

Novel Therapeutic Modalities to Address Nondrugable Protein Interaction Targets

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Small molecule drugs are relatively effective in working on ‘drugable’ targets such as GPCRs, ion channels, kinases, proteases, etc but ineffective at blocking protein–protein interactions that represent an emerging class of ‘nondrugable’ central nervous system (CNS) targets. This article provides an overview of novel therapeutic modalities such as biologics (in particular antibodies) and emerging oligonucleotide therapeutics such as antisense, small-interfering RNA, and aptamers. Their key properties, overall strengths and limitations, and their utility as tools for target validation are presented. In addition, issues with regard to CNS targets as it relates to the blood–brain barrier penetration are discussed. Finally, examples of their application as therapeutics for the treatment of pain and some neurological disorders such as Alzheimer’s disease, multiple sclerosis, Huntington’s disease, and Parkinson’s disease are provided.

Neuropsychopharmacology Reviews (2009) **34**, 142–158; doi:10.1038/npp.2008.115; published online 27 August 2008

Keywords: antibodies; aptamers; antisense; protein–protein interaction; siRNA; target validation

INTRODUCTION

Small molecule drugs have dominated central nervous system (CNS) therapeutic markets for decades. Although small molecules are relatively effective in working at ‘drugable’ targets, such as G-protein-coupled receptors, ion channels, kinases, and proteases, they are ineffective at blocking protein–protein interactions that represent the emerging class of ‘nondrugable’ targets. The implementation of genomic and genetic technologies has led to the identification and validation in cellular and animal models of several promising protein–protein interaction targets for the treatment of neurological and psychiatric disorders. The birth of the biotechnology industry has led to introduction of several new drug categories, including recombinant proteins and therapeutic antibodies in the 1980s and 1990s and the introduction of emerging oligonucleotide therapeutics such as antisense, small-interfering RNA (siRNA), and aptamers over the past decade. Although the relative growth rate of small molecule drugs continues to decline, biologics, and in particular recombinant proteins and monoclonal antibodies (mAbs) have revolutionized the treatment of metabolic and immunological disorders, infectious diseases,

and cancer and represent the fastest growing product segment of the pharmaceutical industry over the next decade. In this article, we will provide an overview of some of the tools such as antibodies, aptamers, antisense, and siRNA that are available to address protein interaction targets and discuss their utility for the validation of both drugable and nondrugable targets. Although an in-depth review of these novel therapeutic modalities is beyond the scope of this article, we will provide you with examples of their application for the treatment of pain and some neurological disorders such as Alzheimer’s disease (AD), multiple sclerosis (MS), Huntington’s disease, and Parkinson’s disease to name a few.

TOOLS TO ADDRESS NONDRUGABLE TARGETS

The cellular points of intervention of different therapeutic modalities are shown in Figure 1. A comparison of several of the key properties of antibodies, aptamers, antisense, and siRNA is summarized in Table 1 and detailed in the sections below.

Antibodies

Since the first therapeutic antibody for a cancer indication was approved by the FDA for marketing in 1997, antibodies have become increasingly important in the fight against cancer, autoimmune, and infectious diseases. To date, 22

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Received 18 April 2008; revised 21 June 2008; accepted 28 June 2008

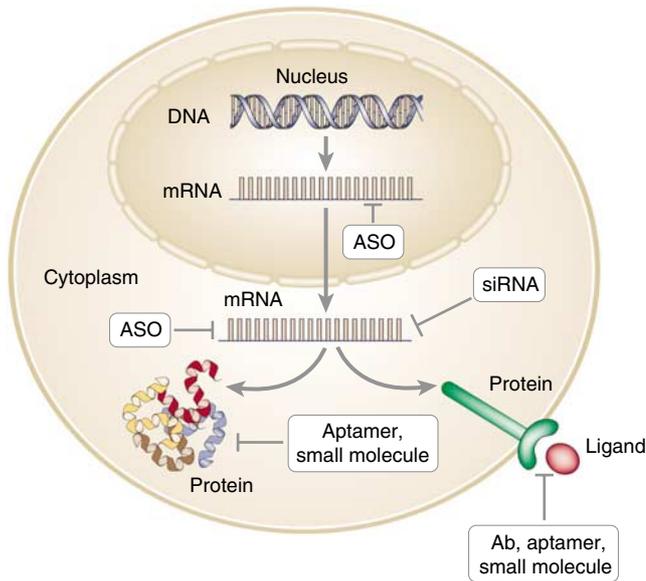


Figure 1. Different therapeutic modalities intervene in different parts of the cell. Antibody (Ab) therapeutics intervene on extracellular proteins, by binding to receptors or ligands to modulate their effects, either by inhibition or stimulation. Aptamers and small molecules bind to extracellular and intracellular proteins to alter their function. Both antagonist and agonist effects are possible on targets such as ion channels, neurotransmitter receptors, and kinases. siRNAs and antisense (ASOs) act further upstream at the mRNA level in the cytoplasm and nucleus, respectively, lowering levels of the corresponding protein.

unconjugated and 5 conjugated mAbs have been approved for these indications. This number will rapidly escalate as there are over 300 mAbs in preclinical and clinical development with the sale of antibodies expected to reach 48% of all therapeutic protein sales by 2009 (Evaluate-Pharma, 2008; Filpula, 2007; Wang and He, 2007).

Antibodies provide excellent affinity and specificity of target recognition. Furthermore, because of their relatively large size, *in vivo* stability, and ability to be sequestered from the blood by interacting with FcRn receptors on endothelial cells, they can have extended pharmacokinetic half-lives (days to weeks) (Daugherty and Mrsny, 2006). The drawbacks for mAbs as therapeutics have included the time and expense of production, tendency to illicit immune responses (ie neutralizing antibodies), inability to access intracellular targets, difficulty in tuning pharmacokinetic characteristics post-production, and formulation issues. The development of new methods and technical innovations has partially alleviated some of these concerns.

Monoclonal antibodies were originally exclusively produced by the hybridoma method in which hybrid cells are produced by the fusion of antibody-producing lymphocytes, acquired from animals after *in vivo* immunization, with a tumor cell. This method was laborious, costly, and produced mAbs prone to illicit a strong and detrimental immune response against murine mAb sequences from

TABLE 1: Properties of Antibody and Oligonucleotide Therapeutics

Property	Antibody	Aptamer	Antisense	siRNA
Targets	Extracellular	Extra- and intra-cellular	Extra- and intra-cellular	Extra- and intra-cellular
Cellular site of action	Extracellular	Extra- and intra-cellular	Intracellular	Intracellular
Potency	pM – nM	pM – nM	High	pM; takes advantage of an endogenous catalytic process
Selectivity	High	High	High	High
Manufacture	Biologics; complex scalability	Chemical; scalable	Chemical; scalable	Chemical; scalable
Cost of goods	High	Moderate	Moderate	Moderate
MW	180,000 Da	10,000 Da 50,000 Da (PEG)	7,000 Da	14,000 Da
Plasma half-life	Days to weeks	Minutes to days	Minutes to days	Minutes to days
Blood-brain barrier penetration	Limited	Limited	Limited	Limited
Routes of administration	Primarily IV or SC	IV, IM, SC, IVT, topical, INH, IT, ICV, IPC	IV, IM, SC, IVT, topical, INH, IT, ICV, IPC	IV, IM, SC, IVT, topical, INH, IT, ICV, IPC
Immunogenicity	Moderate	No	No	No
Immune stimulation	No	No evidence, but can be “designed” in or out	Yes; TLR9; can be minimized	Yes; TLR7; can be “designed” out
Target validation	Extracellular targets	Intra- and extra-cellular targets	Intra- and extra-cellular targets	Intra- and extra-cellular targets
Shelf-life	Limited; requires refrigeration	Generally stable	Generally stable	Generally stable

Abbreviations: ICV – intra-cerebroventricular, IM – intramuscular, INH – inhalation, IPC – intraparenchymal (CNS), IT – intrathecal, IV – intravenous, IVT – intravitreal, PEG – polyethylene glycol, SC – subcutaneous

human patients. This issue has been largely overcome by chimerization and humanization of murine antibodies, although even humanized mAbs show at least a low incidence of immunogenicity (Filpula, 2007). Furthermore, using transgenic mice expressing repertoires of human antibody genes has simplified the production of humanized antibodies from mice (Lonberg, 2005). Concurrently, a variety of methods have been developed to discover humanized antibodies by wholly *in vitro* means. Such methods rely on the screening of large (eg $> 10^9$) antibody libraries using display technologies with phage, cell-surface, and ribosomal scaffolds (Wark and Hudson, 2006). Although still laborious, display methods potentially increase throughput and provide the ability to tune antibodies subsequent to the initial selection.

Antibodies are composed of distinct modules consisting of different functions. For instance, IgG antibodies are bivalent (Figure 2); containing two proteolytically defined variable Fab fragments that consist of two variable domains (V_H and V_L) and are capable of binding antigen and one constant Fc fragment that is responsible for recruiting immune cells to targeted tissue and subsequent antibody-dependent cellular cytotoxicity or complement-dependent cytotoxicity. This modular architecture is conducive to recombinant manipulation toward producing therapeutics with altered properties, especially circulating half-lives (Figure 2; Holliger and Hudson, 2005). For example, removal of the Fc module (single-chain antibodies or scFvs) will shorten serum half-life and eliminate potential toxic side effects by inappropriate activation of immune cells. This modality has been clinically validated with pexelizumab, the scFvs anti-C5 complement inhibitor, in trials against acute myocardial infarction (Carrier *et al*, 2006). Also, recombinant linking of three or more Fab fragments (tribodies, tetrabodies, etc) can increase avidity and possibly serum half-life. The linked Fab fragments can be directed to distinct molecular targets as was done with an anti-CEA/CD3 bispecific diabody in an effort to stimulate the immune systems attack of colon carcinoma cells (Holliger and Manzke, 1999).

Other antibody engineering methods have been developed to deliver drugs/toxins to specific tissues such as has

been done with radioisotopes (Huhlov and Chester, 2004) or chemotherapeutics (Lillo *et al*, 2004). The reverse can also be done to engineer antibodies in a way allowing their delivery to specific sites. One example is to chemically couple an antibody to a pH-sensitive polymer that functions to disrupt membranes at acidic pH. This allows an internalized antibody to dissociate from endosomal vesicles and be released into the cytoplasm (Lackey and Press, 2002). Another particularly elegant example is the construction of a trifunctional fusion antibody as a therapeutic for AD. This tribody contains three modules. The first domain binds the human insulin receptor. The insulin receptor mediates the uptake of insulin by the brain and thus this module allows the entire tribody to enter the brain. The second domain is composed of an scFv antibody that binds the amyloid- β peptide ($A\beta$). Previous work has shown that anti- $A\beta$ antibodies injected into the brain of AD transgenic mice were capable of inducing the rapid clearance of preexisting amyloid plaques. The final module of the tribody contains the CH2-CH3 interface of the human Fc receptor. IgG molecules are rapidly exported from the brain to blood across the blood-brain barrier via the FcR. With these three binding modules, the $A\beta$ antibody was thus capable of (1) crossing the blood-brain barrier into the brain (influx), (2) disaggregating amyloid plaques by binding of $A\beta$, and (3) removal of the $A\beta$ by efflux across the blood-brain barrier back to the blood (Boado *et al*, 2007).

Aptamers

Aptamers are nonnaturally occurring oligonucleotides that bind their targets with high affinity and have the capacity to inhibit protein-protein interactions with potencies similar to what is observed with antibodies (Gold *et al*, 1995; James, 2000). The general process for the identification and optimization of aptamers is shown in Figure 3. Aptamers are generated by the SELEX (systematic evolution of ligands by exponential enrichment) process where high-affinity ligands are isolated from highly diverse (10^{13} – 10^{15}) starting pools by repeated rounds of affinity capture and amplification (Tuerk and Gold, 1990). SELEX can be conducted with nucleic acid pools that are highly stabilized to nucleases

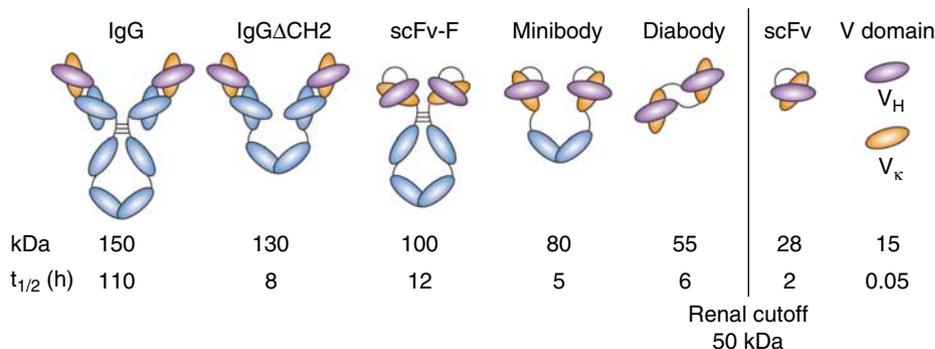


Figure 2. IgG and alternate engineered antibody structures and their effect on circulating half-life ($t_{1/2}$). Schematic representation of different antibody formats with approximate molecular weights and serum half-lives (β -phase). The classic full-length IgG is shown on the left with various and progressively more radical recombinant versions to the right (From: Holliger and Hudson, *Nat Biotechnol* 23: 1126–1136, 2005).

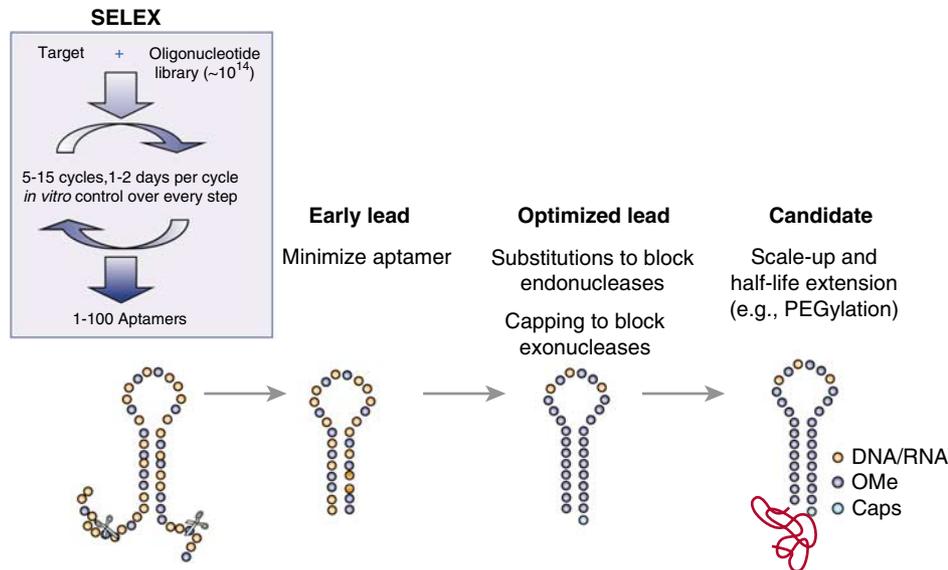


Figure 3. Aptamer discovery process. Schematic representation of the aptamer identification and optimization process. Aptamers are generated by an *in vitro* selection process referred to as SELEX (systematic evolution of ligands by exponential). Hit aptamers with the desired *in vitro* functionality are minimized to a length suitable for chemical manufacture at large scale. Chemical modifications to the minimized hit aptamer are introduced to optimize potency and stability in serum. High molecular weight polyethylene glycol (PEG) is then appended to the aptamer reduce or prevent renal filtration and extend aptamer half-life *in vivo*.

relative to DNA and RNA (Lin *et al*, 1994; Burmeister *et al*, 2005, 2006). The advantage of using such pools for SELEX is that the resulting aptamers bind their targets with high affinity and may be significantly more stable in biological fluids such as plasma than aptamers selected with DNA or RNA. Each oligonucleotide in the SELEX pool typically contains a 30–40 nucleotide region of randomly generated sequence flanked by 15–20 nucleotide 5'- and 3'- primer binding sites, so aptamer hits emerging from a SELEX experiment are typically 60–80 nucleotides in length. To facilitate efficient manufacture at a scale sufficient for therapeutic development and commercialization, aptamers approximately 40 nucleotides long or shorter are desired. Accordingly, the next step in the discovery process, referred to as 'minimization', involves identifying the core element within the hit aptamer sequence that binds the target and retains the desired functionality. Subsequently, aptamers may be optimized for potency and nuclease stability by introducing chemical modifications in the ribose ring, at the phosphodiester linkage or on the base moiety at specific residues (Yang and Gorenstein, 2004; Eaton *et al*, 1997). Aptamer residence time *in vivo* may be increased by appending high molecular weight polyethylene glycol (PEG) moieties to the aptamer termini (Watson *et al*, 2000). By modulating both the composition of the aptamer backbone and the overall molecular weight, the pharmacokinetic properties of an aptamer may be tuned to meet the needs of a particular therapeutic setting (Healy *et al*, 2004; Wilson and Keefe, 2006). For example, pegaptanib (an anti-vascular endothelial growth factor (VEGF) inhibitor for the treatment of age-related macular degeneration), the first aptamer to be approved for therapeutic use, comprises both a nuclease stable aptamer core and a 40kDa PEG and is administered once every 6 weeks (Chapman and Beckey, 2006; Ng *et al*, 2006).

Aptamer antagonists have been characterized for a wide range of proteins (reviewed in Cload *et al*, 2006), including cytokines (Rhodes *et al*, 2000, 2001; Ninichuk *et al*, 2008), growth factors (Bock *et al*, 1992; Jellinek *et al*, 1995; Pagratis *et al*, 1997; Ruckman *et al*, 1998; Rusconi *et al*, 2002), proteases, immunoglobulins (Wiegand *et al*, 1996), and cell adhesion molecules (Jenison *et al*, 1998; Watson *et al*, 2000). SELEX may be conducted against targets presented on the surface of cells (reviewed in Shamah *et al*, 2008), which allows the opportunity for direct selection of aptamer agonists as well. The first example of an aptamer agonist, targeting the co-stimulatory receptor 4-1BB, with demonstrated efficacy *in vivo* was recently reported (McNamara *et al*, 2008).

Although pegaptanib is currently the only approved aptamer, several aptamers are currently in clinical development. For example, phase 1 clinical data for an anti-Factor IXa aptamer (Chan *et al*, 2008) and an anti-von Willebrand Factor (vWF) aptamer (Gilbert *et al*, 2007) were recently reported. The anti-vWF aptamer acts through a mechanism unlikely to be achieved by a small molecule, namely blockade of the interaction between the A1 domain of vWF and platelets, a key step in the initiation of platelet aggregation. Given their high affinity and specificity, ability to block protein:protein interactions, aptamers present a promising modality for addressing nondrugable targets.

Antisense

Antisense oligodeoxynucleotides (ASOs) are single-stranded oligodeoxynucleotides of approximately 15–25 nucleotides in length that reduce the synthesis of the target protein. In contrast to antibodies and aptamers that bind to

proteins to inhibit their function, ASOs exert their effects further upstream, reducing gene expression at the mRNA level (Figure 1). As a result, ASOs can reduce the gene expression of both extracellular and intracellular targets. After entering the cell and binding to targeted mRNA, ASOs downmodulate gene expression by enabling the RNase-H-mediated degradation of the target mRNA or physically blocking translation of the target mRNA. As chemical modifications hinder RNase-H cleavage, such ASO modifications can be leveraged to favor the physical blockade mechanism. Recently, in a novel and very interesting application of RNA-based strategies, ASOs have been used to alter pre-mRNA processing by targeting motifs that involve in splicing (Lacerra *et al*, 2000; Lim and Hertel, 2001; Mann *et al*, 2001; Lu *et al*, 2005; Madocsai *et al*, 2005; Alter *et al*, 2006).

To identify ASOs with high selectivity and potency, careful *in vitro* selection with multiple cell lines to assess efficacy at the target RNA and/or protein level, and confirmation of mechanism of action and specificity by evaluation of closely related isotypes are critical (Crooke, 2004). Further efficacy and specificity analyses *in vivo* are then required, along with chemical modifications to promote stability and facilitate delivery. Phosphorothioate modifications have been widely used for achieving increased stability of ASOs *in vivo* against nucleases present in serum. After intravenous administration, these phosphorothioate-modified ASOs are rapidly distributed to peripheral tissues (distribution half-life of 0.4 h; Cossum *et al*, 1993), accumulating preferentially in organs such as the liver, kidney, and bone marrow. Other routes of administration such as subcutaneous, intradermal, intravitreal, topical, and inhalation have also been used successfully for *in vivo* delivery of ASOs to peripheral organs and tissues (Crooke, 2004; Cossum *et al*, 1994). A consideration with repeat dosing of phosphorothioate-modified ASOs is immune stimulation, as evidenced by splenomegaly and mononuclear cell infiltration in organs such as the liver and kidney (Henry *et al*, 2000). This is particularly pronounced with oligonucleotides containing a CpG motif that activates the Toll-receptor 9 (TLR9), a member of the Toll-receptor family involved in recognition of pathogen-associated molecular patterns and an innate immune cellular response. Stimulation of TLR9 results in an increased production of proinflammatory cytokines, and activation of adaptive immunity. By avoiding these CpG motifs, the proinflammatory response can be minimized. This immune stimulation can be reduced further with chemical modifications such as 5-methyl cytosine and 2'-methoxyethyl (Henry *et al*, 2000) but not completely eliminated (Senn *et al*, 2005). Although delivery formulations such as nanoparticles are being investigated for systemic delivery of ASO to nonneural targets, simple ASO formulations in saline appear to be sufficient for uptake into neurons and activity *in vivo*. In the nervous system, ASOs have been applied successfully with direct CNS administration in preclinical

studies to inhibit multiple molecular targets of interest, including ion channels (Porreca *et al*, 1999), neurotransmitter receptors (Wahlestedt *et al*, 1993), growth factors (Porritt *et al*, 2005), signaling molecules (Sakai *et al*, 2007), and enzymes (Smith *et al*, 2006).

Although ASOs represent a useful tool for elucidation of target function in the nervous system, the more than 60 clinical trials of ASOs to date include only one neurological disorder—MS. Vitravene (fomivirsin) was the first antisense drug to receive FDA approval and has been marketed for cytomegalovirus retinitis since 1998. Thus far, the clinical settings in which ASO drug candidates have been tested comprise mostly oncology indications, as well as hypercholesterolemia, diabetes, MS, HIV, HCV, Crohn's disease, ulcerative colitis, rheumatoid arthritis, psoriasis, and asthma. However, an ASO targeting superoxide dismutase 1 (SOD1) has been reported to lower production of mutant SOD1 and prolong survival after intraventricular infusion in a transgenic rat model of ALS (Smith *et al*, 2006). This ASO, ISIS 333611, is in preclinical development at ISIS and may enter clinical trials for the treatment of ALS in the near future.

siRNA

RNA interference (RNAi) is a naturally occurring cellular process of post-transcriptional gene silencing, discovered in 1998 by Fire and Mello (Fire *et al*, 1998). Synthetic siRNAs are double-stranded RNAs of 21–25 nucleotides in length that harness this naturally occurring pathway. In contrast to antibodies and aptamers that bind to proteins to inhibit their function, siRNAs (like ASOs) exert their effects further upstream, reducing gene expression at the mRNA level (Figure 1). Consequently, siRNAs offer a powerful solution for inhibiting disease targets from all molecular classes, including those that are difficult to modulate selectively with traditional pharmaceutical approaches employing small molecules or proteins. In RNAi, the target mRNA is enzymatically cleaved, leading to decreased levels of the corresponding protein, with specificity being a key feature of the RNAi mechanism. Synthetic siRNAs leverage the naturally occurring process of RNAi in a consistent and predictable manner with regard to extent and duration of action. In the cytoplasm, siRNAs are processed to single guide (antisense) and passenger (sense) strands (Matranga *et al*, 2005; Rand *et al*, 2005), and incorporated into RNA-induced silencing complexes (RISC). The RISC complex seeks out and binds the target mRNA of complementary sequence, resulting in the cleavage of the target mRNA by the enzyme Argonaut 2 in the RISC complex (Hammond *et al*, 2001). As siRNAs harness an endogenous catalytic mechanism, present in all cells, the potency of siRNAs is extremely high with typical siRNAs exhibiting picomolar EC₅₀'s *in vitro*.

siRNAs have been used successfully for elucidating the *in vitro* and *in vivo* function of numerous molecular targets of interest. With careful sequence design and selection, and

appropriate chemical modification, siRNAs are not only potent, but also highly specific for the target of interest, stable *in vivo*, and do not elicit immune responses (Bumcrot *et al*, 2006). Standard sequence design comprises alignment of mRNAs from rodent, monkey and human, and identification of all possible conserved target regions. If sufficient homology exists across species, then multiple siRNA sequences can be screened *in vitro* for subsequent testing in preclinical models and potential clinical advancement. These possible target regions are then subjected to a BLAST-like analysis, to select sequences that are unlikely to result in 'off-targeting' due to silencing of genes that have partial homology to the gene of interest, especially in critical portions of the target region such as the 'seed' region (Jackson *et al*, 2006). These selected sequences, predicted to be specific for the gene of interest, are then tested empirically *in vitro* to identify the most potent sequences from dose-response studies in cell culture systems. Another mechanism whereby siRNA duplexes can induce unintended effects is through stimulation of the innate immune system. It has been demonstrated that siRNA duplexes with certain sequence motifs can activate TLR7 (Hornung *et al*, 2005). However, with the use of appropriate chemical modification, the motifs can be inactivated, eliminating immune stimulation without compromising silencing activity (Judge *et al*, 2005). These same chemical modifications, 2'-O-methyl substitutions, together with phosphorothioate modifications at the 3' end, provide the additional benefit of stability against endo- and exonucleases, respectively, that are present in biological fluids (Choung *et al*, 2006). Thus, with appropriate design, selection, and chemical modification, highly potent and specific siRNAs can be identified for *in vivo* studies.

As siRNA, antisense, and aptamers are all oligonucleotide based, one can envision combining them to generate bifunctional molecules with enhanced properties. In particular, aptamers that target proteins on the cell surface may be useful as delivery agents for siRNA or antisense oligonucleotides. Recently, an anti-prostate-specific membrane antigen (PSMA) aptamer fused to an anti-PLK1 siRNA was shown to specifically reduce levels of PLK1 in LNCaP cells expressing PSMA on their surfaces (McNamara *et al*, 2006). In addition, the aptamer-siRNA chimera inhibited tumor growth in a xenograft model. As more aptamers that specifically mediate cell delivery are identified, these aptamer-siRNA (ASO) chimeras may find broad utility.

In vivo studies in the nervous system with siRNAs have included inhibition of well-known molecular targets such as the dopamine and serotonin transporters (Thakker *et al*, 2004, 2005), P2X3 (Dorn *et al*, 2004), δ -opioid receptor (Luo *et al*, 2005), and huntingtin (Wang *et al*, 2005; DiFiglia *et al*, 2007). In these studies, only the expected phenotypic effects were reported, supporting the specificity of siRNAs *in vivo*. Moreover, the use of siRNAs as tools for target validation is practical. For neuronal gene silencing *in vivo*, siRNA simply formulated in saline and delivered directly to the CNS by intrathecal, intraventricular, or intraparenchymal

administration achieves neuronal uptake and inhibition of the gene of interest. Although transfection reagents such as iFECT have been used to enhance local delivery *in vivo* (Hassani *et al*, 2005; Luo *et al*, 2005; Tan *et al*, 2005; Wang *et al*, 2005; Kumar *et al*, 2006; Salahpour *et al*, 2007), such reagents are not absolutely required for neuronal uptake and have an associated risk of cytotoxicity.

A consideration for clinical translation is that for chronic treatment of neurodegenerative diseases, continuous or repeated long-term infusion of siRNA may be required, which is achievable with implantable infusion devices (see below). Although systemic delivery of siRNA to the CNS would be simpler from a clinical standpoint, the blood-brain barrier represents a formidable hurdle for successful delivery. Recently, Kumar *et al* (2007) have reported silencing in the CNS after intravenous administration of siRNA complexed to the rabies virus glycoprotein peptide, which was proposed to enable delivery across the blood-brain barrier and neuronal uptake by interaction with the nicotinic acetylcholine receptor. Xia *et al* (2007) have reported successful delivery to intracranial tumors in rats, using intravenous administration of a biotinylated siRNA attached by a biotin-streptavidin linker to a transferrin receptor antibody. If these results can be optimized for drug development, then a significant step would be taken toward turning siRNAs into CNS therapeutics.

For therapeutic application, not only is the identification of specific and potent siRNAs against any molecular target of interest rapid and efficient, but also synthesis of siRNAs in large quantities is straightforward and siRNAs are highly stable upon storage and during handling. Presently, there are five clinical trials in progress with siRNA candidate therapeutics. An siRNA targeting respiratory syncytial virus (RSV) for the treatment of RSV infection is being tested with local administration by Alnylam Pharmaceuticals. Three clinical trials, by OPKO Health, Allergan/Merck, and Quark/Pfizer, target the wet form of macular degeneration, using intravitreal injection of siRNA against VEGF, VEGF-R1, and RTP801i-14, respectively. The first systemic siRNA clinical trial recently commenced with Quark Pharmaceuticals targeting p53 for the treatment of acute renal failure. Additional clinical programs with siRNA are likely to commence in the near future. Not only do siRNAs represent a powerful tool for inhibiting any molecular target of interest for elucidation of target function, but also the exciting potential of siRNAs for therapeutic intervention is being revealed in ongoing clinical trials.

TARGET VALIDATION

Target validation is the process of determining if a particular protein is directly or indirectly involved in propagating a particular disease and is therefore an appropriate target for new drugs. The most straightforward method for accomplishing this task is to inhibit the protein's function in the context of a relevant disease

model. Target validation methods attack a gene's function at every level of expression—DNA, RNA, and protein (Figure 1). Knockout technologies attack gene expression at the DNA level and have been invaluable for the target validation of many proteins. However, constructing gene knockouts in mice can be laborious and the results can be difficult to interpret due to differences between mouse and human biology. Also knockout mice sometimes show a phenotype that is inconsistent with extensive *in vitro* research (Pich and Epping-Jordan, 1998) possibly due to the action of compensating genes or developmental anomalies.

RNAi and antisense are two target validation approaches that inhibit gene expression at the level of RNA. The strengths and limitations of RNAi, siRNA, and antisense have been described in detail in the preceding sections. Although these techniques which target expression at the DNA or RNA levels are powerful in validating targets of interest, they may have very different effects from drugs that act at the protein level. As proteins often either have more than one function or take part in numerous multi-protein complexes, deletion of the protein through altering mRNA levels can lead to the disruption of numerous pathways. In contrast, antibodies and aptamers target a specific protein–protein interaction.

Validating targets at the protein level is more direct and closely mimics the action of commonly used therapeutics such as small molecule drugs and biologics including antibodies. Antibodies, which bind proteins with high affinity and specificity, are a powerful way to inactivate protein targets and have been used against hundreds of extracellular targets (Lichtlen *et al*, 2002). However, antibodies are too large to traverse the cell membrane to attack intracellular targets. To circumvent this liability scFv antibodies have been expressed intracellularly (Lo *et al*, 2008). Intracellular antibodies have been named 'intrabodies'. Because intrabodies have disulfide bonds that are important for maintaining the proper structure there has been some difficulty in achieving proper folding in the reducing intracellular environment. Auf der Maur *et al* (2004) ameliorated this problem by using an *in vivo* screening method to find scFv sequences that are more stable. These and other advances in intrabody technology have allowed the practical use of intrabodies for target validation. For instance, Mclear *et al* (2008) used intrabodies to suppress neurodegeneration in photoreceptors in a fly model of Huntington's disease and found that the intrabodies were more effective in suppressing this aspect of the disease than Hsp70 overexpression.

Similar to antibodies, nucleic acid aptamers bind to the target at the protein level and have been used extensively for target validation purposes (Pendergrast *et al*, 2005). Aptamers have many attributes that make them suitable for target validation. They bind with high affinity and also exhibit a degree of specificity that allows them to distinguish between closely related molecules. Like small molecule therapeutics their dosage can be easily adjusted and they exhibit good dose-dependent pharmacokinetic and

pharmacodynamic responses. Importantly, an aptamer can inhibit target function *in vivo* by blocking or knocking out a single domain of a protein while leaving the remainder of the protein functional. Small molecule drugs are also known to have this capability (Aramburu *et al*, 1999). An example of the use of aptamers for validation of an extracellular target *in vivo* is a series of experiments using an anti-platelet-derived growth factor (PDGF)-B aptamer (Pietras *et al*, 2001, 2002). PDGF isoforms are potent mitogens and have been implicated in the abnormally high level of interstitial fluid pressure (IFP) seen in human tumors. This elevated pressure is not conducive to the penetration of drugs into the tumors. PDGF-B inhibition using an aptamer led to decreased IFP levels and concomitantly increased levels of chemotherapeutic agents within the tumor (Pietras *et al*, 2001). Furthermore, the high-affinity aptamer that was specific to PDGF-B and not PDGF-A was able to decrease IFP in a KAT-4 xenograft model and enhance Taxol uptake and efficacy (Pietras *et al*, 2002).

Aptamers, being nucleic acids, are more amenable for use against intracellular targets than antibodies as they can be easily delivered intracellularly by standard transfection techniques or synthesized *in vivo* with the appropriate expression vectors. Furthermore, they do not need cysteine bonds to fold properly as do protein inhibitors. Some intracellular targets may be refractory to either siRNA and/or aptamer knockout. However, as these methods are both nucleic acid based and act at different levels of expression it was theorized that the two methods could be easily combined and that the combination would be more effective than either method alone. Indeed, Chan *et al* (2006) knocked down a well-characterized intracellular protein's activity, NFkB, by expressing either aptamers or siRNAs on novel vectors driven by U6 and HIV-1 promoter sequences. The aptamers were encoded into transcripts known to form stable stem and loop structures that localize within the cytoplasm. The best construct placed the aptamer within the natural stable RNA aptamer 7SL with an artificial terminator sequence added on the 3' end (Figure 4a). This construct diminished NFkB's activity by 64%, about the same as the best siRNA tested (Figure 4b). Interestingly, expression of both aptamers and siRNAs simultaneously suppressed NFkB activity better than either method alone (up to 90%; Figure 4b). The exquisite specificity of aptamers is further exemplified by their ability to discriminate between phosphorylated *vs* nonphosphorylated kinases (Seiwert *et al*, 2000) with greater than 10,000-fold specificity suggesting the potential for validating the post-translational active state of a protein toward a particular phenotype.

ADDRESSING THE BLOOD–BRAIN BARRIER

With few exceptions, small molecules have been the mainstay for the treatment of psychiatric and neurological disorders as well as in pain management. In addition to the

blood–brain barrier (eg $A\beta$ neutralization approaches in the periphery or limited access to the brain; pain targets in the periphery), addressing diseases with a compromised blood–brain barrier such as MS or direct delivery of the therapeutic to brain using implantable delivery devices.

The feasibility of delivering proteins across the blood–brain barrier has been demonstrated using antibody and endocytotic pathways targeting transferrin receptors and low-density lipoprotein receptors that are highly concentrated on brain blood vessels. Carrier systems involving mAbs directed against the transferrin receptor have been utilized to transport large proteins such as nerve growth factor (NGF) across the blood–brain barrier (Granholtm *et al*, 1994). More recently, Spencer and Verma (2007) reported the use of the lentivirus vector system to deliver the lysosomal enzyme glucocerebrosidase and a secreted form of GFP to the neurons and astrocytes in the CNS using a fusion of the low-density lipoprotein receptor-binding domain of the apolipoprotein B to the targeted protein.

Implantable infusion systems for direct delivery of therapeutics to the CNS are being used for intrathecal administration of pain medications and have also been used in clinical trials for intraputamenal administration of a therapeutic candidate for Parkinson's disease. The Medtronic SynchroMed infusion system is an implantable, programmable drug infusion system consisting of a pump, catheter, and programmer that is approved for intrathecal delivery of therapeutics such as Prialt and morphine for the treatment of chronic, intractable pain. This infusion system was also used recently in clinical trials of glial cell line-derived neurotrophic factor (GDNF) for the treatment of Parkinson's disease (Patel *et al*, 2005; Lang *et al*, 2006; Slevin *et al*, 2006) where GDNF was continuously infused for 6 months to 2 years. Although differing efficacy results

were obtained in these studies, the trials clearly provide a precedent for the use of implantable infusion systems for direct intraparenchymal delivery of therapeutics to the CNS. Such infusion systems will be valuable for direct CNS delivery of novel therapeutics such as proteins, antibodies, and oligonucleotides. Alnylam Pharmaceuticals and Medtronic are currently co-developing a therapeutic for the treatment of Huntington's disease that is a drug–device combination consisting of an siRNA targeting huntingtin and an implantable infusion system.

EXAMPLES OF CNS DISORDERS ADDRESSED BY NOVEL THERAPEUTIC MODALITIES

Alzheimer's Disease: β -Amyloid Neutralization Approaches

AD, the major cause of dementia in the older population, affects approximately 5.1 million people in the United States (Alzheimer's Disease Facts and Figures 2007, Alzheimer's Association, www.alz.org). Approved drugs for AD, such as the acetylcholinesterase inhibitors Aricept (donepezil) and Exelon (rivastigmine), alleviate symptoms of the disease such as memory loss but none currently on the market impacts the underlying disease progression. AD is characterized by the presence neuritic plaques composed primarily of the 42 and 40 amino-acid $A\beta$ peptide, products of protease cleavage of the amyloid precursor protein (APP) by the γ - and β -secretases (Selkoe, 2001; Selkoe and Schenk, 2003).

Animal models (murine) such as the PDAPP mouse that expresses an APP mutant (V717F), which results in an increase in the production of $A\beta_{(1-42)}$, recapitulate AD-like

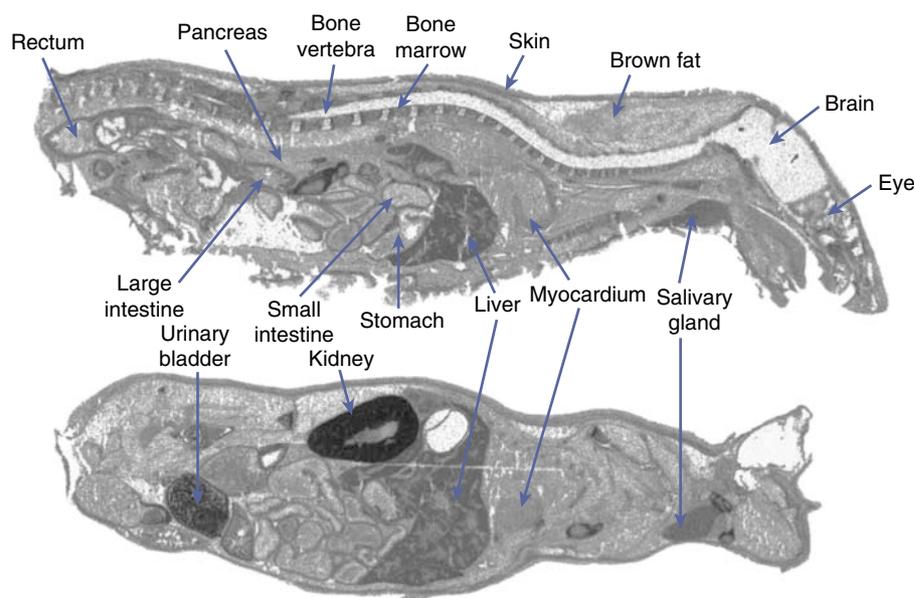


Figure 5. Aptamer distribution. Quantitative whole-body autoradiograph images of a CD-1 mouse 3 h after single intravenous bolus administration of an unconjugated ^3H -labeled aptamer (Cload *et al*, 2006).

plaque pathology, and the concomitant behavioral aspects of the disease (Games *et al*, 1995). These mouse models have been used extensively to study potential anti- $A\beta$ therapies for AD. In an early study, Schenk *et al* (1999) demonstrated that immunization of young PDAPP mice with $A\beta_{(1-42)}$ peptide resulted in production of anti- $A\beta$ antibodies and prevented the development of $A\beta$ plaques, dystrophic neurites, and astrocytosis that are characteristic of untreated animals and AD patients. Importantly, immunization of older animals with established plaques yielded similar results.

Several groups have studied the impact of passive immunization of AD mice with anti- $A\beta$ antibodies (Figure 6). Bard *et al* (2000) have raised anti- $A\beta$ mAbs using peptides representing the N-terminal through the C-terminal residues of $A\beta_{(1-42)}$ as immunogens and evaluated their effects on pathology in a mouse model of AD. Peripheral administration of antibodies to the N terminus of $A\beta_{(1-42)}$ for 6 months dramatically reduced plaque burden in PDAPP mice. Cryostat sections of the brain stained positively with fluorescently labeled anti-mouse antibodies, suggesting that the anti- $A\beta$ antibodies were able to enter the brain. In an *ex vivo* assay, primary microglial cells were cultured with sections of brain from PDAPP or human AD patients' brains in the presence of various anti- $A\beta$ antibodies. Antibodies that had caused plaque reduction *in vivo* also induced plaque phagocytosis. Further characterization suggested that the ability of the antibody to bind to $A\beta$ plaques rather than soluble $A\beta$ correlated with induction of plaque phagocytosis *ex vivo* and reduction of plaque burden *in vivo*. A $F(ab')_2$ fragment of one efficacious antibody, 3D6, was able to bind plaque but did not induce phagocytosis, suggesting an Fc-receptor-mediated

mechanism. In contrast, Bacskai *et al* (2002) demonstrated that the $F(ab')_2$ fragment of 3D6 was as effective as the full-length antibody at removing plaque deposits from the brains of Tg2576 APP transgenic mice (Holcomb *et al*, 1998) after topical administration to the surface of the brain. The behavioral impact of $A\beta$ plaque reduction was assessed in PDAPP mice treated with peripherally administered 10D5, a plaque-clearing antibody (Bard *et al*, 2000); these animals showed improved spatial learning and reduction of $A\beta$ fibrils in the hippocampus (Hartman *et al*, 2005).

Antibodies raised against the C terminus of $A\beta_{(1-42)}$ have also demonstrated efficacy in AD transgenic mouse models. Wilcock *et al* (2004a) administered an anti- $A\beta$ antibody RN2286 (raised against $A\beta_{28-40}$) for 1, 2, or 3 months to older Tg2576 APP mice to study the impact of treatment on animals with existing plaques. Antibody binding to plaque was detected, and treatment with RN2286 resulted in a reduction in plaque load, microglial activation at early time points, and an increase in circulating $A\beta$ levels (Wilcock *et al*, 2004a). In a separate study, weekly peripheral administration of RN2286 to Tg2576 APP mice resulted in complete reversal of learning and memory deficits observed in untreated Tg2576 mice as measured by radial-arm water-maze testing (Wilcock *et al*, 2004b). The requirement for intact antibody effector function to the plaque-clearing action of anti-C-terminal antibodies has been investigated. The impact of treatment with native and deglycosylated forms of 2H6, an antibody to the C terminus of $A\beta$, has been studied in the Tg2576 mouse AD model (Carty *et al*, 2006). Intracranial administration with native 2H6 resulted in activation of microglia and upregulation of $Fc\gamma$ RII and $Fc\gamma$ RIII whereas similar treatment with the deglycosylated antibody, which does not bind with high affinity to the Fc

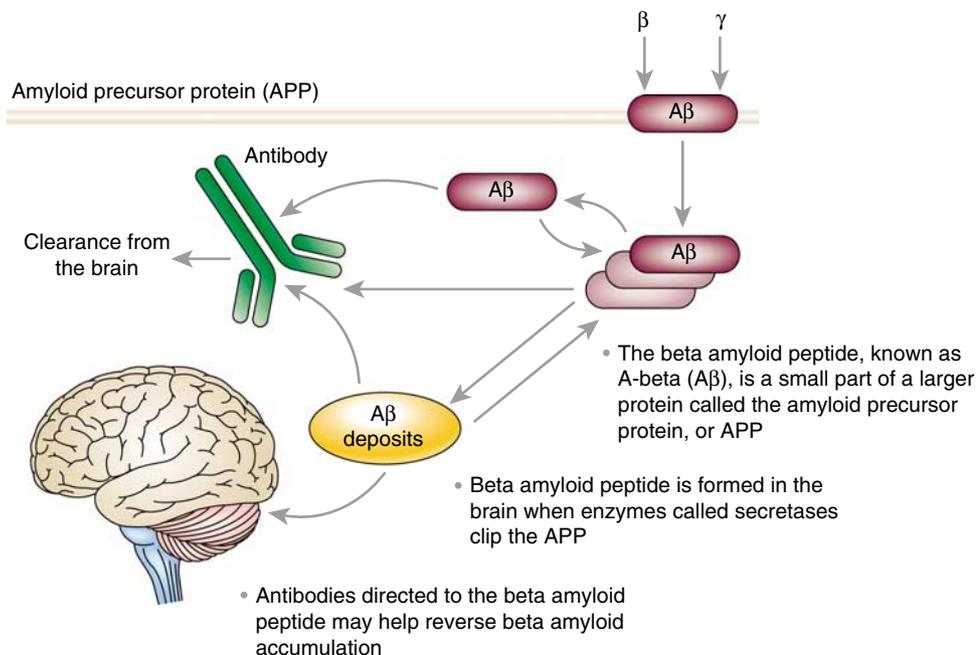


Figure 6. Amyloid- β immunotherapy approach. Anti- $A\beta$ antibodies may prevent or reverse $A\beta$ accumulation in the brain. (http://www.elan.com/Images/FINAL%20Slide%20%231%20MOA_tcm3-6288.pdf).

receptors, did not result in microglial activation or upregulation of the receptors. In both cases, treatment resulted in clearance of compact and diffuse plaques from the hippocampus and frontal cortex, which suggests that although phagocytosis may be involved in plaque clearance, it may not be required.

The results described above suggest that treatment with anti- $A\beta$ antibodies may lower plaque burden through an active mechanism within the brain involving microglial-mediated phagocytosis or perhaps disintegration of amyloid plaques initiated by antibody binding. A second series of papers suggest that agents that bind and sequester soluble $A\beta$ may show promise as well, although acting through a different mechanism. DeMattos *et al* (2001) showed that peripheral administration of m266, an anti- $A\beta$ antibody against amino acids 13–28 of $A\beta$ (Seubert *et al*, 1992) to PDAPP mice between the ages of 4 and 9 months reduced plaque burden in the hippocampus. In addition, m266 resulted in a 1000-fold increase in the plasma concentration of $A\beta$ within 24 h after administration. m266 binds with high affinity to soluble (nonfibrillar) $A\beta$ peptide and no binding of m266 to plaques was observed. Furthermore, the measured flux of the m266 antibody into the CNS was not sufficient to account for the increase in plasma $A\beta$. The authors propose that m266 shifts the equilibrium of $A\beta$ between the CNS and the periphery, causing the accumulation of $A\beta$ in the plasma. In a separate study, 24-month-old PDAPP mice treated with m266 for 6 weeks showed memory deficit reversals in object recognition and hole board learning and memory tasks, although their $A\beta$ burden in the brain was not decreased (Dodart *et al*, 2002). Interestingly, acute treatment of 11-month-old PDAPP mice with m266 resulted in a dose-dependent improvement in object recognition that correlated with an increase of soluble $A\beta$ in the plasma. Building upon that intriguing result, Bales *et al* (2006) showed that soluble $A\beta$ binds to ChT-1, resulting in impairment of cholinergic neurotransmission in PDAPP mice. Treatment with m266 quickly restored hippocampal ACh release to wild-type levels and restored the impaired habituation response observed in PDAPP mice to wild-type levels.

Many patients with AD exhibit cerebral amyloid angiopathy (CAA), which refers to amyloid deposits in the cerebral vasculature. CAA is observed in murine AD models (Calhoun *et al*, 1999), and in some cases treatment with anti- $A\beta$ antibodies increases the extent of CAA-associated microhemorrhages. DeMattos *et al* (2001) demonstrated that antibody binding to CAA-associated plaques correlated with CAA-associated hemorrhage. 3D6 and 10D5 that target the N terminus of $A\beta$, but not m266, which targets the central domain, bind CAA-associated plaques and increase the extent of microhemorrhage in the PDAPP AD mouse model (Racke *et al*, 2005). Separately, it was shown that treatment of Tg2576 mice with RN2286, which targets the C-terminal domain of $A\beta$ and binds plaque (Wilcock *et al*, 2004a), results in CAA-associated microhemorrhage (Wilcock *et al*, 2004b). The results described above (Carty *et al*, 2006) suggest that use of deglycosylated anti- $A\beta$

antibodies for passive immunization may be expected to reduce the incidence of CAA-associated microhemorrhage.

Both active and passive anti- $A\beta$ immunization strategies have moved into clinical development for the treatment of AD. The first trial, involving immunization of preaggregated synthetic $A\beta_{1-42}$ (AN1742; Wyeth/Elan) in combination with the adjuvant QS-21, was terminated after patients developed meningoencephalitis (Gilman *et al*, 2005; Orgogozo *et al*, 2003). Post-mortem analysis of patients revealed findings similar to those observed after immunization of PDAPP mice, including reduction in plaque load, association of collapsed plaques with microglia, and small hemorrhages (Ferrer *et al*, 2004; Nicoll *et al*, 2003). Further characterization suggested that changes in formulation between the Phase 1 and Phase 2 studies were associated in a shift in the patient T-cell response from Th2 to the proinflammatory Th1 response to the C terminus of $A\beta_{1-42}$ (Pride *et al*, 2008). A second clinical trial using a new immunogen consisting the N-terminal domain of $A\beta$ and a protein conjugate (ACC-001; Wyeth/Elan) is currently recruiting for Phase 2 studies (www.clinicaltrials.gov). Three of the anti- $A\beta$ antibodies (3D6, RN1219, and m266) described above are also in clinical development. The furthest advanced is bapineuzumab (humanized 3D6; Elan/Wyeth), which is in Phase 3, whereas LY2062430 (humanized m266; Eli Lilly) is in Phase 2 and PF04360365 (humanized RN2286; Pfizer) is currently in Phase 1.

Both aptamer and siRNA approaches to $A\beta$ neutralization/reduction have been described, although the work is at an earlier stage than the antibody-based programs. Ylera *et al* (2002) reported an aptamer selected against immobilized $A\beta_{1-40}$ that binds to fibrils consisting $A\beta_{1-40}$ with fairly high affinity (~ 40 nM). As yet, no functional data regarding, for example, fibril dissociation, have been reported. $A\beta$ peptide is produced by the sequential cleavage of the APP by β - and γ -secretases. APP transgenic mice with β -secretase (BACE1) deficiency are rescued from memory deficits (Ohno *et al*, 2004), suggesting that reducing $A\beta$ load by inhibiting BACE1 may be a viable therapeutic approach. Treatment of APP mice with anti-BACE1 siRNA via a lentiviral expression system reduced the level of $A\beta_{1-42}$ in the hippocampus. In addition, the siRNA-treated mice performed similarly to nontransgenic control mice in the Morris water maze (Singer *et al*, 2005).

Multiple Sclerosis: Anti-Interleukin-23

Multiple sclerosis is a chronic, progressive, demyelinating disease of the CNS. MS affects over 400,000 people in the United States and may affect 2.5 million people worldwide. The pathological hallmark of the disease is white matter lesions associated with focal infiltration of mononuclear cells with variable degrees of demyelination, axonal loss, and gliosis. The susceptibility to MS is thought to be multifactorial and it is generally considered to be an autoimmune Th1-mediated disease directed against myelin antigens. Interleukin-23 (IL-23) is a member of the

interleukin-12 (IL-12) family of proteins, which are predominant cytokines for triggering Th1-mediated (auto-immune) responses. IL-23 and IL-12 are heterodimeric proteins that contain an identical p40 subunit and a unique p19 (IL-23) or p35 (IL-12) subunit (Figure 7). Both IL-12 and IL-23 are produced by macrophages and dendritic cells and both cytokines affect T cells by binding to heterodimeric receptors. IL-12 acts primarily on naive T cells and promotes their differentiation into Th-1 cells that produce interferon- γ (IFN- γ). IL-23 acts primarily on Th-17 cells and is thought to stimulate their proliferation and production of IL-17 (reviewed in McKenzie *et al*, 2006).

A large amount of data have been generated linking IL-23 activity to autoimmune disease in the CNS. Mice with the p40 or the p19 subunits knocked out are highly resistant to experimental autoimmune encephalomyelitis (EAE), the mouse model for MS, whereas p35 knockout mice are, if anything, more susceptible to the disease (Figure 7; Cua *et al*, 2003). Moreover, intracerebral, but not intravenous, injection of IL-23 gene transfer vectors partially rescued disease susceptibility. Together, these data suggest that CNS-generated IL-23 but not IL-12 is important for EAE development. This hypothesis is supported by data showing that the IFN- γ is not required for EAE and that interleukin-17 (IL-17) knockout mice are also resistant (Ferber *et al*, 1996). Furthermore, Langrish *et al* (2005) showed that *in vitro* generated IL-23-dependent CD4+ T cells are highly pathogenic and essential for the establishment of CNS autoimmunity. This same pattern of resistance was seen in these knockout mice for other autoimmune disease models for rheumatoid arthritis (Murphy *et al*, 2003) and inflammatory bowel disease (Yen *et al*, 2006).

Antibody experiments validate IL-23 as a therapeutic target for MS. Neutralizing antibodies against IL-17 can ameliorate mouse EAE (Hofstetter *et al*, 2005) and anti-p19

(IL-23-selective) antibodies can prevent EAE induction and even reverse established disease (Chen *et al*, 2006). Furthermore, anti-p40 (IL-12/IL-23) antibodies have shown efficacy whether administered before or after onset of disease in a relatively new *Callithrix jacchus* (nonhuman primate marmoset) EAE model (Hart *et al*, 2008) and have been localized to lesions in the brain (Brok *et al*, 2002). In humans, overexpression of p40 mRNA (van Boxel-Dezaire *et al*, 1999) and p19 mRNA (Li *et al*, 2007) is observed in the CNS of MS patients and lower baseline levels of p40 mRNA predict clinical responsiveness to IFN- β treatment (van Boxel-Dezaire *et al*, 2000).

Anti-p40 antibodies, which eliminate both IL-23 and IL-12 activity, have been clinically validated for other autoimmune diseases. CNTO-1275 (ustekinumab), an anti-p40 mAb developed by Centocor Corp., has shown efficacy in both Phase 1 and Phase 2 trials for psoriasis. In the Phase 2 trial, 81% of patients receiving four weekly 90 mg doses achieved at least 75% improvement in their psoriasis, as measured by the Psoriasis Area Severity Index (PASI 75; Krueger *et al*, 2007). In addition, 52% achieved PASI 90, and 20% achieved PASI 100 (total skin clearance). Another anti-p40 antibody, ABT-874, showed similar results, with 90% of patients who received four weekly 200 mg subcutaneous injections achieving PASI 75 (Kimball *et al*, 2008). ABT-874 has also been reported to show efficacy in a Phase 2 trial for Crohn's disease (Mannon *et al*, 2004). Both antibodies were well tolerated in these short-term trials. It remains to be seen if these drugs will have efficacy against MS. So far one trial with CNTO-1275 in MS was conducted but the results have not been reported.

Aside from antibodies, short, stabilized anti-IL-12/23 aptamers have been reported (Burmeister *et al*, 2006). Several of these aptamers bind with low nanomolar affinities and inhibit IL-23- and IL-12-dependent STAT3 activation on activated human primary T cells. One of these aptamers has been truncated to 36 nucleotides and still retains activity. Thus, it shows great potential for development into a therapeutic for autoimmune diseases, including MS. However, because IL-12 is critical for immune responses, especially to intracellular pathogens (Altare *et al*, 1998) and not critical for the development of mouse EAE, it may be preferable to specifically target IL-23 over IL-12. To that end, aptamers selective for IL-23 have been recently developed (Pendergrast *et al*, 2006). One of these is able to inhibit IL-23-dependent IFN- γ release from primary human T cells over 100-fold more efficiently than IL-12-dependent IFN- γ release from the same type of cells. This same aptamer can also completely inhibit human IL-23-dependent release of IL-17 from mouse splenocyte cells (data not shown).

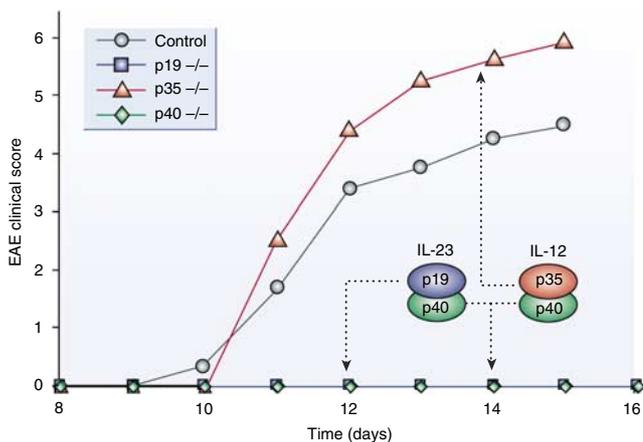


Figure 7 Interleukin-23 is the more relevant p40-containing cytokine for experimental autoimmune encephalomyelitis (EAE) (animal model of multiple sclerosis). Clinical EAE score over time (days) for wild-type controls (filled black circles), p40^{-/-} (filled green diamonds; IL-12/IL-23 knockout), p35^{-/-} (closed red triangles; IL-12 knockout), and p19^{-/-} (closed blue squares; IL-23 knockout) mice. (From: Cua *et al*, *Nature* 421:744–748, 2003).

Huntington's Disease: Huntingtin Inhibition

Huntington's disease is an inherited neurodegenerative disorder, caused by expansion of the CAG repeat region within the huntingtin gene (The Huntington's Disease

Collaborative Research Group, 1993). Clinical symptoms are characterized by involuntary movement, cognitive decline, and behavioral abnormalities; underlying these symptoms are dysfunction and degeneration of striatal and cortical neurons. Despite the identification of the mutated gene 15 years ago, there is no disease-modifying therapy available to the patient population. Functional studies on the mutated huntingtin protein and its molecular interactions with other genes and proteins have implicated huntingtin in multiple critical cellular processes including transcriptional regulation, axonal transport, apoptosis, and cell signaling. Transgenic mice expressing the mutant human huntingtin gene with expanded CAG repeat regions recapitulate molecular, cellular, and behavioral facets of Huntington's disease, providing essential systems for studying pathological mechanisms and testing therapeutic candidates (Mangiarini *et al*, 1996; Slow *et al*, 2003). However, to date, huntingtin has represented a nondrugable target and the key molecular partners in pathogenesis remain under question. Studies in mouse models with regulatable transgenes (Yamamoto *et al*, 2000) and virally delivered RNAi (Harper *et al*, 2005; Rodriguez-Lebron *et al*, 2005) have demonstrated that downmodulating huntingtin represents an important strategy with potential therapeutic benefit. Both unconjugated and cholesterol-conjugated siRNAs targeting huntingtin have been shown to be bioactive *in vivo*. Ge *et al* (2006; unpublished data) reported neuronal uptake of labeled siRNAs in the rodent striatum as well as huntingtin mRNA suppression after intrastriatal siRNA infusion. In a recent report, a cholesterol-conjugated siRNA was demonstrated to ameliorate neuropathology and behavioral abnormalities in a mouse model of Huntington's disease after intrastriatal administration (DiFiglia *et al*, 2007). Importantly, these studies have demonstrated that complete downmodulation of mutant huntingtin is not required for improvement of neuropathology or behavioral deficits; partial downmodulation of mutant huntingtin appears to be sufficient for significant benefit. In a recent report (McBride *et al*, 2008), shRNA expression systems that resulted in very high levels of two antisense RNAs appeared to be toxic to the mouse brain, whereas a third sequence that resulted in much lower levels of antisense RNA but similar silencing was well tolerated. Although the generalizability of these results remains to be determined, these observations illustrate the challenges of a gene therapy approach where consistency and predictability of shRNA expression levels can be problematic. In contrast, direct CNS delivery of synthetic siRNAs provides the needed degree of control over drug levels. McBride *et al* (2008) also report that the mouse brain can tolerate >50% reduction of huntingtin mRNA for at least 4 months, suggesting that the partial downmodulation of huntingtin that appears sufficient for significant benefit based on mouse models also is likely to be well tolerated. Both siRNA and antisense approaches are being pursued to develop drug candidates for the treatment of Huntington's disease, with direct CNS

delivery. These therapeutic strategies represent novel and promising approaches that have been enabled by a unique feature of siRNA and antisense molecules—disease intervention at the mRNA level to reduce the level of a pathogenic protein, in the absence of an identified interaction to inhibit.

Parkinson's Disease: α -Synuclein Inhibition

Parkinson's disease is a neurodegenerative disorder characterized by resting tremor, rigidity, bradykinesia, and postural instability. The primary cells affected are dopaminergic nigrostriatal neurons. Current therapies primarily focus on dopamine replacement and alleviate symptoms but do not alter the underlying progression of disease. The recent clinical trial of GDNF as a disease-modifying therapeutic candidate did not provide sufficient benefit to continue clinical development. However, this trial provided an important landmark for direct intraparenchymal delivery of CNS therapeutics and for evaluation of a disease-modifying drug candidate. SNCA (α -synuclein) has been linked genetically to Parkinson's disease in familial Parkinson's disease, not only in the context of mutant SNCA (Polymeropoulos *et al*, 1997) but also with respect to overexpression of wild-type SNCA resulting from SNCA locus multiplication (Singleton *et al*, 2003; Chartier-Harlin *et al*, 2004; Ibáñez *et al*, 2004). Recently, the important discovery that SNCA is a susceptibility gene for sporadic Parkinson's disease provided further support for SNCA as a key target for therapeutic intervention (Maraganore *et al*, 2006). Taken together, these reports suggest that SNCA downmodulation represents a useful potential strategy for the treatment of Parkinson's disease, and that partial downmodulation of SNCA would be sufficient for therapeutic benefit. Farrer *et al* (2007; unpublished data) recently reported SNCA mRNA suppression after direct CNS infusion of siRNA-targeting SNCA. As with huntingtin, disease intervention at the SNCA mRNA level represents a novel and promising approach that is enabled by direct CNS delivery of RNA therapeutic modalities for reduction of a pathogenic protein.

Pain Management: Nerve Growth Factor and Artemin

Chronic pain represents a debilitating condition that severely impacts quality of life. Unlike acute pain, which has a protective role and is transient, chronic pain, such as neuropathic pain, persists and serves no useful purpose. Current therapies, predominantly consisting of small molecules targeting neurotransmitter receptors and ion channels, are inadequate in most patients and palliative rather than curative, with side effects representing significant limitations. The molecular pathways underlying chronic pain are complex and continue to be elucidated. The recent discovery that neurotrophic factors such as NGF and artemin are important in chronic pain suggests that

these pathways may serve as novel intervention points for disease-modifying therapy. Although NGF was originally identified as a key neurotrophin required for the survival of pain sensory neurons, subsequent studies implicated excessive NGF in inflammatory pain as well as possibly neuropathic pain (Lewin *et al*, 1994; Woolf *et al*, 1994; McMahon *et al*, 1995; Koltzenburg *et al*, 1999). NGF levels rise dramatically in damaged or inflamed tissue, and enhanced retrograde transport of NGF occurs in sensory neurons, leading to increased levels of substance P and CGRP, which can contribute to hyperalgesia. The pivotal role of NGF in mediating inflammatory hyperalgesia was subsequently established in animal models of chronic inflammation with TrkA-IgG fusion protein and anti-NGF antiserum. On the basis of these and other data, a clinical trial was conducted with RI624, a humanized anti-NGF antibody, for the treatment of osteoarthritis pain.

Another strategy for the treatment of chronic pain is supplementation with artemin, a member of the GDNF family. Like GDNF, artemin supports the survival of cultured sensory neurons (Baloh *et al*, 1998a,b). However, the distribution pattern of artemin's receptor, GFR α 3, is relatively restricted, being expressed predominantly in adult rodent and human nociceptive sensory neurons (Orozco *et al*, 2001; Yang *et al*, 2006), suggesting that artemin would specifically modulate the pain sensory system in the adult. This was established in the rat spinal nerve ligation model where behavioral, neurochemical, and histochemical changes associated with neuropathic pain were reversed with systemic administration of artemin (Gardell *et al*, 2003). More recently, expression of GFR α 3 immunoreactivity has also been reported in other tissues, including a subset of epithelial cells in the digestive and reproductive systems; the functional consequences of this expression remain to be determined (Yang *et al*, 2006). In contrast to small molecule drugs targeting individual ion channels or neurotransmitter receptors, these protein-based therapeutic strategies represent novel and promising approaches for the treatment of chronic pain that have the potential to impact multiple facets of pathology.

SUMMARY AND CONCLUSIONS

Therapeutic antibodies and oligonucleotide therapeutics such as antisense, siRNA, and aptamers represent new classes of drugs to inhibit 'nondrugable' targets, including the emerging class of protein-protein interaction targets. They represent excellent tools to validate targets by knocking out gene expression at the level of mRNA (antisense, RNAi, and siRNA) or protein (antibodies and aptamers). These novel therapeutic modalities (*vs* small molecules) have the distinct advantage of disrupting protein-protein interaction but have some limitations in their ADME properties with regard to limited oral absorption and their limited ability to cross the blood-brain barrier. Their utility for effective treatment of CNS

disorders is reviewed in the context of their limited access of relevant targets outside the blood-brain barrier (eg A β neutralization approaches for the treatment of AD; pain targets such as NGF and artemin in the periphery), addressing diseases with a compromised blood-brain barrier such as MS (anti-IL-23 approaches) or direct delivery of the therapeutic to brain using implantable delivery devices (inhibition of huntingtin and SNCA for the treatment of HD and PD, respectively). Broader application of these novel therapeutic modalities for the treatment of neurological and psychiatric disorders is envisioned as new technologies are developed to overcome the current limitations.

DISCLOSURES AND ACKNOWLEDGEMENTS

We thank Eric Gornstein for helpful research.

The authors are full-time employees of Archemix Corporation, a company focused on therapeutic aptamers (Errol B De Souza, Sharon T Cload, and P Shannon Pendergrast) and Alnylam Pharmaceuticals, a company focused on siRNA development (Dinah WY Sah). In addition, Errol B De Souza is on the following boards of directors or scientific advisory boards (Bionomics Ltd, IDEXX Laboratories, National Institute of Mental Health Board of Scientific Counselors, Palatin Technologies, and Targate Inc.).

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