

The notorious R.N.A. in the spotlight - drug or target for the treatment of disease

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ABSTRACT

mRNA is an attractive drug target for therapeutic interventions. In this review we highlight the current state, clinical trials, and developments in antisense therapy, including the classical approaches like RNaseH-dependent oligomers, splice-switching oligomers, aptamers, and therapeutic RNA interference. Furthermore, we provide an overview on emerging concepts for using RNA in therapeutic settings including protein replacement by in-vitro-transcribed mRNAs, mRNA as vaccines and anti-allergic drugs. Finally, we give a brief outlook on early-stage RNA repair approaches that apply endogenous or engineered proteins in combination with short RNAs or chemically stabilized oligomers for the re-programming of point mutations, RNA modifications, and frame shift mutations directly on the endogenous mRNA.

Abbreviations: ASO, Antisense oligonucleotide; CD, Cluster of differentiation; CFTR, Cystic fibrosis transmembrane conductance regulator; CRISPR/Cas9, Clustered regularly interspaced short palindromic repeats/CRISPR-associated 9; FDA, US Food and Drug Administration; GalNAc, N-acetyl galactosamine; IVT-mRNA, In-vitro transcribed mRNA; MHC, Major histocompatibility complex; miRNA, microRNA; MOE, 2'-O-methoxyethyl; mRNA, messenger RNA; ψ , pseudouridine; PS, Phosphothioate; RNAi, RNA interference; siRNA, Short interfering RNA; SSO, Splice-switching oligonucleotide; SMN2, Survival of motor neuron 2; TALEN, Transcription activator-like effector nuclease; TLR, Toll-like receptor; $T_H1/2$ cell, Type 1/2 T helper cell; T_R1 , Type 1 regulatory T cell; VEGF, Vascular endothelial growth factor; VEGFR-1, Vascular endothelial growth factor receptor 1; ZFN, Zinc finger nuclease

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Introduction

During the last 15 y the diverse roles of RNA in regular but also pathological cellular processes became increasingly clear. RNA is not only a short-lived messenger and part of the translational machinery but RNA contributes significantly to the regulation and diversification of the genetic information. There is now increasing insight into the mechanistic role of defective RNA processing, including (alternative) splicing, modification, translation, and decay for the etiology of various diseases.^{1–4} However, not only mis-regulation and defective processing cause disease, but even RNA species themselves can initiate disease processes independent of their protein-coding function. Nucleotide repeat diseases are typical examples.⁵ To employ this new mechanistic knowledge and to translate it into therapy requires drugs that reliably target nucleic acids in a sequence-specific manner. However, there are only few small molecule drugs that target nucleic acids and those are limited in their capacity of sequence addressing. In contrast, oligonucleotide analogs provide a basis for the rational design of highly sequence-specific drugs to target virtually any cellular nucleic acid in a specific manner.⁶ Classical drugs like small molecules target enzymes and receptors to block or alter their specific functions. In contrast, the interference at the nucleic acid level would allow to manipulate the transcriptome and the proteome itself. This is

not limited to the simple up- or down-regulation of target gene expression. Most appealing is the possibility of actively creating new transcript and protein isoforms with altered properties and functions, for instance by re-programming a protein-coding stretch, or by altering splice sites, modification patterns, polyadenylation states, miRNA binding sites, etc.⁷ Affecting the cell by targeting its nucleic acids clearly enlarges the scope of currently available therapeutic interventions including the causal treatment of some genetic diseases.

However, already short oligonucleotides have unfavorable pharmacological properties. They are hydrophilic, polyanionic macromolecules that can hardly overcome cellular membranes, are unstable against RNases, and suffer from rapid renal clearance.⁸ This leads to short half-life and low bioavailability. Furthermore, adverse toxic effects may appear that include immune-reactions and off-target binding to non-targeted cellular nucleic acids. Together, oligonucleotide drugs are often characterized by low efficacy and high toxicity which strongly limits their clinical application.⁶ During the last decades, medicinal chemists have put enormous effort into the development of new chemistries that improve lifetime, delivery, potency, and efficacy of the drugs while reducing their toxicity and immunogenicity. These new chemistries are now approaching clinical trials and will hopefully pave the way for

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the broad clinical application of oligonucleotide drugs. An overview on recent developments in oligonucleotide medicinal chemistry can be found elsewhere.^{6,7}

In principle, interference with the genetic information could be achieved permanently at the DNA- or transiently at the RNA-level. In this review we will focus on the RNA-level. Even though novel approaches for genome engineering are currently keenly explored,⁹ we believe that it would be foolish to carelessly discard the RNA alternative. With respect to ethical issues and safety aspects, the transient and thus reversible nature of RNA manipulation could turn out as a blessing in disguise. Both, the therapeutic effects and the potential adverse effects, are likely to be tunable. Furthermore, manipulations are conceivable that are inaccessible or difficult to realize on the genome level per se. This includes amino acid changes or transcript level changes that would kill a cell if they are permanently enforced. Potentially lethal interventions on kinases, apoptosis

factors, transcription or translation factors could be realized on the RNA-level suddenly, transiently or partially to obtain a therapeutic effect, for instance. Manipulation at the RNA-level might also be much more efficient compared to HDR-dependent genomic knock-in, which remained persistently inefficient in vivo, in particular in postmitotic tissues like the brain.⁹ For many genetic diseases, which are caused by loss-of-function mutations, a patient would benefit more from a drug that can restore a small fraction (like 5%) of functional gene product in a large fraction of a the tissue than from a drug that can restore full gene function (100%) but only in a small fraction of the tissue. A typical example is cystic fibrosis.¹⁰

In this review we will first update on recent developments in the classical approaches, like RNaseH-dependent decay, chemically stabilized oligonucleotides that target mRNAs to induce splice-switching, aptamers, and the knock-down via RNAi (Fig. 1). After painful years of repeated relapse one seems to

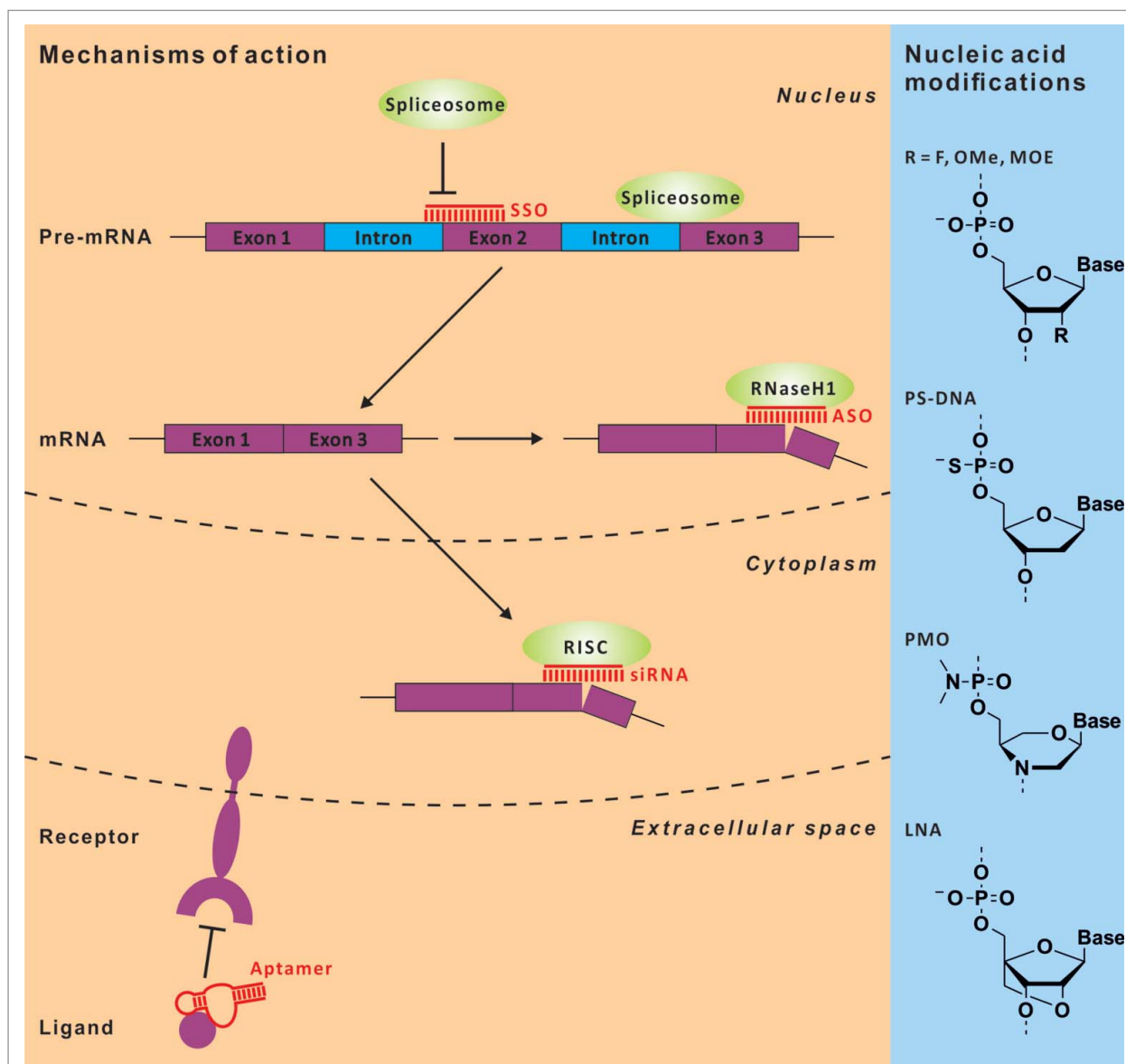


Figure 1. Chemically stabilized, short oligonucleotides can employ various mechanisms for their therapeutic effects ranging from blocking ligand – receptor binding, RNA degradation via RISC or RNaseH(1) recruitment, and alteration of splicing. The classical modes of action are shown on the left panel, a small section of typically used chemical backbone modifications are depicted on the right.

have learned the lessons and have now substantially improved the effectiveness of such drugs. For instance, in 2015 therapeutic RNAi was demonstrated in a relevant monkey model by subcutaneous administration of a chemically stabilized siRNA that partially knocks down antithrombin in the monkey's liver.¹¹ The problem of delivery and toxicity seems to be solved, at least for simple oligonucleotide drugs and for some organs, and allows therapeutic intervention with an affordable amount of the drug under compliant administration routes. Consequently, the number of promising clinical phase II and III studies has increased during the last few years (see Table 1).

Every new discovery in RNA function and regulation offers a starting point to develop novel therapies. After its discovery in 1998 we now find numerous drug candidates in clinical studies that apply the RNAi mechanism (Table 1).¹² In the second part of this review we highlight emerging concepts that are still in the pre-clinical or very early clinical exploration stage but that have the potential to become medicines of the future. This includes therapeutic mRNAs, mRNAs as vaccine, and RNA repair approaches. The latter apply endogenous or engineered enzymes to repair, re-program, or modify a target RNA at a specific site in order to provoke a therapeutically relevant effect (Fig. 2).

Update on established approaches

RNaseH-dependent antisense oligonucleotides

Oligonucleotides working through an RNaseH-dependent cleavage mechanism are the oldest class of antisense oligonucleotides (ASO). They are extensively explored and represent the largest class of nucleic acid analog drugs in clinical trials. RNaseH-dependent ASOs are short DNA oligomers targeting mRNA. Once the DNA-oligo/mRNA heteroduplex is formed, human RNaseH1 binds to it and catalyzes RNA cleavage under release of the intact DNA oligomer.¹³

Medicinal chemists have undertaken great efforts to improve ASO design regarding nuclease resistance, circulation half-life, target affinity (potency), and tissue specificity. The first ASOs tested in clinical trials, also referred to as 1st generation ASOs, have been modified by oxygen-to-sulfur substitutions in the phosphate backbone. ASOs with such a phosphothioate (PS) backbone show enhanced nuclease resistance and prolonged plasma half-life due to non-specific binding to plasma proteins preventing them from rapid renal filtration. However, numerous toxicities were also associated with that type of modification.⁶ In 1998, fomivirsen was the first FDA-approved ASO and was applied for the treatment of human cytomegalovirus-induced retinitis in HIV patients.¹⁴⁻¹⁶ Marketed as Vitravene, the 21 nt PS-oligonucleotide was administered by intravitreal injection to target the immediate early region 2 of the viral mRNA. Since the approval of fomivirsen, several ASOs belonging to the 1st generation are under clinical review. For instance, targeting the mRNA of intercellular adhesion molecule 1 and the insulin receptor substrate 1 are advanced in the treatment of pouchitis^{17,18} and vascular disorders in the eye,¹⁹⁻²² respectively. The RNaseH-mediated degradation of Akt-1 mRNA to impede tumor proliferation²³ is currently tested for clinical application.²⁴⁻²⁶

Due to the early success with 1st generation ASO, further medicinal chemistry was explored to improve half-life and potency of the drugs in order to reduce the administered dose, the application frequency, the costs, and to minimize adverse effects.²⁷ This resulted in the 2nd generation ASOs, also referred to as gapmers. A typical gapmer is a 20 nt oligonucleotide comprising a PS backbone and 5 flanking 2'-O-methoxyethyl (MOE) groups at both termini. Due to the unmodified internal DNA gap, such ASOs remain good substrates for RNaseH, whereas the terminal MOE modifications increase nuclease resistance and enhances the binding of the ASO to the target mRNA.²⁸ 2nd generation ASOs entered clinical trials for various therapeutic applications. The most prominent representative of the 2nd generation is the MOE gapmer mipomersen as the second FDA-approved RNaseH-dependent ASO. The compound targets apolipoprotein B-100 mRNA and is subcutaneously administered to treat familiar hypercholesterolemia. The genetic disorder is caused by the loss of low-density lipoprotein (LDL) receptor function leading to high LDL cholesterol plasma concentration and early cardiovascular disease. Phase III trials had demonstrated an efficient decrease of LDL cholesterol by lowering ApoB-100 amount in patients obtaining mipomersen.²⁹⁻³¹ The treatment obviously profited from the general pharmacokinetics of systemically administered ASOs which preferably accumulate in the liver where ApoB-100 synthesis takes place.⁸ Recently, an RNase-dependent ASO³² has reached clinical phase III to reduce transthyretin expression in patients suffering from familial amyloid polyneuropathy.³³⁻³⁵ Chemotherapy combined with RNaseH-mediated degradation of clusterin mRNA is a potential therapeutic option in the treatment of prostate³⁶⁻³⁸ and lung cancer.^{39,40}

Generation 2.5 ASO are derived from the traditional gapmer design. For this, the MOE modifications are replaced by 2',4'-constrained ethyl (cEt) bridges in the flanking nucleotides. It was found that cEt-modified oligonucleotides provide the same superior target affinity, but increased nuclease resistance as compared to locked nucleic acid (LNA)-containing oligonucleotides.⁴¹ One of the generation 2.5 ASOs targets the mRNA of signal transducer and activator of transcription 3⁴² and is currently tested for the treatment of various cancer types.⁴³⁻⁴⁶

Most recently, a new chemistry has been developed that strongly increases the liver-specific uptake of oligonucleotide drugs, including ASO and siRNA therapeutics. For this, ASOs⁴⁷ and siRNAs⁴⁸ are conjugated with triantennary N-acetyl galactosamine (GalNAc₃). GalNAc₃ mediates liver-specific uptake through the asialoglycoprotein receptor (ASGPR) that is exclusively expressed on hepatocytes. Marketed as ligand-conjugated antisense (LICA) technology (Ionis Pharmaceuticals), it could be shown that the conjugation increases the potency of MOE gapmers up to 10-fold for inhibiting the expression of hepatic genes in mice.⁴⁹ When using a GalNAc₃-conjugated cEt gapmer, the RNaseH-mediated mRNA degradation was enhanced around 60-fold as compared to the corresponding 2nd generation MOE ASO. Additionally, Ionis Pharmaceuticals announced that its LICA drug targeting apolipoprotein(a) was 30-fold more potent in a phase I study than the unconjugated MOE gapmer.^{50,51}

Table 1. Overview on the most recent and advanced clinical trials in the corresponding fields not claiming completeness. For RNAi, failed early trials are also listed that have been terminated in 2009 or before.

Table 1: Clinical Trials using RNA-Therapeutics*						
RNaseH-dependent Antisense Oligonucleotides						
Compound	Application Route	Target	Indication	Company / Initiator	Phase / Status	Trial ID / Reference
Fomivirsen (Vitravene® , ISIS 2922)	IVT	IE2	CMV retinitis	Ionis Pharmaceuticals /Ciba Vision	FDA Approved	14-16
Mipomersen (KYNAMRO® , ISIS 301012)	SC	ApoB-100	HoFH	Ionis Pharmaceuticals / Genzyme	FDA Approved	29-31
Custirsen (OGX-011)	IV	Clusterin	Prostate cancer	Ionis Pharmaceuticals / OncoGenex Technologies	III - Completed	NCT01188187 ³⁶
Custirsen (OGX-011)	IV	Clusterin	Prostate cancer	Ionis Pharmaceuticals / OncoGenex Technologies	III - Active	NCT01578655 ³⁷
Custirsen (OGX-011)	IV	Clusterin	Lung cancer	Ionis Pharmaceuticals / OncoGenex Technologies	III - Recruiting	NCT01630733 ⁴⁰
Aganirsen (GS-101)	Eye drops	IRS-1	Corneal neovascularization in graft patients	Gene Signal	III - Completed	EudraCT 2008-005388-33 ²¹
Aganirsen (GS-101)	Eye drops	IRS-1	Ischemic central retinal vein occlusion	Gene Signal	II/III - Recruiting	EudraCT 2014-000239-18 ²²
Alicaforsen (ISIS 2302)	Enema	ICAM-1	Pouchitis	Ionis Pharmaceuticals / Atlantic Healthcare	III - Recruiting	NCT02525523 ¹⁸
IONIS-TTRRx (ISIS 420915)	SC	TTR	FAP, TRR Amyloidosis	Ionis Pharmaceuticals / GlaxoSmithKline	II/III - Active	NCT01737398 ³³
IONIS-TTRRx (ISIS 420915)	SC	TTR	FAP, TRR Amyloidosis	Ionis Pharmaceuticals / GlaxoSmithKline	III - Enrolling by invitation	NCT02175004 ³⁴
RX-0210 (Arhexin®)	IV	Akt-1	Pancreatic cancer	Rexahn Pharmaceuticals	II - Completed	NCT01028495 ²⁴
RX-0210 (Arhexin®)	IV	Akt-1	Renal cell cancer	Rexahn Pharmaceuticals	IB/II - Recruiting	NCT02089334 ²⁵
IONIS-STAT3-2.5Rx (AZD9150, ISIS 481464)	IV	STAT3	Gastrointestinal or ovarian cancer	Ionis Pharmaceuticals / AstraZeneca	II - Recruiting	NCT02417753 ⁴³
IONIS-STAT3-2.5Rx (AZD9150, ISIS 481464)	IV	STAT3	Metastatic Squamous Cell Carcinoma of the Head and Neck	Ionis Pharmaceuticals / AstraZeneca	IB/II - Recruiting	NCT02499328 ⁴⁴
IONIS-STAT3-2.5Rx (AZD9150, ISIS 481464)	IV	STAT3	Lymphoma	Ionis Pharmaceuticals / AstraZeneca	I/II - Active	NCT01563302 ⁴⁵
IONIS-STAT3-2.5Rx (AZD9150, ISIS 481464)	IV	STAT3	Lymphoma	Ionis Pharmaceuticals / AstraZeneca	IB - Not yet recruiting	NCT02549651 ⁴⁶
IONIS-APO(a)-LRx (ISIS 681257)	SC	ApoA	Elevated Lipoprotein(a)	Ionis Pharmaceutical / Akcea Therapeutics	I - Completed	NCT02414594 ⁵¹
Splice-switching Oligonucleotides						
Compound	Application Route	Target	Indication	Company / Initiator	Phase / Status	Trial ID / Reference
PRO051/GSK2402968 (drisapersen/ kyndrisa®)	SC	Dystrophin exon 51 (skipping)	DMD	BioMarin Pharmaceuticals (developed by Prosenza Therapeutics)/GlaxoSmithKline)	III - Recruiting	NCT01803412 ⁵⁶
PRO051/GSK2402968 (drisapersen/ kyndrisa®)	SC	Dystrophin exon 51 (skipping)	DMD	BioMarin Pharmaceuticals (developed by Prosenza Therapeutics)/GlaxoSmithKline)	IIIB - Enrolling by invitation	NCT02636686 ⁵⁷
PRO044/BMN 044	SC/IV	Dystrophin exon 44 (skipping)	DMD	BioMarin Pharmaceuticals (developed by Prosenza Therapeutics)/GlaxoSmithKline)	II - Enrolling by invitation	NCT02329769 ²⁴⁵
PRO045/BMN 045	SC	Dystrophin exon 45 (skipping)	DMD	BioMarin Pharmaceuticals (developed by Prosenza Therapeutics)	IIIB - Active	NCT01826474 ²⁴⁶
PRO053/BMN 053	SC/IV	Dystrophin exon 53 (skipping)	DMD	BioMarin Pharmaceuticals (developed by Prosenza Therapeutics)	I/II - Active	NCT01957059 ²⁴⁷

Compound	Application Route	Target	Indication	Aptamers Indication	Company / Initiator	Phase / Status	Trial ID / Reference
AVI-4685 (eteplirsen)	IV	Dystrophin exon 51 (skipping)	DMD	DMD	Sarepta Therapeutics	III - Recruiting	NCT02255552 ⁵⁹
SRP-4045	IV	Dystrophin exon 45 (skipping)	DMD	DMD	Sarepta Therapeutics	III - Not yet recruiting	NCT02500381 ²⁴⁸
SRP-4053	IV	Dystrophin exon 53 (skipping)	DMD	DMD	Sarepta Therapeutics	III - Not yet recruiting	NCT02500381 ²⁴⁸
DS-5141b	SC	Dystrophin exon 45 (skipping)	DMD	DMD	Daiichi Sankyo / Orphan Disease Treatment Institute	I/II - Recruiting	NCT02667483 ⁶⁵
NS-065/NCNP-01	IV	Dystrophin exon 53 (skipping)	DMD	DMD	Nippon Shinyaku / National Center of Neurology and Psychiatry	II	not yet registered ⁶²
IONIS-SMNRx / ISIS 396443 / ASO-10-27 (nusinersen)	IT	SNM2 exon 7 (retention)	SMA	SMA	Ionis Pharmaceuticals / Biogen	III - Recruiting	NCT02193074 ⁷⁰
IONIS-SMNRx / ISIS396443 / ASO-10-27 (nusinersen)	IT	SNM2 exon 7 (retention)	SMA	SMA	Ionis Pharmaceuticals / Biogen	III - Active	NCT02292537 ⁷¹
Compound	Application Route	Target	Indication	Aptamers Indication	Company / Initiator	Phase / Status	Trial ID / Reference
Pegaptanib sodium (Macugen [®])	IVT	VEGF	AMD	AMD	Evetech Pharmaceuticals, Pfizer	FDA - Approved	^{75,76}
RB006 (pegnivacogin) as component of REG1	IV	Coagulation factor IXa	CAD	CAD	Regado Bioscience	III - Terminated	NCT01848106 ⁷⁸
E10030 (Fovista [®])	IVT	PDGF	AMD	AMD	Ophthotech	III - Recruiting	NCT01944839 ⁸⁰ NCT01940887 ⁸¹
ARC1905 (Zimura [®])	IVT	complement component C5	AMD	AMD	Ophthotech	I/III - Recruiting	NCT02686658 ⁸³
NOX-A12 (olaptesed pegol)	IV	CXCL12/SDF-1	MM	MM	Noxzon Pharma	IIA - Completed	NCT01521533 ⁸⁸
NOX-A12 (olaptesed pegol)	IV	CXCL12/SDF-1	CLL	CLL	Noxzon Pharma	IIA - Active	NCT01486797 ⁸⁹
NOX-E36 (emapticap pegol)	IV	CCL2/MCP-1	DM2, Albuminuria	DM2, Albuminuria	Noxzon Pharma	IIA - Completed	NCT01547897 ⁹¹
NOX-H94 (lexaptapid pegol)	IV	Hepcidin	Anemia	Anemia	Noxzon Pharma	IIA - Completed	NCT01691040 ⁹²
Compound	Application Route	Target	Indication	RNA Interference	Company / Initiator	Phase / Status	Trial ID / Reference
Bevasiranib	IVT	VEGF	Macular Degeneration	Macular Degeneration	OPKO Health	III - Terminated	NCT00499590 ¹⁰¹
AGN 211745	IVT	VEGFR-1	CNV, AMD	CNV, AMD	Allergan	II - Terminated	NCT00395057 ¹⁰⁴
ISNP	IVT	p53	AKI	AKI	Quark Pharmaceuticals	I - Terminated	NCT00683553 ¹⁰⁰
PF-04523655	IVT	DDIT4	Diabetic Retinopathy, Diabetes Complications	Diabetic Retinopathy, Diabetes Complications	Quark Pharmaceuticals	II - Terminated	NCT00701181 ¹⁰²
PRO-040201	IV	ApoB	Hypercholesterolemia	Hypercholesterolemia	Arbutus Biopharma	I - Terminated	NCT00927459 ¹⁰³
ND-L02-s0201	IV	HSP47	Hepatic Fibrosis	Hepatic Fibrosis	Nitto Denko	I - Ongoing	NCT02227459 ¹¹⁴
ALN-AS1	SC	ALAS1	AIP	AIP	Alnylam Pharmaceuticals	I - Recruiting	NCT02452372 ¹¹⁶
pbi-shRNA STMN1 LP	IT	STMN1	Advanced / Metastatic Cancer, Solid Tumors	Advanced / Metastatic Cancer, Solid Tumors	Gradalis	I - Ongoing	NCT01505153 ¹¹⁹
Bamosiran	Eye drops	ADRB2	OAG, Ocular Hypertension	OAG, Ocular Hypertension	Sylentis	II - Completed	NCT02250612 ¹²¹
ARB-001467	IV	Three sites of the HBV genome	Chronic Hepatitis B	Chronic Hepatitis B	Arbutus Biopharma	II - Recruiting	NCT02631096 ¹¹⁵
ALN-TTRsc/ reusiran	SC	TTR	FAC	FAC	Alnylam Pharmaceuticals	III - Recruiting	NCT02319005 ¹¹⁷
ALN-A13	SC	AT	Hemophilia	Hemophilia	Alnylam Pharmaceuticals	I - Recruiting	NCT02035605 ¹¹⁸
siRNA-transfected PBMCs (APN401)	IV	Cbl-b	Several types of cancer	Several types of cancer	Comprehensive Cancer Center of Wake Forest University	I - Recruiting	NCT02166255 ¹²⁰
QPI-1007	IVT	CASP2	NAION	NAION	Quark Pharmaceuticals	II/III - Recruiting	NCT02341560 ¹²²
MRX34	IV	miRNA34a	Several types of cancer	Several types of cancer	Mirna Therapeutics	I - Recruiting	NCT01829971 ¹²⁸
Targoirs	IV	miRNA16	Malignant Pleural Mesothelioma	Malignant Pleural Mesothelioma	University of Sydney	I - Recruiting	NCT02369198 ¹²⁷
AZD4076	SC	miRNA103/107	NASH	NASH	AstraZeneca	I - Recruiting	NCT02612662 ²⁴⁹
MiraVirsen	SC	miRNA122	Hepatitis C	Hepatitis C	Santaris Pharma A/S	II - Ongoing	NCT02031133 ¹²⁵

(continued)

Table 1. (Continued)

Compound	Application Route	Target	mRNA Therapy		Company / Initiator	Phase / Status	Trial ID / Reference
			Indication	Indication			
CV9104	ID	—	Prostate Cancer	Prostate Cancer	CureVac	II - Ongoing	NCT01817738 ²⁰⁸
CV9202	ID	—	Stage IV NSCLC	Stage IV NSCLC	CureVac	I - Ongoing	NCT01915524 ²⁰⁶
CV7201	IM	—	Rabies	Rabies	CureVac	I - Ongoing	NCT02241135 ²⁰⁴
CV9104	ID	—	Prostate Carcinoma	Prostate Carcinoma	CureVac	II - Ongoing	NCT02140138 ²⁰⁹
IVAC_W_bre1_uid	Not available	—	TNBC	TNBC	BioNtech	I - Not yet Recruiting	NCT02316457 ²⁵⁰
IVAC_W_bre1_uid/IVAC_M_uid	—	—	—	—	—	—	—
IVAC MUTANOME RBL001/RBL002	IN	—	Melanoma	Melanoma	BioNtech	I - Ongoing	NCT02035956 ²¹⁰
Lipo-MERIT	IV	—	Melanoma	Melanoma	BioNtech	I - Recruiting	NCT02410733 ¹⁷²
AGS-003	ID	—	mRCC	mRCC	Argos Therapeutics	III - Ongoing	NCT01582672 ¹⁹⁶
Dendritic cell vaccine (plus temozolomide chemotherapy)	ID	—	GBM	GBM	University Hospital, Antwerp	II - Recruiting	NCT02649582 ²⁰⁰
Dendritic cell vaccine (plus chemotherapy)	IV	—	MPM	MPM	University Hospital, Antwerp	II - Recruiting	NCT02649829 ²⁰¹
pp65 Dendritic cell vaccine	SC	—	GBM	GBM	University of Florida	II - Not yet Recruiting	NCT02465268 ²⁰²
ZFN Modified CD4+ T Cells	Infusion	CCR5	HIV	HIV	University of Pennsylvania	I - Recruiting	NCT02388594 ²⁵¹

Abbreviations of targets, indications and administration routes: ADRB2: Adrenergic receptor $\beta 2$, AIP: Acute intermittent porphyria, AKI: Acute kidney injury, ALAS: ALA-synthase, AMD: Age-related macular degeneration, ApoA/B: Apolipoprotein A/B, ApoB-100: Apolipoprotein B-100, AT: Antithrombin, CAD: Coronary artery disease, CASP2: Caspase 2, Cbl-b: Casitas B-lineage lymphoma proto-oncogene B, CCL2/MCP-1: CC-chemokine ligand 2/monocyte chemoattractant protein 1, CMV: Cytomegalovirus, CNV: Choroid neovascularization, CLL: Chronic lymphocytic leukemia, CXCL-12: C-X-C motif chemokine ligand 12, DDIT4: DNA damage-inducible transcript 4, DMD: Duchenne muscular dystrophy, FAP: Familial amyloid polyneuropathy, FAC: Familial amyloidotic cardiomyopathy, GBM: Glioblastoma multiforme, HSP47: Heat shock protein 47, HoFH: Homozygous familial hypercholesterolemia, HIV: Human immunodeficiency virus, ICAM-1: Intercellular adhesion molecule 1, ID: intra-dermal, IE2: Immediate early 2 protein, IM: intra-muscular, IN: intra-nodal, IRS-1: Insulin receptor substrate 1, IT: intra-tumoral, IV: intra-venous, IVT: intra-vitreous, MPM: Malignant pleural mesothelioma, mRCC: Metastatic renal cell carcinoma, MM: Multiple myeloma, Akt-1: Murine thymoma viral oncogene homolog 1, NASH: Non-alcoholic steatohepatitis, NAION: Nonarteritic anterior ischemic optic neuropathy, NSCLC: Non-small cell lung cancer, OAG: Open angle glaucoma, PDGF: Platelet-derived growth factor, SDF-1: Stromal cell-derived factor 1, STAT3: Signal transducer and activator of transcription 3, SMA: Spinal muscular atrophy, STMN1: Stathmin-1, SC: Sub-Cutaneous, TTR: Transthyretin, DM2: Type 2 diabetes mellitus, VEGF: Vascular endothelial growth factor, VEGFR-1: Vascular endothelial growth factor receptor 1.

Splice-switching oligonucleotides

Pre-mRNA is matured during a complex nuclear process called splicing that removes the introns (non-coding sequences) and joins the exons (coding sequences). By applying alternative splice sites and by occasional inclusion or exclusion of exons and introns, multiple protein variants are derived from one gene (alternative splicing). Several diseases are related to aberrant RNA-splicing leading to non-functional proteins, and great efforts have been undertaken to develop antisense oligonucleotides, referred to as splice-switching oligonucleotides (SSOs) that manipulate splicing. Therapeutic SSOs promoting exon skipping and exon retention for the treatment for Duchenne muscular dystrophy (DMD) and spinal muscular atrophy (SMA) are currently evaluated in clinical trials.⁷

Dystrophin, the protein encoded by the DMD gene, is crucial for the integrity of muscle tissue.⁵² In rare cases, newborn males harbor a defect dystrophin gene on their X chromosome. The patients suffer from successive muscle wasting resulting in a premature death due to respiratory or cardiac failure. In most cases, the loss-of-protein-function results from exonic out-of-frame deletions. In many cases the reading frame can be restored by skipping the aberrant exon by addressing a SSO to an internal exonic splicing enhancer.⁵³ The resulting truncated dystrophin protein retains partial function and gives the less severe Becker muscular dystrophy phenotype.⁵⁴ Several SSOs have been developed that are clinically evaluated for the skipping of exons 44, 45, 51, and 53, including drisapersen and eteplirsen (Table 1). Recently, both companies submitted new drug applications for their lead compounds drisapersen⁵⁵⁻⁵⁷ and eteplirsen,^{58,59} both amenable to exon 51 skipping. In case of drisapersen, the FDA rejected the application due to major concerns about the efficacy and safety of the drug.⁶⁰ The high dosage required led to severe adverse effects including renal and vascular injury. To improve efficacy and safety other SSO chemistries might be more successful. Whereas drisapersen is a 20 nt 2'-O-methoxy phosphorothioate RNA analog, eteplirsen is a 30 nt phosphorodiamidate oligomer, a so-called morpholino. The final decision on the efficacy and safety evaluation by the FDA is still pending for eteplirsen. Additionally, a new, morpholino-based SSO for exon 53 skipping is currently under clinical evaluation (NS-065/NCNP-01).^{61,62} For the future, we can hope in new chemistries. A SSO that relies on 2',4'-C-ethylene-bridged nucleosides (ENA oligonucleotides)⁶³ which mediate nuclease resistance and improved binding affinity to RNA has now entered a clinical phase I/II trial for the treatment of DMD (DS-4151b).^{64,65}

Spinal muscular atrophy (SMA) is a rare genetic disorder caused by survival of motor neuron 1 (SMN1) gene mutations.⁶⁶ Infant patients affected by this disease suffer from the loss of motor neurons and associated muscle wasting. However, there is a therapeutic approach by activating the SMN2 gene, which is almost identical to SMN1, but a single mutation in a splicing enhancer strongly prevents the inclusion of exon 7 resulting in an unstable protein unable to replace the lost SMN1 function.⁶⁷ In a mouse model, a highly potent 2'-O-methoxyethyl PS SSO for exon 7 retention in SMN2 was identified (IONIS-SMN_{Rx}).⁶⁸ The drug is injected in the spinal cord ensuring the direct delivery to the affected motor neurons without the need to cross the blood-brain barrier. After promising clinical phase II results regarding efficacy

and safety of the drug candidate,⁶⁹ two phase III trials were recently initiated for evaluating IONIS-SMN_{Rx}.^{70,71}

Although the SSO design remains challenging, several new therapeutic applications were successfully validated in preclinical studies.⁷² Possible drug approvals of eteplirsen or IONIS-SMN_{Rx} in the near future could eventually proof the feasibility of the splice-modulating antisense oligonucleotide approach.

Aptamers

Aptamers are 20 – 100 nt long oligomers that adopt complex three dimensional structures that allow them to interact potently and specifically with various proteins typically achieving nM- to pM binding affinities.⁷³ They are readily obtained in an iterative laboratory evolution procedure called SELEX (systematic evolution of ligands by exponential enrichment).⁷⁴ Currently, aptamers are mainly targeting extracellular structures such as plasma proteins and cell surface receptors thus avoiding the problem of intracellular delivery. Hence, aptamers are comparable in many aspects to antibodies, however, aptamers are much smaller, can penetrate tissues deeper, are chemically synthesized to highest purity and homogeneity and differ in their toxicity and immunogenicity profile. To improve their plasma life-time and to adjust their toxicity, aptamers are typically chemically stabilized (2'-OMe, 2'-F, 3' inverted dT) and PEGylated.

In 2004, the first (and until today the only) aptamer, Macugen, was approved by the FDA for clinical therapy of AMD (age-related macular degeneration). The 27-nt chemically stabilized RNA oligomer is directed against the vascular endothelial growth factor (isoform 165) and blocks VEGF-receptor-induced neovascularization.^{75,76} After achieving its highest sales in 2010, it has now almost entirely been displaced by antibodies (Ranibizumab and Bevacizumab, for instance) which can bind additional VEGF isoforms besides VEGF-165 and thus benefit for their poorer specificity compared to the aptamer. After this early breakthrough with Macugen, numerous aptamers have been explored in clinical settings. However, some programs suffered very unfortunate setbacks at late clinical trial states, like the aptamer-containing anticoagulation system REG1 which was terminated in 2014 in a phase III study due to unexpected toxicity / immunogenicity issues (Table 1).^{77,78}

Currently, several aptamers for the local treatment of eye diseases are in late clinic trials (II and III), for instance the aptamers Fovista⁷⁹⁻⁸¹ and Zimura,^{82,83} which target PDFG (it is a growth factor) and C5, respectively. In combination with VEGF inhibitors they might find application in the treatment of AMD in the near future. To overcome the prevalent problems with toxicity and immunogenicity, NOXXON Pharma develops so-called Spiegelmer therapeutics.⁸⁴ These drugs apply stereochemically inverted nucleotides based on L-ribose instead of the natural D-ribose, can be evolved via SELEX, and are suggested to be resistant against nucleases⁸⁵ and invisible for the immune system.⁸⁶ Currently, 3 Spiegelmer aptamers⁸⁶⁻⁹² are in clinical phase II studies (Table 1).

Therapeutic RNAi

RNA interference (RNAi) is a mechanism of posttranscriptional gene regulation that was discovered in 1998.¹² RNAi can

interfere with gene expression in various ways including the degradation of a specific mRNA target via endonucleolytic cleavage, or via recruitment of deadenylation / decapping enzymes, but it can also positively affect the stability and translation of a specific mRNA. The mechanistic details that lead to the respective responses are still under exploration. In principle, a dsRNA that is introduced into the cytoplasm is processed by the RNase Dicer into ~22 bp RNA duplexes and loaded onto the endonuclease Argonaute-2 (Ago-2). Ago-2 slices the passenger strand of the RNA-duplex and applies the remaining guide strand for sequence-specific mRNA-targeting.⁹³ While short interfering RNAs (siRNAs) are fully complementary to their target mRNA and promote cleavage (knock-down), microRNAs (miRNAs) contain bulges and loops that prohibit slicing by Ago-2, but alter the stability and translational activity of the target.⁹⁴

Allowing the selective knock-down of genes in cell culture and animal-models, RNAi quickly became a valuable tool in basic biology.⁹⁵⁻⁹⁷ In parallel a race started to exploit the RNAi mechanism for therapeutic purposes and several big pharma companies, like Merck, Roche, and Pfizer made large investments that resulted in the first clinical trials in 2004, already 6 y after the discovery of RNAi.^{98,99} However, in the aftermath those early trials mostly failed due to strong innate immune reactions and/or lack of patients' benefit, and in the consequence big pharma left RNAi again.¹⁰⁰⁻¹⁰⁴ In the 18 y since its discovery the field of therapeutic RNAi went from enthusiastic interest over despondence and back again, resulting in a re-assessment of the technological obstacles and more realistic expectations for clinical trials. This has been accompanied by commentary elsewhere.^{105,106}

However, after recent successes in clinical trials, showing the efficacy of RNAi therapeutics to reduce transthyretin¹⁰⁷ and PCSK9¹⁰⁸ in patients, the interest in RNAi is currently growing and even big pharma including Sanofi and Roche started to invest again.⁹⁸ The initial drawbacks in clinical trials were mostly related to the low efficacy of the drugs, off-target issues and immune-related toxicity.¹⁰⁹ Off-target effects include immune-reactions induced by the siRNA/miRNA precursors, and up- and downregulation of non-target mRNAs due to saturation of the RNAi machinery and off-target binding of the siRNA.¹¹⁰ There is now increasing success in tackling all those issues. Current innovations include chemical modification / sequence optimization of siRNAs and its precursors, and new solutions to the delivery problem. The latter include various forms of (lipid) nanoparticles and bioconjugates. The details of this progress are comprehensively reviewed elsewhere.¹¹⁰⁻¹¹³ Briefly, clinical trials seem more successful when they are confined to readily accessible organs like the liver, cancer, and immune-privileged areas like the eye.¹¹⁴⁻¹²² Whereas the eye is a good target for naked siRNAs, treatment of the liver benefited from lipid-based nanoparticles and the above-mentioned GalNac₃ conjugates.¹¹⁶ In particular the GalNac₃ approach has significantly improved the efficacy of siRNA-conjugates, allowing now the weekly administration of liver-targeting siRNA via subcutaneous injection in non-human primates to knock-down antithrombin to clinically relevant levels.¹¹ Notable in this approach is that it allows to knockdown an essential protein (like antithrombin) in a tunable and reversible manner,

whereas the permanent knock-out of antithrombin (for instance at the DNA-level) is lethal.¹¹ Overall, more than 20 siRNA drugs in various formulations are in clinical trials now (up to phase III, Table 1).¹²³ RNAi-therapy clearly has the potential to tackle currently undruggable diseases and to appear in the clinics soon.

The therapeutic use of the miRNA-related mechanism (not applying the slicing activity of Ago2) is still in its infancy. Attractive is the possibility of manipulating larger networks of genes simultaneously in both, a negative and positive manner.¹²⁴ This might become interesting for the treatment of complex diseases like cancer. On the other hand, endogenous miRNAs are involved in many cellular processes and their manipulation could also be disease-relevant. The knockdown of miRNA 122 with antisense oligonucleotides was shown to interfere with hepatitis C virus progression and is currently in phase II clinical studies.¹²⁵ As the hepatitis virus seems to require the endogenous miRNA for its functioning the knockdown of this host-specific factor is particularly promising as the virus cannot adapt easily by evolution.¹²⁶ Other miRNAs that are linked to cancer like miRNA 16 and 34a are also targeted with ASOs and are currently in clinical trials phase I.^{127,128}

Emerging concepts for therapy

Therapeutic mRNA

For a long time it has been believed that only short, chemically stabilized oligonucleotides are suitable as drugs. However, long (protein-encoding) mRNAs have recently proven their enormous therapeutic potential. Protein replacement experiments were first performed in the early 1990ties with naked mRNA in mice and rats.^{129,130} Even though replacement experiments were successful to some degree, there have been massive problems related to the well-known RNA-dependent immune-stimulation through interferon-I (IFN-I) and a generally low translation efficiency.^{131,132}

However, during the last 15 years, our mechanistic understanding of the immune-stimulatory effect of RNA has substantially improved. This was due to the discovery of RNA sensors including the Toll-like receptors (TLR) 3, 7, 8, Melanoma differentiation-associated protein 5 (MDA-5), Retinoic acid inducible gene I (RIG-I), as well as various RNA helicases.¹³³ Besides the activation of the innate immune response under release of the respective signaling molecules we have also learned how these RNA-sensing events are directly linked to the general repression of mRNA translation in the affected cells. Among others, general translation repression is mediated by phosphorylation of translation initiation factor 2 α via protein kinase R activation.^{134,135} In the worst case, IFN-I activates 2'-5'-adenylate synthase and RNaseL and leads to apoptosis.¹³⁶

RNA replacement strategies aim to achieve high translation levels under minimal immune stimulation. Both can be achieved by designing mRNAs that evade RNA-sensing. The following strategies turned out as particularly successful.

- a) Chemically modified pyrimidine nucleotides like pseudouridine (ψ), 2-thiouridine (s2U), and 5-methylcytidine (m5C) are incorporated into mRNAs during in-vitro-transcription to minimize recognition by RNA

sensors.¹³⁷ Substitution of uridine by pseudouridine was shown to diminish recognition by TLR-3, -7, -8, and RIG-I.^{137,138} To fine-tune effects on translation efficiency, nucleotide analogs are often mixed with their natural counterparts. The extent to which these modifications may induce mistranslation is yet unknown.¹³⁹

- b) Rigorous purification of the mRNA product from unincorporated nucleoside triphosphates, small abortive transcripts, remaining DNA templates, and in particular dsRNA via HPLC (High performance liquid chromatography) was shown to dramatically reduce immunogenicity of the transcripts and can increase the translation 10- to 1000-fold.^{140,141}
- c) Synthetic cap analog structures like ARCA (anti-reverse-cap-analog) can further decrease immune response and improve translation. In contrast to older cap analogs, ARCA is always incorporated in correct orientation.^{142,143} A new ARCA variant contains a phosphothioate that resists enzymatic decapping and can increase the half-life of the mRNA.¹⁴⁴
- d) Computational sequence design allows to reduce the number of particularly immune-stimulatory nucleotides and combinations (like UW, with W = A or U).¹⁴⁵⁻¹⁴⁷ Furthermore, transcript stability can be optimized by the introduction of 3'-UTRs (or some elements) taken from other mammalian or viral genes as well as addition of Poly(A)-tails.¹⁴⁸⁻¹⁵³

The RNA replacement strategy is particularly advantageous when a transient, burst-like expression of a protein is desired. Typical examples for the latter are the epigenetic re-programming (induced pluripotency), wound healing, and genome editing. In this sense, in-vitro transcribed mRNA (IVT-mRNAs) has been used to deliver a) human bone morphogenetic protein 2 (hBMP-2) to support bone regeneration in rats; to deliver b) the transcription factor mix that induces pluripotency; and to deliver c) vascular endothelial growth factor-A (VEGF-A) into a mouse model for myocardial infarction resulting in an improved heart function and enhanced survival.¹⁵⁴⁻¹⁶⁰ Furthermore, IVT-mRNAs have been successful in the delivery of surfactant protein B in deficient mice, and in the delivery of murine erythropoietin to increase the hematocrit.^{138,161}

IVT-mRNA could turn out as a valuable tool for genome editing. Genome editing holds great promise for the treatment of various diseases by a permanent repair of a gene via a site-directed knock-in or knockout.¹⁶² However, the respective nucleases that induce the required double-strand DNA breaks including ZFNs, Talens, and CRISPR/Cas, should not be persistently expressed as this would dramatically increase the chance of off-target genome editing.⁹ Consequently, its delivery as an mRNA is beneficial compared to a DNA vector and also circumvents the typical safety risks of viral and non-viral DNA-based methods like genomic insertion and antivector immunogenicity. Encoding of genome editing tools via IVT-mRNAs has already been widely used to generate transgenic animals.¹⁶³⁻¹⁶⁹ In a proof-of-concept study, gene function was restored via homology-directed promoter exchange in a surfactant-B-deficient mouse model by in-vivo-delivery of the ZFN in form of an IVT-mRNA. However, this required the additional delivery of the repair template (with the promoter) in

form of an AAV6 (Adeno-associated-virus serotype 6).¹⁷⁰ Successful promoter exchange was demonstrated and resulted in a prolonged life of the treated mice. IVT-mRNA encoded Talen have been used successfully to disrupt the CCR5 (CC chemokine receptor type 5) gene via non-homologous-end-joining in the T-cell line PM1. As the loss of CCR5 function confers resistance toward R5-tropic HIV-1 infection, side-directed nucleases are promising to target this infectious disease.¹⁷¹ An initial clinical phase I study is currently starting.¹⁷² As IVT-mRNA is a young field, this study represents the first clinical study that uses IVT-mRNAs, but more are likely to follow soon.

mRNA can have many advantages over DNA vectors to deliver therapeutic proteins. Besides its transient nature, we want note that mRNA is very well and quickly translated in postmitotic cells that are difficult to transfect with DNA vectors. mRNA also works independent of a promoter, but this can potentially limit its application if tissue-specificity is required. However, we know from various studies that there is a large number of regulatory elements, typically in the 3'-UTR, including miRNA binding sites, stabilizing and destabilizing elements that could allow to manipulate the expression of an IVT-mRNA in a tissue-specific manner in the future.^{94,173}

Oligonucleotides for vaccination and desensitization

As indicated above, very successful strategies haven't been developed to evade the RNA-sensing event and to trick the innate immune system. However, inducing a specific immune response can be highly desired. Thus the recent knowledge on the immune stimulation by RNA can be used for the latter. Currently, the classical vaccination is based on the delivery of inactivated or living viruses, virus-like particles, or antigenic peptides. While the antigenic peptides require additional vaccination adjuvants like alum salts, the other entities contain sufficient pathogen-associated-molecular-patterns (PAMPs) in form of proteins, nucleic-acids, and lipopolysaccharides. These PAMPs are detected by pattern-recognition-receptors (including the above-mentioned RNA sensors) and induce the release of type-I interferons, pro-inflammatory cytokines, and chemokines. This is reviewed in-depth elsewhere.^{174,175} Short peptide fragments are then presented to the immune system via MHC-complexes on dendritic cells and other antigen presenting cells.¹⁷⁶ This process finally induces a humoral as well as cellular immune response of the adaptive immune system.

The presented antigens are mainly protein-derived peptides. This opens the intriguing possibility to deliver antigens for MHC-presentation encoded as IVT-mRNAs under simultaneous induction of the necessary innate and adaptive immune stimulation as the IVT-mRNA itself can function as PAMP. By doing so, it is well conceivable to create specific immune responses not only against viruses and bacteria, but also against cancer cells or for allergy treatment.¹⁷⁷⁻¹⁸¹ The design of such mRNA-based vaccines would be highly rational, fast, cheap, and could be done in a personalized manner, for instance against the specific transcriptome of a patient-specific cancer.¹⁸² IVT-mRNA vaccines would be faster available as the generation of virus-particles (and similar entities) would be circumvented. Lyophilized mRNA vaccines can be stored at 37 °C for several weeks.¹⁸³ This allows the transport of vaccines into

regions that cannot provide an uninterrupted cold chain. The safety-profile could also be better compared to DNA-based methods (insertion mutagenesis, low efficiency) or virus-like entities (therapy-induced virus-specific humoral immune response).¹⁸⁴⁻¹⁸⁶ Again, also for vaccination, the transient nature of RNA expression is beneficial, as a low-level, long-term expression of an antigen might induce tolerance.¹⁸⁷

Two major IVT-mRNA-based vaccination strategies are currently explored: the ex-vivo and the in-vivo approach. The first, which was earlier developed, is based on the ex-vivo pulsing of allogenic (= patient-derived) dendritic cells with antigen-encoding mRNA, which allows the redirection of the adaptive immune system to target cancer or virus-infected cells. The feasibility and safety of this method was proven in pre- and clinical trials focused on HIV and various cancer types. However, personalized ex-vivo therapies require time-consuming and expensive individualized manufacturing processes which currently limit their broad clinical application.¹⁸⁸⁻¹⁹⁵ Nevertheless, further clinical trials up to phase III are currently running.¹⁹⁶⁻²⁰²

Even though cumbersome, the ex-vivo strategy allows to optimize and control mRNA transfection and immune stimulation more carefully. The in-vivo approach, however, is potentially more simple and elegant, but encounters additional problems. Whereas all IVT-mRNA strategies require stable and highly translatable transcripts, the in-vivo strategy requires additionally the immune-stimulatory effect that counteracts translation. It was found that complexation of IVT-mRNA with protamine enhances immunogenicity via TLR-7 activation and simultaneously improves stability, however, with the downside of low antigen expression.²⁰³ Anyway, a combination of protamine-complexed IVT-mRNA together with naked IVT-mRNA of the same sequence turned out to satisfy both needs at the same time: high translation efficiency and immune stimulation. Those self-adjuncting mRNAs are currently in phase I and II clinical trials against prostate cancer, late stage

lung cancer, and rabies; pre-clinical trials against influenza have been performed.^{183,186,204-209} We wish to mention that also other approaches that apply naked or formulated IVT-mRNAs are in clinical trials, for instance for targeting other cancer entities.^{172,210} Furthermore, non-coding RNA can also be used as a vaccination adjuvant replacing the classical alum salts as adjuvant of protein- or peptide-based vaccines.²¹¹

Currently, IVT-mRNA are expensive therapies. On one hand, the GMP (Good manufacturing practice) production of IVT-mRNA in large scale is not yet fully established, but CureVac has announced significant progress here.²¹² On the other hand the potency of IVT-mRNA could be further improved by assisted delivery via lipid-nanoparticles, polymeric nanoparticles, gold nanoparticles, among others, as reviewed elsewhere.²¹³ Furthermore, there are promising attempts to develop self-replicating RNA-vaccines that apply viral RNA-dependent RNA-polymerases (from α -virus) to produce the RNA vaccine from a dilute IVT-mRNA template.²¹⁴⁻²¹⁶ However, there are safety concerns related to the control of the replication process and the tolerance against the viral RNA-polymerase, but the strategy is still in the pre-clinical exploration phase.²¹⁷

Finally, mRNA vaccines could also be used in allergy treatment to desensitize the immune system against a specific antigen. Desensitization against type-I allergies is typically accomplished through repeated intra-dermal, intra-nodal, or sub-lingual application of allergens. Whereas a strong Immunglobuline E and CD8⁺ T-cell responses is intended during vaccination, desensitization aims to change the T_{H1}/T_{R1} to T_{H2} cell ratio toward T_{H1}/T_{R1} to fine-tune the immune response and to induce tolerance.¹⁷⁸ Application of low-dose IVT-mRNA could be used for that purpose, and there is pre-clinical data that prove efficacy and suggest a long-term protective effect.²¹⁸ One can expect first clinical trials to start within the next few years. Applying mRNA as an anti-allergic vaccine has several advantages compared to the classical allergen extract (like standardized cat extract) or DNA-based vaccines.^{219,220} IVT-mRNA is

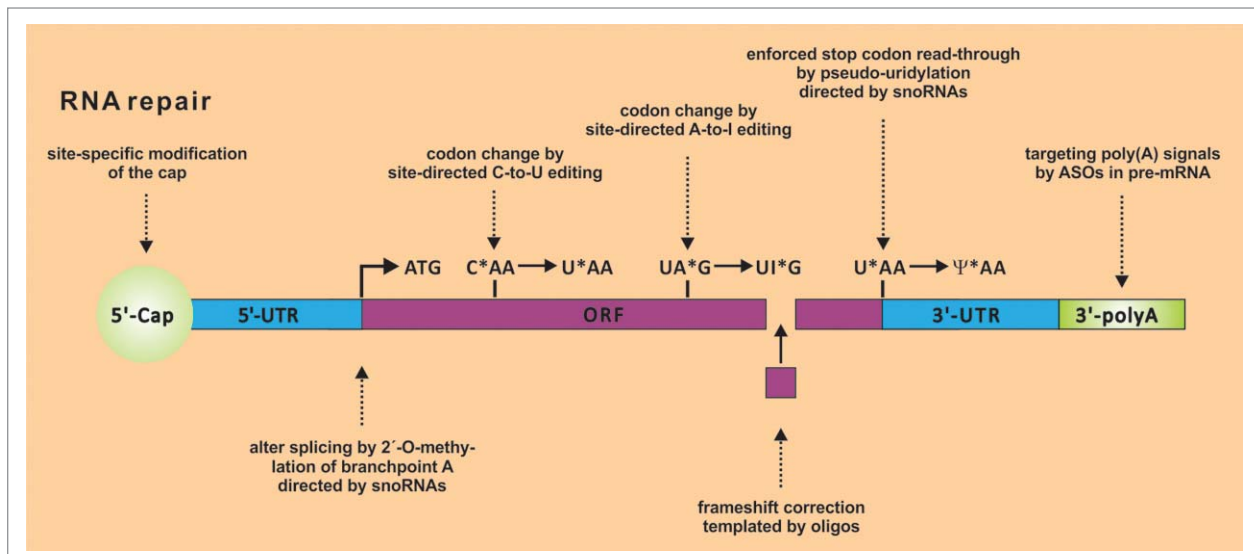


Figure 2. Overview on selected enzymatic processes that could be harnessed to restore gene function by repairing or re-programming mRNA site-specifically. Site-directed A-to-I editing, 2'-O-methylation, pseudouridylation, and frameshift correction via expression or administration of short guideRNAs has already been demonstrated. Many other processes are conceivable and currently under exploration.

obtained in a defined and highly pure state thus avoiding unintended antigens that can be included in allergen extracts.^{221,222} DNA-based allergy treatment on the other hand suffers from the above mentioned safety concerns and thus harbors disproportional risk in the context of a preventative therapy.

RNA repair

Besides the manipulation of splicing, most interventions on the RNA-level aim to destroy or block their endogenous targets. Strategies to restore the function of an RNA that is corrupted by missense, nonsense or frameshift mutation, or by defective processing are rare. In case of loss-of-function mutations, the administration of a therapeutic mRNA to replace the non-functional variant might solve the problem, as discussed above. However, this is only feasible with a small number of therapeutic mRNAs that can be translated under low control of translation level and tissues specificity. Indeed, many transcripts are tightly regulated with respect to their dose and tissue specificity and come as a mixture of various isoforms due to alternative promoter usage, alternative splicing, alternative polyadenylation and alternative posttranscriptional modification. Such transcript variants may differ in their function, localization, stability, etc. To address this variety in an mRNA replacement strategy seems impractical. A better alternative would be the repair of the endogenously expressed but defective RNA transcript, a strategy, we call RNA repair.

Very recently, we and others have engineered artificial RNA-guided editing machineries that allow to re-program genetic information at the RNA level.²²³⁻²²⁵ For this, adenosine-to-inosine (A-to-I) RNA editing enzymes^{226,227} are directed toward specific sites on selected transcripts and allow for the precise posttranscriptional manipulation of the genetic information. The manipulation results from the fact that inosine is biochemically interpreted as guanosine. Thus, formal A-to-G conversions become accessible, in a highly site-specific manner. The specificity comes from the guide-RNA that addresses the editing enzymes and can be readily programmed in rational way, simply by applying Watson-Crick pairing rules.²²⁸ Even though only A-to-G mutations are accessible the scope of manipulations is large. Twelve out of the 20 canonical amino acids can be manipulated, comprising almost all of the polar ones which are essential for protein function.²²³ Furthermore, START and STOP codon, splice elements, polyadenylation signals, and viral RNA are potential targets.^{226,227} We and others have shown that such strategies work inside mammalian cell culture²²⁹ and even in a simple organism²³⁰ and allow the repair of disease-relevant genes, like the CFTR mRNA.²²⁵

Other people have recently shown the possibility of redirecting snoRNA-guided RNA modification machineries, like the 2'-O-methylation²³¹ and the pseudouridylation machinery.¹³⁹ The first modification allows interference with splicing, the second allows the read-through of premature STOP codons. Mammalian cells harbor a plethora of RNA modifying and processing enzymes. There is no need to restrict ourselves to the usage of nucleases, like RISC, RNaseH, and RNaseP.²³² Just to give a few examples, there are RNA editing and modifying enzymes inside the cell that can change nucleotides (A-to-I, C-

to-U²³³, U-to-ψ, A-to-m6A,²³⁴ and many more for the tRNAs²³⁵), that add the cap²³⁶ and the poly(A)-tail,²³⁷ RNAs can be precisely processed, for instance by the CCA-adding enzymes,^{238,239} TUTases,²⁴⁰ etc.²⁴¹ Thus, even complex repair processes are conceivable, including the repair of insertion and deletion mutations at the RNA-level. In this respect, we want to recall a largely overseen work from 2004, done by Paul Zamecnik, the pioneer of antisense therapy, in his early nineties shortly before he passed away. He demonstrated the possibility of repairing the terrible Δ508 deletion mutation in the CFTR gene, the main cause of cystic fibrosis, simply by administration of 2 chemically stabilized RNA oligomers.²⁴² In cell culture, the efficiency of mRNA repair was sufficient to restore the chloride channel function. Unfortunately, he was unable to elucidate the mechanism, but he could clearly demonstrate the repair to take place at the mRNA. Such a complex repair requires a concerted nuclease, ligase (and polymerase) activity at a specific site on an mRNA molecule. In summary, it seems that numerous endogenous enzymes stand ready inside the cell for RNA repair processes. We just have to learn how to make use of them.^{243,244} If successful, one can establish novel platforms for therapeutic intervention.

Conclusions

While splice-switching oligomers and aptamers are still struggling on their ways to the clinic, major progress has been made for RNaseH-dependent ASOs and for therapeutic RNAi with chemically stabilized siRNAs. This is due to the development of new chemistries that improve efficacy and delivery of the drugs to some specific organs. An impressive example is the development of the GalNAC₃ conjugation that clearly improves liver targeting and might allow for the administration of siRNA and ASO by subcutaneous administration in the future. However, overcoming problems with delivery and efficacy remains elusive for many organs and will require massive basic research in the future.

Among the emerging approaches, the usage of in-vitro-transcribed mRNA for protein replacement and vaccination has made impressive progress. This was mainly due to the tailored suppression or harnessing of the RNA-induced immune response by chemical modification and formulation. The approach has the potential to find wide application in the clinics whenever a transient, burst-like expression is advantageous. The RNA repair approach is still in its infancy, but we believe that the harnessing of artificial and in particular endogenous RNA repair proteins might enable new therapies, complementing the above-mentioned classical RNA-based and the approaching genome editing methods, and being superior to the latter with respect to safety and ethical issues.

Overall, the progress during last years is impressive. The increasing number of clinical trials for various approaches makes us feel optimistic that numerous nucleic-acid-based drugs will soon find their ways to the patients to enable novel therapies.

Disclosure of potential conflicts of interest

No potential conflicts of interest were disclosed.

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