to those elements specifically identified. Thus, as a non-limiting example, a reference to "A and/or B", when used in conjunction with open-ended language such as "comprising" can refer, in one embodiment, to A only (optionally including elements other than B); in another embodiment, to B only (optionally including elements other than A); in yet another embodiment, to both A and B (optionally including other elements); etc.

[0130] As used herein in the specification and in the claims, "or" should be understood to have the same meaning as "and/or" as defined above. For example, when separating items in a list, "or" or "and/or" shall be interpreted as being inclusive, i.e., the inclusion of at least one, but also including

more than one, of a number or list of elements, and, optionally, additional unlisted items. Only terms clearly indicated to the contrary, such as "only one of" or "exactly one of," or, when used in the claims, "consisting of," will refer to the inclusion of exactly one element of a number or list of elements. In general, the term "or" as used herein shall only be interpreted as indicating exclusive alternatives (i.e. "one or the other but not both") when preceded by terms of exclusivity, such as "either," "one of," "only one of," or "exactly one of." "Consisting essentially of," when used in the claims, shall have its ordinary meaning as used in the field of patent law.

[0131] As used herein in the specification and in the claims, the phrase "at least one," in reference to a list of one or more elements, should be understood to mean at least one element selected from any one or more of the elements in the list of elements, but not necessarily including at least one of each and every element specifically listed within the list of elements and not excluding any combinations of elements in the list of elements. This definition also allows that elements may optionally be present other than the elements specifically identified within the list of elements to which the phrase "at least one" refers, whether related or unrelated to those elements specifically identified. Thus, as a non-limiting example, "at least one of A and B" (or, equivalently, "at least one of A or B," or, equivalently "at least one of A and/or B") can refer, in one embodiment, to at least one, optionally including more than one, A, with no B present (and optionally including elements other than B); in another embodiment, to at least one, optionally including more than one, B, with no A present (and optionally including elements other than A); in yet another embodiment, to at least one, optionally including more than one, A, and at least one, optionally including more than one, B (and optionally including other elements); etc.

[0132] It should also be understood that, unless clearly indicated to the contrary, in any methods claimed herein that include more than one step or act, the order of the steps or acts of the method is not necessarily limited to the order in which the steps or acts of the method are recited.

[0133] In the claims, as well as in the specification above, all transitional phrases such as "comprising," "including," "carrying," "having," "containing," "involving," "holding," "composed of," and the like are to be understood to be open-ended, i.e., to mean including but not limited to. Only the transitional phrases "consisting of" and "consisting essentially of' shall be closed or semi-closed transitional phrases, respectively, as set forth in the United States Patent Office Manual of Patent Examining Procedures, Section 2111.03. It should be appreciated that embodiments described in this document using an open-ended transitional phrase (e.g., "comprising") are also contemplated, in alternative embodiments, as "consisting of" and "consisting essentially of" the feature described by the open-ended transitional phrase. For example, if the disclosure describes "a composition comprising A and B", the disclosure also contemplates the alternative embodiments "a composition consisting of A and B" and "a composition consisting essentially of A and B".

SEQUENCE LISTING

<160> NUMBER OF SEQ ID NOS: 11 <210> SEQ ID NO 1 <211> LENGTH: 68 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEOUENCE: 1 aaccuuuggg agaugcacau auuugugaag ccacagauga aauaugugca ucucccaaag 60 uuuuuquu 68 <210> SEQ ID NO 2 <211> LENGTH: 68 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEOUENCE: 2 aacaaaaacu uugggagaug cacauauuuc aucuguggcu ucacaaauau gugcaucucc 60 caaagguu 68 <210> SEQ ID NO 3 <211> LENGTH: 21 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide

16

<400> SEQUENCE: 3	
aaauaugugc aucucccaaa g	21
<210> SEQ ID NO 4 <211> LENGTH: 21 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide	
<400> SEQUENCE: 4	
cuuugggaga ugcacauauu u	21
<210> SEQ ID NO 5 <211> LENGTH: 738 <212> TYPE: PRT <213> ORGANISM: Adeno-associated virus 8	
<400> SEQUENCE: 5	
Met Ala Ala Asp Gly Tyr Leu Pro Asp Trp Leu Glu Asp Asn Leu Ser 1 5 10 15	
Glu Gly Ile Arg Glu Trp Trp Ala Leu Lys Pro Gly Ala Pro Lys Pro 20 25 30	
Lys Ala Asn Gln Gln Lys Gln Asp Asp Gly Arg Gly Leu Val Leu Pro 35 40 45	
Gly Tyr Lys Tyr Leu Gly Pro Phe Asn Gly Leu Asp Lys Gly Glu Pro 50 55 60	
Val Asn Ala Ala Asp Ala Ala Ala Leu Glu His Asp Lys Ala Tyr Asp 65 70 75 80	
Gln Gln Leu Gln Ala Gly Asp Asn Pro Tyr Leu Arg Tyr Asn His Ala 85 90 95	
Asp Ala Glu Phe Gln Glu Arg Leu Gln Glu Asp Thr Ser Phe Gly Gly 100 105 110	
Asn Leu Gly Arg Ala Val Phe Gln Ala Lys Lys Arg Val Leu Glu Pro 115 120 125	
Leu Gly Leu Val Glu Glu Gly Ala Lys Thr Ala Pro Gly Lys Lys Arg 130 135 140	
Pro Val Glu Pro Ser Pro Gln Arg Ser Pro Asp Ser Ser Thr Gly Ile145150155160	
Gly Lys Lys Gly Gln Gln Pro Ala Arg Lys Arg Leu Asn Phe Gly Gln 165 170 175	
Thr Gly Asp Ser Glu Ser Val Pro Asp Pro Gln Pro Leu Gly Glu Pro 180 185 190	
Pro Ala Ala Pro Ser Gly Val Gly Pro Asn Thr Met Ala Ala Gly Gly 195 200 205	
Gly Ala Pro Met Ala Asp Asn Asn Glu Gly Ala Asp Gly Val Gly Ser 210 215 220	
Ser Ser Gly Asn Trp His Cys Asp Ser Thr Trp Leu Gly Asp Arg Val 225 230 235 240	
Ile Thr Thr Ser Thr Arg Thr Trp Ala Leu Pro Thr Tyr Asn Asn His 245 250 255	
Leu Tyr Lys Gln Ile Ser Asn Gly Thr Ser Gly Gly Ala Thr Asn Asp 260 265 270	
Asn Thr Tyr Phe Gly Tyr Ser Thr Pro Trp Gly Tyr Phe Asp Phe Asn	

-continued

		275					280					285			
Arg	Phe 290	His	Сув	His	Phe	Ser 295	Pro	Arg	Asp	Trp	Gln 300	Arg	Leu	Ile	Asn
Asn 305	Asn	Trp	Gly	Phe	Arg 310	Pro	Lys	Arg	Leu	Ser 315	Phe	Lys	Leu	Phe	Asn 320
Ile	Gln	Val	Lys	Glu 325	Val	Thr	Gln	Asn	Glu 330	Gly	Thr	Lys	Thr	Ile 335	Ala
Asn	Asn	Leu	Thr 340	Ser	Thr	Ile	Gln	Val 345	Phe	Thr	Asp	Ser	Glu 350	Tyr	Gln
Leu	Pro	Tyr 355	Val	Leu	Gly	Ser	Ala 360	His	Gln	Gly	Сүз	Leu 365	Pro	Pro	Phe
Pro	Ala 370	Asp	Val	Phe	Met	Ile 375	Pro	Gln	Tyr	Gly	Tyr 380	Leu	Thr	Leu	Asn
Asn 385	Gly	Ser	Gln	Ala	Val 390	Gly	Arg	Ser	Ser	Phe 395	Tyr	Суз	Leu	Glu	Tyr 400
Phe	Pro	Ser	Gln	Met 405	Leu	Arg	Thr	Gly	Asn 410	Asn	Phe	Gln	Phe	Thr 415	Tyr
Thr	Phe	Glu	Asp 420	Val	Pro	Phe	His	Ser 425	Ser	Tyr	Ala	His	Ser 430	Gln	Ser
Leu	Asp	Arg 435	Leu	Met	Asn	Pro	Leu 440	Ile	Asp	Gln	Tyr	Leu 445	Tyr	Tyr	Leu
Ser	Arg 450	Thr	Gln	Thr	Thr	Gly 455	Gly	Thr	Ala	Asn	Thr 460	Gln	Thr	Leu	Gly
Phe 465	Ser	Gln	Gly	Gly	Pro 470	Asn	Thr	Met	Ala	Asn 475	Gln	Ala	Lys	Asn	Trp 480
Leu	Pro	Gly	Pro	Cys 485	Tyr	Arg	Gln	Gln	Arg 490	Val	Ser	Thr	Thr	Thr 495	Gly
Gln	Asn	Asn	Asn 500	Ser	Asn	Phe	Ala	Trp 505	Thr	Ala	Gly	Thr	Lys 510	Tyr	His
Leu	Asn	Gly 515	Arg	Asn	Ser	Leu	Ala 520	Asn	Pro	Gly	Ile	Ala 525	Met	Ala	Thr
His	Lys 530	Asp	Asp	Glu	Glu	Arg 535	Phe	Phe	Pro	Ser	Asn 540	Gly	Ile	Leu	Ile
Phe 545	Gly	Lys	Gln	Asn	Ala 550	Ala	Arg	Asp	Asn	Ala 555	Asp	Tyr	Ser	Asp	Val 560
Met	Leu	Thr	Ser	Glu 565	Glu	Glu	Ile	Lys	Thr 570	Thr	Asn	Pro	Val	Ala 575	Thr
Glu	Glu	Tyr	Gly 580	Ile	Val	Ala	Asp	Asn 585	Leu	Gln	Gln	Gln	Asn 590	Thr	Ala
Pro	Gln	Ile 595	Gly	Thr	Val	Asn	Ser 600	Gln	Gly	Ala	Leu	Pro 605	Gly	Met	Val
Trp	Gln 610	Asn	Arg	Aap	Val	Tyr 615	Leu	Gln	Gly	Pro	Ile 620	Trp	Ala	Lys	Ile
Pro 625	His	Thr	Asp	Gly	Asn 630	Phe	His	Pro	Ser	Pro 635	Leu	Met	Gly	Gly	Phe 640
Gly	Leu	Lys	His	Pro 645	Pro	Pro	Gln	Ile	Leu 650	Ile	Lys	Asn	Thr	Pro 655	Val
Pro	Ala	Asp	Pro 660	Pro	Thr	Thr	Phe	Asn 665	Gln	Ser	Lys	Leu	Asn 670	Ser	Phe
Ile	Thr	Gln 675	Tyr	Ser	Thr	Gly	Gln 680	Val	Ser	Val	Glu	Ile 685	Glu	Trp	Glu

-continued

18

Leu Gln Lys Glu Asn Ser Lys Arg Trp Asn Pro Glu Ile Gln Tyr Thr 690 695 700 Ser Asn Tyr Tyr Lys Ser Thr Ser Val Asp Phe Ala Val Asn Thr Glu 705 710 715 720 Gly Val Tyr Ser Glu Pro Arg Pro Ile Gly Thr Arg Tyr Leu Thr Arg 730 725 735 Asn Leu <210> SEQ ID NO 6 <211> LENGTH: 1524 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 6 cgcgccactt ggcctgcctc cgtcccgccg cgccacttcg cctgcctccg tcccccgccc 60 geogegeeat geotgtggee ggeteggage tgeogegeeg geoettgeee eccgeegeae 120 aqqaqcqqqa cqccqaqccq cqtccqccqc acqqqqaqct qcaqtacctq qqqcaqatcc 180 aacacateet eegetgegge gteaggaagg acgaeegeae gggeaeegge accetgtegg 240 300 tattcqqcat qcaqqcqcqc tacaqcctqa qaqatqaatt ccctctqctq acaaccaaac gtgtgttctg gaagggtgtt ttggaggagt tgctgtggtt tatcaaggga tccacaaatg 360 ctaaagagct gtcttccaag ggagtgaaaa tctgggatgc caatggatcc cgagactttt 420 tggacageet gggattetee accagagaag aaggggaett gggeeeagtt tatggettee 480 agtggaggca ttttgggggca gaatacagag atatggaatc agattattca ggacagggag 540 ttgaccaact gcaaagagtg attgacacca tcaaaaccaa ccctgacgac agaagaatca 600 tcatgtgcgc ttggaatcca agagatcttc ctctgatggc gctgcctcca tgccatgccc 660 tctgccagtt ctatgtggtg aacagtgagc tgtcctgcca gctgtaccag agatcgggag 720 acatgggcct cggtgtgcct ttcaacatcg ccagctacgc cctgctcacg tacatgattg 780 cgcacatcac gggcctgaag ccaggtgact ttatacacac tttgggagat gcacatattt 840 acctgaatca catcgagcca ctgaaaattc agcttcagcg agaacccaga cctttcccaa 900 ageteaggat tettegaaaa gttgagaaaa ttgatgaett caaagetgaa gaettteaga 960 ttgaagggta caatccgcat ccaactatta aaatggaaat ggctgtttag ggtgctttca 1020 aaggagettg aaggatattg teagtettta ggggttggge tggatgeega ggtaaaagtt 1080 ctttttgctc taaaagaaga aggaactagg tcaaaaatct gtccgtgacc tatcagttat 1140 taatttttaa ggatgttgcc actggcaaat gtaactgtgc cagttctttc cataataaaa 1200 ggctttgagt taactcactg agggtatctg acaatgctga ggttatgaac aaagtgagga 1260 gaatgaaatg tatgtgctct tagcaaaaac atgtatgtgc atttcaatcc cacgtactta 1320 taaagaaggt tggtgaattt cacaagctat ttttggaata tttttagaat attttaagaa 1380 tttcacaagc tattccctca aatctgaggg agctgagtaa caccatcgat catgatgtag 1440 agtgtggtta tgaactttat agttgtttta tatgttgcta taataaagaa gtgttctgca 1500 ttcgtaaaaa aaaaaaaaaa aaaa 1524

<210> SEQ ID NO 7

-continued

19

<211> LENGTH: 1569 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polypucleotide	
ADD. CECHENCE. 7	
	60
gggggege ggaagggte elgeeacege geeacligge elgeeleegt eeegeegege	100
saertegeet geereegtee eeegeeegee gegeeatgee tgtggeegge teggagetge	120
rgegeeggee ellgeeceee geegeacagg agegggaege egageeget eegeegeacg	180
gggagetgea gtaeetgggg cagateeaae acateeteeg etgeggegte aggaaggaeg	240
acegeacggg caeeggeace eigieggiat teggeatgea ggegegetae ageetgagag	300
atgaatteee tetgetgaea accaaacgtg tgttetggaa gggtgttttg gaggagttge	360
tgtggtttat caagggatee acaaatgeta aagagetgte tteeaaggga gtgaaaatet	420
gggatgecaa tggateeega gaettittgg acageetggg atteteeaee agagaagaag	480
gggacttggg cocagtttat ggottocagt ggaggcattt tgggggcagaa tacagagata	540
tggaatcaga ttattcagga cagggagttg accaactgca aagagtgatt gacaccatca	600
aaaccaaccc tgacgacaga agaatcatca tgtgcgcttg gaatccaaga gatcttcctc	660
tgatggeget geeteeatge eatgeeetet geeagtteta tgtggtgaae agtgagetgt	720
cetgecaget gtaccagaga tegggagaea tgggeetegg tgtgeettte aacategeea	780
getaegeeet geteaegtae atgattgege acateaeggg eetgaageea ggtgaettta	840
tacacacttt gggagatgca catatttacc tgaatcacat cgagccactg aaaattcagc	900
ttcagcgaga acccagacct ttcccaaagc tcaggattct tcgaaaagtt gagaaaattg	960
atgacttcaa agctgaagac tttcagattg aagggtacaa tccgcatcca actattaaaa	1020
tggaaatggc tgtttagggt gctttcaaag gagctcgaag gatattgtca gtctttaggg	1080
yttgggctgg atgccgaggt aaaagttett tttgetetaa aagaaaaagg aactaggtea	1140
aaaatotgto ogtgaootat cagttattaa tttttaagga tgttgooact ggoaaatgta	1200
actgtgccag ttctttccat aataaaaggc tttgagttaa ctcactgagg gtatctgaca	1260
atgetgaggt tatgaacaaa gtgaggagaa tgaaatgtat gtgetettag caaaaacatg	1320
tatgtgcatt tcaatcccac gtacttataa agaaggttgg tgaatttcac aagctatttt	1380
tggaatattt ttagaatatt ttaagaattt cacaagctat tccctcaaat ctgagggagc	1440
tgagtaacac catcgatcat gatgtagagt gtggttatga actttaaagt tatagttgtt	1500
ttatatgttg ctataataaa gaagtgttct gcattcgcca aaaaaaaaaa	1560
aaaaaaaa	1569
<210> SEQ ID NO 8 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide	
<400> SEQUENCE: 8	
atgeetgtgg eeggetegga getgeegege eggeeettge eeeeegeege acaggagegg	60
gacgeegage egegteegee geaeggggag etgeagtaee tggggeagat eeaaeaeate	120

20

-continued

ctccgctgcg						
	gcgtcaggaa	ggacgaccgc	acgggcaccg	gcaccctgtc	ggtattcggc	180
atgcaggcgc	gctacagcct	gagagatgaa	ttccctctgc	tgacaaccaa	acgtgtgttc	240
tggaagggtg	ttttggagga	gttgctgtgg	tttatcaagg	gatccacaaa	tgctaaagag	300
ctgtcttcca	agggagtgaa	aatctgggat	gccaatggat	cccgagactt	tttggacagc	360
ctgggattct	ccaccagaga	agaagggggac	ttgggcccag	tttatggctt	ccagtggagg	420
cattttgggg	cagaatacag	agatatggaa	tcagattatt	caggacaggg	agttgaccaa	480
ctgcaaagag	tgattgacac	catcaaaacc	aaccctgacg	acagaagaat	catcatgtgc	540
gcttggaatc	caagagatct	tcctctgatg	gcgctgcctc	catgccatgc	cctctgccag	600
ttctatgtgg	tgaacagtga	gctgtcctgc	cagctgtacc	agagatcggg	agacatgggc	660
ctcggtgtgc	ctttcaacat	cgccagctac	gccctgctca	cgtacatgat	tgcgcacatc	720
acgggcctga	agccaggtga	ctttatacac	actttgggag	atgcacatat	ttacctgaat	780
cacatcgagc	cactgaaaat	tcagcttcag	cgagaaccca	gacctttccc	aaagctcagg	840
attcttcgaa	aagttgagaa	aattgatgac	ttcaaagctg	aagactttca	gattgaaggg	900
tacaatccgc	atccaactat	taaaatggaa	atggetgttt	tg		942
<210> SEQ 3 <211> LENG" <212> TYPE <213> ORGAN <220> FEATU <223> OTHEN	D NO 9 CH: 942 DNA HISM: Artif: JRE: NFORMATIC	icial Sequer DN: Syntheti	nce ic Polynucle	eotide		
<400> SEQU	ENCE: 9					
atgcctgtgg	ccggctcgga	gctgccgcgc	cggcccttgc	cccccgccgc	acaggagcgg	60
gacgccgagc	cgcgtccgcc	acacaaaaaa	ctocantacc			
		geacgggggag	eegeageaee	tggggcagat	ccaacacatc	120
ctccgctgcg	gcgtcaggaa	ggacgaccgc	acgggcaccg	tggggcagat gcaccctgtc	ccaacacatc ggtattcggc	120 180
ctccgctgcg atgcaggcgc	gcgtcaggaa gctacagcct	ggacgaccgc gagagatgaa	acgggcaccg ttccctctgc	tggggcagat gcaccctgtc tgacaaccaa	ccaacacatc ggtattcggc acgtgtgttc	120 180 240
ctccgctgcg atgcaggcgc tggaagggtg	gcgtcaggaa gctacagcct ttttggagga	ggacgaccgc gagagatgaa gttgctgtgg	acgggcaccg ttccctctgc tttatcaagg	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag	120 180 240 300
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca	gcgtcaggaa gctacagcct ttttggagga agggagtgaa	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat	acgggcaccg ttccctctgc tttatcaagg gccaatggat	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc	120 180 240 360
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaaggggac	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg	120 180 240 300 360 420
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct catttgggg	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga cagaatacag	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaaggggac agatatggaa	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag tcagattatt	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt caggacaggg	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg agttgaccaa	120 180 240 360 420 480
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct cattttgggg ctgcaaagag	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga cagaatacag tgattgacac	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaaggggac agatatggaa catcaaaacc	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag tcagattatt aaccctgacg	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt caggacaggg acagaagaat	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg agttgaccaa catcatgtgc	120 180 240 300 360 420 480 540
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct catttgggg ctgcaaagag gcttggaatc	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga cagaatacag tgattgacac caagagatct	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaagggggac agatatggaa catcaaaacc tcctctgatg	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag tcagattatt aaccctgacg gcgctgcctc	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt caggacaggg acagaagaat catgccatgc	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg agttgaccaa catcatgtgc cctctgccag	120 180 240 300 360 420 480 540
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct catttgggg ctgcaaagag gcttggaatc ttctatgtgg	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga cagaatacag tgattgacac caagagatct tgaacagtga	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaaggggac agatatggaa catcaaaacc tcctctgatg gctgtcctgc	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag tcagattatt aaccctgacg gcgctgcctc cagctgtacc	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt caggacaggg acagaagaat catgccatgc	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg agttgaccaa catcatgtgc cctctgccag agacatgggc	120 180 240 300 360 420 480 540 600
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct catttgggg ctgcaaagag gcttggaatc ttctatgtgg ctcggtgtgc	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga cagaatacag tgattgacac caagagatct tgaacagtga ctttcaacat	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaaggggac agatatggaa catcaaaacc tcctctgatg gctgtcctgc cgccagctac	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag tcagattatt aaccctgacg gcgctgcctc cagctgtacc gccctgctca	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt caggacaggg acagaagaat catgccatgc	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg agttgaccaa catcatgtgc cctctgccag agacatgggc tgcgcacatc	120 180 240 300 360 420 480 540 600 660
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct catttgggg gctggaatc ttctagtgg ctcggtgtgc acgggcctga	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga cagaatacag tgattgacac caagagatct tgaacagtga ctttcaacat agccaggtga	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaagggggac agatatggaa catcaaaacc tcctctgatg gctgtcctgc cgccagctac ctttatacac	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag tcagattatt aaccctgacg gcgctgcctc cagctgtacc gccctgctca actttgggag	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt caggacaggg acagaagaat catgccatgc	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg agttgaccaa catcatgtgc cctctgccag agacatgggc tgcgcacatc ttacctgaat	120 180 240 300 360 420 480 540 600 720 780
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct catttgggg gcttggaatc ttctatgtgg ctcggtgtgc acgggcctga cacatcgagc	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga tgattgacac caagagatct tgaacagtga ctttcaacat agccaggtga cactgaaaat	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaaggggac agatatggaa catcaaaacc tcctctgatg gctgtcctgc cgccagctac ctttatacac	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag tcagattatt aaccctgacg gcgctgcctc cagctgtacc gccctgctca actttgggag cgagaaccca	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt caggacaggg acagaagaat catgccatgc	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg agttgaccaa catcatgtgc cctctgccag agacatgggc tgcgcacatc ttacctgaat aaagctcagg	120 180 240 300 360 420 480 540 600 660 720 780 840
ctccgctgcg atgcaggcgc tggaagggtg ctgtcttcca ctgggattct catttgggg gctggaatc ttctatgtgg ctcggtgtgc acgggcctga cacatcgagc attcttcgaa	gcgtcaggaa gctacagcct ttttggagga agggagtgaa ccaccagaga tgattgacac caagagatct tgaacagtga ctttcaacat agccaggtga cactgaaaat aagttgagaa	ggacgaccgc gagagatgaa gttgctgtgg aatctgggat agaagggggac agatatggaa catcaaaacc tcctctgatg gctgtcctgc cgccagctac ctttatacac tcagcttcag aattgatgac	acgggcaccg ttccctctgc tttatcaagg gccaatggat ttgggcccag tcagattatt aaccctgacg gcgctgctcc cagctgtacc gccctgctca actttgggag cgagaaccca ttcaaagctg	tggggcagat gcaccctgtc tgacaaccaa gatccacaaa cccgagactt tttatggctt caggacaggg acagaagaat catgccatgc	ccaacacatc ggtattcggc acgtgtgttc tgctaaagag tttggacagc ccagtggagg agttgaccaa catcatgtgc cctctgccag agacatgggc tgcgcacatc ttacctgaat aaagctcagg	120 180 240 300 360 420 480 540 600 720 780 840 900

<210> SEQ ID NO 10 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE:

-continued

<223> OTHER INFORMATION: Synthetic Polynucleotide	
<400> SEQUENCE: 10	
atgeetgtgg eeggetegga getgeegege eggeeettge eeeeegeege acaggagegg	60
gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc	120
ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc	180
atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc	240
tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag	300
ctgtcttcca agggagtgaa aatctgggat gccaatggat cccgagactt tttggacagc	360
ctgggattct ccaccagaga agaagggggac ttggggcccag tttatggctt ccagtggagg	420
cattttgggg cagaatacag agatatggaa tcagattatt caggacaggg agttgaccaa	480
ctgcaaagag tgattgacac catcaaaacc aaccctgacg acagaagaat catcatgtgc	540
gcttggaatc caagagatct tcctctgatg gcgctgcctc catgccatgc	600
ttctatgtgg tgaacagtga gctgtcctgc cagctgtacc agagatcggg agacatgggc	660
ctcggtgtgc ctttcaacat cgccagctac gccctgctca cgtacatgat tgcgcacatc	720
acgggcctga agccaggtga ctttatacac actttgggag atgcacatat ttacctgaat	780
cacatcgagc cactgaaaat tcagcttcag cgagaaccca gacctttccc aaagctcagg	840
attettegaa aagttgagaa aattgatgae tteaaagetg aagaetttea gattgaaggg	900
tacaatccgc atccaactat taaaatggaa atggctgttt tg	942
-2105 SEO ID NO 11	
<210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide	
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11</pre>	
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg</pre>	60
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccgcgtccgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc</pre>	60 120
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc</pre>	60 120 180
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc</pre>	60 120 180 240
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag</pre>	60 120 180 240 300
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag ctgtcttcca agggagtgaa aatctgggat gccaatggat cccgagact tttggacagc</pre>	60 120 180 240 300 360
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag ctgtcttcca agggagtgaa aatctgggat gccaatggat cccagagctt tttggacagc ctgggattct ccaccagaga agaaggggac ttgggcccag tttatggctt ccagtggagg</pre>	60 120 180 240 300 360 420
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag ctgtcttcca agggagtgaa aatctgggat gccaatggat cccgagactt tttggacagc ctgggattct ccaccagaga agaaggggac ttgggcccag tttatggctt ccagtggagg cattttgggg cagaatacag agaatatggaa tccagattatt caggacaggg agttgaccaa</pre>	60 120 180 240 300 360 420 480
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag ctgtcttcca agggagtgaa aatctgggat gccaatggat cccagagatt tttggacagc ctgggattct ccaccagaga agaaggggac ttgggcccag tttatggctt ccagtggagg cattttgggg cagaatacag agatatggaa tcagattatt caggacaggg agttgaccaa ctgcaaagag tgattgacac catcaaaacc aaccctgacg acagaagaat catcatgtgc</pre>	60 120 180 240 360 420 480 540
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacgggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag ctgtcttcca agggagtgaa aatctgggat gccaatggat cccagagatt tttggacagc ctgggattct ccaccagaga agaaggggac ttgggcccag tttatggctt ccagtggagg cattttgggg cagaatacag agatatggaa tcagattatt caggacagg agttgaccaa ctgcaaagag tgattgacac catcaaaacc aaccctgacg acagaagaat catcatgtgc gcttggaatc caagagatct tcctctgatg gcgctgcctc catgccatgc</pre>	60 120 180 240 300 360 420 480 540
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag ctgtcttcca agggagtgaa aatctgggat gccaatggat cccagagat tttggacagc ctgggattct ccaccagaga agaaggggac ttgggcccag tttatggctt ccagtggagg cattttgggg cagaatacag agatatggaa tccgattatt caggacaggg agttgaccaa ctgcaaagag tgattgacac catcaaaacc aaccctgacg acagagaat catcatgtgc gcttggaatc caagagatct tcctctgatg gcgctgcctc catgccatgc</pre>	60 120 180 240 360 420 480 540 600 660
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctccgca gcaccggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccga acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagaagtgaa ttccctctgc tgacaaccaa acgtggttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag ctgtcttcca agggagtgaa aatctgggat gccaatggat cccagacagg gagattet ccaaccagaga agaaggggac ttgggcccag tttatggct ccagtggagg cattttgggg cagaatacag agatatggaa tcagattatt caggacaggg agttgaccaa ctgcaaagag tgattgacac catcaaaacc aaccctgacg acagaagaat catcatgtgc gcttggaatc caagagatct tcctctgatg gcgctgcctc catgccatg ccttgccag ttctatgtgg tgaacagtga gctgtcctgc cagctgtacc agagatcgg agacatgggc ctcggtgtgc ctttcaacat cgccagctac gccctgctca cgtacatgat tgcgcacatc tccggtgtgc ctttcaacat cgccagctac gccctgcta cgtacatgat tgcgcacatc tccggtgtgc ctttcaacat cgccagctac gccctgcta cgtacatgat tgcgcacatc</pre>	60 120 180 240 300 360 420 480 540 600 660
<pre><210> SEQ ID NO 11 <211> LENGTH: 942 <212> TYPE: DNA <213> ORGANISM: Artificial Sequence <220> FEATURE: <223> OTHER INFORMATION: Synthetic Polynucleotide <400> SEQUENCE: 11 atgcctgtgg ccggctcgga gctgccgcgc cggcccttgc cccccgccgc acaggagcgg gacgccgagc cgcgtccgcc gcacggggag ctgcagtacc tggggcagat ccaacacatc ctccgctgcg gcgtcaggaa ggacgaccgc acgggcaccg gcaccctgtc ggtattcggc atgcaggcgc gctacagcct gagagatgaa ttccctctgc tgacaaccaa acgtgtgttc tggaagggtg ttttggagga gttgctgtgg tttatcaagg gatccacaaa tgctaaagag ctgtcttcca agggagtgaa aatctgggat gccaatggat cccagtaggag ctgggattct ccaccagaga agaaggggac ttgggcccag tttatggct ccagtggagg ctgggattct ccaccagaga agaagggac ttgggcccag tttatggct ccagtggagg ctgtcaaagag tgattgacac catcaaaacc aaccctgacg acagaagaat catcatgtgc gcttggaatc caagagatct tcctctgatg gcgctgcctc catgccatgc</pre>	60 120 180 240 300 360 420 480 540 600 660 720 780

-continued

attcttcgaa	aagttgagaa	aattgatgac	ttcaaagctg	aagactttca	gattgaaggg	900
tacaatccgc	atccaactat	taaaatggaa	atggctgttt	ag		942

What is claimed is:

1. An recombinant adeno-associated viral (rAAV) particle comprising a nucleic acid vector that comprises (a) a heterologous nucleic acid region comprising a sequence that encodes an interfering RNA that comprises a region of complementarity with a thymidylate synthase mRNA and (b) inverted terminal repeat (ITR) sequences flanking the heterologous nucleic acid region.

2. The rAAV particle of claim **1**, wherein the interfering RNA is a small hairpin RNA (shRNA) or a microRNA.

3. The rAAV particle of claim **1** or **2**, wherein the region of complementarity is 100% complementary to at least 8 contiguous nucleotides of any of SEQ ID NOs: 6-11.

4. The rAAV particle of claim **1** or **2**, wherein the interfering RNA comprises at least 8 contiguous nucleotides of SEQ ID NOs: 1, 2, 3 and/or 4.

5. The rAAV particle of any one of claims **1** to **4**, wherein the particle is an AAV8 particle or a modified AAV8 particle.

6. The rAAV particle of claim 5, wherein the modified AAV8 particle comprises an AAV8 capsid protein comprising a Y275F, Y447F, or Y733F mutation, or any combination thereof.

7. The rAAV particle of claim 6, wherein the AAV8 capsid protein comprises both a Y447F and Y733F mutation.

8. The rAAV particle of any one of claims **1** to **7**, wherein the interfering RNA is under expression control of a promoter sequence.

9. The rAAV particle of claim **8**, wherein the interfering RNA is under expression control of an RNA polymerase III promoter.

10. The rAAV particle of claim 8, wherein the interfering RNA is under expression control of a pancreas-specific promoter, optionally wherein the pancreas-specific promoter is an islet-specific promoter, further optionally wherein the islet-specific promoter is an insulin promoter.

11. A composition comprising the rAAV particle of any one of claims 1-10.

12. The composition of claim **11**, further comprising one or more physiologically acceptable carriers and/or one or more physiologically acceptable adjuvants.

13. A method of decreasing thymidylate synthase expression in a cell, the method comprising administering to the cell the rAAV particle of any one of claims 1 to 10 or the composition of claim 11 or 12.

14. The method of claim 13, wherein the cell is a cell of the pancreas.

15. A method of decreasing thymidylate synthase expression in a subject, the method comprising administering to the subject the rAAV particle of any one of claims 1 to 10 or the composition of claim 11 or 12.

16. The method of claim **15**, wherein the administration results in delivery of the rAAV particle to the pancreas.

17. A method of treating pancreatic cancer in a subject, the method comprising administering to the subject the rAAV particle of any one of claims 1 to 10 or the composition of claim 11 or 12.

18. The method of claim **17**, wherein the pancreatic cancer is islet cell carcinoma.

19. The method of any one of claims **15-18**, wherein the subject is a human subject.

20. A synthetic ribonucleic acid (RNA) molecule comprising:

a sense strand of sequence AACCUUUGGGAGAUGCA-CAUAUUUGUGAAGCCACAGAUGAAAUAU-GUGCAUCUC CCAAAGUUUUUUGUU (SEQ ID NO: 1) and an antisense strand of sequence AACAAAAACUUUGGGAGAUGCACAUAUUU-CAUCUGUGGCUUCACAAAUAUGUGC AUCUC-CCAAAGGUU (SEQ ID NO: 2).

21. The synthetic RNA molecule of claim **20**, wherein the RNA is a small hairpin RNA (shRNA).

22. A shRNA of having a targeted sequence that comprises RNA of sequence AAAUAUGUGCAUCUC-CCAAAG (SEQ ID NO: 3) or RNA of sequence CUUUGGGAGAUGCACAUAUUU (SEQ ID NO: 4).

23. The synthetic RNA of any one of claims **20-22**, further comprising an unpaired overhang sequence at the 5' and/or 3' end.

24. The synthetic RNA of claim 23, wherein the unpaired overhang sequence comprises a sequence of repeating bases.

25. The synthetic RNA of claim **24**, wherein the sequence of repeating bases comprises repeating uracil (U) bases.

26. The synthetic RNA of claim **25**, wherein the unpaired overhang sequence is UU.

27. A composition comprising the synthetic RNA of any one of claims 20-26.

28. The composition of claim **27**, further comprising one or more physiologically acceptable carriers and/or one or more physiologically acceptable adjuvants.

* * * * *