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Supplemental Information

Locked Nucleic Acid Gappers and Conjugates

Potently Silence *ADAM33*, an Asthma-Associated

Metalloprotease with Nuclear-Localized mRNA

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Locked Nucleic Acid gapmers and conjugates induce potent silencing of *ADAM33*, an asthma-associated metalloprotease with nuclear-localized mRNA

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Supporting Information:

Supporting Figure S1: Comparison of unmodified and fully modified duplex siRNA

Supporting Figure S2: qRT-PCR time course results, gymnotic delivery

Supporting Tables S1–S5: Sequences and mass spectrometry data

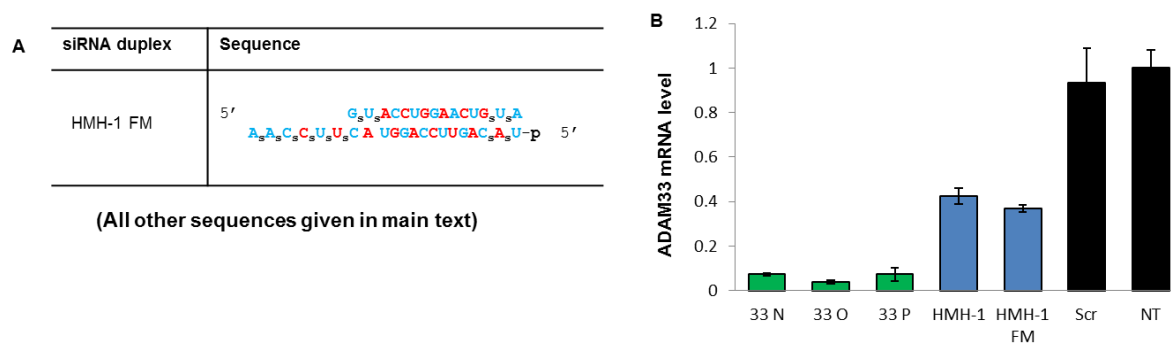


Figure S1. Silencing by a fully chemically modified duplex siRNA. **(A)** Sequence and modification pattern for the fully modified siRNA. Sequence modification codes are 's' (phosphorothioate linkage); red: 2'F-RNA; blue: 2'OMe-RNA; p (5'-phosphate). Passenger strand is listed on top and guide strand underneath. **(B)** qPCR results showing ADAM33 level 3 days after transfection. Oligonucleotides were delivered with Lipofectamine RNAiMAX at 50 nM and normalized to a non-treated control. Error bars represent standard deviation of independent biological duplicates. Green bars are LNA gapmers, Blue bars are native or modified duplex siRNAs, Black bars are negative controls.

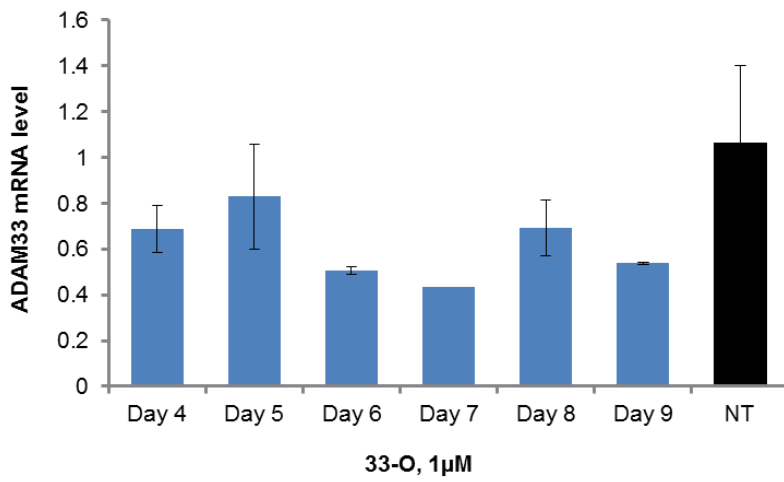


Figure S2. qRT-PCR time course results of gymnotic delivery at 1 μM concentration of ASO **33-O**. Oligonucleotides are delivered at 1 μM dose and normalized to a non-treated control. Error bars are standard deviation of technical replicates. The degree of knockdown is intentionally chosen to be suboptimal (by using 1 μM rather than 3 μM ASO) to allow better discrimination of any differences in efficacy.

Table S1. siRNA sequences designed to target *ADAM33* mRNA, with mass spectrometry data. S represents sense strand, AS represents anti-sense strand.

siRNA	Sequence Uppercase: RNA; Lowercase: DNA For the bottom row only: 2'-F-RNA ; 2'-O-methyl-RNA ; p: 5' phosphate; 's': phosphorothioate	Calculated mass (g/mol)	Found mass (g/mol)
HMH-1	5' GGAAGUACCUGGAACUGUA tt CCUUCAUGGACCUUGACAU 5'	S: 6124.7 AS: 6567.0	S: 6123.8 AS: 6565.8
HMH-2	5' GGUGAGAGGUAGCUCCUAA tt CCACUCUCCAUCGAGGAUU 5'	S: 6140.7 AS: 6566.0	S: 6139.8 AS: 6564.9
HP--1	5' AGAAAGACAUGUUGGCUAU tt UCUUUCUGUACAACCGAUA 5'	S: 6109.7 AS: 6552.0	S: 6108.7 AS: 6550.8
HP- 2	5' GGGAGAUGCUCUUGGAAAC tt CCCUCUACGAGUACCUUUG 5'	S: 6163.8 AS: 6543.0	S: 6162.7 AS: 6542.7
HP-3	5' UGCUUGAGCUGGAGAAGAA tt ACGAACUCGACCUCUUCUU 5'	S: 6164.8 AS: 6527.0	S: 6163.7 AS: 6525.7
HP-4	5' UGGUGAACUCUGCGGGAGA tt ACCACUUGAGACGCCUCU 5'	S: 6156.7 AS: 6565.0	S: 6155.7 AS: 6564.8
HP-5	5' CCCAACACACGGAUCAUU tt tt GGGUUGGUGUGCCUAGUAA 5'	S: 6572.1 AS: 6743.1	S: 6570.9 AS: 6742.9
HP- 6	5' UGGCCCUGUGCAGGCUGAA tt tt ACCGGGACACGUCCGACUU 5'	S: 6701.1 AS: 6644.1	S: 6701.0 AS: 6643.0
HP- 7	5' AGUCCAGAUGCCAAGAUGC tt tt UCAGGUCUACGGUUCUAGG 5'	S: 6652.1 AS: 6663.0	S: 6652.0 AS: 6662.9
HP- 8	5' CCAGACGUUUACCUACUGG tt tt GGUCUGCAAUUGGAUGACC 5'	S: 6606.0 AS: 6709.1	S: 6605.9 AS: 6708.0
HP- 9	5' AGGGCGCCACAGUGGGCCU tt tt UCCCGCGGUGUCACCCGGA 5'	S: 6739.1 AS: 6636.1	S: 6739.0 AS: 6636.0
HP-10	5' GAUCAAGUCCAGAUGCCAA tt tt CUAGUUCAGGUCUACGGUU 5'	S: 6676.1 AS: 6624.0	S: 6675.0 AS: 6623.8
HP- 11	5' UAGCAACCAUAACUGCCAC tt tt AUCGUUGGUAUUGACGGUG 5'	S: 6596.1 AS: 6704.0	S: 6594.9 AS: 6703.9
Scr	5' AGUGGAGGGCGCCUGCCAC tt tt UCACCUCCCGCGGACGGUG 5'	S: 6739.1 AS: 6636.1	S: 6739.05 AS: 6634.99
HMH-1 FM	5' G_s U_s ACCUGGAACUG_s U_s A_s A_s A_s C_s C_s U_s U_s C_s A_s UGGACCUUGAC_s A_s U_s-p 5'	S: 4977.3 AS: 7010.7	S: 4977.7 AS: 7009.8

Table S2. ss-siRNA sequences designed to target *ADAM33* mRNA, with mass spectrometry data.

ss-siRNA	Sequence (5'-3') ' _s ': phosphorothioate; LNA; 2'-F-RNA; 2'-O-methyl-RNA; MOE; P: 5' phosphate	Calculated mass (g/mol)	Found mass (g/mol)
ssi-HMH-1a	P-U _s ACAGUCCAGG _s U _s A _s C _s U _s U _s C _s C _s A _s A	7131.0	7131.6
ssi-HP-2	P-G _s UUUCCAUGAGC _s A _s U _s C _s U _s C _s C _s C _s A _s A	7107.0	7106.6
ssi-HP-3	P-U _s UCUUCUCCAGC _s U _s C _s A _s A _s G _s C _s A _s A _s A	7091.0	7090.6
ssi-HMH-1b	P-U _s A _s CA _s GU _s UC _s CA _s GG _s UA _s C _s U _s U _s C _s C _s A _s A	7196.0	7195.7
ssi-HMH-1c	P-U _s A _s CA _s GU _s UC _s CA _s GG _s UA _s C _s U _s U _s C _s C _s A _s A	7108.1	7107.7
ssi-HMH-1d	P-U _s A _s CA _s GU _s UC _s CA _s GG _s UA _s C _s U _s U _s C _s C _s A _s A	7104.1	7103.6

Table S3. LNA gapmer ASOs designed to target h*ADAM33* mRNA, and LNA controls, with mass spectrometry data.

LNA gapmer	Sequence (5'-3') -' _s ': phosphorothioate; LNA; Lowercase: DNA; HOP: Hexadecyloxypropyl	Calculated mass (g/mol)	Found mass (g/mol)
33-G	T _s G _s A _s t _s C _s C _s g _s t _s g _s t _s g _s g _s T _s T _s G	5023.0	5023.1
33-H	A _s T _s G _s a _s t _s C _s C _s g _s t _s g _s t _s g _s G _s T _s T	5007.0	5007.1
33-I	A _s A _s T _s g _s a _s t _s C _s C _s g _s t _s g _s t _s G _s GG _s T	5016.0	5016.1
33-J	C _s A _s A _s t _s g _s a _s t _s C _s C _s g _s t _s g _s T _s GG _s G	5015.0	5015.0
33-L	T _s G _s T _s C _s a _s t _s g _s g _s t _s t _s t _s t _s C _s A _s G	4996.0	4996.0
33-M	G _s G _s T _s g _s t _s C _s a _s t _s g _s g _s t _s T _s T _s C	5012.0	5012.1
33-N	A _s G _s G _s t _s g _s t _s C _s a _s t _s g _s g _s t _s T _s T _s T	5022.0	5022.0
33-O	T _s C _s A _s t _s t _s t _s t _s a _s g _s g _s a _s g _s C _s T _s A	5003.0	5003.1
33-P	T _s T _s C _s a _s t _s t _s t _s t _s a _s g _s g _s a _s G _s C _s T	4994.0	4994.1
33-Q	T _s G _s T _s t _s C _s a _s t _s t _s t _s a _s g _s G _s A _s G	5006.0	5006.0
33-R	T _s C _s C _s g _s t _s g _s g _s a _s a _s t _s t _s G _s C _s A	5027.0	5027.1
Ina ctrl	A _s T _s T _s t _s t _s a _s t _s t _s C _s g _s g _s a _s G _s C _s T	4980.0	4980.4
Ina scr2	A _s A _s C _s a _s ^m C _s g _s t _s ^m C _s t _s a _s t _s a _s C _s G _s C _s (^m C is 5-methylcytosine)	4984.1	4983.5

Table S4. Biostable and biocleavable LNA gapmer ASOs designed to target hADAM33 mRNA, with mass spectrometry data.

Conjugated LNA gapmer	Sequence (5'-3') ‘ _s ’: phosphorothioate; LNA; Lowercase: DNA; HOP: Hexadecyloxypropyl	Calculated mass (g/mol)	Found mass (g/mol)
33-N biostable	HOP _s A _s G _s G _s t _s g _s t _s c _s a _s g _s g _s t _s T _s T _s T	5400.2	5399.7
33-O biostable	HOP _s T _s C _s A _s t _s t _s t _s t _s a _s g _s g _s a _s g _s C _s T _s A	5364.2	5364.8
33-P biostable	HOP _s T _s T _s C _s a _s t _s t _s t _s t _s a _s g _s g _s a _s G _s C _s T	5355.2	5355.8
33-N biocleavable	HOP t t t t t t A _s G _s G _s t _s g _s t _s c _s a _s g _s g _s t _s T _s T _s T	7209.4	7208.8
33-O biocleavable	HOP t t t t t t T _s C _s A _s t _s t _s t _s t _s a _s g _s g _s a _s g _s C _s T _s A	7190.5	7189.3
33-P biocleavable	HOP t t t t t t T _s T _s C _s a _s t _s t _s t _s t _s a _s g _s g _s a _s G _s C _s T	7181.5	7181.0

Table S5. LNA gapmer ASOs designed to target mAdam33 mRNA, with mass spectrometry data.

LNA gapmer	Sequence (5'-3') ‘ _s ’: phosphorothioate; LNA; Lowercase: DNA	Calculated mass (g/mol)	Found mass (g/mol)
m33-G	(same as human 33-G sequence, Table S3)	5023.0	5022.1
m33-N	A _s G _s G _s c _s a _s t _s c _s t _s c _s g _s g _s t _s T _s T _s G	4991.9	4991.4
m33-P	T _s A _s A _s g _s c _s t _s c _s a _s g _s a _s g _s t _s T _s C _s G	4999.0	4998.4
m33-Q	G _s G _s T _s a _s a _s g _s c _s t _s c _s a _s g _s a _s G _s T _s T	5025.0	5024.4
m33-R	T _s C _s T _s a _s t _s g _s a _s c _s a _s a _s c _s a _s G _s C _s T	4957.3	4956.5