

Full-length article

Design and screening of antisense oligodeoxynucleotides against PAI-1 mRNA in endothelial cells *in vitro*¹

Qi-sheng JIANG, Sheng-qi WANG²

Beijing Institute of Radiation Medicine, Beijing 100850, China

Key words

plasminogen activator inhibitor 1; antisense oligodeoxynucleotide; endothelial cells

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² Correspondence to Prof Sheng-qi WANG. Phn/Fax 86-10-6693-2211. E-mail sqwang@nic.bmi.ac.cn

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Abstract

Aim: To design and screen antisense oligodeoxynucleotides (ASODNs), which inhibit type-1 plasminogen activator inhibitor (PAI-1) expression in human umbilical vein endothelial cells (HUVEC) *in vitro*. **Methods:** Twenty seven ASODNs against different sites of PAI-1 mRNA were designed and transfected to HUVEC by lipofectin *in vitro*. The effects of ASODNs on PAI-1 antigen, PAI-1 activity and PAI-1 mRNA expression were detected by ELISA, amidolytical assay and RT-PCR, respectively. **Results:** Transforming growth factor β_1 (TGF- β_1)-treated HUVEC increased the expression of PAI-1 compared with the normal HUVEC. Five among twenty seven designed ASODNs were effective in inhibiting the increase in PAI-1 antigen and PAI-1 activity in a dose-dependent manner after 48-h transfection. In particular, ASODN 14 (AO14) exhibited the best inhibitory effect. The control sequences of AO14, including sense, scramble, and mismatch sequences, did not significantly inhibit PAI-1 activity. It was revealed that the inhibitory efficacy of AO14 was in a sequence-specific manner. RT-PCR showed that ASODN 1, 7, 8, 14, and 15 decreased PAI-1 mRNA expression induced by TGF- β_1 and AO14 showed the best inhibitory effect. **Conclusion:** ASODN 1, 7, 8, 14, and 15, among twenty seven designed ASODNs against PAI-1 mRNA, significantly decreased PAI-1 antigen and PAI-1 activity induced by TGF- β_1 in a dose-dependent manner in HUVEC *in vitro*. AO14 showed the best inhibitory effect on PAI-1 expression in a sequence-specific manner. The results of RT-PCR indicated that inhibitory effects of ASODNs on PAI-1 biosynthesis occurred at the mRNA level. Four among five effective target sites of ASODNs located at the translation initiation site or within the translation area of PAI-1 mRNA, suggesting that these sites may be promising sites for the design of effective ASODNs.

Introduction

As more and more gene sequences, mostly of unknown biological function, become available through genome sequencing efforts, antisense oligonucleotides (ASODNs) are widely used for the elucidation of gene and protein function and as therapeutic agents in clinical trials^[1,2]. But not all ASODN are equally effective in their ability to inhibit gene expression and protein synthesis^[3,4]. To date, the screening of multiple sequences has been the most familiar way to identify effective antisense sequences.

Type-1 plasminogen activator inhibitor (PAI-1) is the

physiological inhibitor of both tissue-type (t-PA) and urokinase-type plasminogen activator and plays an important role in the process of fibrinolysis and thrombus formation. PAI-1 is expressed in vascular endothelial cells and smooth muscle cells. Overexpression of PAI-1 is strongly associated with life-threatening thrombotic diseases^[5–8] in atherosclerosis, myocardial infarction, deep-vein thrombosis and gram negative sepsis.

The aim of this study is to design some ASODNs that hybridize to various target sites of PAI-1 mRNA and evaluate their inhibitory effect on PAI-1 expression in cultured human umbilical vein endothelial cells (HUVEC), which will

then provide a basis for further screening of thrombolytic ASODN.

Materials and methods

ASODNs design and modification PAI-1 mRNA reported by Ginsburg *et al* (Genebank accession: M16006)^[9] was used as a target sequence for design of ASODNs. The secondary structure of PAI-1 mRNA was simulated by software of RNA-structure 3.6^[10]. Twenty seven antisense sequences targeted to different sites of PAI-1 mRNA were designed (Table 1), including two against the initiation site, twelve against the translation area and thirteen against the 3' side. The ASODNs were synthesized by a DNA synthesizer (ABI 3900, Weiterstadt, Germany) and modified by phosphoramidite solid-phase approach^[11]. The sulphurization step was performed by means of *bis*(*O,O*-disopropoxy phosphinothioyl) disulfide. After standard cleavage from the support, the protection oligonucleotide was subjected to double reverse phase HPLC purification, followed by Na⁺-ion exchange.

Their purity was routinely controlled by polyacrylamide gel electrophoresis (PAGE) and was not lower than 95%.

Cell culture and ASODNs delivery HUVEC was kindly provided by Dr Xiao NAN (Surgery Institute of Third Military Medical University, Chongqing, China). HUVEC were grown to confluence on 48 wells of fibronectin-coated culture plates with DMEM medium supplemented with 10% fetal calf serum (Hyclone, Logan, UT, USA), 100 U/mL penicillin and 100 µg/mL streptomycin for 48 h before transfection of ASODNs. The media was replaced with serum-free media containing 10 µg/mL lipofectin (Invitrogen, Carlsbad, CA, USA) and various concentrations of ASODNs (0.0, 0.1, 0.25, 0.5, and 1.0 µmol/L), and were incubated for 6 h at 37 °C. Cells were stimulated by transforming growth factor β₁ (TGF-β₁, Roche Diagnostics, Mannheim, Germany) with a concentration of 0.5 ng/mL (for increasing PAI-1 concentration in media, as normal concentration is extremely low) for 48 h before collecting media. The control group was treated only with TGF-β₁ 0.5 ng/mL without the use of ASODNs. The HUVEC were used for total RNA extraction and the media were stored

Table 1. Sequences of twenty seven ASODNs targeted to PAI-1 mRNA.

ASODN No	Target site	nt	Sequence (5'-3')
1 (AO1)	67-86	20	GAC ATC TGC ATC CTG AAG TT
2 (AO2)	70-89	20	GGA GAC ATC TGC ATC CTG AA
3 (AO3)	147-166	20	CGT AGG ATG GGG GAT GGT GC
4 (AO4)	199-218	20	TGC GCC ACCTGC TGA AAC AC
5 (AO5)	320-339	20	CTT GAA TCC CAT AGC TGC TT
6 (AO6)	363-382	20	TGT ACA GAT GCC GGA GGG CG
7 (AO7)	489-508	20	TGA CCG TGC TCC GGAACA GC
8 (AO8)	559-578	20	TTT GTG TGT GTC TTC ACC CA
9 (AO9)	641-660	20	GTT GAA GTA GAG GGC ATT CA
10 (AO10)	756-775	20	TAT AGT TFA ACT TGT TGG TC
11 (AO11)	862-881	20	GGC ACC TCT TTT TCA TAA GG
12 (AO12)	1113-1132	20	CGT TCA CCT CGA TCT TCA CT
13 (AO13)	1188-1207	20	CCA TGA TGA TCT CCT CGG GG
14 (AO14)	1246-1265	20	CCC ATG AAA AGG ACT GTT CC
15 (AO15)	1297-1316	20	TTT GTC CCA GAT GAA GGC GT
16 (AO16)	1275-1294	20	TTC CCC AGG GTC AGG GTT CC
17 (AO17)	1538-1557	20	AGGCGT CAC CGT CTG GTT TG
18 (AO18)	1789-1808	20	TGC CAC AGT GGA CTC TGA GA
19 (AO19)	1886-1905	20	GGG GAG GGA GAT GGC CAG GC
20 (AO20)	2059-2078	20	ATA TGA TAA ATA TTT AGG TA
21 (AO21)	2176-2195	20	GGC TGG ACT TCC TGA GAT AC
22 (AO22)	2450-2469	20	AAA GTT CTG TCC TGG TAG GT
23 (AO23)	2483-2502	20	CCA ATG CGG CTG TGA GTC AC
24 (AO24)	2595-2614	20	AAA GAT TAT CTA AGG TAG TT
25 (AO25)	1084-1103	20	CCC CTC CCC ACC AAG AGA TT
26 (AO26)	53-72	20	GAA GTT CAC AGA GGT GCC TT
27 (AO27)	57-76	20	TCC TGA AGT TCT CAG AGG TA

at -70 °C until assay for PAI-1 antigen and activity.

PAI-1 antigen assay PAI-1 antigen in conditioned medium was determined by specific enzyme-linked immunosorbent assay (ELISA)^[12] kits (Diagnostic Stago, American Bioproducts, Parsippany, New Jersey, USA). In brief, 50 μ L of media was incubated with PAI-1 monoclonal antibodies that precoated the flat bottom of 96-well cell culture plates (Corning Laboratories, Corning, NY, USA) for 1 h at 37 °C. Peroxidase conjugated anti-PAI-1 antibody 0.1 mL was added to all wells and incubated for 1 h at 37 °C. Tetramethylbenzidine (TMB) substrate 0.1 mL was added to all wells and incubated for 10 min at room temperature. Stop solution 0.1 mL was added to each well. The PAI-1 Ag in the media was measured by reading the absorbance at 450 nm and calculated against standard regression line.

PAI-1 activity assay PAI-1 activity was measured by amidolytical assay^[13]. One unit PAI-1 activity was defined as the amount of PAI that inhibited one international unit of human single-chain t-PA.

PAI-1 mRNA expression by RT-PCR The primers were designed by computer assistance according to the gene bank. PAI-1: forward, 5'-CGGAGCACGGTCAAGCAAGTG-3'; reverse, 5'-GTTGAGGGCAGAGAGAGGCGC-3', the size of amplified fragment is 401 bp. Internal control GAPDH: forward, 5'-CCATGGAGAAGGCTGGGG-3'; reverse, 5'-CAAAGTTGTCA-TGGATGACC-3'; the size of amplified fragment is 195 bp. Total RNA from each sample was isolated using TRIzol solution (Invitrogen). Total RNA was quantified with the ratio of absorption values of RNA samples at 260 nm and 280 nm. For each sample, 4 μ g of total RNA was reverse transcribed into the first strand of cDNA in a 20- μ L reaction system at 37 °C for 50 min. Then polymerase chain reaction was performed from the synthesized cDNA in a 50 μ L solution containing 3 μ L of cDNA, 1 μ L of 25 mmol primers (up-stream and down-stream) of PAI-1, 0.5 μ L of 25 mmol primers (up-stream and down-stream) of GAPDH, 10 mmol dNTP 1 μ L, 25 mmol MgCl₂ 4 μ L, 10 \times buffer 5 μ L, 0.3 μ L of Taq DNA

polymerase (Qiagen, Valencia, CA, USA). Amplification was performed in a thermal cycler (Bio-Rad, Alfred Nobel Drive Hercules, CA, USA) under the following conditions: 26 cycles of denaturation at 94 °C for 50 s, annealing at 59 °C for 45 s, extension at 70 °C for 40 s, followed by a final extension for 5 min. PCR product 10 μ L was electrophoresed on a 2% agarose gel, and stained with EB. The PAI-1 mRNA level in each sample was semi-quantified by comparing the intensities of PAI-1 mRNA band with those of the internal control GAPDH band.

Statistics assay Data were shown as mean \pm SD. Statistical analysis was performed by analysis of variance (ANOVA). The level of statistical significance was chosen as $P < 0.05$.

Results

Effects of ASODNs on PAI-1 Ag After 48-h transfection, in comparison to control group (290.0 \pm 57.2 ng/mL), all of the ASODNs did not significantly inhibit PAI-1 Ag at a concentration of 0.1 μ mol/L. ASODN 1, 7, 14, and 15 could significantly decrease PAI-1 Ag at concentrations of 0.25, 0.5 and 1.0 μ mol/L (inhibition rate: 25.9%–78.7%). ASODN 8 could also remarkably inhibit PAI-1 Ag at concentration of 0.5 and 1.0 μ mol/L (inhibition rate: 60.8%, 65.1%, respectively). The inhibitory effect was in a dose-independent manner (Table 2). These data suggested that five among twenty seven designed ASODNs significantly inhibited PAI-1 Ag expression and AO14 exhibited the best inhibitory effect.

Effects of ASODNs on PAI-1 activity After 48-h transfection, compared to control group (15.0 \times 10⁻² \pm 1.5 \times 10⁻² AU/mL), at a concentration of 0.1 μ mol/L, all of the ASODNs did not significantly inhibit PAI-1 activity. At 0.25 μ mol/L, ASODN 1, 14, and 15 significantly reduced PAI-1 activity (inhibition rate: 41.3%, 40.0%, 33.3%, respectively). At 0.5 and 1.0 μ mol/L, ASODN 1, 7, 8, 14, and 15 remarkably decreased PAI-1 activity (inhibition rate: 40.0%–52.7%). These data (Table 3) suggested that five ASODNs that

Table 2. Effects of ASODNs on PAI-1 antigen (ng/mL) in HUVEC after 48-h transfection *in vitro*. $n=3$. Mean \pm SD. ^b $P < 0.05$, ^c $P < 0.01$ vs control.

ASODNs	Concentration of ASODNs/ μ mol \cdot L ⁻¹				
	0 (C)	0.1	0.25	0.5	1.0
AO1	290.0 \pm 57.2	302.7 \pm 34.5	150.6 \pm 18.8 ^b	109.3 \pm 14.0 ^b	91.3 \pm 11.6 ^c
AO7	290.0 \pm 57.2	343.6 \pm 33.0	215.0 \pm 29.1 ^b	131.0 \pm 27.2 ^b	103.0 \pm 22.0 ^c
AO8	290.0 \pm 57.2	316.3 \pm 29.6	218.0 \pm 29.7	113.6 \pm 11.6 ^b	101.3 \pm 16.8 ^c
AO14	290.0 \pm 57.2	330.3 \pm 34.7	137.5 \pm 18.2 ^b	91.0 \pm 18.6 ^c	61.7 \pm 14.4 ^c
AO15	290.0 \pm 57.2	292.3 \pm 18.2	134.7 \pm 21.2 ^b	120.0 \pm 21.6 ^b	104.3 \pm 23.7 ^c

Table 3. Effects of ASODNs on PAI-1 activity ($\times 10^{-2} \text{AU}\cdot\text{mL}^{-1}$) in HUVEC after 48-h transfection *in vitro*. $n=3$. Mean \pm SD. ^b $P<0.05$ vs control, ^c $P<0.01$ vs control.

ASODNs	Concentration of ASODNs/ $\mu\text{mol}\cdot\text{L}^{-1}$				
	0 (C)	0.1	0.25	0.5	1.0
AO1	15.0 \pm 1.5	12.0 \pm 2.6	8.8 \pm 1.5 ^b	9.0 \pm 1.0 ^b	9.0 \pm 1.2 ^c
AO7	15.0 \pm 1.5	13.0 \pm 2.5	10.0 \pm 1.7	9.0 \pm 1.1 ^b	8.1 \pm 1.1 ^c
AO8	15.0 \pm 1.5	13.0 \pm 2.0	11.1 \pm 3.1	9.0 \pm 1.1 ^b	9.0 \pm 0.6 ^c
AO14	15.0 \pm 1.5	11.1 \pm 0.6	9.0 \pm 2.1 ^b	8.0 \pm 1.7 ^c	7.1 \pm 0.6 ^c
AO15	15.0 \pm 1.5	12.1 \pm 2.5	10.0 \pm 1.1 ^b	8.1 \pm 1.2 ^c	7.8 \pm 0.6 ^c

inhibited PAI-1 Ag could also inhibit PAI-1 activity and AO14 showed the best inhibitory effect.

Effects of ASODNs on expression of PAI-1 mRNA After 48-h transfection, at a concentration of 1.0 $\mu\text{mol/L}$, ASODN 1, 7, 8, 14, and 15 significantly decreased PAI-1 mRNA expression induced by TGF- β_1 . The inhibition rates were 25.3%, 43.3%, 40.0%, 86.3%, 61.4%, respectively, when compared with the control (Figure 1).

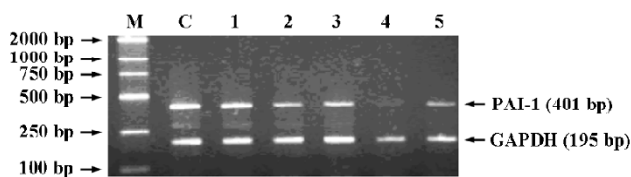


Figure 1. PAI-1 expression in HUVEC after transfection of 1.0 $\mu\text{mol/L}$ ASODNs. M and C are representative of marker DL2000 and control, respectively. Lane 1: ASODN 1; Lane 2: ASODN 7; Lane 3: ASODN 8; Lane 4: ASODN 14; Lane 5: ASODN 15. The expression of PAI-1 mRNA was significantly decreased in five ASODN groups after transfection for 48 h and AO14 showed the best inhibitory effect compared with control.

Specific inhibitory effects of ASODNs on PAI-1 activity

Positive results of decrease PAI-1 expression by five ASODNs in cultured endothelial cells prompted us to investigate further the specificity of action of the ASODNs.

Because AO14, among twenty seven designed ASODNs, was confirmed to have the best efficiency in inhibiting PAI-1 mRNA, PAI-1 Ag and PAI-1 activity, we designed and synthesized corresponding control oligonucleotides of AO14, including sense (AO28), scrambled (AO29), and an oligonucleotide with two mismatches (AO30). Their nucleotide sequences are listed in Table 4. Transfection (1.0 $\mu\text{mol/L}$) of these control oligonucleotides to HUVEC by lipofectin was under the same experimental condition as AO14. The results showed that sense, scrambled, and mismatched oligonucleotides of AO14 did not have an inhibitive effect on PAI-1 activity after 48-h of transfection (Table 4). This suggests that the inhibitory effect of AO14 on PAI-1 activity was indeed of specific antisense origin.

Discussion

Antisense strategy is a novel approach for inhibiting target gene expression by ASODNs complementary to preselected regions of mRNA by Watson-Crick base pairing. So ASODNs have been proposed as potential therapeutic agents for inhibiting candidate genes that may account for specific diseases^[1,2]. But not every site of mRNA sequence is accessible to hybridization with ASODNs. A major obstacle in employing ASODN is the election of target sites within nucleotide sequences for effective inhibition of ex-

Table 4. Effects of AO14 and its control oligonucleotides (1.0 $\mu\text{mol/L}$) on PAI-1 activity ($\times 10^{-2} \text{AU}\cdot\text{mL}^{-1}$) in HUVEC after 48-h transfection *in vitro*. $n=3$. Mean \pm SD. ^c $P<0.01$ vs control.

Oligonucleotides	Description	Nucleotide Sequence (5'-3')	0 h	48 h
AO14	ASODN	CCC ATG AAA AGG ACT GTT CC	15.0 \pm 1.5	7.1 \pm 0.6 ^c
AO28	Sense	GGA ACA GTC CTT TTC ATG GG	15.0 \pm 1.5	12.6 \pm 2.2
AO29	Scrambled	CCC ATG AGA A AT G AC GTT CC	15.0 \pm 1.5	14.1 \pm 2.9
AO30	Two mismatches	CCC ATG ATA AGG AGT GTT CC	15.0 \pm 1.5	13.8 \pm 1.6

pression^[14]. Several reports suggest that the most effective target sites are either the 5'-nontranslated sequences or the ATG start site for translation^[15]. In the present study, we designed twenty seven ASODNs (20 nt) that are complementary to PAI-1 mRNA local sites, including the initiation site, translation field and 3' side, and transfected them respectively into cultured HUVEC by lipofectin. The results showed that five among twenty seven designed ASODNs significantly decreased PAI-1 mRNA, PAI-1 antigen and PAI-1 activity. The effective target sites were located as follows: 1 (total 2) at initiation site, 3 (total 12) at translation field and 1 (total 13) at 3' side. The results showed that the efficiency of ASODNs against the initiation site (1/2) and translation field (3/12) were higher than 3' side (1/13) and 5' side (0/2), which showed that not only the initiation site but also the translation field were promising sites for ASODNs design (because the 5' side of PAI-1 mRNA is too short, we designed only two ASODNs there).

PAI-1 is one of the most important factors in the pathogenesis of thrombosis^[5-8]. Overexpression of PAI-1 is associated with thrombotic disease in atherosclerosis, myocardial infarction, and gram negative sepsis. A similar effect was also observed in transgenic mice, indicating that individuals with elevated concentrations of the inhibitor in their blood tend to be at risk of developing thrombotic problems. However, some researchers have shown that neutralization of plasma PAI-1 by monoclonal antibody^[16], small molecular weight inhibitors^[7,17] or synthetic peptides^[18] would remarkably increase fibrinolysis and protect against thrombus formation. In the present study, twenty seven ASODNs against PAI-1 mRNA were designed and transfected to HUVEC *in vitro*. The results showed that ASODN 1, 7, 8, 14, and 15 were effective in inhibiting the increase in PAI-1 antigen and PAI-1 activity, which was induced by TGF- β_1 after 48 h of transfection, and AO14 showed the best inhibitory effect. In addition to determination of the protein level under suppression of the translation process by antisense constructs, RT-PCR of mRNA for PAI-1 was performed for mRNAs isolated from HUVEC untreated (control) and treated with ASODN 1, 7, 8, 14, and 15. Only cells treated with ASODN 1, 7, 8, 14, and 15 showed a decrease in PAI-1 mRNA. These results indicate that inhibition of PAI-1 biosynthesis occurs at the mRNA level. Among five effective ASODNs, AO14 was found to be the most efficient inhibitor of PAI-1 synthesis in cultured HUVEC. For further determination of the antisense mechanism of sequence-specific ASODN, the control sequences of AO14, including sense, scramble, and mismatch sequences (Table 4), were tested under the same experimental conditions. The results indicated that the con-

trol sequences of AO14 did not significantly inhibit PAI-1 activity. This shows that the inhibitory efficacy of AO14 was not only in a dose-dependent manner but also in a sequence-specific manner in HUVEC *in vitro*.

The remarkable advantage of antisense strategy in specificity of action suggests that the prevention and treatment of thrombotic disease by ASODNs may be a potential method that is worthy of further exploration.

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