siRNA relieves chronic neuropathic pain

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ABSTRACT

Double stranded, short interfering RNAs (siRNA) of 21-22 nt length initiate a sequence-specific, posttrancriptional gene silencing in animals and plants known as RNA interference (RNAi). Here we show that RNAi can block a pathophysiological pain response and provide relief from neuropathic pain in a rat disease model by down regulating an endogenous, neuronally expressed gene. Rats, intrathecally infused with a 21 nt siRNA perfectly complementary to the pain-related cation-channel P2X₃, showed diminished pain responses compared to missense (MS) siRNA-treated and untreated controls in models of both agonist-evoked pain and chronic neuropathic pain. This form of delivery caused no adverse effects in any of the animals receiving P2X₃ siRNA, MS siRNA or vehicle. Molecular analysis of tissues revealed that P2X₃ mRNA expressed in dorsal root ganglia, and P2X₃ protein translocated into the dorsal horn of the spinal cord, were significantly diminished. These observations open a path toward use of siRNA as a genetic tool for drug target validation in the mammalian central nervous system, as well as for proof of concept studies and as therapeutic agents in man.

INTRODUCTION

RNAi is a naturally occurring mechanism for regulating gene expression which has been observed in several model organisms and is mediated by double stranded RNA (dsRNA) (1–3). Previously, RNAi has been observed and applied in plants (4), in invertebrates (1,2), in mammalian cell culture (5,6) and in early mouse development (7–9). Recently, siRNAs, either administered directly or provided via a variety of expression constructs including virus vectors, have been used *in vivo* to demonstrate efficacy in inhibition of expression of several disease-related targets including VEGF (10,11), Caspase-8 (12), hepatitis B (13,14) and C viruses (15), β -catenin (16), TNF- α (17), AGRP (18), β -glucuronidase (19) and the endogenous Fas gene expressed in adult mouse liver (20).

The intracellular presence of dsRNA homologous to a gene results in post-transcriptional gene silencing via induced sequence-specific degradation of the corresponding mRNA. Mechanistic studies have shown that Dicer, a ribonuclease III, specifically cleaves long dsRNA into short 21-22-nucleotide (nt) dsRNA (siRNA) carrying 2-3-nt 3'-overhangs on both sense and antisense strands (21-23). These siRNAs subsequently serve as guide sequences that instruct a multicomponent nuclease (the RISC complex) to destroy complementary mRNA (23,24). Unlike plants and invertebrates, mammalian cells respond to dsRNAs that are greater than 30 bp in length. These responses include interferon production (25) and activation of two pathways: PKR (26) and 2'-5' polyadenylic acid (27). The former phosphorylates $eIF2\alpha$, leading to a global shutdown of protein synthesis, and the latter activates RNaseL, mediating non-specific degradation of cellular mRNA. Activation of such mechanisms has been circumvented in mammalian cell culture by using siRNAs directly (5).

These observations indicate that it will now be possible to down regulate adult mammalian genes on a broader scale in vivo with RNAi, as is already done systematically in Caenorhabditis elegans (28,29), and enable the comprehensive study, proof of concept and validation of the roles of putative disease genes in whole mammalian organisms. Although the field is advancing rapidly, inhibition of a pathophysiological condition upon treatment with an siRNA has, to date, only been reported for concavalin A-induced liver damage (20) to prevent sepsis (17). siRNAs have been applied successfully to down regulate gene expression in the central nervous system (CNS) (18); however, no experiments have yet been reported demonstrating a direct effect of an siRNA on a pathophysiological disease phenotype in the CNS. In the work described here, we have examined the efficacy of a 21-nt siRNA bearing 3'-overhangs of two phosphorothioated (PS) 2'-O-(2-methoxyethyl)-ribonucleotides (MOE) (30), targeting $P2X_3$ in vivo, using a system we had previously established for administration of P2X₃ antisense oligonucleotides (ASO) (31) in the CNS, in which we measured effects of specific intrathecally administered oligonucleotides in a chronic neuropathic pain model. The siRNA sequence used in this

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study was chosen based upon the optimal ASO sequence selected in previous studies (32) (indicating that the site is available for oligonucleotide targeting) and which, after thorough testing in vitro, was shown to be even more potent than the original ASO (33). The purpose of our study was twofold. Firstly, our goal was to evaluate the roles of putative pain genes in pain models in order to identify novel drug targets. Secondly, our plan was to test to see if siRNAs could be used therapeutically to alleviate a disease phenotype. We chose P2X₃ as our first target for testing the principle of siRNA since we had already proven its role in pain via ASO treatment in vivo and had already established all the conditions for optimal delivery of oligonucleotides. Studies on P2X₃ knockout mice also indicated that this gene plays a role in inflammatory pain signalling (34,35). This system, therefore, provided the ideal means for testing the efficacy of siRNAs in vivo in an adult animal disease model in the CNS. Administration of these potent P2X₃ siRNAs in vivo resulted in a dramatic relief of pain as indicated by a significant inhibition in the neuropathic pain response, while exhibiting no apparent toxicity or non-specific side effects. Furthermore, in addition to being safer, siRNA was also more efficacious when compared with the administration of equivalent doses of ASO.

MATERIALS AND METHODS

siRNA and ASO compounds

The ASO compound has been previously published (31). $P2X_3$ siRNA sequences were synthesized (5'-UCACUCGGCUG-GAUGGAGUtt and 5'-ACUCCAUCCAGCCGAGUGAag), together with a corresponding 4-nt missense (MS) control siRNA sequence (5'-UCACUGCGCUCGAUGCAGUtt and 5'-ACUGCAUCGAGCGCAGUGAag) where lower case letters represent phosphorothioated ribonucleotides carrying a 2'-O-(2-methoxyethyl)-group (30).

Details of siRNA synthesis

Synthetic oligoribonucleotides were prepared using standard TOM-phosphoramidite chemistry (Xeragon AG) on an OligoPilot II synthesizer (Amersham Pharmacia Biotech) at 180 µmol scale. Phosphoramidites were dissolved in acetonitrile at 0.2 M concentration, mixed in a 1:1 ratio with a 0.2 M solution of benzimidazolium triflate in acetonitrile for coupling over 5 min. A first capping was made using standard capping reagents. Sulfurization was made by using a 0.05 M of solution N-ethyl,N-phenyl-5-amino-1,2,4-dithiazol-3thione for 2 min. Oxidation was made by a 0.5 M tbutylhydroxyperoxide in dichloromethane for 2 min. A second capping was performed after oxidation or sulfurization. Oligonucleotides were detritylated for the following coupling by 2% dichloroacetic acid in dichloroethane. Upon completion of the sequences, the support-bound compounds were cleaved and base and phosphodiester deprotected as 'Trityl-on' material by a Methylamine solution (41% aqueous methylamine/33% ethanolic methylamine 1:1 v/v) at 35°C for 6 h. Resulting suspensions were lyophilized to dryness. 2'-O-silyl groups were removed upon treatment with 1 M tetrabutylammonium fluoride (10 min at 50°C and 6 h at 35°C). The obtained crude solutions were directly purified by RP-HPLC. The purified detritylated compounds were analysed by capillary gel electrophoresis for purity and quantified by UV according to their extinction coefficient at 260 nM. Identity was checked by electrospray mass spectrometry (5'-UCA-CUCGGCUGGAUGGAGUtst: $M_{calc} = 6866.5$ Da, $M_{meas} = 6826.5$ Da; 5'-ACUCCAUCCAGCCGAGUGAasg: $M_{calc} = 6826.5$ Da, $M_{meas} = 6830.5$ Da; 5'-UCACU<u>GCGCUC</u>-GAUGCAGUtst: $M_{calc} = 6784.4$ Da, $M_{meas} = 6784.0$ Da; 5'-ACUGCAUCGAGCGAGUGAasg: $M_{calc} = 6904.5$ Da, $M_{meas} = 6903.5$ Da).

siRNA annealing

For annealing of siRNA, 1 mM single strands were incubated in isotonic buffer (100 mM potassium acetate, 30 mM HEPES–KOH, 2 mM magnesium acetate, 26 mM NaCl, pH 7.4 at 37°C) for 5 min at 90°C followed by 1 h at 37°C. These conditions differ slightly from the original conditions described by Elbashir *et al.* (5): higher concentrations of oligonucleotides were used for annealing, which was confirmed by gel analysis; denaturation at 90°C was performed for 5 min instead of 1 min; NaCl was added to assure a nonirritating, isotonic solution for *in vivo* testing. At the time of annealing, both strands were fully deprotected.

Animal models

All experiments were carried out according to Home Office (UK) guidelines and with approval of the local Novartis Animal Welfare and Ethics Committee. The partial ligation model and infusion of siRNA, MS siRNA and ASO into naïve and neuropathic rats via minipump as well as behavioural testing for mechanical hyperalgesia and allodynia were performed as described previously (31,36). The intrathecal dose of 400 µg/day P2X₃ siRNA and MS siRNA or 180 µg/ day P2X₃ ASO was continuously infused over the course of 6–7 days. Each animal was tested in a blind random order. Statistical significance of mechanical hyperalgesia and allodynia data were analysed with ANOVA followed by Tukey's HSD test, **P* < 0.05, ***P* < 0.005, ****P* < 0.001.

Molecular analysis

Fixed, cryopreserved spinal cord was dissected and treated with an immunohistochemistry protocol described previously (31) with one difference: as a control, biotinylated IB4 (Sigma, Buchs, Switzerland; 1:35) was used instead of anti-CGRP streptavidin-TRITC followed by (Jackson ImmunoResearch, West Grove, PA; 1:200). Dorsal root ganglia (DRG) from non-perfused rats were snap-frozen and RNA was isolated separately for each animal by crushing the frozen DRG with mortar and pestle and syringing 10 times to fragment genomic DNA. The RNA was then purified with RNeasy Columns (Qiagen, Basel, Switzerland). 50 ng of each sample was analysed three times with the ABI PRISM[™] 7700 Sequence Detector (PE Applied Biosystems, Foster City, CA) as described previously (31). Values were then normalized to β -actin analysis done in parallel on the same samples. Statistics: ANOVA followed by Tukey's HSD test, *P < 0.05.

RESULTS AND DISCUSSION

PS-MOE modifications are known to offer superior stability against nucleases when incorporated into ASO and have proven particularly useful in *in vivo* studies (31). Used as 3'-

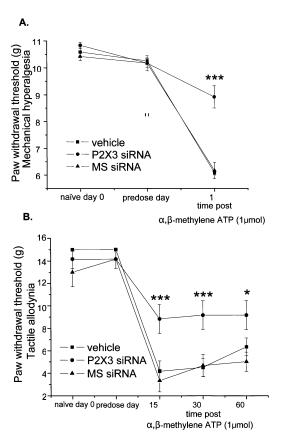


Figure 1. P2X₃ siRNA reduced agonist-induced pain *in vivo*. Significant inhibition of α , β -methylene-ATP evoked (**A**) mechanical hyperalgesia and (**B**) tactile allodynia in the hind paw of naïve rats after 6-day intrathecal treatment with P2X₃ siRNA compared to MS siRNA or vehicle. PWTs were measured prior to cannulation on day 0, and on day 6 before and after intraplantar injection of 1 µmol α , β -methylene-ATP (*n*/group = 6).

overhangs in siRNAs, these RNA analogues have been shown not to adversely affect the RNAi mechanism in cell culture experiments compared to standard siRNAs carrying deoxythymidine dimers (dTdT) (33). Furthermore, this chemical modification, 2'-MOE at two 3' terminal nucleotides of each strand linked via a phosphorothioate, has been shown to be just as potent in vitro as native dsRNA (33) and was included as a chemical protection against possible in vivo nuclease degradation. P2X₃ is an ATP-gated cation channel, expressed in a restricted fashion in the cell bodies of sensory neurons and their projections. Evidence for its involvement in pain is based on the regulation of P2X₃ message expression in animal models of chronic pain and altered pain-related responses in knockout mice (34,35). More direct proof was provided by our demonstration of an inhibition of hyperalgesia in a model of neuropathic pain by ASO delivery, a finding which has been independently confirmed (37) and is further supported by the recent report of analgesic activity of a small molecule antagonist of P2X₃ in a variety of animal models of pain (38,39). Since we have already established a role for P2X₃ in a rat model of neuropathic pain, and have delineated the optimal conditions for delivery of oligonucleotides in this model, this target was chosen to test the principle of the utility of siRNA in validating potential targets in disease states in vivo.

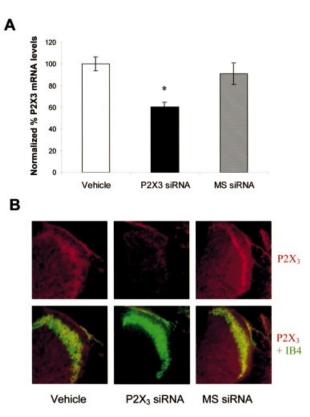


Figure 2. Molecular down regulation of P2X₃ RNA and protein upon treatment with siRNA. (**A**) P2X₃ mRNA levels in lumbar L4/L5 DRG, normalized to β -actin, were significantly reduced in P2X₃ siRNA- versus MS siRNA- and vehicle-treated rats (*n*/group = 3). (**B**) P2X₃ immunoreactivity in lamina II of the spinal cord was visibly reduced in siRNA-treated animals (top panel). Co-immunostaining with neuron-specific isolectin B4 (green) showed that equivalent regions of each spinal cord were analysed (*n*/group = 3).

P2X₃ siRNA sequences, complementary to the same region of the gene targeted in our earlier ASO study (31), together with an appropriate mismatch control were synthesized. The reagents that we employed in this study were thoroughly characterized in vitro for their ability to inhibit P2X₃ at mRNA, protein and functional levels (33). In preliminary experiments, varying doses of siRNAs were delivered intrathecally into the spinal fluid of naïve rats (n/group = 3)via an indwelling cannula attached to an osmotic minipump (data not shown). This work established that doses up to 400 ug/day did not elicit any signs of neurotoxicity (hind limb paralysis, vocalization or anatomical damage to the spinal cord). This dose, approximately equivalent to the ASO dose evaluated in our previous study (31), was used in all subsequent studies, resulting in significant and specific behavioural effects. Thus, following intrathecal delivery of P2X₃ siRNA for 6 days in naïve rats, an intraplantar injection of the P2X₃ agonist α , β -methylene ATP (1 µmol) into the left hindpaw resulted in a diminished level of mechanical hyperalgesia (hypersensitivity to a noxious pressure stimulus of ~ 100 g) and tactile allodynia (hypersensitivity to a normally non-noxious pressure stimulus of ~15-20 g), as measured by a reduction in paw withdrawal thresholds (PWTs) (P < 0.001, Fig. 1) compared to vehicle- and MS siRNA-treated rats. Molecular analysis using tissues isolated from the rats

revealed that the down regulation of $P2X_3$ mRNA levels in DRG and protein levels in the dorsal horn of the spinal cord correlated with the decrease in pain response in $P2X_3$ siRNA-treated rats (Fig. 2). Quantification of $P2X_3$ mRNA from DRG using quantitative RT–PCR, the most sensitive method of mRNA quantitation, demonstrated 40% down regulation of the target for $P2X_3$ siRNA-treated animals compared to MS siRNA and vehicle controls. The incomplete knock-down of target mRNA may be a result of residual mRNA in the cell nucleus, or may simply reflect an incomplete transfection of DRGs. $P2X_3$ protein was analysed immunohistochemically at its site of action in the entire dorsal horn of lumbar spinal cord tissue slice and appeared visibly down regulated in lamina II in the $P2X_3$ siRNA group compared to MS siRNA and vehicle-treated controls (Fig. 2B).

The significant reduction of mechanical hyperalgesia and tactile allodynia observed in the agonist-induced pain model described above was subsequently confirmed in the partial sciatic ligation model of neuropathic pain (40). Within 2 days of initiation of treatment, P2X₃ siRNA, but not MS siRNA or vehicle, produced a significant reduction (P < 0.001) in tactile allodynia which was maintained for the duration of the siRNA administration (Fig. 3A). In the same experiment, administration of P2X₃ ASO at a dose of 180 μ g/day, in contrast, did not reduce allodynia, confirming results from our earlier study (31). The cannulation process itself does not cause any change in nociceptive indices as indicated by the PWTs of the naïve vehicle group. In the contralateral paws, where the nerve was not ligated (Fig. 3B), all groups exhibited similar PWTs to the naïve controls. However, intraplantar application of α,β methylene ATP on the final day of the experiment evoked tactile allodynia, measurable 15, 30 and 60 min post-agonist injection in vehicle-, ASO- and MS siRNA-treated groups. In contrast, the allodynic response was significantly (P < 0.001)diminished in the P2X₃ siRNA-treated group, demonstrating a functional down regulation of the $P2X_3$ channel (Fig. 3B). Taken together with the effects observed with the match and mismatch siRNAs (sequences of which do not exist in any other P2X family member (e.g. P2X1 or P2X2) and the fact that the P2X₃ isoform is the primary α , β -methylene-ATP sensitive P2X target expressed in DRG neurons (36), confines the functional effects of this siRNA to inhibition of P2X₃. When the same animals were tested for mechanical hyperalgesia, a significant reversal of PWTs was observed in neuropathic paws 1 day after initiating treatment, and this was maintained for the duration of the application of P2X₃ siRNA and ASO. However, the double-stranded P2X₃ siRNA (400 μ g/day) was more effective than the ASO at approximately equimolar concentrations (180 µg/day). Reversal of mechanical hyperalgesia was not seen in the control groups (Fig. 4A), supporting a specific down regulation of P2X₃ receptors. Mechanical hyperalgesia was not observed in the contralateral paws of any group until the agonist α , β -methylene ATP was injected on the last day of the experiment, whereupon vehicleand MS siRNA-treated groups exhibited a considerable decrease in PWTs (Fig. 4B). Remarkably, P2X₃ siRNA treatment completely suppressed the hyperalgesic response of α , β -methylene ATP, whereas P2X₃ ASO diminished it by more than 50%, both confirming functional down regulation of the channel (Fig. 4B). Both measurements of the hyperalgesic response (Fig. 4A and B) confirm that the P2X₃ siRNA reagent

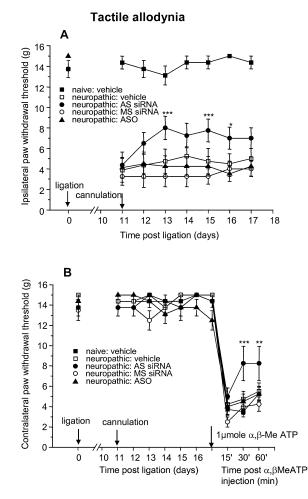


Figure 3. P2X₃ siRNA reduced tactile allodynia in a neuropathic pain model. (A) Tactile allodynia was significantly inhibited in the neuropathic paw (partially ligated sciatic nerve) in P2X₃ siRNA- versus MS siRNA-, P2X₃ ASO- and vehicle-treated rats. An additional naïve (non-ligated) group that received vehicle only controlled for the cannulation process. (B) PWTs in the contralateral paws of all groups were the same as naïve animals. Upon agonist injection (1 µmol α,β -methylene-ATP) on the last day of the experiment (day 17), all control groups and the ASO group showed agonist-induced tactile allodynia when assayed at 15, 30 and 60 min post injection. P2X₃ siRNA significantly diminished agonist-induced tactile allodynia (*n*/group = 8).

is considerably more effective at reducing pain than the analogous ASO and this difference may be one possible explanation as to why the $P2X_3$ siRNA is more potent than the analogous ASO in the allodynia model measurements.

An important observation was the absence of any apparent toxicity due to intrathecal administration of siRNAs at the doses described in these studies. There were no animals lost; no hind leg paralysis was observed, and no signs of discomfort were exhibited throughout the experiments. The rats were carefully observed during the experiments, both during the behavioural studies and during dissection of the DRG and spinal cord tissue. Although the siRNA was administered intrathecally directly into the spinal fluid, there were no observable behavioural changes and no obvious signs of spinal inflammation upon dissection.

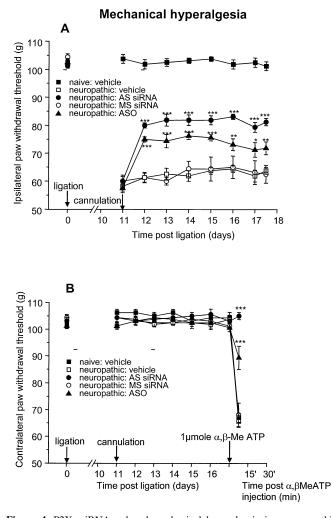


Figure 4. P2X₃ siRNA reduced mechanical hyperalgesia in a neuropathic pain model. (**A**) Mechanical hyperalgesia was significantly inhibited in the neuropathic paw (partially ligated sciatic nerve) in P2X₃ siRNA- and in P2X₃ ASO- versus MS siRNA- and vehicle-treated rats. As a control, an additional naïve vehicle group had not been ligated and showed normal pain perception. (**B**) PWTs in the contralateral paws of all groups was the same as normal animals. Upon agonist injection (1 µmol α,β -methylene-ATP) on the last day of the experiment (day 17), all groups showed agonist-induced mechanical hyperalgesia at 15 min after injection, except for the P2X₃ siRNA- and the ASO-treated groups. P2X₃ siRNA completely suppressed agonist-induced mechanical hyperalgesia (*n*/group = 8).

In summary, we have shown in an agonist-induced pain model, that siRNA produced a significant behavioural reduction of mechanical hyperalgesia and allodynia, which correlated with a molecular down regulation of P2X₃ in DRG and spinal cord. This behavioural effect was subsequently reproduced in a disease model of chronic neuropathic pain. The reversal of pain was not complete, possibly due to the partial down regulation of P2X₃, or more likely, the involvement of proteins other than P2X₃ in neuropathic pain.

However, it has been already demonstrated that partial inhibition of $P2X_3$ with ASO treatment produces significant behavioural effects (31,37). In general, the behavioural effects produced by application of $P2X_3$ siRNA were similar to those observed after administration of an equivalent dose of $P2X_3$ ASO; however, an important difference was the reversal of

allodynia by the former, an effect that we did not observe upon ASO treatment. Both the knock-out animals and the siRNAtreated animals clearly show an important role of P2X₃ in pain behaviour. The experiments are complementary, as P2X₃ knock-out animals were tested for sensitivity to warming stimuli and bladder response, whereas the siRNA-treated animals were assessed in agonist-induced and neuropathic pain models. Concerning neuropathic pain, the effects observed using these siRNAs correlate very well with those from two recently published studies characterizing a low molecular weight inhibitor of P2X₃ and P2X_{2/3} (38,39).

In comparison to previous reports, our results are novel in three important respects: firstly, an endogenous diseaserelated neuronal gene in the nervous system has been targeted. Secondly, siRNA was delivered intrathecally to the spinal fluid, a delivery method which is commonly used for clinical treatment of severe pain. Thirdly, and most importantly, the readout was a measurable pathophysiological response in a relevant and well-defined disease model. These data have immediate implications in the study of gene function in that the mechanism is sequence specific, rapid and scalable, allowing one to envisage multiparallel analyses on collections of genes for *in vivo* target validation in drug discovery.

Finally, considering its effectiveness *in vivo* and relative lack of toxicity even at a high concentration, these data indicate that siRNA could be used extensively to test the involvement of potential novel targets in disease processes, both in animal models and in man. This is particularly attractive where drugs or small molecule compounds directed at these targets are not available. On a more speculative level there is the additional possibility that RNAi can be used itself as a therapy for certain disease states where delivery and target protein knockdown can be optimized.

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