

A small molecule that binds Hedgehog and blocks its signaling in human cells

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Small-molecule inhibition of extracellular proteins that activate membrane receptors has proven to be extremely challenging. Diversity-oriented synthesis and small-molecule microarrays enabled the discovery of robotnikinin, a small molecule that binds the extracellular Sonic hedgehog (Shh) protein and blocks Shh signaling in cell lines, human primary keratinocytes and a synthetic model of human skin. Shh pathway activity is rescued by small-molecule agonists of Smoothed, which functions immediately downstream of the Shh receptor Patched.

Sonic hedgehog (*Shh*), the most widely characterized of the Hedgehog homologs, is essential for proper embryonic development¹⁻³. The *Shh* pathway involves the autocleavage of full-length Shh into an active 20-kDa N-terminal fragment (ShhN), which binds to its 12-pass transmembrane receptor, Patched (Ptc1), reversing its inhibitory effect on Smoothed (Smo). One effect of this de-repression is the activation of Gli transcription factors, which regulate the transcription of target genes that include *Gli1* and *Ptch1* (ref. 1).

Several synthetic and natural small-molecule modulators of Smo, an apparent member of the family of G protein-coupled receptors (GPCRs), have been discovered using cell-based phenotypic screens⁴. *Shh* signaling antagonists that bind to Smo include cyclopamine (**3**), SANT1 (**4**) and Cur-61414 (**5**)^{5,6}. *Shh* signaling agonists that bind to Smo include the synthetic small molecules purmorphamine (**6**) and Hh-Ag1.2 (SAG; **7**)^{7,8}. Furthermore, small molecules that inhibit Shh signaling downstream of Smo, GANT61 (**8**) and GANT58 (**9**), have also been reported (Supplementary Fig. 1

online)⁹. The discovery of small-molecule modulators of *Shh* signaling provides an avenue to regulate the activity of a pathway implicated in medulloblastoma, basal cell carcinoma (BCC), pancreatic cancer, prostate cancer and developmental disorders^{2,10,11}. Phase 1 clinical trials in BCC and pancreatic cancer involving the Smo antagonists GDC-0449 and IPI-926 (NCT00607724 and NCT00761696 at <http://clinicaltrials.gov>, respectively) are currently underway^{12,13}.

None of the reported synthetic Shh pathway inhibitors are known to target the Shh protein itself. To our knowledge, all reported examples of discoveries of small-molecule Shh signaling modulators resulted from the use of cell-based phenotypic assays. Target-based discovery of modulators of Shh signaling was expected to provide a complementary approach.

Small-molecule microarray (SMM)-based screens have enabled the discovery of small molecules that bind target proteins of interest and modulate the cellular functions of their targets¹⁴⁻¹⁶. In this system, small molecules have been linked covalently to a glass surface and screened for binding to either a purified protein or an epitope-tagged protein in a complex cell lysate^{17,18}. Here, we report a screen of bacterially expressed ShhN using SMMs containing a collection of approximately 10,000 diversity-oriented synthesis (DOS) compounds and natural products arrayed on a single microscope slide¹⁹.

In the ShhN SMM screen, some structurally related macrocycles emerged as noteworthy assay positives. A representative macrocycle, **1** (Fig. 1a), was retested for binding to ShhN via surface plasmon resonance (SPR) (Fig. 1b). The compound exhibited binding to ShhN in a concentration-dependant fashion with a K_D of 9 μ M, determined

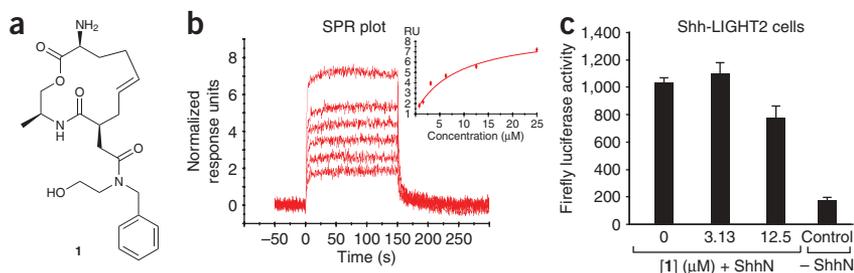


Figure 1 Characterization of SMM hit **1**. (a) Structure of **1**. (b) SPR plot of **1** binding to purified ShhN. The plot shows normalized response units (RUs) on the y axis and time (s) on the x axis. The concentrations plotted are 0.78 μ M, 1.56 μ M, 3.13 μ M, 6.25 μ M, 12.5 μ M and 25 μ M, in order of increasing normalized RUs. (c) Luminescence plots for a Gli-dependent firefly luciferase reporter gene assay of **1** at the indicated concentrations. ShhN represents a positive control for medium containing ShhN palmitoylated at the N terminus. The assays were performed at 0.25% (v/v) DMSO. Each value represents the average of five experiments, with the error bar denoting the s.d.

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by fitting steady state data. To our knowledge, this is the first reported discovery of a small molecule capable of binding to the ShhN protein.

We examined the activity of **1** in Shh-LIGHT2 cells (American Type Culture Collection)²⁰, which is an NIH3T3 cell line with a Gli-dependent firefly luciferase reporter. These cells have been widely used to demonstrate the efficacy of Shh pathway inhibitors (cyclopamine) and activators (purmorphamine and SAG)^{4,6,21}. Shh pathway activity was inferred by measuring firefly luciferase levels after a 30 h incubation with compound in the presence of N-palmitoylated ShhN. The compound exhibited moderate Shh pathway inhibition (**Fig. 1c**) and did not demonstrate cytotoxicity at any of the experimental concentrations based on a cell titer viability assay run in parallel (see **Supplementary Methods** online). This raised the possibility that the ShhN binding was related to the moderate Shh pathway inhibition.

As part of an ongoing structure-activity relationship effort, we identified a 12-membered macrocycle, which we have named robotnikinin (**2**), that showed increased ShhN binding and displayed a substantially longer dissociation time than **1**. Based on SPR experiments, robotnikinin (**Fig. 2a**) demonstrated ShhN binding capacity at concentrations between 1.56 μ M and 25 μ M, with a K_d of 3.1 μ M derived from kinetic data (**Fig. 2b**). When this compound was tested in an NIH3T3 line transformed with a Gli-luciferase construct, or Shh-LIGHT2 cells¹¹, it showed concentration-dependent inhibition of ShhN-induced pathway activation (**Fig. 2c**). Additionally, at concentrations above 30 μ M, the macrocycle exhibited inhibition comparable to that observed upon treatment with 6.25 μ M cyclopamine in Shh-LIGHT2 cells. No substantial cytotoxicity was observed as judged by cell titer measurements using a cell viability assay.

To explore further the potential mechanism of Shh pathway inhibition involving direct perturbation of the ShhN protein complex,

the same compounds were tested in a *Ptch1*^{-/-} cell line derived from mouse embryos lacking *Ptch1* function. The cell line had both *Ptch1* alleles replaced with a β -galactosidase (β -gal) reporter²¹. Because *Ptch1* inhibits Hh pathway activation by repressing Smo function and is also a target gene, removing both *Ptch1* alleles results in constitutive pathway activation and β -gal expression. Small-molecule pathway inhibitors that act downstream of *Ptch1* remain active in this cell line. In the *Ptch1*^{-/-} cell line, Shh pathway activity is proportional to the β -gal levels observed after 30 h of incubation with compound. No substantial difference was observed when the *Ptch1*^{-/-} cell line was treated with N-palmitoylated ShhN or low serum-containing culture medium, which confirms that with the *Ptch1* receptor absent, the Shh pathway is constitutively activated and ShhN does not increase pathway activation²¹. Previous studies have demonstrated that cyclopamine, whose target (Smo) is downstream of *Ptch1* (refs. 6,22), is effective at ablating β -gal reporter activity in this cell line. In our study, treatment with 6.6 μ M cyclopamine resulted in substantial pathway inhibition. In contrast, no pathway inhibition was observed using robotnikinin at any of the concentrations tested after normalizing luminescence data for cell titer (**Supplementary Fig. 2a** online). These results support a model in which this small molecule inhibits the Shh pathway upstream of *Ptch1* in Shh-LIGHT2 cells.

Our model predicts that treatment of Shh-LIGHT2 cells with the Smo agonists purmorphamine and SAG would override the inhibitory effect of robotnikinin given that Smo functions downstream of Shh/*Ptch1*. When we tested the model by co-administering 3.6 μ M purmorphamine in addition to various concentrations of robotnikinin, virtually all of the inhibitory effect was eliminated. This effect was recapitulated when 100 nM SAG was co-administered (**Fig. 2c**). Next, we sought to test our model in the context of a different cell

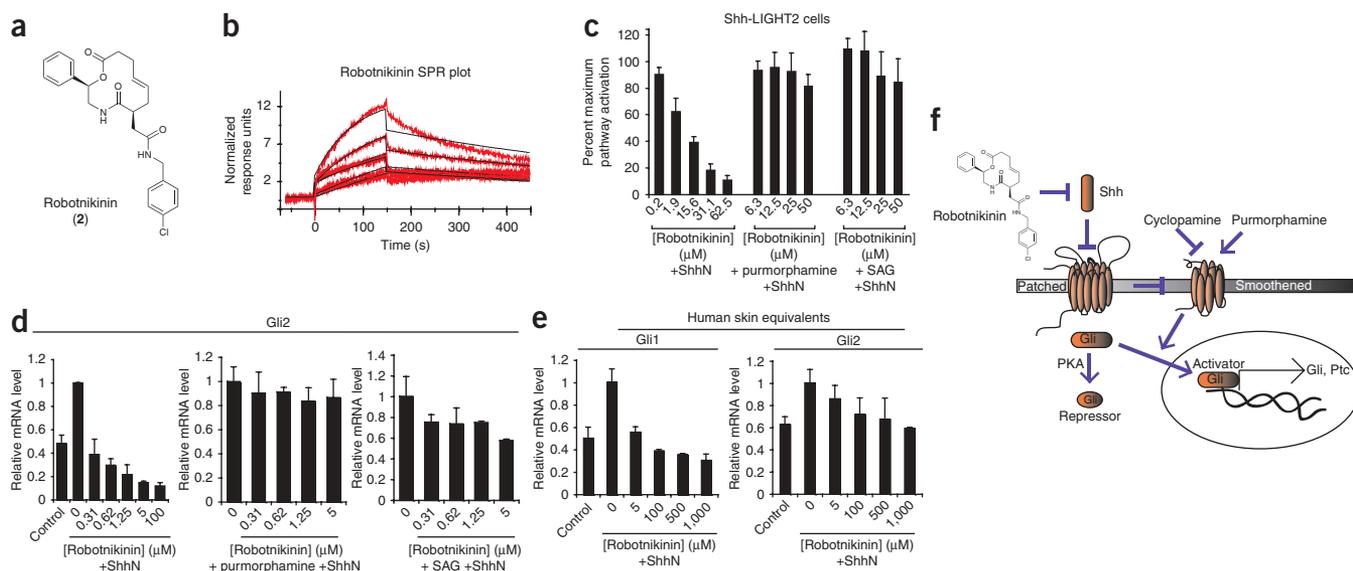


Figure 2 Robotnikinin. **(a)** The structure of robotnikinin, which is the compound that resulted from follow-up chemistry efforts to optimize potency. **(b)** SPR curve of robotnikinin showing concentration-dependent binding to purified ShhN. Normalized RUs are plotted over a time course. The concentrations plotted are 1.56 μ M, 3.13 μ M, 6.25 μ M, 12.5 μ M and 25 μ M, in order of increasing Rus. **(c)** Inhibition of Gli signaling by robotnikinin in Shh-LIGHT2 cells stimulated with medium containing ShhN palmitoylated at the N terminus, relative to 6.25 μ M cyclopamine (a small-molecule inhibitor of Smoothened). Shh-LIGHT2 cells stimulated with N-palmitoylated ShhN along with 3.6 μ M purmorphamine or 100 nM SAG (small-molecule activator of Smoothened) showed negligible inhibition at the indicated concentrations of inhibitor. **(d)** Robotnikinin lowers levels of endogenous Gli2 mRNA (analyzed by qPCR) in primary human keratinocytes in a dose-dependent manner; this effect is blocked by the co-administration of Smo agonists. Note that there is some Gli expression in the absence of exogenous Shh due to the presence of a basal amount of Shh in the growth medium. **(e)** When analyzed by qPCR, synthetic human skin displayed Gli1 and Gli2 transcriptional repression in the presence of varying concentrations of robotnikinin. **(f)** Robotnikinin inhibits the induction of the Shh pathway. Our experiments support a mechanism involving inhibition of the actions of Shh, either directly or indirectly by interfering with a precursor complex. All error bars show s.d.

line, without reliance on a reporter gene construct. C3H10T1/2 cells are an immortalized mouse mesenchymal stem cell line that differentiate to osteoblasts upon treatment with N-palmitoylated ShhN, with alkaline phosphatase as a reliable marker of the transformation^{8,23}. Robotnikinin showed dose-dependent inhibition of alkaline phosphatase induction in the cell line in the presence of ShhN, but no detectable inhibition when 3.6 μ M purmorphamine was co-administered (**Supplementary Fig. 2b**). These data indicate that the inhibition and epistatic rescue of robotnikinin was not dependent on the Shh-LIGHT2 cell line. Consistent with its mode of action upstream of Smo and in contrast with other small-molecule modulators of Shh signaling, we also determined that at concentrations up to 25 μ M, robotnikinin does not compete for Smo binding with BODIPY-cyclopamine in a Smo-overexpressing human embryonic kidney (HEK) 293 cell line (**Supplementary Fig. 2e**).

To obtain direct evidence of the ability of robotnikinin to block Shh signaling, we investigated its actions on two types of human primary cells on endogenous levels of transcripts resulting from Shh-induced genes. The first study used real-time PCR (qPCR) to measure quantitatively levels of mRNA for the Shh-induced transcription factors Gli1 and Gli2 in primary human keratinocytes. When primary human keratinocytes were stimulated with N-palmitoylated ShhN and incubated with robotnikinin, Gli2 and Gli1 transcription was markedly repressed by 30 h (**Fig. 2d** and **Supplementary Fig. 2c**). Most Gli2 transcription was abolished in the presence of 100 μ M robotnikinin at the 30 h time point. When Smo agonists purmorphamine and SAG were co-administered with robotnikinin, an active Shh pathway phenotype was observed with a loss of response to robotnikinin, as evidenced by Gli2 and Gli1 mRNA levels (**Fig. 2d** and **Supplementary Fig. 2c**).

A second study used synthetic skin that was prepared by first populating dehydrated collagen matrix, itself derived from human skin grafts, with primary human keratinocytes. Next, several dermal layers were formed by culturing over an extended period. Following incubation with robotnikinin, the primary human synthetic skin tissue was analyzed by qPCR for levels of Gli1 and Gli2 transcripts. These experiments revealed that the tissue displayed reduced levels of Gli1 and Gli2 mRNA while remaining histologically normal (**Fig. 2e** and **Supplementary Fig. 2e**). Robotnikinin was thus found to retain its activity in human-derived tissue.

We have described to our knowledge the first example of a small molecule that binds to purified ShhN protein. Robotnikinin inhibits Shh signaling in a concentration-dependent manner but exhibits no inhibitory activity in a cell line lacking the Ptc1 receptor, does not compete with cyclopamine-Smo interactions and does not exhibit an inhibitory effect in the presence of the well-characterized Smo agonists purmorphamine and SAG. Robotnikinin displays substantial repression of Shh-induced Gli1 and Gli2 in primary human skin cells and human-derived skin tissue. In light of the ShhN-binding properties of the macrocycle and the results of our epistasis analyses (the macrocycle's lack of substantial Shh pathway inhibition, using Gli activity as a surrogate for pathway activity in the *Ptc1*^{-/-} cell line, and the ability of two agonists of the downstream Smo to override its effects), we propose a new mechanism of action involving direct targeting of the ShhN protein complex (**Fig. 2f**). Recent evidence has indicated that Hh signaling is facilitated by HhN binding partners Ihog, Boi and

heparin in *Drosophila melanogaster*, and Shh binding partners (Ihog orthologs) Cdo and Boc in vertebrates^{24,25}. The data presented here suggest that robotnikinin interferes with the ability of the ShhN protein complex to relay its signal efficiently to Ptc1. Small molecule-mediated disruption of protein-protein interactions involving extracellular growth and differentiation factors with their receptors is in general challenging, yet the process described here led directly to a success. The discovery of robotnikinin has provided a powerful small-molecule probe of an important signaling pathway involving a step in the pathway not previously accessible to small-molecule modulation. We believe that robotnikinin will be especially valuable as a probe of diseases associated with aberrant Shh pathway activity.

Note: Supplementary information and chemical compound information is available on the Nature Chemical Biology website.

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COMPETING INTERESTS STATEMENT

The authors declare competing financial interests: details accompany the full-text HTML version of the paper at <http://www.nature.com/naturechemicalbiology/>.

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