



Mesyl phosphoramidate backbone modified antisense oligonucleotides targeting miR-21 with enhanced in vivo therapeutic potency

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Contributed by Sidney Altman, October 16, 2020 (sent for review August 3, 2020; reviewed by Sebastien Lyonnais and Asako Yamayoshi)

The design of modified oligonucleotides that combine in one molecule several therapeutically beneficial properties still poses a major challenge. Recently a new type of modified mesyl phosphoramidate (or μ -) oligonucleotide was described that demonstrates high affinity to RNA, exceptional nuclease resistance, efficient recruitment of RNase H, and potent inhibition of key carcinogenesis processes in vitro. Herein, using a xenograft mouse tumor model, it was demonstrated that microRNA miR-21-targeted μ -oligonucleotides administered in complex with folate-containing liposomes dramatically inhibit primary tumor growth via long-term down-regulation of miR-21 in tumors and increase in biosynthesis of miR-21-regulated tumor suppressor proteins. This antitumoral effect is superior to the effect of the corresponding phosphorothioate. Peritumoral administration of μ -oligonucleotide results in its rapid distribution and efficient accumulation in the tumor. Blood biochemistry and morphometric studies of internal organs revealed no pronounced toxicity of μ -oligonucleotides. This new oligonucleotide class provides a powerful tool for antisense technology.

nucleolar stress and fragmentation, up-regulation of P21, and activation of caspases, leading to large-scale apoptosis (4, 5, 8).

To improve clinical performance of oligonucleotides, attempts are made to design hybrid molecules that combine in one structure several modifications. However, the preparation of mixmers complicates synthesis and increases the cost of such compounds. Design of new modified oligonucleotides that would combine in one oligomer the maximum number of therapeutic capabilities remains as relevant as ever.

Oligodeoxynucleotides containing an aromatic sulfonyl phosphoramidate group were first obtained over a decade ago but never studied as potential antisense compounds (9). Recently, we synthesized oligodeoxynucleotides with the tosyl phosphoramidate group and showed that their duplex with RNA was destabilized relative to the native DNA:RNA duplex, presumably due to the bulkiness of the tosyl group at the internucleotidic position (10). This has prompted us to investigate a novel DNA analog that

antisense oligonucleotide | DNA modification | mesyl oligonucleotide | oncogenic microRNA | phosphorothioate

Significance

Sequence-specific targeting of disease-associated RNAs by new types of chemically modified antisense oligonucleotides with improved physicochemical and biological properties is still in high demand. miRNA-addressing mesyl phosphoramidate oligonucleotides represent prospective therapeutics since they reveal good bioavailability in tumor tissue, potent and long-lasting antitumor effect, high targeted specificity associated with reactivation of tumor suppressor proteins, and physiological safety.

Antisense-based RNA targeting approaches laid the foundation for one of the powerful drug discovery platforms (1). Antisense oligonucleotides (ASOs) are short synthetic nucleic acids designed to inhibit the expression of target RNA by sequence-specific recognition.

The first generation modified ASOs are well-known phosphorothioate oligodeoxynucleotides (PS ASOs) (1, 2). Despite decades of efforts, no other modification has been developed combining a number of important properties, such as high nuclease resistance, the ability to recruit RNase H, and enhanced protein binding facilitating ASOs uptake by the cells and accumulation in peripheral tissues (3). The PS modification currently forms the backbone of the main ASOs in clinical trials, combined with other modifications of the second or third generation, such as 2'-O-methoxyethyl (MOE), locked nucleic acid (LNA), or constrained ethyl (cEt), stabilizing duplex interactions (the gapmer oligonucleotides) (2). Despite the number of benefits, one of the significant drawbacks of PS ASOs that limit its clinical utility is substantial toxicity (4–7). It was shown that toxicity of some MOE, LNA, or cEt PS gapmer ASOs arises from enhanced binding to cellular proteins, resulting in RNase H1-dependent nucleolar mislocalization of paraspeckle proteins,

Author contributions: O.A.P., V.V.V., D.A.S., and M.A.Z. designed research; O.A.P., S.K.G.M., A.V.S., I.A.S., D.V.G., and A.A.F. performed research; E.A.B., M.A.M., E.V.S., and D.A.S. contributed new reagents/analytic tools; O.A.P., S.K.G.M., A.V.S., I.A.S., D.V.G., M.A.M., M.J.A.W., S.A., and M.A.Z. analyzed data; O.A.P., S.K.G.M., A.V.S., V.V.V., S.A., D.A.S., and M.A.Z. wrote the paper; M.A.M. synthesized folate containing liposomes; and E.V.S. synthesized folate lipoconjugate and prepared folate containing liposomes.

Reviewers: S.L., CNRS Montpellier France; and A.Y., Nagasaki University.

The authors declare no competing interest.

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This article contains supporting information online at <https://www.pnas.org/lookup/suppl/doi:10.1073/pnas.2016158117/-DCSupplemental>.

First published December 7, 2020.

incorporates the mesyl (methanesulfonyl) phosphoramidate group (μ -modification), which has the smallest one-carbon side chain (Fig. 1). Oligodeoxynucleotides with a few μ -groups replacing the phosphodiester (PO) group indeed showed improved RNA binding over tosyl analogs (11). Next, ASOs substituting μ -modifications for natural PO groups at every internucleotide position were designed to inhibit oncogenic microRNAs (miRNAs) (12), in particular, miR-21, which is recognized as a potent oncogene and causes global dysregulation of gene expression in malignant cells (13–17). μ -oligonucleotides demonstrated high affinity to RNA, exceptional resistance to nucleases, and potent recruitment of RNase H, with increased efficiency of catalytic miRNA degradation as compared to PS ASOs. The biological potential of μ ASOs was shown in cell culture and contributes to the efficient and specific inhibition of key carcinogenic processes in vitro (12). The present investigation focuses on the study of therapeutically important characteristics of the miR-21-targeted μ ASOs as potential antitumor agents in vivo: biodistribution, specific and off-target effects, kinetics of tumor suppression, and assessment of the main criteria of systemic toxicity. Herein, the enhanced in vivo potency of μ ASOs compared with PS ASOs indicates that this oligonucleotide class represents a highly-potent tool of antisense technology.

Results

Structure of Chemically Modified ASOs. In this work, 22-mer oligodeoxyribonucleotides complementary to mature miR-21 (18) and containing μ - or PS-modifications of all internucleotide bonds, termed as μ -miR-21-ON and PS-miR-21-ON, respectively, were studied (for sequences and structures, see Fig. 1 and *Materials and Methods*). As a control, fully modified μ - or PS-scrambled oligodeoxyribonucleotides (μ -Scr-ON and PS-Scr-ON) of the same length, with a sequence that had no homology in the mammalian genome, were used.

Kinetics of miRNA Down-Regulation In Vitro by μ and PS ASOs. To develop the optimal regimen of antisense therapy with μ ASOs in vivo, the kinetics of miRNA silencing by μ ASOs (100 nM) was investigated in melanoma B16 cells that combine the main malignant features—aggressiveness, rapid proliferation, and high metastatic potential. Lipofectamine 2000-mediated transfection of B16 cells with μ -miR-21-ON causes a 6-d-lasting decrease in the miR-21 level, with maximum down-regulation observed at 72 h (*SI Appendix, Fig. S1*). Similar kinetics of miRNA down-regulation was also evidenced for μ ASOs targeted to oncogenic miRNAs miR-155 and miR-17: The lowest level of miRNAs

accounting for an 80 to 90% decrease was reached by 72 h and was preserved up to 144 h (*SI Appendix, Fig. S1*). It should be emphasized that no unspecific down-regulation of other miRNAs, for instance, miR-155 and miR-17, was observed after cell treatment with μ -miR-21-ON (*SI Appendix, Fig. S1*). In turn, PS-miR-21-ON promoted no more than a twofold decrease in the level of miR-21 observed by 48 h, with the total duration of inhibition being less than 72 h (*SI Appendix, Fig. S1*).

Biodistribution of μ and PS ASOs in Tumor-Bearing Mice. To study the biodistribution and therapeutic potential of μ ASOs, a xenograft model of human epidermoid carcinoma KB-8-5 cells in severe combined immunodeficient (SCID) mice was chosen. The use of this model permits one to assess a direct antitumor effect and minimize the interference of the immunity engagement in response to oligonucleotide therapy.

To study biodistribution and conduct an initial analysis of pharmacokinetic parameters of the developed oligonucleotides, KB-8-5 tumor-bearing SCID mice were peritumorally or intravenously (i.v.) injected with 40 μ g of μ - or PS-Cy5.5-modified oligonucleotides. Folate-containing cationic liposomes F shown to efficiently deliver small interfering RNA (siRNA) to KB-8-5 tumors characterized by high abundance of folate receptors were used as a delivery vehicle (19).

Data analysis showed that, after peritumoral administration, both oligonucleotides accumulated mainly in the kidneys and tumor tissue (Fig. 2). The content of μ ASO and PS ASO in the kidneys after 4 h was 71% and 36% of the total amounts of the therapeutics retained in the body (Fig. 2B). Four hours after peritumoral administration, the relative content of oligonucleotides in the tumor was 23% for μ ASO and 63% for PS ASO. By 24 h, the excretion of μ ASO slowed, and its relative content in the tumor increased to 48% (Fig. 2B). The content of PS ASO in tumor tissue by 24 h decreased to 48%, with more active accumulation in the kidneys. As can be seen from the fluorescence imaging of mouse internal organs, most of the PS ASOs remained anchored in the area of administration while the μ -oligonucleotide was distributed throughout the body, providing high accumulation in the tumor tissue (Fig. 2A).

After i.v. administration, the main fraction of both μ and PS ASOs was localized in the kidneys and liver of animals. The content of μ and PS ASOs in the kidneys after 24 h was 90% and 77%, respectively, and 16% of PS ASO accumulated in the liver (Fig. 2A and B). Accumulation of oligonucleotides in tumor tissue after i.v. administration was poor: By 24 h, the relative content of both oligonucleotides did not exceed 2% (Fig. 2B). According to the obtained data, peritumoral injections were used to study the antitumor potential of μ ASOs in vivo.

Confocal microscopy images of tumor node cryosections confirmed the accumulation of μ and PS oligonucleotides in tumor tissue (Fig. 2C and *SI Appendix, Fig. S2*). According to the results of confocal microscopy, both oligonucleotides localize in the intercellular space as well as in the cytoplasm and nucleus of tumor cells 24 h after injection.

Tumor Growth Retardation In Vivo by miR-21-Targeted μ and PS ASOs. The antitumor potential of μ ASOs was studied using a xenograft KB-8-5/SCID model. The miR-21 inhibitory effect of μ ASO in KB-8-5 tumor cells was preliminarily confirmed in vitro by stem-loop PCR (*SI Appendix, Fig. S3*). At day 12, when the primary tumor node was formed, 10 μ g of μ or PS ASOs in complex with cationic liposomes F were peritumorally administered to animals. In total, four injections were given at 3-d intervals (Fig. 3A). This treatment regimen was used based on the data on stability of the oligonucleotides in serum, kinetics of miRNA down-regulation, and biodistribution. Control groups received unloaded liposomes F or Opti-MEM culture medium.

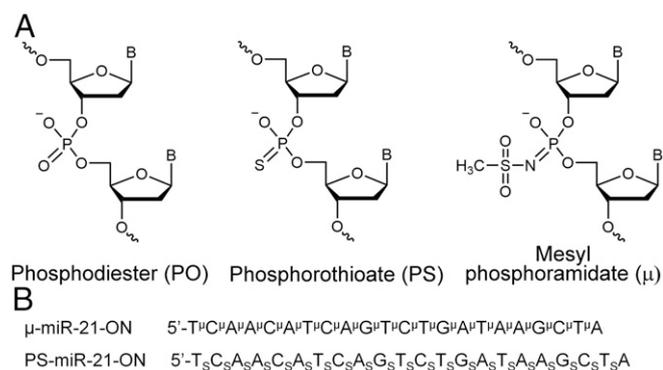


Fig. 1. Structures and sequences of antisense oligonucleotides (ASOs) targeted to miR-21. (A) Structures of DNA phosphate group modifications: phosphodiester (PO), phosphorothioate (PS), and mesyl phosphoramidate (μ). (B) Sequences of oligonucleotides μ -miR-21-ON and PS-miR-21-ON used in the study. Indices μ and S stand for mesyl phosphoramidate group and phosphorothioate group, respectively.

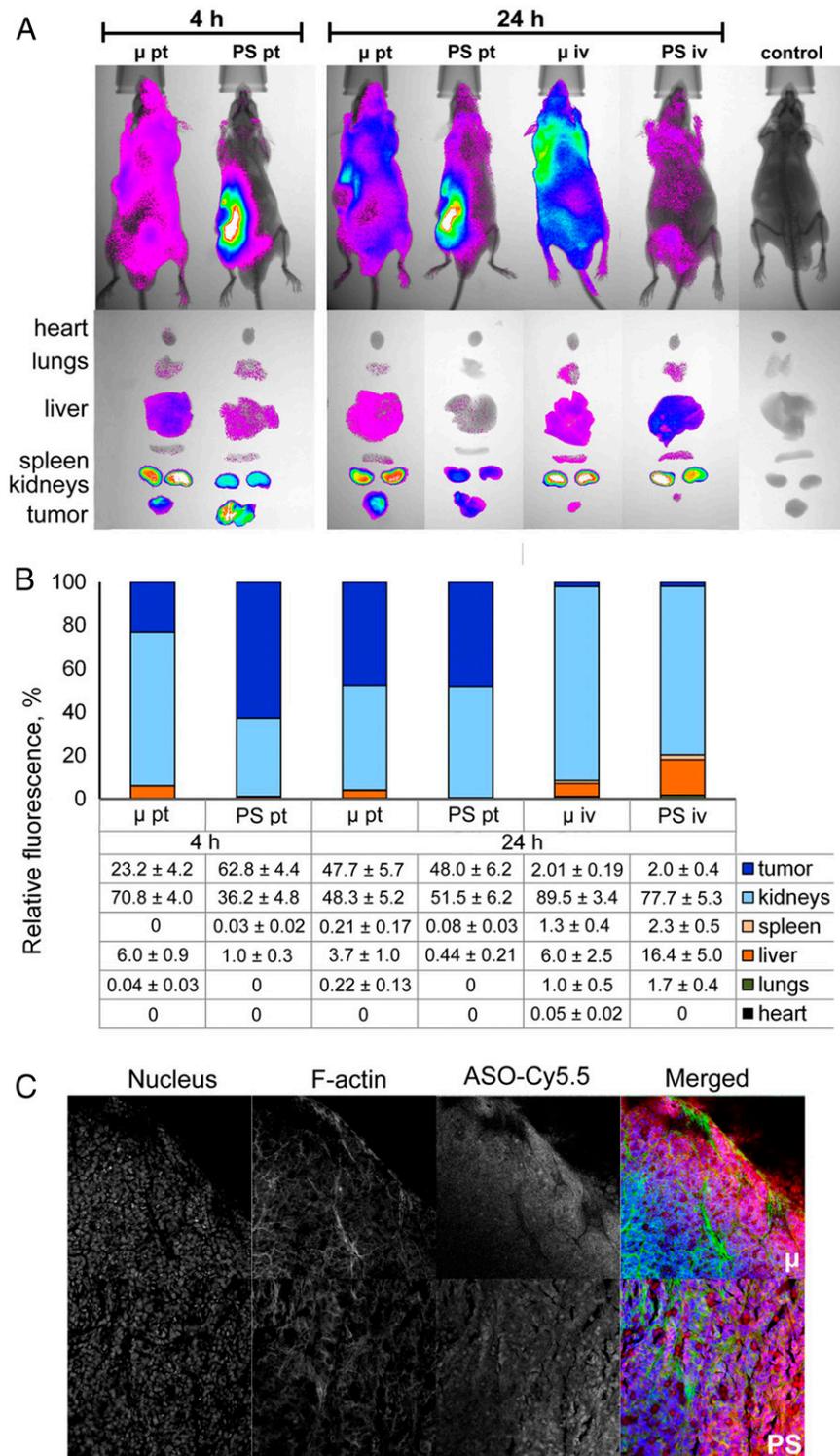


Fig. 2. Biodistribution of Cy5.5-labeled μ and PS ASOs in tumor-bearing SCID mice. (A) Lifetime fluorescence imaging of tumor-bearing SCID mice ($n = 3$ mice per group), dorsal orientation view, and dissected organs 4 and 24 h after peritumoral (pt) and i.v. (iv) injections of 40 μg per mouse of modified oligonucleotides. Control, noninjected mice. (B) Percent biodistribution of modified oligonucleotides after 4 and 24 h after pt and i.v. injections of μ and PS ASOs (mean \pm SE). (C) Confocal microscopic images of cryosections of KB-8-5 tumors from mice 24 h after pt injections of Cy5.5-labeled μ and PS ASOs at a dose of 40 μg per mouse. Three-channel pictures are shown. Merged images represent staining of nucleus with DAPI (blue), staining of actin filaments with phalloidin-TRITC (green), and visualization of oligonucleotides by Cy5.5-labeling (red). Magnification $\times 400$.

Analysis of kinetics of tumor growth revealed that PS-miR-21-ON, despite significant accumulation in tumor tissue, did not promote any statistically reliable retardation of tumor growth,

and its effect was statistically insignificant in comparison with control PS-Scr-ON and unloaded liposomes F (Fig. 3B). In turn, μ -miR-21-ON provided an eightfold reduction of tumor volume

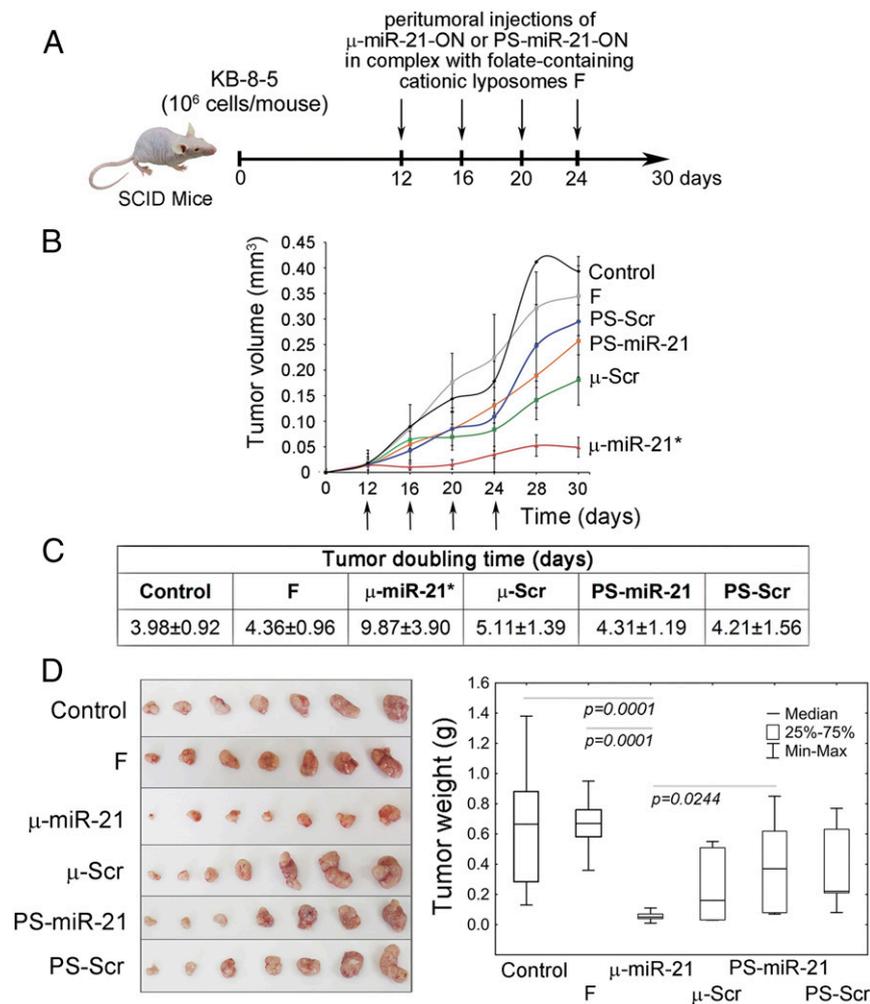


Fig. 3. Antitumor effect of μ -miR-21-ON. (A) Experimental design including engraftment of SCID mice with human epidermoid carcinoma KB-8-5 and peritumoral injections of μ -miR-21-ON, PS-miR-21-ON, or scrambled μ or PS ASOs in complex with folate-containing liposomes F at a dose of 10 μ g per mouse. In total, four injections were made at days 12, 16, 20, and 24 after tumor cell implantation. (B) Kinetics of KB-8-5 tumor growth after treatment with μ -miR-21-ON and PS-miR-21-ON ($n = 7$ mice per group). The days of injections are marked by arrows. An * indicates statistically significant difference of μ -miR-21-ON group from all of the other groups, with $P < 0.05$. (C) Doubling time of KB-8-5 tumors after treatment with oligonucleotides. Statistically reliable difference from all groups is marked by asterisk. (D) Tumor weight at day 30. Data were statistically analyzed using one-way ANOVA with post hoc Tukey test; P value indicates a statistically reliable difference. F, liposomes F.

compared to control, fivefold compared to PS-miR-21-ON, and fourfold compared to μ -Scr-ON (Fig. 3B). Moreover, application of μ -miR-21-ON contributed to a considerable increase in the tumor doubling time that in average was twofold longer than in other groups (Fig. 3C). Comparison of tumor weights showed that the average tumor weight in the μ -miR-21-ON-injected group was up to 12-fold less than in control and about sixfold less compared to μ -Scr-ON (Fig. 3D). In contrast, tumor weight in groups injected with PS ASOs was only twofold less than in the control, with no reliable differences between PS-miR-21-ON and PS-Scr-ON (Fig. 3D).

To confirm that the antitumor effect of μ -miR-21-ON was miR-21-mediated, the levels of miR-21 and its direct protein targets PTEN and PDCD4 in tumor tissue were evaluated at the end of the experiment. μ -miR-21-ON administration caused a 50% decrease in the miR-21 level, compared to control (Fig. 4A), while PS-miR-21-ON administration reduced the level of miR-21 only by 25% (Fig. 4A). It is noteworthy that the observed effect of μ -miR-21-ON was specific: No detectable down-regulation of other miRNAs, such as miR-155 and miR-17 in tumor tissue, was observed (Fig. 4B).

After inhibition of oncogenic miR-21, the activation of the biosynthesis of tumor suppressor proteins, among which PTEN and PDCD4 represent the most well-established targets, might be expected (20, 21). Indeed, the level of these proteins was significantly enhanced after μ -miR-21-ON treatment: 1.5-fold for PDCD4 and 3.5-fold for PTEN, compared to control, while μ -Scr-ON, PS-miR-21-ON, and PS-Scr-ON had no detectable impact on the expression of these proteins (Fig. 4 C and D).

Histological Analysis of KB-8-5 Tumors after Treatment with miR-21-Targeted μ and PS ASOs. Microscopic examination of KB-8-5 carcinomas showed that tumor tissue is represented by polymorphic epidermal cancer cells with acidophilic cytoplasm, a large nucleus with two to three hyperchromic nucleoli, and a high mitotic rate (Fig. 5). Necrotic changes and inflammatory infiltration, represented predominantly by lymphocytes with a small number of neutrophils and macrophages located at the border of unaltered tumor tissue and necrosis, were found in the tumors of all groups (SI Appendix, Table S1). However, in the group treated by μ -miR-21-ON, areas of necrosis and inflammation were much smaller than in other groups (SI Appendix, Table S1).

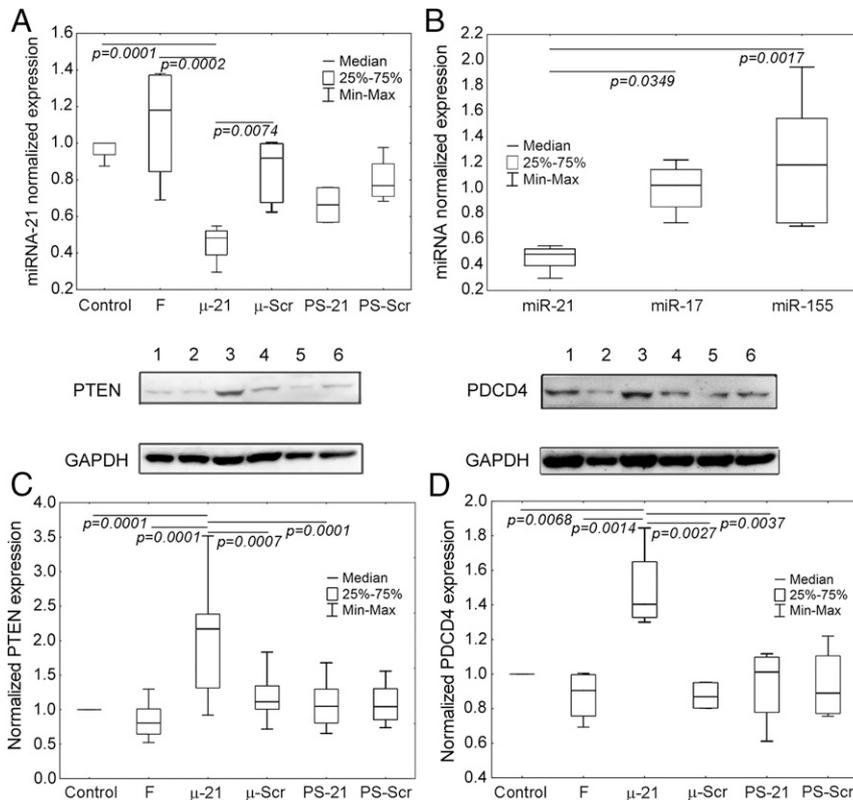


Fig. 4. The level of miRNAs and proteins PTEN and PDCD4 in human epidermoid carcinoma KB-8-5 tissue after administration of μ -miR-21-ON and PS-miR-21-ON. (A) The level of miR-21 in tumor tissue after completion of treatment with μ -miR-21-ON and PS-miR-21-ON. (B) The level of miR-21, miR-155, and miR-17 after treatment with μ -miR-21-ON. Expression of miRNAs was measured by qPCR and normalized to the expression of snRNA U6. (C and D) Expression of tumor suppressor proteins PTEN and PDCD4, respectively, after treatment with μ -miR-21-ON and PS-miR-21-ON measured by Western blot. Protein levels were normalized to the level of GAPDH. Numbers indicate the following groups: 1, control; 2, folate-containing liposomes F; 3, μ -miR-21-ON; 4, μ -Scr-ON; 5, PS-miR-21-ON; 6, Ps-Scr-ON. Data were statistically analyzed using one-way ANOVA with post hoc Tukey test; *P* value indicates a statistically reliable difference.

The smaller size of the primary tumor node in this group and its trophic and blood supply may have significant influence on these parameters.

Histological analysis showed that tumor tissue of μ -miR-21-ON-treated mice was characterized by fourfold reduced mitotic activity, as compared to the control group, and twofold reduced, as compared to groups treated with μ -Scr-ON and PS-miR-21-ON (Fig. 5 A and B). Morphometric analysis of caspase-3 immunohistochemical images demonstrated that nontreated KB-8-5 tumor tissue had a small number of apoptotic cells (Fig. 5 C and D). μ -miR-21-ON treatment led to a 12.8-fold increase in the number of caspase-3 positive cells as compared with the control group ($P = 0.0001$) (Fig. 5D). Administration of oligonucleotides, regardless of target specificity and modification type, led to an increase in the number of apoptotic cells in tumor tissue, but the average number of apoptotic cells in the μ -miR-21-ON-treated group was 1.5-fold higher compared to groups of mice injected with other oligonucleotides. However, due to the increase in the scatter of experimental data, statistically significant differences between the number of caspase-3 positive cells were observed only after the treatment with μ -miR-21-ON and PS-Scr ($P = 0.0005$) (Fig. 5D).

Toxicity of Oligonucleotides. Biochemical blood tests showed that μ ASOs exhibited no toxic effect on the liver and kidneys of mice. No statistically significant differences in blood indicators, such as ALT, ALK, total protein, creatinine, and blood urea nitrogen

(BUN), between the groups (including healthy mice) were found (SI Appendix, Table S2).

The development of a KB-8-5 tumor, per se, has a distinguishable toxic effect on the liver, manifested in an increase in destructive changes (dystrophy plus necrosis) up to 35% from the entire liver parenchyma (SI Appendix, Fig. S4 and Table S3). Administration of ASOs caused no additional detrimental effect on the liver. Moreover, in the groups treated with μ -miR-21-ON or PS-miR-21-ON, a significantly reduced percentage of destructive changes in the liver parenchyma was observed, which reflects a therapeutic effect. Tumor development as well as applied therapy did not affect the regenerative properties of the liver: No changes in numerical density of binuclear hepatocytes were observed (SI Appendix, Table S3).

Morphometric analysis of kidneys showed that, upon tumor development, the normal kidney tissue reduced to about 75% (SI Appendix, Table S4). In the group treated with PS ASOs, some increase in destructive changes in the kidney tissue was observed: 25 to 27% (compared to 23 to 24% in other experimental groups), manifested in dystrophy of the epithelium of the proximal tubules (SI Appendix, Fig. S4 and Table S4). Despite the predominant localization of both μ and PS ASOs in the kidney, signs of dystrophy were most pronounced only in the groups injected with PS ASOs.

Discussion

Over the history of the development of nucleic acid therapeutics, more than two dozen modifications of internucleotidic bonds in

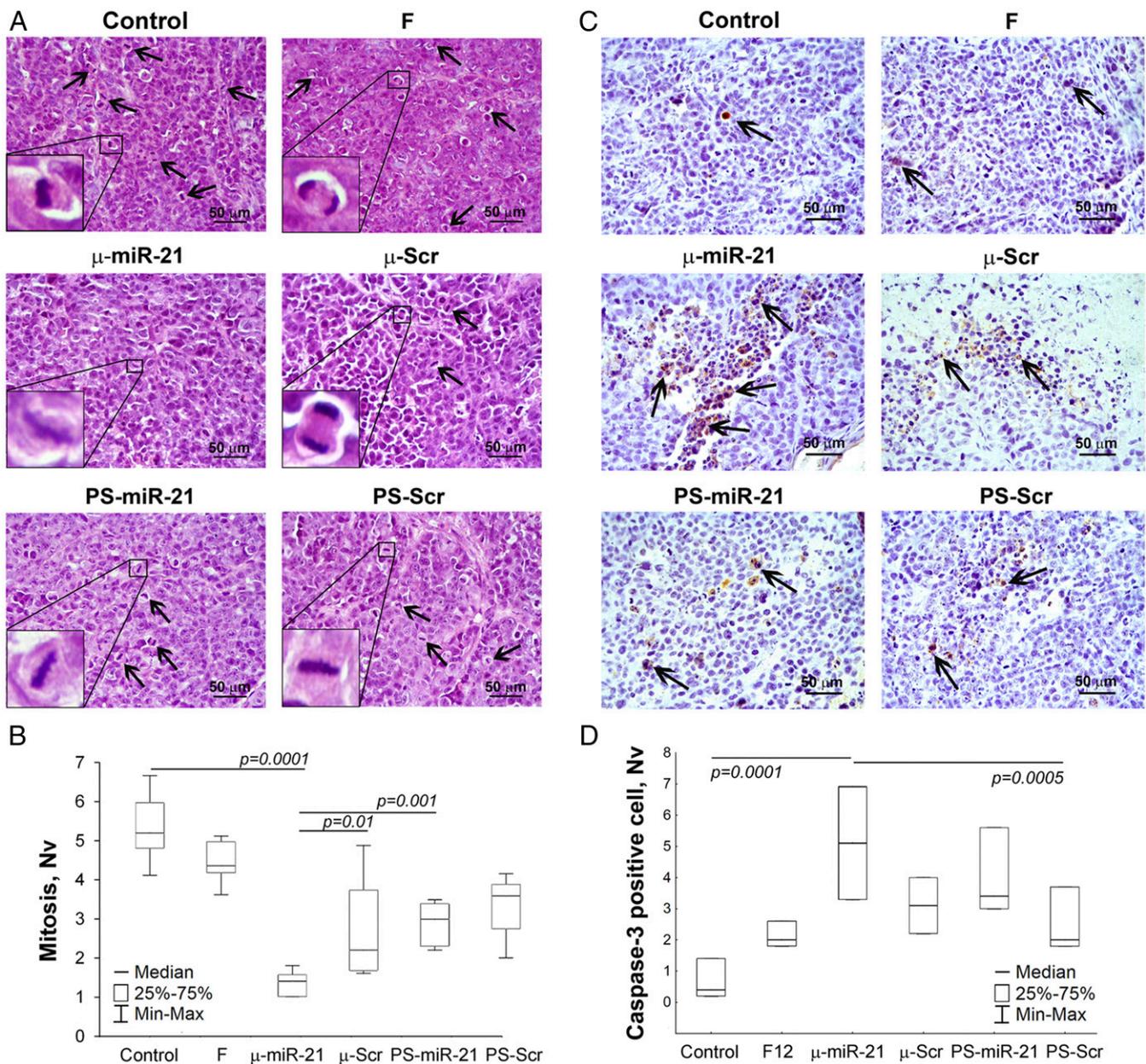


Fig. 5. Mitosis and apoptosis in KB-8-5 tumors after therapy with μ -miR-21-ON and PS-miR-21-ON. (A) Typical images of tumor sections after hematoxylin and eosin staining. Magnification $\times 400$. Bar corresponds to 50 μ m. Mitoses are indicated by arrows. Typical examples of individual mitotic events are shown with magnification $\times 1,000$ in the bottom left corner. (B) Morphometric analysis of tumor tissue with mitosis counting. (C) Typical images of tumor sections after immunohistochemical staining with caspase-3 monoclonal antibodies. Examples of caspase-3-positive cells are indicated by arrows. Magnification $\times 400$. (D) Morphometric analysis of tumor tissue with apoptotic cells counting. Nv $\times 400$. Nv, the numerical density indicating the number of particles in the unit tissue volume. Data were statistically analyzed using one-way ANOVA with post hoc Tukey test; *P* value indicates a statistically reliable difference.

oligonucleotides have been proposed, in which the phosphodiester backbone was replaced by different variants of phosphorothioates, methylphosphonates, phosphoramidates, phosphotriesters, phosphorodithioates, or derivatives with guanidinium groups or *S*-methylthiourea motifs (22–24). Among the proposed modifications, the most successful are phosphorothioates. Although this chemical substitution is endowed with a number of beneficial properties, the presence of serious shortcomings (4, 5, 25–27) has led to a search for new, improved modifications. In this regard, the μ -modification has a number of important favorable properties, including exceptional resistance to nucleases, high affinity for the RNA targets, and faster RNA degradation by RNase H, compared

to PS ASOs (12). The listed biological characteristics of μ ASOs are directly reflected in the duration of functioning within the cell and the strength of silencing effect. The visible effect of μ ASOs lasts for at least 6 d, achieving 80 to 95% down-regulation of miRNA level, while PS ASOs caused only a twofold decrease, and it expired after 1 d. The main *in vitro* and *in vivo* characteristics of μ -oligonucleotides in comparison with phosphorothioate analogs are summarized in Table 1.

An important difference between μ and PS ASO is the ability of phosphorothioates to penetrate inside the cells in a carrier-free mode (28, 29). In the study of antitumor activity of μ -miR-21-ON *in vivo*, both oligomers were enclosed in folate-containing

lipoplexes in order to equalize the conditions of their delivery to tumor cells. The biodistribution data showed that μ ASO rapidly spread throughout the body, ensuring efficient accumulation in tumors, while phosphorothioate oligonucleotides are characterized by anchoring in the area of administration and lower penetration efficiency into the tumor tissue. At 24 h postinjection, both μ and PS ASOs were almost equally distributed between tumor and kidney (Fig. 2).

The study of antitumor potential showed that μ -miR-21-ON provides drastic retardation of tumor growth, up to 90% 3 d after the first injection, which lasted up to 30 d until the end of the experiment. The antitumor effect of μ -miR-21-ON was accompanied by a significant decrease in mitotic activity and stimulation of proapoptotic, but not necrotic, behavior of tumor cells. μ -miR-21-ON administrated to mice specifically decreased the miR-21 level in tumor observed even a week after the last injection; that was not the case for PS-miR-21-ON therapy. Moreover, during this period, a μ -miR-21-ON-mediated increase in expression of tumor suppressor proteins such as PTEN and PDCD4, was maintained. These data demonstrate the specific and potent therapeutic effect of μ -miR-21-ON whereas the effect of PS-miR-21-ON is not so pronounced and unspecific. Earlier, it was noticed that efficient uptake of PS ASOs by cells does not always result in a pharmacologically functional intracellular pathway and therapeutic activity (30, 31).

According to the results of in vivo experiments, it seems that μ -Scr-ON is exerting some effects on tumor cells. The obtained data show that administration of oligonucleotides of any type, regardless of target specificity, leads to a moderate nontargeted effect and an increase in the scatter of experimental parameters within the group. It should be emphasized that there were no statistically significant differences between the effects of μ -Scr-ON and PS-Scr-ON, as well as PS-miR-21-ON. Moreover, our experiments have shown that no decrease in the level of miR-21, miR-17, or miR-155 was detected in response to μ -Scr-ON, thus additionally indicating the nontargeted systemic response to oligonucleotide administration. It cannot be excluded that this nontargeted effect can be not only of negative nature, resulting from general cytotoxicity, but also has a positive character,

providing an immunomodulatory or immunostimulating effect, leading to tumor regression.

The use of targeted folate-equipped cationic liposomes allowed one not only to balance the delivery of μ and PS ASOs into the tumor, but also to reduce the dose of the administered compounds, as well as mitigate their possible toxicity. For instance, in vivo administration of PS ASOs in a carrier-free mode requires doses of 20 mg/kg for the inhibition of tumor growth (32). Application of targeted delivery systems provides the decrease in PS ASO dose to 2 mg/kg (10 μ g per mice), which is sufficient for their effective accumulation in tumor tissue (33). The study of blood biochemistry showed that neither the presence of a tumor, nor the therapy performed, has an effect on the level of liver and kidney enzymes. Moreover, it has been shown that the effect of both μ - and PS ASOs on the tumor process apparently reduces the total destructive changes in the liver tissue.

The other side of the ability of phosphorothioates to circulate in the body for a long time is the off-target effects associated with nonspecific interactions. The use of μ -oligonucleotides with higher therapeutic performance formulated in cationic liposomes or other nanoparticles or conjugated to some chemical moieties that interact with cell-specific receptors for effective tissue delivery seems to be safer and advantageous. In this direction, the use of *N*-acetylgalactosamine (GalNAc) moiety, providing interaction with asialoglycoprotein receptors, conjugated to antisense oligonucleotide was shown to greatly expand the therapeutic opportunities of ASOs (34–36). Conjugation of the GalNAc ligand to gapmer PS ASOs attenuates toxicity and promotes penetration into cells by a productive endocytic pathway that leads to a 30-fold increase in therapeutic potency and at least a 40-fold decrease in the dosage of PS ASOs (35, 37).

The high performance of antisense therapy in vivo is determined to a large extent by the appropriate RNA target choice. Selected in the study, miR-21 represents a crucial regulator of a plethora of cellular processes, such as proliferation, differentiation, apoptosis, migration, etc. (38–41). As a result, inhibition of the aberrant hyperactivity of miR-21 in tumor tissue leads to global changes in cellular signaling pathways and promotes

Table 1. Comparison of biological properties of μ and PS ASOs

Biological properties	Mesyl phosphoramidate (μ) ASO	Phosphorothioate (PS) ASO
In vitro (12)		
Hybridization efficiency in equimolar concentration with miRNA, %	100	46
Stability in 10% fetal bovine serum, $t_{1/2}$, h	>168	96
RNase H cleavage rate, k_{obs} , $10^{-6} \cdot s^{-1}$	28.3 ± 2.2	8.7 ± 0.6
Efficiency of miRNA down-regulation in tumor cells, 72 h, %	75 to 90	—**
Duration of miRNA down-regulation in tumor cells, h	>144	<72
Inhibition of tumor cell migration, 72 h	19-fold	Fivefold
In vivo		
Reduction in tumor growth*		
Average tumor volume	Eightfold	Twofold
Average tumor weight	12-fold	Twofold
Tumor doubling time, d	9.9 ± 2.9	4.3 ± 1.2
Decrease in mitosis	3.9-fold	1.8-fold
Increase in caspase-3 positive cells	12.8-fold	8.5-fold
Efficiency of miRNA down-regulation in tumor tissue, %*	50	25
Increase in tumor suppressor protein expression in tumor tissue, fold*	1.5- to 3.5-fold	—**
Toxicity		
Blood biochemistry	No	No
Liver	No	No
Kidney	No	Moderate

*The effect was measured 7 d after the last administration of ASO.

**No effect was observed.

Appendix, Table S5. PCR amplification was carried out using BioMaster HS-qPCR SYBR Blue mix (Biolabmix) according to the manufacturer's protocol. The obtained qPCR data were analyzed by standard Bio-Rad iQ5 v.2.0 software. For each sample, the threshold cycle (Ct) was determined. Quantitative assessment of the level of transcript representation and relative miRNA expression was performed by comparing the Ct values for miRNA and U6 small nuclear RNA (snRNA) used as a reference.

Western Blot Analysis. To prepare cell lysates from KB-8-5 tumors, tumor tissue specimens were homogenized in radioimmunoprecipitation assay (RIPA) buffer (Thermo Scientific) using an electric homogenizer, followed by centrifugation at +4 °C (12,000 rpm, 20 min). Collected cell lysates were separated in 12.5% sodium dodecyl sulfate (SDS)/PAGE and transferred to a poly(vinylidene difluoride) (PVDF) membrane using a semidry transfer. Western Blot analysis was performed as described earlier (47, 48) using primary antibodies against PTEN (ab154812, 1:800; Abcam), PDCD4 (ab79405, 1:1,000; Abcam), and GAPDH (ab9485, 1:2,000; Abcam), and secondary HRP-conjugated goat anti-rabbit antibodies (ab6721; Abcam).

Histology. Histological analysis was performed similar to ref. 49. Briefly, the specimens of fixed tumors, kidneys, and livers from each animal were dehydrated in ascending ethanols and xylols and embedded in HISTOMIX paraffin (BioVitrum). Paraffin sections (5 μm) were sliced on a Microm HM 355S microtome (Thermo Fisher Scientific) and stained with hematoxylin and eosin, microscopically examined, and scanned. Tumor sections for immunohistochemical (IHC) studies (3 to 4 μm) were deparaffinated and rehydrated; antigen retrieval was carried out after exposure in a microwave oven at 700 W. The samples were incubated with the caspase-3 (ab2302, Abcam) specific antibodies according to the manufacturer's protocol. Next, the sections were incubated with secondary HRP-conjugated antibodies (Spring Bioscience detection system), exposed to 3,3'-diaminobenzidine (DAB) substrate, and stained with Mayer's hematoxylin. Images were obtained using an Axiostar Plus microscope equipped with an Axiocam MRC5 digital camera (Zeiss) at 100x and 400x magnifications. Twenty random fields were studied in each specimen, comprising 140 fields for each group of mice.

Morphometric analysis of tumor, liver, and kidney sections was performed as described in ref. 49 by point counting, using a counting grid which consists of 100 testing points in a testing area equal to $3.2 \times 10^6 \mu\text{m}^2$. Ten to 15 random fields were studied in each specimen, comprising 70 to 100 fields for each group of mice in total during morphometric analyses of liver and kidney, and 20 to 50 random fields were studied in each specimen, comprising 140 to 350 fields for each group of mice in total during morphometric analyses of tumors. Morphometric analysis of tumor tissue included evaluation of the volume densities (Vv, %) of unaltered tumor tissue, lymphoid infiltration, necrosis, total destructive changes (lymphoid infiltration plus necrosis), and numerical density (Nv) of mitoses and caspase-3-positive cells in tumor tissue. The ratio of unaltered tumor tissue to mutilated tumor tissue was estimated.

The volume density (Vv, %) representing the volume fraction of tissue occupied by this compartment was determined from the points lying over this structure and calculated using the following formula: $Vv = (P_{\text{structure}}/P_{\text{test}}) \times 100\%$, where $P_{\text{structure}}$ denotes the number of points over the structure and P_{test} denotes the total number of test points, in this case 100. The numerical density (Nv) indicating the number of particles in the unit tissue volume was evaluated as the number of particles in the square unit, $3.2 \times 10^6 \mu\text{m}^2$ in this case.

Statistics. The data obtained were statistically processed using one-way ANOVA and post hoc Tukey test ($P \leq 0.05$). The statistics package STATISTICA version 10.0 was used for this analysis.

Data Availability. All study data are included in the article and *SI Appendix*.

ACKNOWLEDGMENTS. This work was funded by Russian Science Foundation Grant 19-74-30011, the Russian State-funded budget project of Institute of Chemical Biology and Fundamental Medicine SB RAS Grant AAAA-A17-117020210024-8, and Russian Foundation for Basic Research Grants 18-515-05007 and 18-515-57006. We thank Dr. B. P. Chelobanov for synthesis of Cy5.5-labeled oligonucleotides.

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