Molecular Therapy

Review



Clinical Applications of Single-Stranded Oligonucleotides: Current Landscape of Approved and In-Development Therapeutics

Juergen Scharner¹ and Isabel Aznarez¹

¹Stoke Therapeutics, Inc., Bedford, MA, USA

Single-stranded oligonucleotides have been explored as a therapeutic modality for more than 20 years. Only during the last 5 years have single-stranded oligonucleotides become a modality of choice in the fields of precision medicine and targeted therapeutics. Recently, there have been a number of development efforts involving this modality that have led to treatments for genetic diseases that were once untreatable. This review highlights key applications of single-stranded oligonucleotides that function in a sequence-dependent manner when applied to modulate precursor (pre-)mRNA splicing, gene expression, and immune pathways. These applications have been used to address diseases that range from neurological to muscular to metabolic, as well as to develop vaccines. The wide range of applications denotes the versatility of single-stranded oligonucleotides as a robust therapeutic platform. The focus of this review is centered on approved single-stranded oligonucleotide therapies and the evolution of oligonucleotide therapeutics into novel applications currently in clinical development.

Single-stranded oligonucleotides are synthetic, short, modified RNA or DNA molecules that function in either a sequence-dependent (antisense oligonucleotides [ASOs] and immune stimulatory oligonucleotides [ISOs]) or tertiary structure-dependent manner (e.g., aptamers¹). In this review, we focus our attention on oligonucleotide therapeutics whose activities are dictated by their sequences; ASOs bind to their target by Watson and Crick base-pairing,² while ISOs are recognized and bound by proteins that detect specific motifs.³

Single-stranded oligonucleotide therapeutics have evolved substantially since the use of DNA ASOs were first described in the late 1970s.^{4,5} The year 1998 saw the first ever approved ASO therapy, formivirsen, which consisted of a 21-mer DNA ASO with a phosphorothioate (PS) backbone (Figure 1).⁶ Formivirsen, much like the first ASO applications, was an antiviral molecule, designed to target cytomegalovirus. It was not until 2013 that the second ASO drug, mipomersen, obtained US Food and Drug Administration (FDA) approval to treat homozygous familial hypercholesterolemia (HoFH) via RNase H-mediated degradation of apolipoprotein B (APOB).⁷ Mipomersen represented a new generation of ASOs using new chemical modifications and design that led to more stable (increased nuclease resistance), safer (decreased pro-inflammatory response), and potent (increased RNA binding affinity) compounds with enhanced pharmacokinetic (PK) and pharmacodynamic (PD) properties (Figure 1). During the 15 years between the two approvals, the field of singlestranded oligonucleotide therapeutics matured, leading to an acceleration in the development of this therapeutic modality and an increase in the number of approved drugs. In the last 4 years, six new ASO therapies and one ISO therapy have obtained FDA or European Medicines Agency (EMA) approval. These therapies address a range of genetic diseases across neurological, metabolic/cardiovascular, and muscular conditions as well as potentiate the efficacy of a hepatitis B vaccine. The approved therapies utilize three major types of single-stranded oligonucleotide modalities, that is, gapmers, steric blocking, and CpGs, to modulate gene expression, precursor (pre-)mRNA splicing, and immune pathways. Target organs expanded beyond the eye and liver to the central nervous system (CNS) and muscle. Singlestranded oligonucleotides represent a broad-spectrum therapeutic platform that continues to evolve with innovations in applications, chemistry, and tissue targeting.

Approved Single-Stranded Oligonucleotide Therapeutics Gapmer ASOs

Mechanism of Action. Gapmers are synthetic, single-stranded, and typically 16–20 nt long with a central stretch of 8–10 DNA nt flanked on each side by a stretch or wing of 4–5 nt containing modifications in the sugar ring. These sugar modifications in the wings increase nuclease resistance protecting the ends of the ASO, avoid or reduce an immune response, and increase binding affinity. Commonly, gapmers contain PS bonds throughout to increase stability and plasma protein binding, which improves the PK properties of the gapmer ASO (Figure 1).⁸ With their typical 4-8-4/5-10-5 (wing-DNA-wing) structure, gapmers bind in a sequence-specific manner by Watson and Crick base-pairing to their target RNA and form a stretch of double-stranded RNA-DNA hybrid. These RNA-DNA hybrids mimic endogenous RNA-DNA hybrids that occur naturally during DNA replication, which are recognized and disrupted via RNase H-mediated cleavage of the RNA strand. The cleavage then

Correspondence: Isabel Aznarez, Stoke Therapeutics, Inc., 45 Wiggins Avenue, Bedford, MA 01730, USA.

E-mail: iaznarez@stoketherapeutics.com

https://doi.org/10.1016/j.ymthe.2020.12.022.



Figure 1. Single-Stranded Oligonucleotide Chemistries

Chemical structures of ribose and backbone modifications used in single-stranded oligonucleotides compared to DNA and RNA structures are shown. Modifications highlighted in green are used in approved oligonucleotide drugs. DNA, deoxyribonucleic acid; RNA ribonucleic acid; Me, methyl; MOE, methoxyethyl; LNA, locked nucleic acid; cEt, constrained ethyl; BNA, bridged nucleic acid; PMO, phosphorodiamidate morpholino oligomer; PO, phosphodiester; PS phosphorothioate.

leads to the degradation of the RNA. Leveraging this endogenous mechanism, gapmers cause RNase H-mediated cleavage of the target pre-mRNA or mRNA at the location bound by the stretch of DNA. The target transcript is subsequently degraded, resulting in reductions of the target transcript and corresponding protein (Figure 2). As single-stranded ASOs, gapmers are taken up by cells in the CNS, eye, liver, kidney, adrenal glands, and lungs via a natural endocytic process, and they are subsequently released from endosomes into the cytoplasm, reaching the nucleus by a yet to be fully understood mechanism.³ Gapmers can be used to reduce the level of any protein or specific protein isoforms, as well as toxic proteins that result from a gain-of-function or dominant negative mutation.⁹

Mipomersen. Mipomersen (Kynamro; Ionis Pharmaceuticals/Genzyme) was approved by the FDA in January 2013 to treat HoFH, a rare and serious hereditary condition with high risk for premature coronary heart disease associated with atherosclerosis.^{10,11} HoFH is caused by homozygous or compound heterozygous mutations in one of three genes (*APOB*, *LDLR*, or *PCSK9*), leading to severely elevated low-density lipoprotein cholesterol (LDL-C) levels.¹⁰ Mipomersen is a second-generation gapmer ASO with a 5-10-5 structure, 2'-O-methoxyethyl (2'MOE) modifications in the wings (2'MOE-DNA-2'MOE), and PS backbone (Figure 1). Mipomersen was designed to bind to the coding region of the APOB-100 mRNA (an isoform encoded by the *APOB* gene), and elicit RNase H-mediated degradation of the mRNA and reduce ApoB protein levels (see Figure 2B for the mechanism of action).¹²

PK studies in preclinical species as well as humans showed that mipomersen has a 30-day half-life in plasma and target tissue (liver).⁷ In patients with HoFH, mipomersen is administered systemically via weekly subcutaneous injections at a dose of 200 mg and is indicated as an adjunct therapy to lipid-lowering medications and diet to reduce LDL-C, ApoB, total cholesterol, and non-high-density lipoprotein cholesterol. In humans, PD studies showed a mipomersen-mediated reduction of ApoB protein of 46%, which resulted in a 47% decrease of LDL-C. 7

During clinical trials, safety concerns of hepatotoxicity as determined by increased levels of transaminases were observed leading to a black label warning for mipomersen.¹³ Because of the risk of hepatotoxicity, mipomersen is available only through a restricted program under a risk evaluation and mitigation strategy. Other adverse effects include injection site reaction, fever, flu-like symptoms, and fatigue.¹⁴ Mipomersen was not approved by the EMA based on safety concerns. In 2016, Ionis and Kastle Therapeutics acquired the rights to mipomersen, and in May 2018 the marketing of the drug was discontinued.

Inotersen. Inotersen (Tegsedi; Ionis Pharmaceuticals/Akcea) was approved by the FDA and EMA in 2018 for the treatment of patients with hereditary amyloid transthyretin (ATTR) amyloidosis, a rare systemic disorder that is characterized by progressive peripheral polyneuropathy, cardiomyopathy, nephropathy, and gastrointestinal dysfunction. The disorder is caused by dominant-negative mutations in the *TTR* gene that lead to destabilization of the tetrameric TTR protein complex, causing aggregation of monomers into extracellular amyloid deposits.¹⁵ The design and mechanism of action of inotersen are analogous to mipomersen, and it was designed to bind and elicit RNase H-mediated degradation of mutant and wild-type *TTR* transcripts, resulting in decreased mutant and wild-type TTR proteins (see Figure 2B for the mechanism of action).¹⁶

Analyses of preclinical species and healthy human volunteers dosed with inotersen demonstrated a more than 80% reduction of plasma TTR protein levels, which correlated with a reduction in the *TTR* transcript in the liver (target organ) of preclinical species.¹⁶ In patients with ATTR amyloidosis, inotersen is administered via weekly subcutaneous injection of a recommended dose of 284 mg to treat the polyneuropathy phenotype. Clinical data indicated that inotersen improved the course of the neurological disease as well as the quality



Figure 2. Mechanism of Action of Gapmer ASOs

(A) Expression of an example target gene from transcription through translation of a wild-type or toxic protein. (B) Gapmer ASO with the typical structure of modified chemistry in the wings to protect the ends from nucleases and an internal stretch of DNA that leads to the formation of the RNA-DNA hybrid when bound to the target transcript. In this example, the gapmer ASO binds to an exon in the pre-mRNA and mRNA and recruits RNase H that recognizes the RNA/DNA hybrid and cleaves the RNA. The cleavage triggers RNA degradation, leading to reduction of wild-type or toxic protein levels. Gapmer ASOs can be designed to target other transcript regions, e.g., introns. Even though RNase H is more abundant in the nucleus, RNase H-mediated cleavage of RNA-DNA hybrids can also occur in the cytoplasm.

of life of ATTR amyloidosis patients. During clinical trials, safety concerns included thrombocytopenia (fatal in some patients) and glomerulonephritis, which occurred independently in 3% of patients.¹⁷ Inotersen is the first approved example of a gapmer ASO that directly targets the root cause of a disease.

Volanesorsen. Volanesorsen (Waylivra; Ionis Pharmaceuticals/Akcea) received a conditional market authorization in Europe issued by the EMA in 2019 for the treatment of familial chylomicronemia syndrome (FCS), which is characterized by 10- to 100-fold increased levels of triglycerides compared to the healthy population, which leads to hypertriglyceridemia and recurrent episodes of pancreatitis. The elevated triglycerides result from reduced or lack of lipoprotein lipase activity caused predominantly by recessive mutations in the *LPL* gene.¹⁸ To address the lack of LPL activity, volanesorsen was designed to reduce the levels of *APOC3* mRNA and apolipoprotein C (ApoC)-III protein to decrease plasma triglycerides via an LPL-independent mechanism (see Figure 2B for the mechanism of action).

Volanesorsen is another second-generation gapmer ASO that targets a primarily liver-expressed mRNA. Preclinical studies in rodents and non-human primates showed a significant reduction of ApoC-III and triglyceride plasma levels. Similar results were obtained from the administration of volanesorsen to healthy human subjects, demonstrating a good translation between preclinical species and human clinical studies.¹⁹ In patients suffering from FCS, volanesorsen treatment consists of a once weekly subcutaneous dose of 300 mg in combination with dietary restrictions. Clinical data indicated that 77% of patients treated with volanesorsen have significantly reduced levels of triglyceride in plasma, reaching values lower than 750 mg/dL, well below the 880 mg/dL required to reduce the risk of pancreatitis. In these patients, dietary restriction alone was not sufficient to reduce cholesterol levels. Similar to the other two systemically administered, approved second-generation gapmer ASOs, thrombocytopenia and site of injection reactions were common adverse events.²⁰

Steric-Blocking ASOs

Mechanism of Action. Steric-blocking ASOs are synthetic, fully modified, single-stranded oligonucleotides that typically range from 15 to 30 nt in length and contain chemical modifications in the sugar as well as the backbone (Figure 1). Many of the chemical modifications are similar to gapmer ASOs and are also intended to increase



Figure 3. Mechanism of Action of Nusinersen for the Treatment of Spinal Muscular Atrophy

(A) Region of *SMN2* pre-mRNA containing exons 6, 7, and 8. *SMN2* exon 7 carries a silent single nucleotide change with respect to *SMN1* that causes exon 7 skipping, which leads to an unstable SMN protein. Only 10% of *SMN2* pre-mRNA is properly spliced, resulting in an insufficient level of functional SMN protein to compensate for the loss of the *SMN1* gene. (B) Nusinersen binding to intron 7 of the *SMN2* pre-mRNA and promoting exon 7 inclusion, which leads to increased levels of SMN protein and improved motor neuron function.

nuclease resistance, reduce the immune response, and increase binding affinity. Unlike gapmers, steric-blocking ASOs do not elicit RNase H-mediated degradation of the target RNA, but rather bind to the target RNA via sequence-specific Watson and Crick base-pairing and either hinder the binding of trans-acting factors (e.g., small nuclear RNA [snRNA], microRNA [miRNA], long non-coding RNA [lncRNA], or RNA-binding proteins) to their cognate sequence or prevent the formation of RNA secondary structures.^{2,9} Their first application dates to mid-1980s in which an ASO was designed to target a 3' splice site in the herpes simplex virus 1 pre-mRNA preventing splicing and functioning as an antiviral agent.²¹ Since then, steric blocking ASOs have been used to correct mutation-driven splicing defects and modulate mRNA stability and protein translation.²² Steric-blocking ASOs are thought to be taken up by cells in the CNS, eye, liver, kidney, adrenal glands, lungs, and, to a much lesser extent, to the muscle (see below for a specific example) via the same mechanism as gapmer ASOs.³

Nusinersen. Nusinersen (Spinraza; Ionis Pharmaceuticals/Biogen) was approved by the FDA in December 2016 and the EMA in 2017 for the treatment of spinal muscular atrophy (SMA), a rare neuromuscular disorder that is the most common genetic cause of infant mortality. SMA is caused by recessive mutations in the survival motor neuron 1 (*SMN1*) gene, and the disease severity is mitigated by a copy gene, *SMN2*, which produces low levels of normal SMN protein.²³ *SMN2* exon 7 carries a single nucleotide change with respect to *SMN1* that severely decreases its inclusion in the final mRNA, leading to an unstable SMN protein (Figure 3A).²⁴ Nusinersen is an 18-mer, fully modified 2'MOE-PS (Figure 1) that was designed to bind to position +10 of intron 7 of the *SMN1/2* pre-mRNA and block access of the splicing repressor hnRNPA1 to its cognate intronic splicing silencer (Figure 3B).²⁵ This results in increased levels of *SMN2* exon 7 inclusion, leading to correctly spliced mRNA and increased levels of SMN protein, which compensates for the loss of *SMN1* (Figure 3B).^{26,27}

Preclinical studies in mice indicated that nusinersen has a long-lasting effect on *SMN2* splicing in the brain with continued maximal effect through the 6-month observational period following a single intracerobroventricular injection. In addition, intrathecal (i.t.) injection in non-human primates showed a widespread distribution throughout the spinal cord where motor neurons (target cell) reside.²⁸ PK studies in patients injected i.t. with up to a 9-mg dose of nusinersen showed consistent results with preclinical species and indicated a prolonged half-life of 4–6 months in cerebrospinal fluid (CSF).²⁹ PK/PD analysis in autopsy tissue demonstrated broad distribution of the drug in the spinal cord and brain regions, as well as increases in SMN protein in the spinal cord.³⁰

Clinical trials have successfully demonstrated efficacy in infants and older children treated with nusinersen shown by an increase in motor function and survival.³⁰ Nusinersen is approved to treat infants through adult SMA patients and is administered to patients directly into the CSF via i.t. injection by lumbar puncture at a dose



Figure 4. Mechanism of Action of Eteplirsen for the Treatment of Duchenne Muscular Dystrophy (DMD)

(A) Effect of a deletion of exons 49–50 region in the *DMD* gene that causes a frameshift and leads to the introduction of a premature termination codon in exon 51. The mutant mRNA is degraded in the cytoplasm by nonsense-mediated mRNA decay (NMD), and no dystrophin protein is produced, causing DMD. (B) Eteplirsen binding to exon 51, which prevents its inclusion and restores the frame. The resulting mRNA lacking exons 49–51 is translated to generate an internally truncated dystrophin protein that retains partial function.

albeit truncated dystrophin protein (e.g., inframe deletions) lead to a milder disease presentation known as Becker muscular dystrophy.³³ This genotype-phenotype correlation provided the rationale to skip specific DMD exons to avoid deleterious nonsense and frameshift mutations that cause DMD to restore the frame at the transcript level and produce a Becker-like truncated, partially functional dystrophin.^{34,35} Following this rationale, eteplirsen was designed as a 30-mer phosphorodiamidate morpholino oligomer (PMO) (Figure 1) that binds to exon 51 of the DMD gene, causing the skipping of exon 51 to avoid deleterious mutations. The skipping of exon 51 restores the mRNA frame and leads to the production of an internally truncated, partially functional dystrophin protein (Figure 4B).³⁶ Unlike nusinersen that can treat the vast majority of SMA patients (patients with no SMN2 are very rare), eteplirsen can only address patients who are amenable to

of 12 mg in 5 mL every 4 months after an initial loading dose of 12 mg administered monthly for the first 4 months.³¹ Adverse reactions include lower respiratory infection, upper respiratory infection, and constipation. The label has warnings for thrombocy-topenia and coagulation abnormalities as well as renal toxicity. While these warnings relate to the class of ASOs, they have not been observed for nusinersen.³⁰ In addition to the approved patient population, nusinersen has been further tested in pre-symptomatic infants with mutations in *SMN1* and two to three copies of *SMN2* in the NURTURE trial, and the results underscore the proven efficacy of the treatment and the importance of neonatal diagnosis.³²

Eteplirsen. Eteplirsen (Exondys 51; Sarepta Therapeutics) received an accelerated approval by the FDA in 2016 for the treatment of Duchenne muscular dystrophy (DMD), which is caused by complete loss-of-function mutations in the X-linked *DMD* gene and predominantly affects males. DMD is a fatal disease due to loss of muscle function caused by the lack of dystrophin protein (Figure 4A). Loss of function mutations in the *DMD* gene that result in partially functional exon 51 skipping, which account for 13% of the DMD patient population.^{37,38}

Proof-of-concept studies using DMD patient muscle cells and a humanized mouse model validated the exon skipping approach preclinically.³⁶ PK studies in the mdx mouse model indicated that eteplirsen has a 6-h plasma half-life and is distributed to heart, skeletal, and diaphragm muscles (target tissue), as well as to kidney, where exposure was highest.³⁹ In patients, muscle biopsies and western blot and immunofluorescence analyses detected a small but significant increase of dystrophin.^{40,41} While muscle is not a tissue that is normally targetable by unconjugated ASOs, the membranes of the muscle of DMD patients are leaky, allowing for eteplirsen to be taken up by myofibers.⁴²

Clinical assessment of patients treated with eteplirsen concluded that there was some preservation of the distance walked in 6 min compared to patients in natural history studies. Despite the small increase in dystrophin and the small number of patients in the clinical trials, the FDA granted conditional approval for eteplirsen for the treatment of males with mutations amenable to exon 51 skipping.

A weekly 30 mg/kg dose is administered systemically by intravenous (i.v.) infusion (35–60 min). Adverse reactions include balance disorder and vomiting. The label has no warnings; however, eteplirsen was not approved by the EMA in 2019, citing lack of efficacy. Follow-on studies are still underway.

Golodirsen and Viltolarsen. Golodirsen (Vyondys 53; Sarepta Therapeutics) and viltolarsen (Viltepso; NS Pharma) are treatments for DMD that were granted accelerated approval by the FDA in 2019 and 2020, respectively. The rationale for the development of golodirsen and viltolarsen and their mechanism of action is analogous to eteplirsen. Golodirsen is a 25-mer PMO that was designed to bind to exon 53 of the *DMD* gene and cause the skipping of exon 53 to avoid deleterious loss-of-function frameshifting mutations. Likewise, viltolarsen is a 21-mer PMO that also binds to exon 53 to avoid mutations. In both cases, the skipping of exon 53 restores the frame and leads to the production of an internally truncated, partially functional dystrophin protein (see Figure 4 for an analogous example). Both golodirsen and viltolarsen address 8% of DMD patients.

Golodirsen and viltolarsen are approved for males with mutations amenable to exon 53 skipping and are administered by i.v. infusion (35–60 min) of a weekly 30 mg/kg dose and 80 mg/kg, respectively. The following adverse reactions have been described for golodirsen: hypersensitivity reactions, including rash, pyrexia, pruritus, urticaria, dermatitis, and skin exfoliation, with some requiring treatment. The most common adverse reactions observed in patients treated with viltolarsen include upper respiratory tract infection, injection site reaction, cough, and pyrexia. In both cases, renal toxicity was observed in animals who received golodirsen or viltolarsen in preclinical studies, but not in the clinical studies.^{43,44}

Milasen: N-of-1 ASO Therapy. The advancements and availability of next-generation sequencings have enabled the identification of pathogenic "private" mutations in a wide range of genetic diseases, especially mutations located in flanking and deep intronic sequences. The vast majority of disease-causing intronic mutations lead to aberrant pre-mRNA splicing. Depending on how disruptive these mutations are to the splicing process, their effect could range from 100% faulty mRNA to varied levels of residual properly spliced mRNA. The latter indicates that splicing is not completely "broken" by the mutation and it could be fixed. Given the targeted nature and success of nusinersen in treating SMA by correcting a splicing defect, combined with local delivery, safety profile, and manufacturing ease,⁴⁵ steric blocking ASOs have become the preferred modality for personalized treatment.

The first N-of-1 drug was developed to treat a single patient with neuronal ceroid lipofuscinosis 7 (CLN7), a form of Batten's disease. This condition is a rare and fatal neurodegenerative disease that is characterized by progressive symptoms resulting in blindness, ataxia, seizures, and developmental delay and results from recessive mutations in the *MFSD8* gene. At the onset at 3 years of age, symptoms in a patient were mild, but by her sixth birthday she could barely speak. After a Batten's disease diagnosis was made, genetic screening

revealed a known pathogenic mutation in one allele of the MFSD8 gene. Using whole-genome sequencing, a deep intronic insertion of a SINE-VNTR-Alu (SVA) retrotransposon was uncovered. RT-PCR analysis of the patient's blood cells revealed that the insertion led to the activation of a cryptic 3' splice site in intron 6, resulting in missplicing and consequently the introduction of a premature termination codon. Seven ASOs were designed to target the cryptic 3' splice site and predicted exonic splicing enhancers, of which three led to an increase in proper MFSD8 splicing. As the chemical modifications used in nusinersen proved safe for local delivery in the CNS, the 2'MOE-PS ASO (Figure 1) targeting MFSD8, which was the most efficacious hit, was selected for further evaluation. The ASO was named "milasen" after the patient's first name, Mila. Milasen (Boston Children's Hospital/Mila's Miracle Foundation) is a 22-mer ASO that was shown to alleviate lysosomal dysfunction phenotypes in the patient's derived fibroblast cells.46

Preclinical toxicology studies in rats were conducted to show that i.t. injection of a single ascending dose of milasen had no adverse effects at the lowest dose (0.06 mg), which guided the subsequent dose selection in clinical studies. As the patient's condition continued to deteriorate, an investigational new drug application was filed with the FDA, and in 2018 clinical investigational treatment under an Expanded Access-Investigational New Drug application was initiated. Similar to nusinersen, milasen was administered via i.t. bolus injection by lumbar puncture. The initial dose of 3.5 mg was increased every 2 weeks until it reached 42 mg, which was then administered once every 3 months. The patient was monitored through the first year after the initial injection and no observable serious adverse events were detected. Neurological and neuropsychological assessments at 3 and 6 months after treatment initiation showed stabilization and improvement of Vineland Adaptive Behavior Scales subscores. Seizure frequency and duration decreased by 50% during the follow-up period after treatment initiation compared to pre-treatment measurements.⁴⁶ Based on subsequent patient evaluations, the dose and dosing regimen was modified and the patient is currently receiving a dose of 70 mg every 2 months (T. Yu, personal communication).

Milasen is an example of the potential of steric-blocking ASOs as a modality applied to the rapid development of personalized treatment for individual patients with fatal genetic diseases caused by splicing mutations. The feasibility of development of N-of-1 treatments as demonstrated by milasen underscores the importance of molecular diagnosis, even if no therapies are available at the time of diagnosis.

ISOs

Mechanism of Action. In addition to splicing modulation and target knockdown, single-stranded oligonucleotides are also being used to stimulate the innate immune system in a variety of applications, including vaccine adjuvants and cancer. Unlike ASOs, ISOs do not bind to RNA, but rather they bind to proteins. They mimic molecular signatures in microorganisms known as pathogen-associated molecular patterns (PAMPs) that trigger the innate immune system via recognition by the Toll-like family of receptors (TLRs) expressed in cells such



as antigen-presenting cells (APCs), dendritic cells (DCs), and B cells.⁴⁷ Examples of PAMPs include double-stranded RNA (dsRNA), which can activate TLR3, single-stranded RNA (ssRNA), which can activate TLR7/8, and unmethylated CpG-containing single-stranded DNA (ssDNA) or dsDNA, which can trigger a protective immune response by activating TLR9.48-51 Synthetic single-stranded oligonucleotides containing unmethylated CpG motifs can mimic bacterial unmethylated DNA and cause a similar response. There are at least three structurally distinct classes of CpG oligonucleotides, that is, CpG-A, CpG-B, and CpG-C.^{52,53} The three classes differ in the specific response they trigger, which is influenced by their sequence, chemistry, and higher order structure.^{54,55} CpG-B ASOs are ssDNA sequences with a PS backbone and contain multiple CpG motifs. They primarily promote the production of T helper (Th)1-type cytokines and type I interferons, the maturation of plasmacytoid DCs (pDCs), and the proliferation and activation of B cells, which boost the immune response to coadministered vaccines (Figure 5).^{51–53}

Figure 5. CpG-Containing Oligonucleotides Mediated Immune Cell Stimulation

Internalized CpG-containing ISOs are mainly recognized by TLR9, which activates a complex signaling cascade resulting in the nuclear translocation of transcription factors including AP1, IRF7, and nuclear factor κB (NF-κB). Transcriptional activation of pro-inflammatory genes regulates the maturation of pDCs, the activation and proliferation of B cells, as well as the production of type I interferons and Th1-type cytokines. CpG-mediated immune cell activation, as well as the subsequent secretion of cytokines, stimulates the immune response to the coadministered antigen (not shown), and it produces a more rapid and longer lasting antibody response when compared to alternative vaccine adjuvants. In addition to vaccine adjuvants, single-stranded CpG-containing ISOs are currently also being tested in cancer patients and to modulate the immune response in inflammatory diseases (Table 1).

The first approved vaccine containing a CpG ISO as an adjuvant was Heplisav-B (Dynavax Technologies) in 2017, a vaccine to prevent hepatitis B virus (HBV) infection in adults.^{52,56} Heplisav-B contains 3 mg of CpG 1018, a 22mer DNA oligonucleotide with a uniform PS backbone, which is delivered together with 20 µg of the hepatitis B surface antigen (HbsAg) via intramuscular administration in two doses 4 weeks apart.^{52,56} When compared to Energix-B (GSK), a three-dose hepatitis B vaccine using aluminum hydroxide (alum) as an adjuvant, CpG 1018 induced a significantly improved immune response that was induced more rapidly and lasted longer. In the first phase 2 study comparing Heplisav-B to Energix-B, 79% of Heplisav-B recipients showed a

protective antibody response 4 weeks after the first dose compared to only 12% of Energix-B recipients.⁵⁷ Adverse events, including fatigue and headache, were low and similar in both groups, while mild injection site tenderness was more common in the presence of CpG 1018.⁵⁷ A more rapid induction of protective antibody levels was confirmed in additional studies.^{58–60} Heplisav-B also showed superior long-term effects 50 weeks after the last active dose,⁶⁰ and it produced seroprotection in patients with chronic kidney disease who were hyporesponsive to currently licensed alum-adjuvanted hepatitis B vaccines.^{61,62} CpG 1018 is currently being investigated in a phase 1 clinical trial as an adjuvant in an HIV vaccine (ClinicalTrials.gov: NCT04177355) as well as vaccines against severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) (ClinicalTrials.gov: NCT04450004, NCT04405908, and NCT04487210).

The first nine approved ASO drugs and milasen represent a broad platform with applications in five different organs (Figure 6) and a



Figure 6. Single-Stranded Oligonucleotide Therapies

Summary of approved drugs to date and their target tissue. Milasen is a clinical investigational treatment under an Expanded Access-Investigational New Drug application.

variety of mechanisms of action. These ASO therapies are administered via local (CNS and eye) or systemic delivery to treat or prevent a wide range of genetic diseases or viral infections. Locally administered ASOs in the eye and CNS have an advantage over systemically injected ASOs, given their lower doses that result in a lower concentration of plasma circulating drug and accumulation of drug in the kidneys. Generally, the higher the ASO dose is, the more likely it is to cause adverse effects. In the second part, this review focuses on some of the mitigation strategies that are underway to reduce ASO doses and improve their therapeutic index by increasing the potency and uptake efficiency of ASOs.

Innovations in the Clinic

Beginning with the first DNA oligonucleotides, and continuing through the currently approved single-stranded oligonucleotides (un-conjugated DNA, 2'MOE, and PMO sequences), a great deal of un-

derstanding and innovations in the ASO field have led to improvements in the PK, distribution, and potency, as well as novel applications of this modality of nucleic acid therapeutics. However, there remains potential for further improvements in ASO potency, specificity, and safety, as well as targeted ASO delivery. Some of these innovations, as well as new applications, are currently being tested in the clinic and will be discussed in further detail.

ASO Chemistry

One of the most successful chemical modifications used in single stranded ASOs is the 2'MOE substitution in the sugar ring (Figure 1). It offers increased nuclease resistance compared to DNA, and it forms duplexes with RNA that are \sim 2°C more stable per modification compared to DNA-PS/RNA hybrids.⁶³ Structural studies and molecular dynamics simulations of fully modified MOE-RNA duplexes show that the 2' substitution locks the sugars in a C3'-endo

conformation.^{64,65} Enforcing a C3'-endo sugar pucker improves affinity for RNA, as well as nuclease resistance in bicyclic 2',4'-constrained MOE (cMOE) and 2',4'-constrained ethyl (cEt) modifications (Figure 1), leading to more potent ASOs.⁶⁶ The first 2',4'-constrained gapmer ASOs had locked nucleic acids (LNAs) (Figure 1) in their wings. However, some LNA gapmers have been associated with hepatotoxicity in animals,⁶⁷ which may be linked to increased RNase H-mediated off-target effects⁶⁸ or specific sequence motifs.⁶⁹ A clinical trial for an LNA gapmer targeting PCSK9 was terminated due to renal tubular toxicity observed in test subjects.⁷⁰ Gapmer ASOs with cEt-modified wings have improved potency when compared to 2'MOE, without increased liver toxicity.67,71 Several ASOs containing the new generation 2.5 cEt chemistry are currently being tested in clinical trials (Table 1). One of them is AZD9150 (danvatirsen; IONIS-STAT3-2.5Rx), a 16-mer gapmer with 3-nt cEt wings targeting STAT3 to treat multiple types of cancer (Table 1). Early results from a phase 1b study in patients with diffuse large B cell lymphoma (DLBCL) showed that AZD9150 is tolerated and demonstrated efficacy at a dose of 3 mg/kg in a subset of heavily pretreated patients with DLBCL.⁷²

ASOs with a stereodefined backbone represent another chemistry innovation currently being tested in clinical trials. Therapeutic ASOs contain PS linkages, enhancing metabolic stability as well as protein-binding properties, facilitating ASO distribution and cellular uptake.^{73,74} The use of PS linkages in the ASO backbone introduces a chiral center with two configurations, Sp and Rp.⁷⁵ A k-mer ASO therefore has 2^{k-1} different stereoisomers, each potentially with a different stability and potency profile, making it challenging to comprehensively study various Sp/Rp configurations. Early work using first-generation DNA-PS ASOs showed that the Sp and Rp forms have different biophysical and biological properties affecting melting temperature (Tm), RNase H activating ability, and nuclease resistance.^{76,77} Wan et al.⁷⁵ tested the effect of a uniform DNA gap chirality on a limited number of second-generation MOE and generation 2.5 cEt-containing gapmers. They showed that in vitro, the effect of Rp ASOs was comparable to their stereorandom counterpart, while Sp ASOs had a lower Tm and improved endonuclease resistance. In vivo, Rp and Sp compounds were both quickly metabolized, resulting in low activity, which led to the conclusion that a mixture of Sp and Rp is required to balance RNase H activity and nuclease resistance.⁷⁵ A screen of stereopure gapmer compounds with one or two Rp and Sp linkages at specific positions did not yield any improvement in potency. However, combining fixed Rp/ Sp linkages with a 2'MOE at gap position 2 resulted in a dramatic improvement of the therapeutic index.⁷⁸ Synthesis of stereopure ASOs, pioneered by Stec et al.,79 was traditionally limited to research use due to low coupling efficiency and difficulties in removing the chiral auxiliary. Advances in the oxazaphospholidine synthesis approach address these traditional hurdles and, while still inefficient compared to phosphoramidite coupling reactions, open the path for stereopure/stereodefined ASOs in clinical applications.^{80–83} The first stereopure compound tested in the clinic was suvodirsen (WVE-210201), an ASO targeting DMD exon 51 in DMD patients. However, the ASO was not efficacious in patients, which resulted in the termination of the phase 2/3 trial (Clinical-Trials.gov: NCT03907072). Other stereodefined compounds are currently being tested in patients with Huntington's disease (HD) in phase 1/2 clinical trials (WVE-120101, ClinicalTrials.gov: NCT03225833; WVE-120102, ClinicalTrials.gov: NCT03225846) (Table 1). In contrast to other HTT-targeting strategies, this technology uses allele-specific knockdown by targeting SNPs linked to the disease-causing CAG-repeat expansions.⁸⁴ While such an approach would require multiple treatment candidates depending on the SNPs present in the HD patient population, it could provide a safer profile than targeting mutant and wild-type HTT non-selectively. Given the immense number of possible stereoisomers for each sequence, it remains to be determined whether the compounds in development to target HTT are the safest and most efficacious. While some progress has been made to identify a stereochemical code for specific sequences,⁸⁰ this uncertainty will likely remain a limitation for the development of stereodefined compounds in the near future.

Ligand-Conjugated ASOs

Naked ASOs systemically delivered *in vivo* are rapidly taken up by liver and kidney.⁸ While ASO accumulation and activity has been detected in a variety of extrahepatic/extra kidney tissues, systemic delivery to specific target organs is relatively inefficient.^{8,85} The development of ASO conjugates has enabled tissue-specific targeting of ASOs and short interfering RNAs (siRNAs). Such conjugates include various receptor ligands, cell-penetrating peptides (CPPs), and antibodies, some of which are already in use in the clinic.^{86–89} In addition to improving the delivery of ASOs, conjugation led to targeted delivery, allowing for dose reduction of ASOs.

One such ligand is triantennary N-acetylgalactosamine (GN3), a sugar moiety binding to the asialoglycoprotein receptor (ASGP-R). The ASGP-R is primarily expressed on hepatocytes, and the binding of its ligand N-acetylgalactosamine (GalNAc) promotes receptormediated endocytosis of the ligand. Leveraging this mechanism, conjugation of ASOs and siRNAs with GN3 promotes the receptormediated uptake of the conjugated therapeutic agent. Upon internalization, the conjugate is metabolized by hydrolysis of phosphodiester bonds between the nucleic acid and GN3, as well as hydrolysis of the amide linkages in GN3.90 The use of GN3 as an siRNA conjugate targeting ALAS1 (givosiran) has been approved by the FDA to treat acute hepatic porphyria.⁸⁸ Several GN3-conjugated gapmers are currently being evaluated in clinical trials to target AGT, ANGPTL3, CFB, GHR, KLKB1, LPA, TMPRSS6, and TTR specifically in the liver (Table 1). In preclinical studies in mice, GN3-conjugated gapmers designed against five different targets showed a 5- to 10-fold greater potency compared to the unconjugated parental gapmer, while being well tolerated. When GN3 was conjugated to a gapmer with higher affinity cEt wings, an ~60-fold potency increase was achieved.⁹⁰ Recently published early results from a clinical trial with a GN3-gapmer targeting APOA (APO(a)-LRx) in patients with hyperlipoproteinemia(a) and cardiovascular disease showed a reduction in lipoprotein(a) levels

www.moleculartherapy.org

Review

Table 1. List of Recent Clinical Trials Using Gapmer and Steric Blocking ASOs and ISOs That Have Completed Trials after January 1, 2020 or Are Currently "Active," "Recruiting," or "Enrolling by Invitation" as Described on ClinicalTrials.gov

	ASO TYPE	МОА	DRUG NAME(S)	DISEASES
111/25	Gapmer	RNAse H degradation of AGT	IONIS-AGT-LRx	Treatment Resistant Hypertension
LIVER	Gapmer	RNAse H degradation of ANGPTI 3	AKCEA-ANGPTL3-LRx: ISIS 703802	Hypertriglyceridemia
	oupiner	invise in degradation of Anon 125	ARCEN ARGI 120 ERX, 1919 / 05002	Type 2 Diabetes Mellitus
				Nonalcoholic Fatty Liver Dicease
	Ganmer	RNAse H degradation of CEP	IONIS.ER.I Ry	Primany Ind. Nenhronathy
	Gapiner	KWASE IT degradation of Crb	IONIS-I D-LIX	Age Belated Magular Degeneration
	Common	DNA U da em detirer of CCCD		Age-Related Macular Degeneration
	Gapmer	RNAse H degradation of GLOR	IONIS-GUGRRX; ISIS 449884	Type 2 Diabetes
	Gapmer	RNAse H degradation of GHR	IONIS-GHR-LRx; ISIS 766720	Acromegaly
	Gapmer	RNAse H degradation of	IONIS-HBVRx	Hepatitis B Viral Infection
		hepatitis B surface antigen		
	Gapmer	RNAse H degradation of KLKB1	IONIS-PKK-LRx; ISIS 721744	Hereditary Angioedema
	Gapmer	RNAse H degradation of LPA	AKCEA-APO(a)-LRx; ISIS 681257	Cardiovascular Disease
	Gapmer	RNAse H degradation of TMPRSS6	IONIS-TMPRSS6-LRx	Beta-thalassemia
	Gapmer	RNAse H degradation of TTR	AKCEA-TTR-LRx	Hereditary Transthyretin-Mediated Amyloid Polyneuropathy
				Transthyretin-Mediated Amyloid Cardiomyopathy
CNS	Steric blocking	NMD exon skipping to increase	STK-001	Dravet Syndrome
	5	full-length SCN1A expression		,
action of	Gapmer	RNAse H degradation of COORF72	BIIB078: IONIS-C9Rx	Amyotrophic Lateral Sclerosis
Contra la	Ganmer	RNAse H degradation of HTT	Tominersen: IONIS-HTTRy: RG6042	Huntington's Disease
SO	Ganmer	RNAse H degradation of	WVF.120101	Huntington's Disease
	Japiner	CND containing mutant //TT	WVL-120101	nunungion a Disease
	Common	DNAss II degradation of	MB/E 120102	Unitington la Disease
	Gapmer	RIVASE H degradation of	WVE-120102	nunungton's Disease
		SNP-containing mutant HTT		
	Gapmer	KNAse H degradation of LRRK2	BIIB094; ION859; IONIS-BIIB7Rx	Parkinson's Disease
	Gapmer	RNAse H degradation of SOD1	Tofersen; BIIB067; Ionis-SOD1Rx	Amyotrophic Lateral Sclerosis
	Gapmer	RNAse H degradation of TAU	IONIS-MAPTRx; BIIB080	Alzheimer's Disease
	Gapmer	RNAse H degradation of UBE3A-ATS	GTX-102	Angelman Syndrome
	Steric blocking	Skipping of cryptic exon activated	Sepofarsen; QR-110	Leber Congenital Amaurosis 10
EYE		by c.2991+1655A>G in CEP290		
	Steric blocking	USH2A exon 13 skipping to remove	QR-421a	Retinitis Pigmentosa
		pathogenic mutations		
	Gapmer	RNAse H degradation of RHO	QR-1123	AD Retinitis Pigmentosa
MUSCLE	Steric blocking	DMD even 51 skipping to restore reading frame	SRP-5051	Duchenne Muscular Dystronby
	Storic blocking	DMD exon 45 skipping to restore reading frame	Casimorson: SPR-4045	Duchenne Muscular Dystrophy
	Steric blocking	DMD exon 45 skipping to restore reading frame		Duchenne Muscular Dystrophy
	Steric blocking	DMD exon 45 skipping to restore reading frame		Duchenne Muscular Dystrophy
	Gapmer	RNASE H degradation of DNM2	DYN 101; IONIS-DNM2-2.5KX	Centronuclear Myopathy
	Gapmer	RNAse H degradation of AR	IONIS-AR-2.5Rx; AZD5312	Prostate Cancer
	Gapmer	RNAse H degradation of STAT3	Danvatirsen; IONIS-STAT3-2.5Rx;	Advanced and Refractory Pancreatic Cancer
			AZD9150	Early Stage and Metastatic NSCLC
				Mismatch Repair Deficient Colorectal Cancer
				Advanced Solid Tumors; Relapsed or Refractory DLBCL
				Relapsed or Refractory Aggressive Non-Hodgkin's Lymphoma
				Muscle Invasive Bladder Cancer
				Metastatic Squamous Cell Carcinoma of Head & Neck
LY WL				
LUNGS	Gapmer	RNAse H degradation of ENAC	IONIS-ENaCRx; IONIS-ENAC-2.5Rx	Cystic Fibrosis
LUNGS		•		
LUNGS				
LUNGS	CpG-A	TLR9 agonist	CMP-001	Malignant Melanoma; Lymph Node Cancer:
IMMUNE CELLS	СрС-А	TLR9 agonist	CMP-001	Malignant Melanoma; Lymph Node Cancer; Lymphoma: NSCLC: Malignant Colorectal Neoplasms
IMMUNE CELLS	СрG-А	TLR9 agonist	CMP-001	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases:
	Срб-А	TLR9 agonist	CMP-001	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Suramour Coll Carrinoma of the Hond and Nork
	CpG-A	TLR9 agonist	CMP-001	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck
IMMUNE CELLS	CpG-A CpG-B	TLR9 agonist	CMP-001 CPG1018; 1018 ISS	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19
IMMUNE CELLS	CpG-A CpG-B CpG-B	TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection
IMMUNE CELLS	CpG-A CpG-B CpG-B CpG-B CpG-B	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease
IMMUNE CELLS	CpG-A CpG-B CpG-B CpG-B CpG-B	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies
IMMUNE CELLS	CpG-B CpG-B CpG-B CpG-B CpG-B CpG-C	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies Advanced or Metastatic Solid Malignancies
IMMUNE CELLS	CpG-A CpG-B CpG-B CpG-B CpG-B CpG-C	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies B-Cell Non-Hodgkin Lymphoma
IMMUNE CELLS	CpG-A CpG-B CpG-B CpG-B CpG-B CpG-C	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies Advanced or Metastatic Solid Malignancies B-Cell Non-Hodgkin Lymphoma Metastatic Pancreatic Adenocarcinoma
IMMUNE CELLS	CpG-A CpG-B CpG-B CpG-B CpG-B CpG-C	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies Advanced or Metastatic Solid Malignancies B-Cell Non-Hodgkin Lymphoma Metastatic Pancreatic Adenocarcinoma Grade 1-3A Follicular Lymphoma; Lymphoma;
IMMUNE CELLS	CpG-B CpG-B CpG-B CpG-B CpG-B CpG-C	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies Advanced or Metastatic Solid Malignancies B-Cell Non-Hodgkin Lymphoma Metastatic Pancreatic Adenocarcinoma Grade 1-3A Follicular Lymphoma; Lymphoma;
IMMUNE CELLS	CpG-A CpG-B CpG-B CpG-B CpG-C CpG-C	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101 Tilsotolimod: IMO-2125	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies Advanced or Metastatic Solid Malignancies B-Cell Non-Hodgkin Lymphoma Metastatic Pancreatic Adenocarcinoma Grade 1-3A Follicular Lymphoma; Iymphoma; Metastatic Melanoma, Head Neck Cancer; Breast Cancer
IMMUNE CELLS	CpG-A CpG-B CpG-B CpG-B CpG-B CpG-C CpG-C	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101 Tilsotolimod; IMO-2125 DV281	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies Advanced or Metastatic Solid Malignancies B-Cell Non-Hodgkin Lymphoma Metastatic Pancreatic Adenocarcinoma Grade 1-3A Follicular Lymphoma; Lymphoma; Metastatic Melanoma, Head Neck Cancer; Breast Cancer Metastatic Melanoma; Solid Tumor Advanced Non Small Cell Lung Cancer
IMMUNE CELLS	CpG-A CpG-B CpG-B CpG-B CpG-B CpG-C CpG-C CpG-C CpG-C	TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101 Tilsotolimod; IMO-2125 DV281 Advasc/ G555 2	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies Advanced or Metastatic Solid Malignancies B-Cell Non-Hodgkin Lymphoma Metastatic Pancreatic Adenocarcinoma Grade 1-3A Follicular Lymphoma; Lymphoma; Metastatic Melanoma, Head Neck Cancer; Breast Cancer Metastatic Melanoma; Solid Tumor Advanced Non Small Cell Lung Cancer
LUNGS OF CELLS	Срб-А Срб-В Срб-В Срб-В Срб-В Срб-В Срб-С Срб-С Срб-С Срб-С Срб-С Срб-С Срб-С Срб-С Срб-С Срб-2	TLR9 agonist TLR9 agonist	CMP-001 CPG1018; 1018 ISS Agatolimod; CPG7909; PF03512676 CPG10104 DUK-CPG-001 SD-101 Tilsotolimod; IMO-2125 DV281 Advax-CpG55.2 Cokinglimod; IMIC9120	Malignant Melanoma; Lymph Node Cancer; Lymphoma; NSCLC; Malignant Colorectal Neoplasms and Liver Metastases; Squamous Cell Carcinoma of the Head and Neck HIV; COVID-19 Anthrax; CMV Infection Hookworm Disease Myeloid Malignancies, Lymphoid Malignancies Advanced or Metastatic Solid Malignancies B-Cell Non-Hodgkin Lymphoma Metastatic Pancreatic Adenocarcinoma Grade 1-3A Follicular Lymphoma; Lymphoma; Metastatic Melanoma, Head Neck Cancer; Breast Cancer Metastatic Melanoma, Head Neck Cancer; Breast Cancer Metastatic Melanoma; Solid Tumor Advanced Non Small Cell Lung Cancer Influenza

CpG indicates unclassified CpG oligonucleotides (A. Krieg, personal communication).



Figure 7. Mechanism of Action of QR-421a, a New Application of Steric-Blocking ASOs

The figure depicts a portion of USH2A pre-mRNA containing exons 12, 13, and 14 (middle) in which exon 13 carries missense (blue sign) or premature termination codon (PTC)-introducing (nonsense or frameshift, red sign) retinitis pigmentosa (RP) mutations. Exon 13 encodes half of laminin epidermal growth factor (EGF)-like domain 4, domains 5, 6, 7, and half of laminin EGF-like domain 8 (light blue rectangles highlighted in red). Normally, missense or PTC-introducing RP mutations lead to non-functional full-length or non-functional truncated usherin, respectively (top). QR-421a (+ASO) promotes skipping of exon 13 (depicted by lines connecting exons 12 and 14), leading to the generation of an usherin protein that lacks half of laminin EGF-like domain 4, as well as domains 5, 6, 7, and half of laminin EGF-like domain 8 (bottom), but maintains proper function. Green rectangle, signal peptide: purple rectangle, laminin G-like domain: orange rectangle, laminin N-terminal domain; light blue rectangles, laminin EGF-like domains; green rectangles, fibronectin type III repeats; yellow circles, laminin G; red rectangle, transmembrane domain; blue circle, PDZ binding motif.

by up to 80% using much lower doses (20, 40, or 60 mg every 4 weeks, 20 mg every 2 weeks, or 20 mg every week) than that of approved unconjugated gapmers, potentially reducing unwanted adverse events such as thrombocytopenia and renal toxicity.⁹¹

While GN3 aims to limit delivery of ASOs directly to hepatocytes, other types of conjugates are being used to broaden ASO distribution in the body. For example, there are three splice-modulating ASOs approved for DMD; however, their efficacy is limited by inefficient uptake of ASOs in skeletal muscle.³⁸ Furthermore, DMD also presents in the heart and diaphragm, two regions where ASO uptake is also limited.92 To achieve better tissue distribution, next-generation DMD-targeting ASOs are covalently linked to CPPs. The use of small cationic peptides to enhance cellular uptake of oligonucleotides was first proposed in 1987.93 CPPs are typically 5-30 aa long and can pass through the cell membrane via energy-dependent and -independent mechanisms without interaction with specific receptors.94,95 While there are only few examples where CPPs increase biological activity of negatively charged ASOs or siRNAs (reviewed in Juliano⁹⁶), CPPs show dramatic effects when linked to neutrally charged peptide nucleic acids (PNAs) and PMOs.⁹⁷ Muscle-specific CPPs typically contain two tandem repeats of a series of arginine (R), 6-aminohexanoic acid (X), and/or beta-alanine (B) residues with or without a short peptide linker.^{98,99} A 14-aa peptide, which is highly potent in the heart, diaphragm, and quadriceps, key muscles in the treatment of DMD, is the B-peptide (RXRRBR)₂XB.⁹⁸ In a dog model of DMD, CPP-conjugated PMOs (PPMOs) rescued dystrophin expression in the myocardium and cardiac Purkinje fibers. PPMOs also improved electrocardiogram abnormalities without apparent toxicity, leading the way toward better therapies for DMD.⁸⁶ PPMOs targeting DMD exons 45 and 51 are currently being tested in patients with DMD (Table 1).

New ASO Applications

Apart from the development of new chemistries and innovations in tissue targeting, new applications of steric-blocking ASOs have also made their way into the clinic. ASO-mediated exon skipping, for example, has traditionally been used to restore the reading frame that was disrupted by frameshifting mutations. A new approach uses an exon skipping strategy to remove retinitis pigmentosa mutations in a constitutive exon of USH2A while maintaining usherin protein function. This strategy was first proposed for diseases caused by mutations in lamin A/C such as Emery-Dreifuss muscular dystrophy.¹⁰⁰ Lamin A/C is an intermediate filament and has an *a*-helical central rod domain. Proteins lacking LMNA exon 5, which encodes part of the central rod domain, localize correctly and perform better in cell-based assays than do proteins harboring dominant-negative missense mutations in exon 5, suggesting that exon skipping is a viable strategy to target missense mutations in exons that encode repetitive domains. Similarly, usherin contains multiple repetitive domains that can be shortened by deleting USH2A exon 13 (exon 12 in the mouse), while maintaining protein function (Figure 7).¹⁰¹ Retinitis pigmentosa patients with pathogenic homozygous or compound heterozygous nonsense and/or missense mutations in USH2A exon 13 might therefore benefit from an exon skipping approach to restore protein function, a strategy currently being tested in the clinic (QR-421a, Table 1).

While skipping of constitutive exons might also be applicable to other diseases, the exon must be in-frame and code for a redundant domain, which has to be assessed via *in vitro* and *in vivo* functional studies. Such applications will therefore likely remain a small niche of therapeutic splice-modulating ASOs.

Another innovative approach, termed TANGO, leverages naturally occurring non-productive alternative splicing events to increase



Figure 8. Mechanism of Action of STK-001, a New Application of Steric-Blocking ASOs

(A) Region of *SCN1A* wild-type pre-mRNA containing non-productive (non-coding) exon X (yellow rectangle) and flanking coding exons (brown rectangles). *SCN1A* premRNA is alternatively spliced such that it generates a non-productive mRNA containing the non-productive exon X, which leads to the introduction of a PTC, and a productive mRNA lacking exon X. Upon export to the cytoplasm, the non-productive mRNA is degraded by nonsense-mediated mRNA decay and the productive mRNA is translated into wild-type Nav1.1 protein. The pre-mRNA carrying DS mutations undergoes the same alternative splicing processing, but the mutant productive mRNA does not produce a functional protein (not shown in the figure), leading to haploinsufficiency of Nav1.1. (B) STK-001 (ASO) binding to the non-productive exon X of the *SCN1A* wild-type and mutant (not shown) pre-mRNA and promotes exon X skipping, which leads to a reduction in non-productive mRNA and increased levels of productive mRNA and wild-type Nav1.1 protein to near normal levels. STK-001 leverages the wild-type gene copy to compensate for the loss-of-function mutant alleles in DS patients.

gene expression.¹⁰² These events lead to the introduction of a premature termination codon (PTC) in the mRNA, inducing nonsensemediated mRNA decay (NMD) and no protein production. ASOmediated prevention of non-productive alternative splicing increased productive mRNA and protein levels in vitro and in vivo.¹⁰² A proposed therapeutic application focused on upregulating productive mRNA and protein in autosomal dominant haploinsufficient diseases, such that upregulation of the wild-type gene copy could compensate for the loss of the mutant copy and restore protein levels. An example of this application (STK-001, Table 1) is currently being tested in clinical trials for the treatment of Dravet syndrome (DS), an autosomal dominant haploinsufficiency (Figure 8). Preclinical studies of STK-001 in a DS mouse model demonstrated that preventing the inclusion of a non-productive exon in the Scn1a gene led to increased productive mRNA and restoration of full-length fully functional Nav1.1 protein levels. Nav1.1 protein restoration significantly increased survival and reduced seizures in DS mice.¹⁰³ Unlike other splicing modulating ASOs in the clinic that address specific mutations

and only a subset of the patient population, the TANGO therapeutic approach leverages the wild-type gene copy in the context of autosomal dominant haploinsufficiencies and therefore has the potential to address all patients with heterozygous loss-of-function mutations.

Single-stranded oligonucleotide therapeutics represent a platform of targeted and selective medicines that have been used to address severe genetic diseases that were once untreatable. With the rise of wholeexome and genome sequencing and the multigene panel mutation screening, molecular diagnosis of genetic diseases is routinely performed leading to the identification of disease-associated genes and earlier diagnosis. Continued advances in chemistry and ligand conjugation has led to the development of new generations of ASO drugs that are more potent and therefore safer. Targeted delivery of ASOs together with new ASO applications are expanding the breadth of the ASO therapeutic platform and have paved the way for a future in which ASO therapeutics become a common modality to treat genetic diseases and beyond.

ACKNOWLEDGMENTS

This work was supported by Stoke Therapeutics. We would like to acknowledge Jennifer Fairman from Fairman Studios for the design of the figures. We would like to thank Arthur Krieg for advice on immunostimulatory oligonucleotides. We would also like to thank the reviewers for insightful comments, and Nga Tong who provided editorial support for this review.

DECLARATION OF INTERESTS

I.A. and J.S. are employees and hold shares of Stoke Therapeutics.

REFERENCES

- 1. Zhu, G., and Chen, X. (2018). Aptamer-based targeted therapy. Adv. Drug Deliv. Rev. 134, 65–78.
- Kole, R., Krainer, A.R., and Altman, S. (2012). RNA therapeutics: beyond RNA interference and antisense oligonucleotides. Nat. Rev. Drug Discov. 11, 125–140.
- Koller, E., Vincent, T.M., Chappell, A., De, S., Manoharan, M., and Bennett, C.F. (2011). Mechanisms of single-stranded phosphorothioate modified antisense oligonucleotide accumulation in hepatocytes. Nucleic Acids Res. 39, 4795–4807.
- Stephenson, M.L., and Zamecnik, P.C. (1978). Inhibition of Rous sarcoma viral RNA translation by a specific oligodeoxyribonucleotide. Proc. Natl. Acad. Sci. USA 75, 285–288.
- Zamecnik, P.C., and Stephenson, M.L. (1978). Inhibition of Rous sarcoma virus replication and cell transformation by a specific oligodeoxynucleotide. Proc. Natl. Acad. Sci. USA 75, 280–284.
- Roehr, B. (1998). Fomivirsen approved for CMV retinitis. J. Int. Assoc. Physicians AIDS Care 4, 14–16.
- Geary, R.S., Baker, B.F., and Crooke, S.T. (2015). Clinical and preclinical pharmacokinetics and pharmacodynamics of mipomersen (Kynamro®): a second-generation antisense oligonucleotide inhibitor of apolipoprotein B. Clin. Pharmacokinet. 54, 133–146.
- Geary, R.S., Norris, D., Yu, R., and Bennett, C.F. (2015). Pharmacokinetics, biodistribution and cell uptake of antisense oligonucleotides. Adv. Drug Deliv. Rev. 87, 46–51.
- Bennett, C.F., and Swayze, E.E. (2010). RNA targeting therapeutics: molecular mechanisms of antisense oligonucleotides as a therapeutic platform. Annu. Rev. Pharmacol. Toxicol. 50, 259–293.
- 10. Goldberg, A.C., Hopkins, P.N., Toth, P.P., Ballantyne, C.M., Rader, D.J., Robinson, J.G., Daniels, S.R., Gidding, S.S., de Ferranti, S.D., Ito, M.K., et al.; National Lipid Association Expert Panel on Familial Hypercholesterolemia (2011). Familial hyper-cholesterolemia: screening, diagnosis and management of pediatric and adult patients: clinical guidance from the National Lipid Association Expert Panel on Familial Hypercholesterolemia. J. Clin. Lipidol. 5 (3, Suppl), S1–S8.
- Rader, D.J., Cohen, J., and Hobbs, H.H. (2003). Monogenic hypercholesterolemia: new insights in pathogenesis and treatment. J. Clin. Invest. 111, 1795–1803.
- Crooke, R.M., Graham, M.J., Lemonidis, K.M., Whipple, C.P., Koo, S., and Perera, R.J. (2005). An apolipoprotein B antisense oligonucleotide lowers LDL cholesterol in hyperlipidemic mice without causing hepatic steatosis. J. Lipid Res. 46, 872–884.
- Genzyme. (2013). Kynamro: full prescribing information, https://www.accessdata. fda.gov/drugsatfda_docs/label/2013/203568s000lbl.pdf.
- Genzyme. (2013). Kynamro® (mipomersen sodium) injection prescribing information, https://www.accessdata.fda.gov/drugsatfda_docs/label/2019/203568s011lbl.pdf.
- Sekijima, Y. (2015). Transthyretin (ATTR) amyloidosis: clinical spectrum, molecular pathogenesis and disease-modifying treatments. J. Neurol. Neurosurg. Psychiatry 86, 1036–1043.
- 16. Ackermann, E.J., Guo, S., Benson, M.D., Booten, S., Freier, S., Hughes, S.G., Kim, T.W., Jesse Kwoh, T., Matson, J., Norris, D., et al. (2016). Suppressing transthyretin production in mice, monkeys and humans using 2nd-generation antisense oligonucleotides. Amyloid 23, 148–157.

- Benson, M.D., Waddington-Cruz, M., Berk, J.L., Polydefkis, M., Dyck, P.J., Wang, A.K., Planté-Bordeneuve, V., Barroso, F.A., Merlini, G., Obici, L., et al. (2018). Inotersen treatment for patients with hereditary transthyretin amyloidosis. N. Engl. J. Med. 379, 22–31.
- Johansen, C.T., Kathiresan, S., and Hegele, R.A. (2011). Genetic determinants of plasma triglycerides. J. Lipid Res. 52, 189–206.
- 19. Graham, M.J., Lee, R.G., Bell, T.A., 3rd, Fu, W., Mullick, A.E., Alexander, V.J., Singleton, W., Viney, N., Geary, R., Su, J., et al. (2013). Antisense oligonucleotide inhibition of apolipoprotein C-III reduces plasma triglycerides in rodents, nonhuman primates, and humans. Circ. Res. 112, 1479–1490.
- 20. Witztum, J.L., Gaudet, D., Freedman, S.D., Alexander, V.J., Digenio, A., Williams, K.R., Yang, Q., Hughes, S.G., Geary, R.S., Arca, M., et al. (2019). Volanesorsen and triglyceride levels in familial chylomicronemia syndrome. N. Engl. J. Med. 381, 531–542.
- 21. Smith, C.C., Aurelian, L., Reddy, M.P., Miller, P.S., and Ts'o, P.O. (1986). Antiviral effect of an oligo(nucleoside methylphosphonate) complementary to the splice junction of herpes simplex virus type 1 immediate early pre-mRNAs 4 and 5. Proc. Natl. Acad. Sci. USA 83, 2787–2791.
- 22. Lundin, K.E., Gissberg, O., and Smith, C.I. (2015). Oligonucleotide therapies: the past and the present. Hum. Gene Ther. *26*, 475–485.
- 23. Feldkötter, M., Schwarzer, V., Wirth, R., Wienker, T.F., and Wirth, B. (2002). Quantitative analyses of SMN1 and SMN2 based on real-time LightCycler PCR: fast and highly reliable carrier testing and prediction of severity of spinal muscular atrophy. Am. J. Hum. Genet. 70, 358–368.
- 24. Monani, U.R., Lorson, C.L., Parsons, D.W., Prior, T.W., Androphy, E.J., Burghes, A.H., and McPherson, J.D. (1999). A single nucleotide difference that alters splicing patterns distinguishes the SMA gene SMN1 from the copy gene SMN2. Hum. Mol. Genet. 8, 1177–1183.
- 25. Hua, Y., Vickers, T.A., Okunola, H.L., Bennett, C.F., and Krainer, A.R. (2008). Antisense masking of an hnRNP A1/A2 intronic splicing silencer corrects SMN2 splicing in transgenic mice. Am. J. Hum. Genet. 82, 834–848.
- 26. Hua, Y., Sahashi, K., Rigo, F., Hung, G., Horev, G., Bennett, C.F., and Krainer, A.R. (2011). Peripheral SMN restoration is essential for long-term rescue of a severe spinal muscular atrophy mouse model. Nature 478, 123–126.
- 27. Passini, M.A., Bu, J., Richards, A.M., Kinnecom, C., Sardi, S.P., Stanek, L.M., Hua, Y., Rigo, F., Matson, J., Hung, G., et al. (2011). Antisense oligonucleotides delivered to the mouse CNS ameliorate symptoms of severe spinal muscular atrophy. Sci. Transl. Med. 3, 72ra18.
- 28. Rigo, F., Chun, S.J., Norris, D.A., Hung, G., Lee, S., Matson, J., Fey, R.A., Gaus, H., Hua, Y., Grundy, J.S., et al. (2014). Pharmacology of a central nervous system delivered 2'-O-methoxyethyl-modified survival of motor neuron splicing oligonucleotide in mice and nonhuman primates. J. Pharmacol. Exp. Ther. 350, 46–55.
- 29. Chiriboga, C.A., Swoboda, K.J., Darras, B.T., Iannaccone, S.T., Montes, J., De Vivo, D.C., Norris, D.A., Bennett, C.F., and Bishop, K.M. (2016). Results from a phase 1 study of nusinersen (ISIS-SMN_{Rx}) in children with spinal muscular atrophy. Neurology 86, 890–897.
- 30. Finkel, R.S., Chiriboga, C.A., Vajsar, J., Day, J.W., Montes, J., De Vivo, D.C., Yamashita, M., Rigo, F., Hung, G., Schneider, E., et al. (2016). Treatment of infantile-onset spinal muscular atrophy with nusinersen: a phase 2, open-label, dose-escalation study. Lancet 388, 3017–3026.
- US Food and Drug Administration (2016). FDA approves first drug for spinal muscular dystrophy, https://www.fda.gov/news-events/press-announcements/fdaapproves-first-drug-spinal-muscular-atrophy.
- 32. De Vivo, D.C., Bertini, E., Swoboda, K.J., Hwu, W.L., Crawford, T.O., Finkel, R.S., Kirschner, J., Kuntz, N.L., Parsons, J.A., Ryan, M.M., et al.; NURTURE Study Group (2019). Nusinersen initiated in infants during the presymptomatic stage of spinal muscular atrophy: Interim efficacy and safety results from the phase 2 NURTURE study. Neuromuscul. Disord. 29, 842–856.
- 33. Koenig, M., Beggs, A.H., Moyer, M., Scherpf, S., Heindrich, K., Bettecken, T., Meng, G., Müller, C.R., Lindlöf, M., Kaariainen, H., et al. (1989). The molecular basis for Duchenne versus Becker muscular dystrophy: correlation of severity with type of deletion. Am. J. Hum. Genet. 45, 498–506.

- 34. Aartsma-Rus, A., Bremmer-Bout, M., Janson, A.A., den Dunnen, J.T., van Ommen, G.J., and van Deutekom, J.C. (2002). Targeted exon skipping as a potential gene correction therapy for Duchenne muscular dystrophy. Neuromuscul. Disord. 12 (Suppl 1), S71–S77.
- 35. Wilton, S.D., Lloyd, F., Carville, K., Fletcher, S., Honeyman, K., Agrawal, S., and Kole, R. (1999). Specific removal of the nonsense mutation from the mdx dystrophin mRNA using antisense oligonucleotides. Neuromuscul. Disord. 9, 330–338.
- 36. Arechavala-Gomeza, V., Graham, I.R., Popplewell, L.J., Adams, A.M., Aartsma-Rus, A., Kinali, M., Morgan, J.E., van Deutekom, J.C., Wilton, S.D., Dickson, G., and Muntoni, F. (2007). Comparative analysis of antisense oligonucleotide sequences for targeted skipping of exon 51 during dystrophin pre-mRNA splicing in human muscle. Hum. Gene Ther. 18, 798–810.
- 37. Aartsma-Rus, A., Straub, V., Hemmings, R., Haas, M., Schlosser-Weber, G., Stoyanova-Beninska, V., Mercuri, E., Muntoni, F., Sepodes, B., Vroom, E., and Balabanov, P. (2017). Development of exon skipping therapies for Duchenne muscular dystrophy: a critical review and a perspective on the outstanding issues. Nucleic Acid Ther. 27, 251–259.
- Aartsma-Rus, A., and Krieg, A.M. (2017). FDA approves eteplirsen for Duchenne muscular dystrophy: the next chapter in the eteplirsen saga. Nucleic Acid Ther. 27, 1–3.
- Sarepta Therapeutics (2016). Eteplirsen briefing document NDA 206488, https:// www.fda.gov/media/121650/download.
- 40. Charleston, J.S., Schnell, F.J., Dworzak, J., Donoghue, C., Lewis, S., Chen, L., Young, G.D., Milici, A.J., Voss, J., DeAlwis, U., et al. (2018). Eteplirsen treatment for Duchenne muscular dystrophy: exon skipping and dystrophin production. Neurology 90, e2146–e2154.
- 41. Mendell, J.R., Goemans, N., Lowes, L.P., Alfano, L.N., Berry, K., Shao, J., Kaye, E.M., and Mercuri, E.; Eteplirsen Study Group and Telethon Foundation DMD Italian Network (2016). Longitudinal effect of eteplirsen versus historical control on ambulation in Duchenne muscular dystrophy. Ann. Neurol. 79, 257–271.
- 42. Hoffman, E.P., Bronson, A., Levin, A.A., Takeda, S., Yokota, T., Baudy, A.R., and Connor, E.M. (2011). Restoring dystrophin expression in Duchenne muscular dystrophy muscle progress in exon skipping and stop codon read through. Am. J. Pathol. 179, 12–22.
- 43. US Food and Drug Administration (2019). FDA grants accelerated approval to first targeted treatment for rare Duchenne muscular dystrophy mutation, https://www. fda.gov/news-events/press-announcements/fda-grants-accelerated-approval-firsttargeted-treatment-rare-duchenne-muscular-dystrophy-mutation.
- 44. US Food and Drug Administration (2020). FDA approves targeted treatment for rare Duchenne muscular dystrophy mutation, https://www.fda.gov/news-events/ press-announcements/fda-approves-targeted-treatment-rare-duchenne-musculardystrophy-mutation.
- Levin, A.A. (2019). Treating disease at the RNA level with oligonucleotides. N. Engl. J. Med. 380, 57–70.
- 46. Kim, J., Hu, C., Moufawad El Achkar, C., Black, L.E., Douville, J., Larson, A., Pendergast, M.K., Goldkind, S.F., Lee, E.A., Kuniholm, A., et al. (2019). Patientcustomized oligonucleotide therapy for a rare genetic disease. N. Engl. J. Med. 381, 1644–1652.
- Medzhitov, R., and Janeway, C.A., Jr. (1997). Innate immunity: the virtues of a nonclonal system of recognition. Cell 91, 295–298.
- 48. Hong, D., Kurzrock, R., Kim, Y., Woessner, R., Younes, A., Nemunaitis, J., Fowler, N., Zhou, T., Schmidt, J., Jo, M., et al. (2015). AZD9150, a next-generation antisense oligonucleotide inhibitor of *STAT3* with early evidence of clinical activity in lymphoma and lung cancer. Sci. Transl. Med. 7, 314ra185.
- **49.** Krieg, A.M. (2002). CpG motifs in bacterial DNA and their immune effects. Annu. Rev. Immunol. *20*, 709–760.
- 50. Takeshita, F., Leifer, C.A., Gursel, I., Ishii, K.J., Takeshita, S., Gursel, M., and Klinman, D.M. (2001). Cutting edge: role of Toll-like receptor 9 in CpG DNAinduced activation of human cells. J. Immunol. *167*, 3555–3558.
- Kayraklioglu, N., Horuluoglu, B., and Klinman, D.M. (2021). CpG oligonucleotides as vaccine adjuvants. Methods Mol. Biol. 2197, 51–85.

- Campbell, J.D. (2017). Development of the CpG adjuvant 1018: a case study. Methods Mol. Biol. 1494, 15–27.
- Klinman, D.M. (2004). Immunotherapeutic uses of CpG oligodeoxynucleotides. Nat. Rev. Immunol. 4, 249–258.
- 54. Martinson, J.A., Tenorio, A.R., Montoya, C.J., Al-Harthi, L., Gichinga, C.N., Krieg, A.M., Baum, L.L., and Landay, A.L. (2007). Impact of class A, B and C CpG-oligodeoxynucleotides on in vitro activation of innate immune cells in human immunodeficiency virus-1 infected individuals. Immunology 120, 526–535.
- 55. Verthelyi, D., Ishii, K.J., Gursel, M., Takeshita, F., and Klinman, D.M. (2001). Human peripheral blood cells differentially recognize and respond to two distinct CPG motifs. J. Immunol. *166*, 2372–2377.
- 56. Kaufman, M.B. (2018). Pharmaceutical approval update. P&T 43, 83-84.
- 57. Halperin, S.A., Dobson, S., McNeil, S., Langley, J.M., Smith, B., McCall-Sani, R., Levitt, D., Nest, G.V., Gennevois, D., and Eiden, J.J. (2006). Comparison of the safety and immunogenicity of hepatitis B virus surface antigen co-administered with an immunostimulatory phosphorothioate oligonucleotide and a licensed hepatitis B vaccine in healthy young adults. Vaccine 24, 20–26.
- 58. Halperin, S.A., Ward, B., Cooper, C., Predy, G., Diaz-Mitoma, F., Dionne, M., Embree, J., McGeer, A., Zickler, P., Moltz, K.H., et al. (2012). Comparison of safety and immunogenicity of two doses of investigational hepatitis B virus surface antigen co-administered with an immunostimulatory phosphorothioate oligodeoxyribonucleotide and three doses of a licensed hepatitis B vaccine in healthy adults 18–55 years of age. Vaccine 30, 2556–2563.
- 59. Heyward, W.L., Kyle, M., Blumenau, J., Davis, M., Reisinger, K., Kabongo, M.L., Bennett, S., Janssen, R.S., Namini, H., and Martin, J.T. (2013). Immunogenicity and safety of an investigational hepatitis B vaccine with a Toll-like receptor 9 agonist adjuvant (HBsAg-1018) compared to a licensed hepatitis B vaccine in healthy adults 40–70 years of age. Vaccine 31, 5300–5305.
- 60. Sablan, B.P., Kim, D.J., Barzaga, N.G., Chow, W.C., Cho, M., Ahn, S.H., Hwang, S.G., Lee, J.H., Namini, H., and Heyward, W.L. (2012). Demonstration of safety and enhanced seroprotection against hepatitis B with investigational HBsAg-1018 ISS vaccine compared to a licensed hepatitis B vaccine. Vaccine 30, 2689–2696.
- 61. Janssen, J.M., Heyward, W.L., Martin, J.T., and Janssen, R.S. (2015). Immunogenicity and safety of an investigational hepatitis B vaccine with a Tolllike receptor 9 agonist adjuvant (HBsAg-1018) compared with a licensed hepatitis B vaccine in patients with chronic kidney disease and type 2 diabetes mellitus. Vaccine 33, 833–837.
- 62. Janssen, R.S., Mangoo-Karim, R., Pergola, P.E., Girndt, M., Namini, H., Rahman, S., Bennett, S.R., Heyward, W.L., and Martin, J.T. (2013). Immunogenicity and safety of an investigational hepatitis B vaccine with a Toll-like receptor 9 agonist adjuvant (HBsAg-1018) compared with a licensed hepatitis B vaccine in patients with chronic kidney disease. Vaccine 31, 5306–5313.
- Manoharan, M. (1999). 2'-Carbohydrate modifications in antisense oligonucleotide therapy: importance of conformation, configuration and conjugation. Biochim. Biophys. Acta 1489, 117–130.
- 64. Teplova, M., Minasov, G., Tereshko, V., Inamati, G.B., Cook, P.D., Manoharan, M., and Egli, M. (1999). Crystal structure and improved antisense properties of 2'-O-(2methoxyethyl)-RNA. Nat. Struct. Biol. 6, 535–539.
- 65. Lind, K.E., Mohan, V., Manoharan, M., and Ferguson, D.M. (1998). Structural characteristics of 2'-O-(2-methoxyethyl)-modified nucleic acids from molecular dynamics simulations. Nucleic Acids Res. 26, 3694–3699.
- 66. Pallan, P.S., Allerson, C.R., Berdeja, A., Seth, P.P., Swayze, E.E., Prakash, T.P., and Egli, M. (2012). Structure and nuclease resistance of 2',4'-constrained 2'-O-methoxyethyl (cMOE) and 2'-O-ethyl (cEt) modified DNAs. Chem. Commun. (Camb.) 48, 8195–8197.
- 67. Swayze, E.E., Siwkowski, A.M., Wancewicz, E.V., Migawa, M.T., Wyrzykiewicz, T.K., Hung, G., Monia, B.P., and Bennett, C.F. (2007). Antisense oligonucleotides containing locked nucleic acid improve potency but cause significant hepatotoxicity in animals. Nucleic Acids Res. 35, 687–700.
- 68. Kasuya, T., Hori, S., Watanabe, A., Nakajima, M., Gahara, Y., Rokushima, M., Yanagimoto, T., and Kugimiya, A. (2016). Ribonuclease H1-dependent hepatotoxicity caused by locked nucleic acid-modified gapmer antisense oligonucleotides. Sci. Rep. 6, 30377.

www.moleculartherapy.org

Review

- 69. Burdick, A.D., Sciabola, S., Mantena, S.R., Hollingshead, B.D., Stanton, R., Warneke, J.A., Zeng, M., Martsen, E., Medvedev, A., Makarov, S.S., et al. (2014). Sequence motifs associated with hepatotoxicity of locked nucleic acid-modified antisense oligonucleotides. Nucleic Acids Res. 42, 4882–4891.
- 70. van Poelgeest, E.P., Hodges, M.R., Moerland, M., Tessier, Y., Levin, A.A., Persson, R., Lindholm, M.W., Dumong Erichsen, K., Ørum, H., Cohen, A.F., and Burggraaf, J. (2015). Antisense-mediated reduction of proprotein convertase subtilisin/kexin type 9 (PCSK9): a first-in-human randomized, placebo-controlled trial. Br. J. Clin. Pharmacol. 80, 1350–1361.
- 71. Seth, P.P., Siwkowski, A., Allerson, C.R., Vasquez, G., Lee, S., Prakash, T.P., Wancewicz, E.V., Witchell, D., and Swayze, E.E. (2009). Short antisense oligonucleotides with novel 2'-4' conformationaly restricted nucleoside analogues show improved potency without increased toxicity in animals. J. Med. Chem. 52, 10–13.
- 72. Reilley, M.J., McCoon, P., Cook, C., Lyne, P., Kurzrock, R., Kim, Y., Woessner, R., Younes, A., Nemunaitis, J., Fowler, N., et al. (2018). STAT3 antisense oligonucleotide AZD9150 in a subset of patients with heavily pretreated lymphoma: results of a phase 1b trial. J. Immunother. Cancer 6, 119.
- Geary, R.S., Yu, R.Z., and Levin, A.A. (2001). Pharmacokinetics of phosphorothioate antisense oligodeoxynucleotides. Curr. Opin. Investig. Drugs 2, 562–573.
- Vosberg, H.P., and Eckstein, F. (1982). Effect of deoxynucleoside phosphorothioates incorporated in DNA on cleavage by restriction enzymes. J. Biol. Chem. 257, 6595– 6599.
- 75. Wan, W.B., Migawa, M.T., Vasquez, G., Murray, H.M., Nichols, J.G., Gaus, H., Berdeja, A., Lee, S., Hart, C.E., Lima, W.F., et al. (2014). Synthesis, biophysical properties and biological activity of second generation antisense oligonucleotides containing chiral phosphorothioate linkages. Nucleic Acids Res. 42, 13456–13468.
- 76. Koziolkiewicz, M., Krakowiak, A., Kwinkowski, M., Boczkowska, M., and Stec, W.J. (1995). Stereodifferentiation—the effect of P chirality of oligo(nucleoside phosphorothioates) on the activity of bacterial RNase H. Nucleic Acids Res. 23, 5000–5005.
- Yu, D., Kandimalla, E.R., Roskey, A., Zhao, Q., Chen, L., Chen, J., and Agrawal, S. (2000). Stereo-enriched phosphorothioate oligodeoxynucleotides: synthesis, biophysical and biological properties. Bioorg. Med. Chem. 8, 275–284.
- 78. Østergaard, M.E., De Hoyos, C.L., Wan, W.B., Shen, W., Low, A., Berdeja, A., Vasquez, G., Murray, S., Migawa, M.T., Liang, X.H., et al. (2020). Understanding the effect of controlling phosphorothioate chirality in the DNA gap on the potency and safety of gapmer antisense oligonucleotides. Nucleic Acids Res. 48, 1691–1700.
- 79. Stec, W.J., Grajkowski, A., Koziolkiewicz, M., and Uznanski, B. (1991). Novel route to oligo(deoxyribonucleoside phosphorothioates). Stereocontrolled synthesis of Pchiral oligo(deoxyribonucleoside phosphorothioates). Nucleic Acids Res. 19, 5883–5888.
- 80. Iwamoto, N., Butler, D.C.D., Svrzikapa, N., Mohapatra, S., Zlatev, I., Sah, D.W.Y., Meena, Standley, S.M., Lu, G., Apponi, L.H., et al. (2017). Control of phosphorothioate stereochemistry substantially increases the efficacy of antisense oligonucleotides. Nat. Biotechnol. 35, 845–851.
- Iwamoto, N., Oka, N., Sato, T., and Wada, T. (2009). Stereocontrolled solid-phase synthesis of oligonucleoside H-phosphonates by an oxazaphospholidine approach. Angew. Chem. Int. Ed. Engl. 48, 496–499.
- 82. Guo, M., Yu, D., Iyer, R.P., and Agrawal, S. (1998). Solid-phase stereoselective synthesis of 2'-O-methyl-oligoribonucleoside phosphorothioates using nucleoside bicyclic oxazaphospholidines. Bioorg. Med. Chem. Lett. 8, 2539–2544.
- 83. Oka, N., Yamamoto, M., Sato, T., and Wada, T. (2008). Solid-phase synthesis of stereoregular oligodeoxyribonucleoside phosphorothioates using bicyclic oxazaphospholidine derivatives as monomer units. J. Am. Chem. Soc. 130, 16031–16037.
- Aslesh, T., and Yokota, T. (2020). Development of antisense oligonucleotide gapmers for the treatment of Huntington's disease. Methods Mol. Biol. 2176, 57–67.
- 85. Hung, G., Xiao, X., Peralta, R., Bhattacharjee, G., Murray, S., Norris, D., Guo, S., and Monia, B.P. (2013). Characterization of target mRNA reduction through in situ RNA hybridization in multiple organ systems following systemic antisense treatment in animals. Nucleic Acid Ther. 23, 369–378.
- 86. Echigoya, Y., Nakamura, A., Nagata, T., Urasawa, N., Lim, K.R.Q., Trieu, N., Panesar, D., Kuraoka, M., Moulton, H.M., Saito, T., et al. (2017). Effects of systemic

multiexon skipping with peptide-conjugated morpholinos in the heart of a dog model of Duchenne muscular dystrophy. Proc. Natl. Acad. Sci. USA *114*, 4213-4218.

- Levin, A.A. (2017). Targeting therapeutic oligonucleotides. N. Engl. J. Med. 376, 86–88.
- 88. Scott, L.J. (2020). Givosiran: first approval. Drugs 80, 335-339.
- 89. Sugo, T., Terada, M., Oikawa, T., Miyata, K., Nishimura, S., Kenjo, E., Ogasawara-Shimizu, M., Makita, Y., Imaichi, S., Murata, S., et al. (2016). Development of antibody-siRNA conjugate targeted to cardiac and skeletal muscles. J. Control. Release 237, 1–13.
- 90. Prakash, T.P., Graham, M.J., Yu, J., Carty, R., Low, A., Chappell, A., Schmidt, K., Zhao, C., Aghajan, M., Murray, H.F., et al. (2014). Targeted delivery of antisense oligonucleotides to hepatocytes using triantennary *N*-acetyl galactosamine improves potency 10-fold in mice. Nucleic Acids Res. 42, 8796–8807.
- 91. Tsimikas, S., Karwatowska-Prokopczuk, E., Gouni-Berthold, I., Tardif, J.C., Baum, S.J., Steinhagen-Thiessen, E., Shapiro, M.D., Stroes, E.S., Moriarty, P.M., Nordestgaard, B.G., et al.; AKCEA-APO(a)-LRx Study Investigators (2020). Lipoprotein(a) reduction in persons with cardiovascular disease. N. Engl. J. Med. 382, 244–255.
- 92. Jirka, S.M., Heemskerk, H., Tanganyika-de Winter, C.L., Muilwijk, D., Pang, K.H., de Visser, P.C., Janson, A., Karnaoukh, T.G., Vermue, R., 't Hoen, P.A., et al. (2014). Peptide conjugation of 2'-O-methyl phosphorothioate antisense oligonucleotides enhances cardiac uptake and exon skipping in mdx mice. Nucleic Acid Ther. 24, 25–36.
- 93. Lemaitre, M., Bayard, B., and Lebleu, B. (1987). Specific antiviral activity of a poly(L-lysine)-conjugated oligodeoxyribonucleotide sequence complementary to vesicular stomatitis virus N protein mRNA initiation site. Proc. Natl. Acad. Sci. USA 84, 648–652.
- **94.** Guidotti, G., Brambilla, L., and Rossi, D. (2017). Cell-penetrating peptides: from basic research to clinics. Trends Pharmacol. Sci. *38*, 406–424.
- Raucher, D., and Ryu, J.S. (2015). Cell-penetrating peptides: strategies for anticancer treatment. Trends Mol. Med. 21, 560–570.
- 96. Juliano, R.L. (2005). Peptide-oligonucleotide conjugates for the delivery of antisense and siRNA. Curr. Opin. Mol. Ther. 7, 132–136.
- 97. Gait, M.J., Arzumanov, A.A., McClorey, G., Godfrey, C., Betts, C., Hammond, S., and Wood, M.J.A. (2019). Cell-penetrating peptide conjugates of steric blocking oligonucleotides as therapeutics for neuromuscular diseases from a historical perspective to current prospects of treatment. Nucleic Acid Ther. 29, 1–12.
- 98. Jearawiriyapaisarn, N., Moulton, H.M., Buckley, B., Roberts, J., Sazani, P., Fucharoen, S., Iversen, P.L., and Kole, R. (2008). Sustained dystrophin expression induced by peptide-conjugated morpholino oligomers in the muscles of mdx mice. Mol. Ther. 16, 1624–1629.
- 99. Yin, H., Saleh, A.F., Betts, C., Camelliti, P., Seow, Y., Ashraf, S., Arzumanov, A., Hammond, S., Merritt, T., Gait, M.J., and Wood, M.J. (2011). Pip5 transduction peptides direct high efficiency oligonucleotide-mediated dystrophin exon skipping in heart and phenotypic correction in mdx mice. Mol. Ther. 19, 1295–1303.
- 100. Scharner, J., Figeac, N., Ellis, J.A., and Zammit, P.S. (2015). Ameliorating pathogenesis by removing an exon containing a missense mutation: a potential exon-skipping therapy for laminopathies. Gene Ther. 22, 503–515.
- 101. Pendse, N.D., Lamas, V., Pawlyk, B.S., Maeder, M.L., Chen, Z.Y., Pierce, E.A., and Liu, Q. (2019). In vivo assessment of potential therapeutic approaches for USH2A-associated diseases. Adv. Exp. Med. Biol. 1185, 91–96.
- 102. Lim, K.H., Han, Z., Jeon, H.Y., Kach, J., Jing, E., Weyn-Vanhentenryck, S., Downs, M., Corrionero, A., Oh, R., Scharner, J., et al. (2020). Antisense oligonucleotide modulation of non-productive alternative splicing upregulates gene expression. Nat. Commun. 11, 3501.
- 103. Han, Z., Chen, C., Christiansen, A., Ji, S., Lin, Q., Anumonwo, C., Liu, C., Leiser, S.C., Meena, Aznarez, I., et al. (2020). Antisense oligonucleotides increase *Scn1a* expression and reduce seizures and SUDEP incidence in a mouse model of Dravet syndrome. Sci. Transl. Med. *12*, eaaz6100.