



Therapeutic antagonism of the neurokinin 1 receptor in endosomes provides sustained pain relief

Alan Hegron^{a,b,c,1}, Chloe J. Peach^{a,b,c,1}, Raquel Tonello^{a,b,c,1}, Philipp Seemann^{d,1}, Shavonne Teng^{a,b,c}, Rocco Latorre^{a,b,c}, Harald Huebner^d, Dorothee Weikert^d, Jeanette Rientjes^e, Nicholas A. Veldhuis^f, Daniel P. Poole^f, Dane D. Jensen^{a,b,c,g}, Alex R. B. Thomsen^{a,b,c}, Brian L. Schmidt^{a,b,c,g}, Wendy L. Imlach^h, Peter Gmeiner^d, and Nigel W. Bunnett^{a,b,c,2}

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The hypothesis that sustained G protein-coupled receptor (GPCR) signaling from endosomes mediates pain is based on studies with endocytosis inhibitors and lipid-conjugated or nanoparticle-encapsulated antagonists targeted to endosomes. GPCR antagonists that reverse sustained endosomal signaling and nociception are needed. However, the criteria for rational design of such compounds are ill-defined. Moreover, the role of natural GPCR variants, which exhibit aberrant signaling and endosomal trafficking, in maintaining pain is unknown. Herein, substance P (SP) was found to evoke clathrin-mediated assembly of endosomal signaling complexes comprising neurokinin 1 receptor (NK₁R), G $\alpha_{q/i}$, and β arrestin-2. Whereas the FDA-approved NK₁R antagonist aprepitant induced a transient disruption of endosomal signals, analogs of netupitant designed to penetrate membranes and persist in acidic endosomes through altered lipophilicity and pK_a caused sustained inhibition of endosomal signals. When injected intrathecally to target spinal NK₁R+ve neurons in knockin mice expressing human NK₁R, aprepitant transiently inhibited nociceptive responses to intraplantar injection of capsaicin. Conversely, netupitant analogs had more potent, efficacious, and sustained antinociceptive effects. Mice expressing C-terminally truncated human NK₁R, corresponding to a natural variant with aberrant signaling and trafficking, displayed attenuated SP-evoked excitation of spinal neurons and blunted nociceptive responses to SP. Thus, sustained antagonism of the NK₁R in endosomes correlates with long-lasting antinociception, and domains within the C-terminus of the NK₁R are necessary for the full pronociceptive actions of SP. The results support the hypothesis that endosomal signaling of GPCRs mediates nociception and provides insight into strategies for antagonizing GPCRs in intracellular locations for the treatment of diverse diseases.

signaling | receptors | endocytosis | pain

G protein-coupled receptors (GPCRs) are seven transmembrane domain receptors for hormones and neurotransmitters. They control homeostatic and disease processes and are therapeutic targets for disease (1). Drug discovery has focused on targeting GPCRs at the cell surface. However, plasma membrane signaling of GPCRs is usually tightly regulated and may not exclusively mediate long-lasting pathology. After binding to extracellular ligands, GPCRs are phosphorylated by GPCR kinases (GRKs) and interact with β arrestins (β ARRs), which uncouple GPCRs from G proteins and desensitize plasma membrane signaling (2, 3). β ARRs also couple GPCRs to clathrin and adaptor protein-2, which leads to dynamin (Dnm)-mediated vesicle scission and mediates receptor endocytosis. Although endocytosis has been considered as a mechanism that terminates GPCR signaling, many GPCRs continue to signal after endocytosis, including most class B GPCRs that interact with β ARRs with sustained kinetics and some class A GPCRs that exhibit transient interactions with β ARRs (4–15). Within endosomes, GPCR, G α , and/or β ARR signaling complexes (signalosomes) generate second messengers and activate kinases in subcellular compartments. Endosomal signaling of GPCRs has been implicated in certain physiological and pathological processes, including hormonal control (4, 7) and pain transmission (9–11, 15–17). Thus, GPCRs in endosomes, rather than at the cell surface, might be the optimal therapeutic target (18).

Painful stimuli provoke the release of substance P (SP) from primary sensory neurons, which stimulates endocytosis of the neurokinin 1 receptor (NK₁R) in second-order spinal neurons and endothelial cells of postcapillary venules (9, 19). NK₁R signals in endosomes mediate sustained activation of spinal neurons and pain transmission (9, 17). The calcitonin-like receptor (15), protease-activated receptor 2 (11, 16), and δ -opioid receptor (10) also signal from endosomes to control pain, which suggests a general role for endosomal GPCR signaling in pain. The hypothesis that endosomal GPCR signaling controls nociception derives from the use of endocytosis inhibitors, which suppress endosomal signaling

Significance

Sustained GPCR signaling from endosomes mediates pathologies, including pain. Optimal therapy requires development of antagonists that penetrate cells, are retained in endosomes, and disrupt endosomal signaling. To enhance membrane penetration and retention in acidified endosomes, analogs of the neurokinin 1 receptor (NK₁R) antagonist netupitant were synthesized with altered lipophilicity and acidity. Lipophilic and acidic analogs antagonized endosomal NK₁R signaling and provided potent, efficacious, and long-lasting pain relief in mice expressing human NK₁R. Mice expressing a truncated NK₁R, corresponding to a naturally occurring variant with aberrant signaling and trafficking, showed diminished nociceptive responses to substance P. The results identify criteria that will facilitate the design of antagonists of endosomal receptors and provide evidence for the contribution of endosomal signaling to disease.

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¹A.H., C.J.P., R.T., and P.S. contributed equally to this work.

²To whom correspondence may be addressed. Email: nwb2@nyu.edu.

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and nociception (9, 11). Preferential blockade of endosomal GPCRs has been achieved by antagonist conjugation to a transmembrane lipid or encapsulation into nanoparticles, which deliver antagonists to endosomes and provide sustained antinociception (9–11, 15, 17, 20–22). Pharmacokinetic considerations may restrict the usefulness of lipidated and nanoparticle-encapsulated GPCR antagonists. The criteria for rational design of small-molecule GPCR antagonists that penetrate plasma and endosomal membranes, disrupt assembly of multiprotein signalosomes in acidic endosomes, and reverse sustained endosomal signaling and nociception are ill-defined. Whether expression of GPCR variants unable to signal from endosomes confers resistance to pain is unknown.

The current study sought to define the criteria for designing GPCR antagonists that disrupt endosomal signalosomes and provide effective pain relief. The NK₁R antagonists aprepitant (AP) and netupitant (NT), which are both FDA approved for treatment of chemotherapy-induced nausea and vomiting (23, 24), were compared with NT analogs with differing lipophilicity (LogP, LogD) and acidity (pKa). Nociception was studied in mice either expressing the human NK₁R, which avoided interspecies differences in NK₁R antagonist potency (25), or a truncated human NK₁R that corresponds to a natural variant with aberrant G protein and β ARR signaling (26–28). The results show that sustained antagonism of the endosomal NK₁R correlates with long-lasting antinociception, and that domains within the C-terminus of the NK₁R are necessary for the full pronociceptive actions of SP.

Results

SP-Induced Assembly of NK₁R, G α , and β ARR Signalosomes in Early Endosomes. Endosomal signaling of GPCRs is thought to entail the assembly of GPCR, G α , and β ARR signalosomes, where G α subunits transduce signals and β ARRs are scaffolds for GPCRs and signaling enzymes. A bioluminescence resonance energy transfer (BRET) assay was developed to simultaneously study recruitment of the NK₁R and isoforms of G α or β ARR to subcellular compartments of HEK293T cells. Conventional BRET is limited to studying the proximity of only two proteins. To surmount this limitation, NanoLuc Binary Technology BRET (NanoBiT-BRET, nbBRET) was used to simultaneously measure the proximity between the NK₁R, an effector (mini (m) G α or β ARR) plus a localization marker (16, 29) (Fig. 1A). mG α proteins are N-terminally truncated G α proteins that diffuse through the cytosol and bind to active conformations of GPCRs (30, 31); mG α_{sq} and mG α_{si} were developed by mutating mG α_s residues to equivalent G α_q and G α_i residues. NbBRET entails a split luciferase assay in which full-length human NK₁R-407 was C-terminally tagged with the natural peptide fragment of NanoLuc (NP, 13 residue NanoBiT fragment), and a plasma membrane (CAAX) or early endosomal (FYVE) marker was tagged with large NanoBiT fragment (LgBiT). Luminescence occurs from a complex between NK₁R-NP and LgBiT-CAAX or LgBiT-FYVE, which serves as an energy donor for fluorophore-tagged Venus-mG α or β ARR-YFP. SP (100 nM) increased nbBRET between NK₁R, mG α_{sq} , or β ARR2 and LgBiT-CAAX or LgBiT-FYVE, consistent with

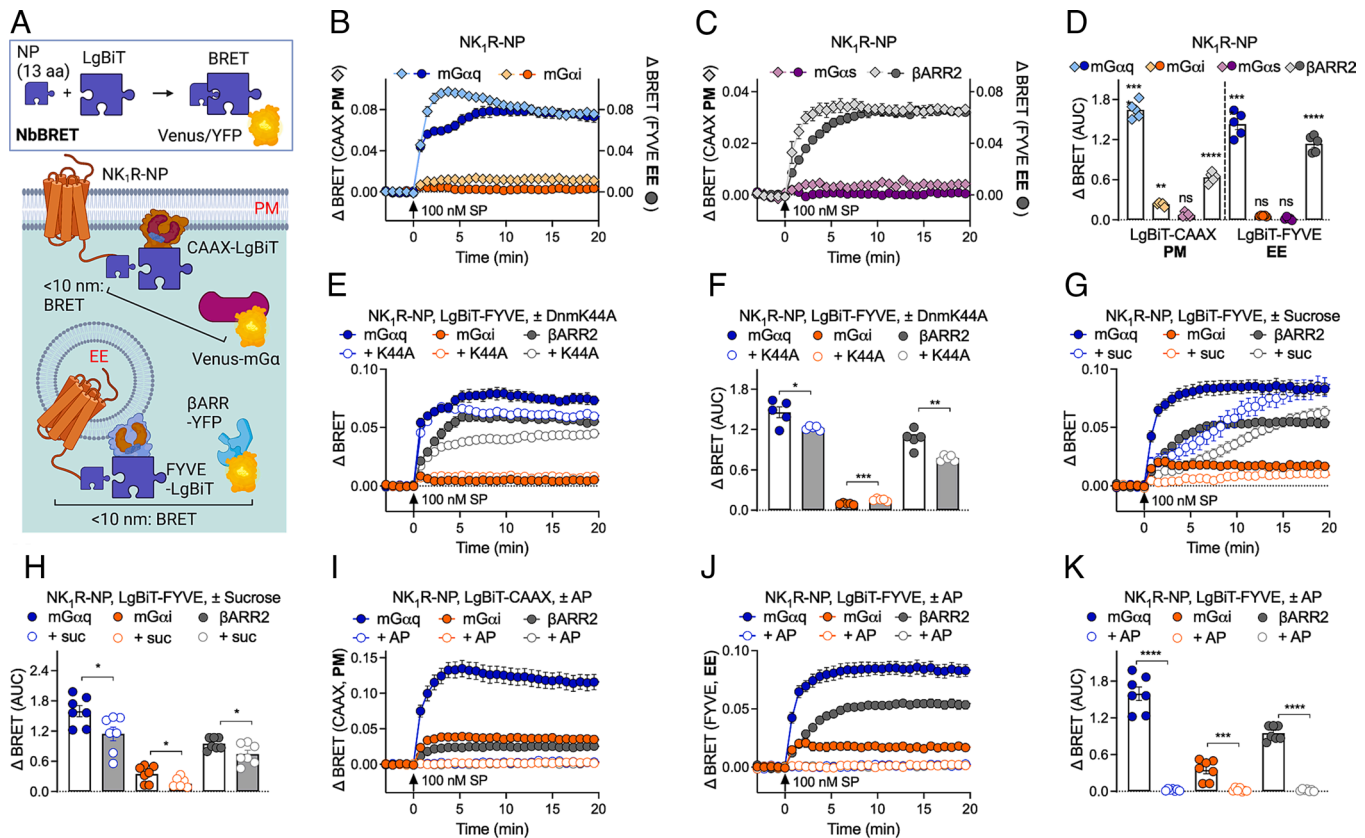


Fig. 1. Assembly of NK₁R signalosomes in HEK293T cells. (A) NanoBiT-BRET (nbBRET) uses a nanoluciferase split into two fragments (natural peptide, NP, and LgBiT) to detect BRET between receptor (NK₁R), mini G α proteins (mG α) or β arrestin (β ARR), and proteins resident to the plasma membrane (PM, CAAX) or early endosome (EE, FYVE). (B–K) Substance P (SP, 100 nM) induced nbBRET between NK₁R-NP, LgBiT-CAAX, or LgBiT-FYVE as well as Venus-mG α_{sq} , Venus-mG α_{si} , Venus-mG α_{sq} , or β ARR2-YFP. (E–H) Effect of dominant-negative dynamin (DnmK44A; E and F) or hypertonic sucrose (0.45 M, 30 min, G and H). (I–K) Effect of aprepitant (AP, 100 nM, 30 min). AUC, area under curve. Mean \pm SEM. N = 5 to 7 independent experiments. **P* < 0.05, ***P* < 0.01, ****P* < 0.001, *****P* < 0.0001. One-way ANOVA with Dunnett's test (D, F, H, and K).

the recruitment of NK₁R, mGα_{sq}, and βARR2 to the plasma membrane and early endosomes (Fig. 1 B–D). SP also stimulated a small increase in nbBRET between NK₁R, mGα_{sq}, and CAAX. Recruitment of mGα_{sq} and βARR2 to early endosomes lagged relative to recruitment to the plasma membrane. Recruitment of NK₁R, mGα_{sq}, or βARR2 to the plasma membrane and endosomes was sustained for at least 20 min. Expression of a dominant-negative mutant of dynamin (DnmK44A), which harbors a mutation in the GTPase domain and selectively inhibits receptor endocytosis without affecting protein exocytosis (32), or pretreatment with hypertonic (0.45 M) sucrose, inhibited the recruitment of NK₁R/mGα_{sq}, NK₁R/mGα_{si}, and NK₁R/βARR2 complexes to early endosomes (Fig. 1 E–H). Preincubation with the NK₁R antagonist AP (100 nM) abolished complex assembly at the plasma membrane and in early endosomes (Fig. 1 I–K).

NK₁R and effector (Gα, βARR) recruitment to intracellular compartments of HEK293T cells was confirmed using conventional BRET assays. To quantify the kinetics of NK₁R trafficking to subcellular compartments, NK₁R-407 tagged with *Renilla* luciferase-8 (Rluc8) was coexpressed with Venus-Kras (plasma membrane marker), Rab5a coupled to tandem *Renilla* green fluorescent protein (tdRGFP, early endosome), Venus-Rab11a (recycling endosome), Venus-Giantin (cis-Golgi network), or YFP-TGN38 (trans-Golgi network) (SI Appendix, Fig. S1A). SP (100 nM) decreased BRET between NK₁R-Rluc8 and Venus-Kras (SI Appendix, Fig. S1 B and C) and increased BRET between NK₁R-Rluc8 and tdRGFP-Rab5a, Venus-Rab11a, and Venus-Giantin, consistent with NK₁R trafficking from the plasma membrane to early and recycling endosomes and the cis-Golgi network (SI Appendix, Fig. S1 D–I). BRET between NK₁R-Rluc8 and YFP-TGN38 transiently decreased and then recovered (SI Appendix, Fig. S1 J and K). DnmK44A inhibited NK₁R-Rluc8 recruitment to early endosomes, recycling endosomes and the cis-Golgi network, implicating clathrin-dependent endocytosis. DnmK44A only partially inhibited SP-induced removal of the NK₁R from the plasma membrane, which may reflect the rapid recycling or mobilization of internalized NK₁R, incomplete disruption of endogenous Dnm, or involvement of additional (non-Dnm) mechanisms of receptor endocytosis. DnmK44A caused the BRET signal between NK₁R-Rluc8 and YFP-TGN38 to further decrease, which may implicate Dnm in maintaining a pool of NK₁R within the trans-Golgi network.

Enhanced bystander BRET (ebBRET), which capitalizes on the affinity of protein pairs tagged with *Renilla* fluorophores to improve sensitivity, was used to confirm the recruitment of mGα and βARR2 to the plasma membrane and early endosomes (SI Appendix, Fig. S2A) (33). EbBRET studies confirmed that SP stimulated recruitment of mGα_{sq}, mGα_{si}, and βARR2 to the plasma membrane and early endosomes (SI Appendix, Fig. S2 B–G). BRET between Gγ-GFP and Rluc2-Gα isoforms was measured to study the effects of SP on proximity of Gγ and Gα. SP decreased BRET between Gγ-GFP and Rluc2-Gα_q or Rluc2-Gα_i, consistent with disassembly of the heterotrimeric Gα, Gβ, and Gγ complex or conformational rearrangement of the activated Gα subunit (SI Appendix, Fig. S2 H and I).

Considered together, these results support the hypothesis that SP induces the recruitment of the NK₁R, mGα_{sq}, and βARR2 to endosomes, where a multiprotein signalosome may transduce intracellular signals.

Design, Synthesis, and Pharmacological Characterization of NT Analogs. Antagonists were designed to optimally inhibit NK₁R signaling in endosomes. The accumulation and retention of GPCR antagonists in endosomes likely depends on their lipophilicity and

charge. Since the lumen of endosomes is acidic (pH 5.5 to 6.0), the degree of ligand protonation may differ between the cytoplasm and endosomal lumen, which could affect endosome membrane penetration and confer enrichment in endosomes and sustained receptor occupancy. Upon receptor binding, the protonation state of the ligand controls association and dissociation and, thus, binding affinity and the kinetic behavior. Furthermore, individual protonation states depending on environment-specific pH values may affect the conformation and, hence, ligand-binding properties of receptors. Ligands were designed with varying lipophilicity (LogP, LogD) and acidity (pKa) to promote accumulation or to take advantage of a potentially altered receptor conformation in endosomes at sufficient concentrations to extend the residence time at the endosomal NK₁R. To determine the criteria for design of antagonists of endosomal GPCRs, a series of structural analogs of the NK₁R antagonist NT (“PS” series) were developed. NT caused a concentration-dependent inhibition of mGα_{sq} recruitment to NK₁R-NP at the plasma membrane and in early endosomes (Fig. 2 A–C). NT had a lower potency in early endosomes (NT, FYVE pIC₅₀ 7.70 ± 0.12, n = 5) compared to the plasma membrane (NT, CAAX pIC₅₀ 8.19 ± 0.17, n = 5; unpaired *t* test, *P* < 0.05).

To better understand NT structure–activity relationships, the weakly basic pyridine nitrogen was replaced with a benzene ring (34) (Fig. 2 D–G). Initial investigations were directed to the synthesis and pharmacological investigation of carba-NT (PS15) with only one basic nitrogen. Further ligand modifications were guided by the crystal structure of the NT/NK₁R complex (PDB ID: 6HLP), which revealed that the N(CH₃) group of the piperazine ring is ideally suited for further modifications (35). Examination of the binding pocket and subsequent docking studies indicated that a replacement of the N(CH₃) group by O or CH₂ is possible, and the introduction of even large N-substituents is well tolerated. Hence, a series of carba-NT derivatives were synthesized to enable introduction of neutral (PS9, PS29), acidic (PS34), weakly basic (PS40), basic (PS21), and strongly basic (PS49) groups (Fig. 2G and Table 1). A maleimide group (PS26) was introduced to enable formation of a covalent bond to the NK₁R. Fluorescent tetramethylrhodamine (TAMRA) derivatives of NT were also prepared (PS67, PS68, PS69). Based on the preparation of NT (36), synthesis started from 2-bromo-4-fluoronitrobenzene to generate a biphenyl system using a Suzuki reaction (SI Appendix). Nucleophilic aromatic substitution allowed introduction of different saturated heterocyclic moieties. Reduction of the nitro group, *N*-methylation, and amide coupling reaction resulted in the introduction of a 1,3-bistrifluoromethylphenyl substituted isobutyric acid moiety and thus, final product formation. To characterize their binding affinity (K_d) at NK₁R, a BRET-based assay was developed using membranes from HEK293T cells expressing human NK₁R tagged at the N terminus with nanoluciferase (Nluc) and TAMRA-NT (SI Appendix, Fig. S3). Binding was measured at normal extracellular pH 7.4 or the acidic pH 6.0 of endosomes. At pH 7.4, TAMRA-NT analogs showed high affinities for Nluc-NK₁R (PS68, K_d 1.9 ± 0.39 nM, n = 12; PS69, K_d 0.81 ± 0.065 nM, n = 21). PS69 was then used as a fluorescent tracer to determine the affinity of SP (K_i 1.4 ± 0.46 μM, n = 6). The orthosteric NK₁R antagonists AP and NT inhibited binding with K_i values of 0.15 nM and 0.91 nM, respectively (Table 1). At pH 7.4, PS analogs retained binding with K_i values ranging from 0.72 nM (PS15) to 110 nM (PS21). Each NT analog also had nanomolar affinity for NK₁R at acidic pH 6.0, thus retaining the ability to antagonize SP/NK₁R binding in acidic endosomes. Most of the compounds showed slightly diminished binding affinities in the acidified environment. An approximately 2-fold and 20-fold increase in affinity was observed for weakly basic

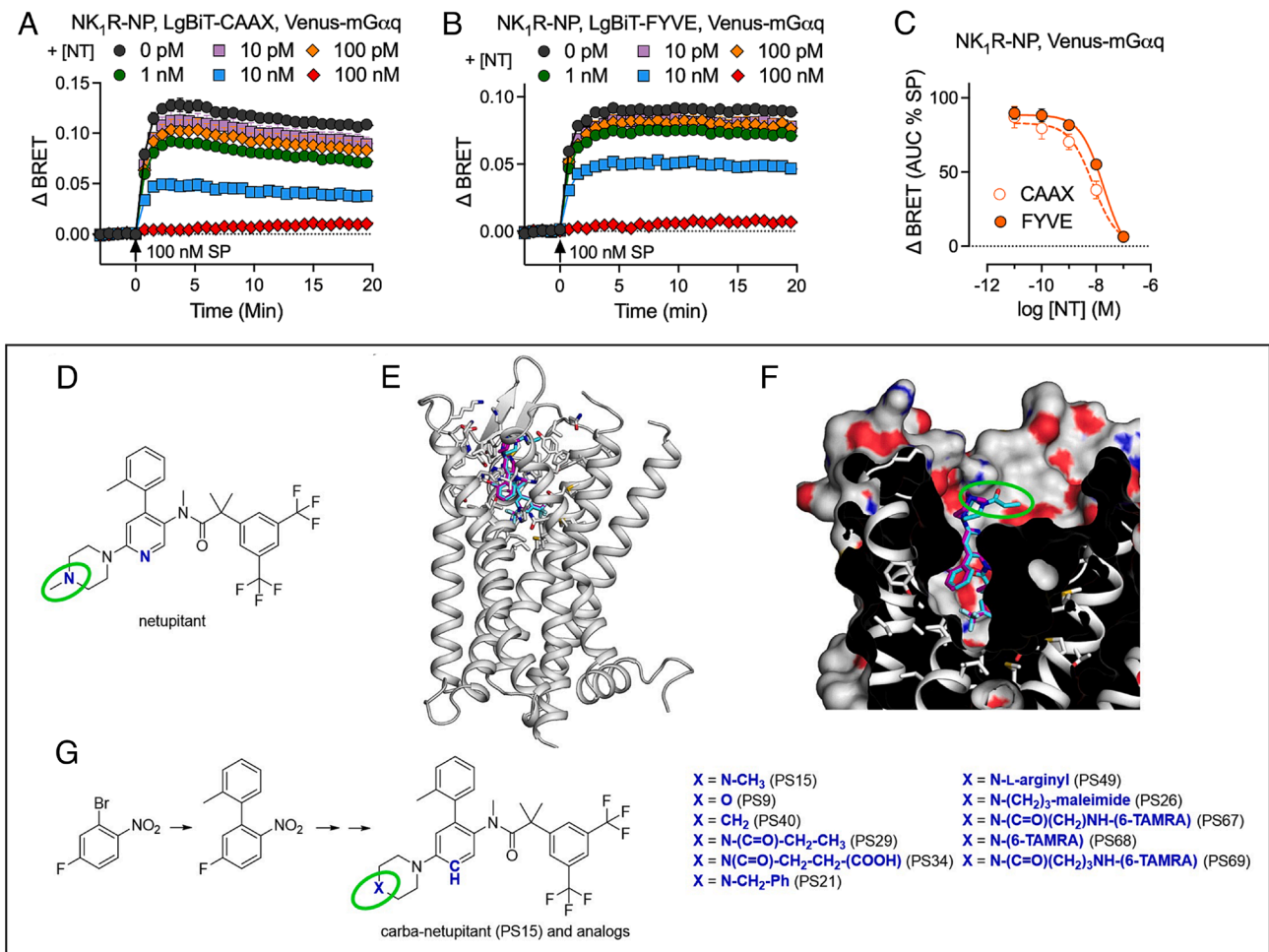


Fig. 2. Development of netupitant analogs. (A–C) Effect of graded concentrations of netupitant (NT, 30 min) on substance P (SP)-induced recruitment of mini $G_{\alpha_{sq}}$ (mG α_q) to the plasma membrane (CAAX, A) or early endosome (FYVE, B) on NanoBIT-BRET (nbBRET) with natural peptide (NP)-tagged NK $_1$ R in HEK293T cells. (D–G) Design and synthesis of NT analogs. (D) Chemical formula of NT. (E) NK $_1$ R crystal structure (PDB ID: 6HLP) in complex with NT (magenta) overlaid with the docked pose of PS29 (blue). Residues are displayed in a range of 8 Å relative to NT and PS29. F. NT (PDB ID: 6HLP) in complex with the NK $_1$ R crystal structure (PDB ID: 6HLP) presented together with the docked pose of PS29. Receptor surface is clipped between the transmembrane helices 4, 5, and 6 to give better insight into the binding pocket and to show the outward reaching piperazine substituent of NT and PS29 (green circle), showing surface atoms and hetero atoms [oxygen (red), nitrogen (blue), sulfur (yellow)]. (G) Synthetic approach toward carba-NT derivatives. Structural changes compared to NT are highlighted in blue. Mean \pm SEM. N = 4 to 9 independent experiments.

arylpiperidine PS40 [K_i (pH 7.4) = 71 nM; K_i (pH 6.0) = 36 nM] and the lipophilic and basic benzylamine PS21 [K_i (pH 7.4) = 110 nM; K_i (pH 6.0) = 4.4 nM], respectively, indicating that protonation affects ligand binding.

Inhibitory Actions of NK $_1$ R Antagonists on SP-Induced Endosomal Signaling. The potency of NT analogs was determined by measuring inhibition of SP-induced inositol 1-phosphate (IP $_1$) accumulation. SP potently stimulated IP $_1$ accumulation (SP, EC $_{50}$ 5.1 \pm 1.1 nM, n = 8). To quantify antagonist potency (IC $_{50}$), HEK293T cells expressing the human NK $_1$ R were preincubated with antagonists (90 min), challenged with SP (10 nM, \sim EC $_{80}$, 90 min), and IP $_1$ accumulation was assessed. AP and NT inhibited SP-stimulated IP $_1$ accumulation with IC $_{50}$ values of 170 nM and 178 nM, respectively (Table 1). PS analogs retained activity with IC $_{50}$ values ranging from 91 nM (PS21) to 407 nM (PS49).

To evaluate whether NK $_1$ R antagonists inhibit endosomal signaling, a Förster resonance energy transfer (FRET) biosensor targeted to the nucleus (Nuc-EKAR) was used to measure nuclear ERK activity, which emanates from NK $_1$ R signaling in endosomes (9). HEK293T cells cotransfected with human NK $_1$ R and NucEKAR were preincubated (30 to 90 min) with graded concentrations of

NK $_1$ R antagonists and challenged with SP (10 nM, \sim EC $_{50}$; *SI Appendix, Fig. S2J*). NT caused a concentration-dependent inhibition of SP-induced nuclear ERK activity (Fig. 3A and Table 1). NT was less potent than AP (Fig. 3B; NT, pIC $_{50}$ 6.59 \pm 0.12; AP, pIC $_{50}$ 6.94 \pm 0.17; n = 4 to 6). PS analogs with different physicochemical properties induced concentration-dependent inhibition of SP-induced activation of nuclear ERK with comparable potency (Fig. 3C and Table 1; pIC $_{50}$ 6.16 to 6.72).

To determine the duration of antagonism (approximating to k_{off}), cells were preincubated with antagonists (30 to 90 min), washed, recovered for 2 or 4 h, and then challenged with SP. Most of the inhibitory effect of AP was lost within 2 h of recovery and is thus transient (Fig. 3D and E and Table 1). In contrast, the inhibitory effects of NT and PS analogs were largely maintained after recovery for 2 h (Fig. 3D and E) and 4 h, respectively (Fig. 3F and *SI Appendix, Fig. S4 A–F*). These results were consistent for two antagonist concentrations (*SI Appendix, Fig. S4 A–F*).

To assess the kinetics with which antagonists reverse endosomal NK $_1$ R signaling (approximating to k_{on}), SP-evoked recruitment of mG α to early endosomes was studied by measuring ebBRET between Rluc8-mG α_{sq} and tdrFP-Rab5a. Cells were incubated

Table 1. Pharmacological properties of NK₁R antagonists

	Compound properties					Binding affinity		Potency of inhibition		Kinetics of inhibition
	Derived [†] , † or calculated ^{‡, §}					BRET (Nluc-NK ₁ R) [¶]		IP ₁ (HTRF)	Nuclear ERK (FRET)	Endosomal miniGα _{sq} (ebbRET)
	pKa [*]	pKa [†]	LogP [§]	LogD _{5.0} [†]	LogD _{7.0} [†]	pK _i at pH 7.4	pK _i at pH 6.0	pIC ₅₀	pIC ₅₀	k (min ⁻¹) for 10 μM
AP (neutral)	n.d.	n.d.	n.d.	n.d.	n.d.	9.82 ± 0.08 (5)	9.27 ± 0.06 (3)	6.77 ± 0.17 (4)	6.94 ± 0.17 (4)	0.15 ± 0.01 (4)
NT (basic)	6.81	6.51	7.22	n.d.	4.97	9.04 ± 0.07 (5)	9.07 ± 0.08 (3)	6.75 ± 0.10 (10)	6.59 ± 0.12 (6)	0.14 ± 0.02 (9)
PS15 (carba-NT)	6.95	7.44	7.54	3.85	5.06	9.14 ± 0.07 (4)	8.59 ± 0.04 (3)	6.50 ± 0.13 (5)	6.38 ± 0.09 (4)	0.23 ± 0.04 (9)
PS9 (neutral)	<3	2.23	7.66	n.d.	3.49	8.84 ± 0.08 (3)	8.45 ± 0.05 (3)	6.87 ± 0.07 (4)	6.29 ± 0.09 (4)	0.11 ± 0.04 (5)
PS40 (weakly basic)	4.31	4.13	8.73	n.d.	n.d.	7.15 ± 0.08 (4)	7.45 ± 0.12 (3)	6.78 ± 0.14 (7)	6.22 ± 0.15 (4)	n.d. – slow kinetics
PS29 (neutral)	<3	1.27	7.37	5.05	4.99	8.73 ± 0.07 (3)	8.31 ± 0.08 (5)	6.84 ± 0.18 (5)	6.46 ± 0.21 (4)	0.14 ± 0.02 (5)
PS21 (basic)	6.17	6.21	9.30	Not trace-able log D >> 5	Not trace-able log D >> 5	6.96 ± 0.02 (3)	8.36 ± 0.07 (3)	7.04 ± 0.11 (4)	6.50 ± 0.25 (4)	n.d. – slow kinetics
PS49 (basic)	>10	11.55	5.45	n.d.	n.d.	8.44 ± 0.08 (4)	7.59 ± 0.08 (6)	6.39 ± 0.09 (5)	6.36 ± 0.16 (5)	0.21 ± 0.01 (7)
PS34 (acidic)	n.d.	6.11	6.41	4.90	3.49	8.74 ± 0.03 (3)	8.69 ± 0.08 (5)	6.75 ± 0.11 (6)	6.16 ± 0.09 (5)	0.11 ± 0.01 (7)
PS26 (covalent)	n.d.	9.16	7.50	n.d.	n.d.	8.64 ± 0.08 (3)	7.91 ± 0.06 (3)	6.68 ± 0.17 (5)	6.72 ± 0.30 (4)	0.11 ± 0.02 (5)

^{*}Free base titrated in 67% MeOH in H₂O (0.01 M HCl).

[†]LogD experiments with distribution between octanol and aqueous buffer solution at pH 5.0 and pH 7.0. n.d. not determined.

[‡]Calculated with Jaguar (Maestro Schroedinger).

[§]Calculated with QikProp (Maestro Schroedinger).

[¶]Determined by competition binding with fluorescent TAMRA-NT (PS69).

with SP (10 nM) for 5 min and then treated with NK₁R antagonists or vehicle (Fig. 3G). SP stimulated recruitment of mGα_{sq} to early endosomes that was maintained for at least 40 min. AP, NT, and PS analogs caused a concentration-dependent inhibition of mGα_{sq} recruitment, reaching equilibrium by 40 min (*SI Appendix, Fig. S5*). At a saturating concentration, PS15 induced a rapid disruption of the endosomal signal within 15 min, with a faster rate than AP, NT, or other analogs (Fig. 3G and H and Table 1).

To evaluate the uptake of NT, HEK293T cells transiently expressing NK₁R-eGFP were incubated with TAMRA-NT (100 nM) and imaged using confocal microscopy. TAMRA-NT was taken up into cells within minutes regardless of NK₁R-eGFP expression (Fig. 3I and *Movie S1*). TAMRA-NT also colocalized with NK₁R-eGFP at the plasma membrane. Preincubation with SP prevented TAMRA-NT binding to NK₁R-eGFP at the plasma membrane (Fig. 3I and *Movie S2*).

Thus, NT analogs penetrate cells and disrupt endosomal NK₁R signaling. Whereas the inhibitory actions of AP are rapidly reversed, NT analogs are long lasting and persist after washing cells to remove ligand.

Generation of Knockin Mice Expressing Full-Length hNK₁R-407 or Truncated hNK₁R-311. Differences in the potency of antagonists between human and mouse NK₁R, including for AP and NT, significantly hamper studies of the NK₁R in preclinical disease models (25). To circumvent this problem, knockin mice were generated in which the mouse NK₁R was replaced by the full-length human NK₁R (407 residues). A naturally occurring variant of the NK₁R has been described in several species, including human, that is truncated at residue 312 and lacks most of the intracellular C-terminal domain (26–28). To examine the function of this variant, knockin mice were generated in which mouse NK₁R was replaced by the truncated human variant (311 residues). This variant binds SP but displays diminished G protein signaling, interaction with βARRs, and internalization (9, 27).

Thus, a comparison of SP-evoked nociception in NK₁R-407 and NK₁R-311 mice could give insight into the contribution of NK₁R endosomal signaling to pain.

Targeting constructs were designed to express tdTomato reporter (to detect NK₁R-expressing cells), a 2A peptide (cleavage would liberate tdTomato), and Flag-NK₁R-407 or Flag-NK₁R-311 (Fig. 4A and *SI Appendix, Fig. S6A*). Constructs were generated by homologous recombination using a 10 kb segment of mouse genomic DNA from a C57 BAC clone. Synthetic cDNAs encoding NK₁R-407 or NK₁R-311 were codon optimized for mouse. tdTomato followed by a 2A peptide was incorporated upstream and a neomycin cassette flanked by FRT sites was incorporated downstream from the NK₁R cDNAs. The cassette was inserted at the ATG site of mouse NK₁R and the first exon was deleted. Constructs were linearized and electroporated in Bruce4 (C57BL/6) ES cells. Screening of 200 clones revealed 8 to 14% ratio of recombination, which was confirmed by southern blotting (Fig. 4B and *SI Appendix, Fig. S6B*). Clones were analyzed using an internal probe annealing on the NEO cassette (Fig. 4C and *SI Appendix, Fig. S6C*). Sequencing of five NK₁R-407 and five NK₁R-311 clones confirmed correct insertion of tdTomato and integrity of FRT sites. Three clones were injected into mouse blastocysts, and male chimeras (black and white) were bred with wild-type C57BL/6 female mice to generate black F1 progeny; two clones gave germline transmission. Expression of NK₁R-407 and NK₁R-311 was confirmed by southern blotting of the heterozygotic F1 generation (Fig. 4D and *SI Appendix, Fig. S4D*) and RT-PCR of knockin mice. F1 mice were backcrossed for >10 generations to generate homozygotic NK₁R-407 and NK₁R-311 mice.

Characterization of hNK₁R-407 and hNK₁R-311 Mice. The expression of the NK₁R was verified in hNK₁R-407, hNK₁R-311, and wild-type mice using PCR and qRT-PCR with primers for mouse (m) or human (h) receptors, including primers specific to hNK₁R-407 but not hNK₁R-311 and primers that would

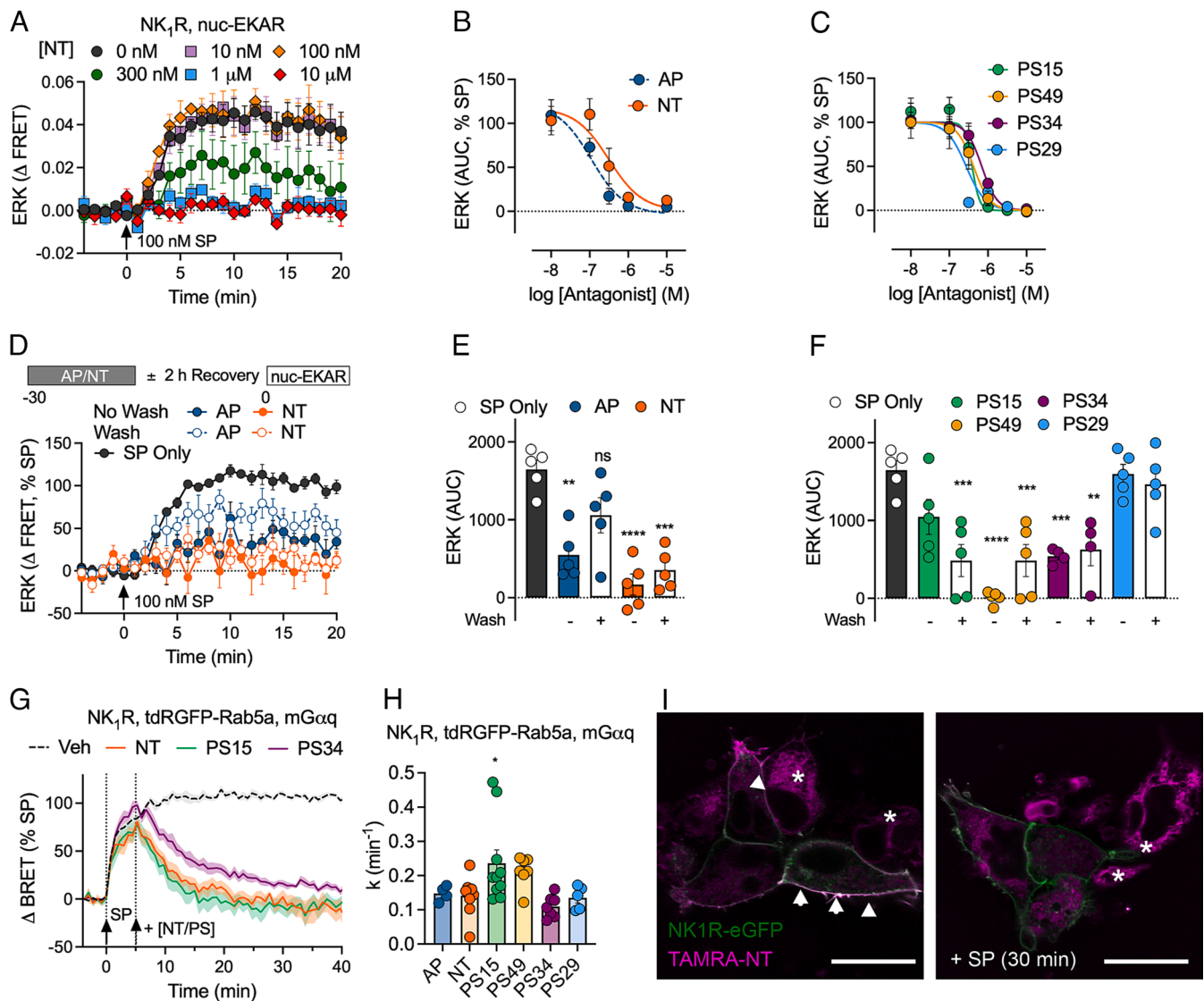


Fig. 3. Antagonism of NK₁R endosomal signaling in HEK293T cells. (A–F) FRET assays of substance P (SP)-induced activation of nuclear ERK activity measured using a nuclear EKAR biosensor. (A–C) Cells were preincubated (30 or 90 min) with graded concentrations of aprepitant (AP), netupitant (NT), PS15, PS29, PS34, or PS49 prior to SP (100 nM) challenge. (D–F) Duration of inhibition following antagonist removal and recovery. Cells were preincubated with antagonist for 30 min (AP, 100 nM; NT, 300 nM) or 90 min (NT analogs, 1 μM), washed, and recovered in agonist-free medium for 2 h (AP, NT) or 4 h (NT analogs; *SI Appendix, Fig. S3*), prior to stimulation with SP (100 nM). (G and H) Enhanced bystander BRET (ebBRET) measuring proximity between mini Gα_{sq} (Venus-mGα_{sq}) and a *Renilla* luciferase-tagged endosomal marker (tdRGFP-Rab5a). Cells were stimulated with SP (100 nM, 5 min), then challenged with vehicle or saturating concentrations of NT, PS15, or PS34 (10 μM). EbBRET was measured for 40 min. Kinetics were fitted as a nonlinear exponential decay from antagonist addition to quantify the rate of antagonism. Mean ± SEM. N = 4 to 9 independent experiments. 1-way ANOVA with Holm-Sidak's (E), Dunnett's (F) test against SP, or Dunnett's against NT (H). (I) Localization of fluorescent TAMRA-NT (PS68, 100 nM, 10 min, magenta) in HEK293T cells expressing NK₁R-eGFP (green), with or without preincubation with unlabeled SP (100 nM, 30 min). Arrow heads denote TAMRA-NT and NK₁R-eGFP colocalization at the plasma membrane. Asterisks denote uptake of TAMRA-NT into cells lacking NK₁R-eGFP. (Scale bar, 20 μm). Representative images, N = 3 experiments.

detect both hNK₁R-407 and hNK₁R-311. Primers specific to hNK₁R-407 detected the receptor in extracts of the spinal cord and colon from hNK₁R-407 mice, but not hNK₁R-311 or wild-type mice (Fig. 5 A–C). Primers common to hNK₁R-407 and hNK₁R-311 detected both receptors at similar levels in the spinal cord and colon of hNK₁R-407 and hNK₁R-311 mice but not in wild-type mice.

RNAScope[®] in situ hybridization was used to localize human and mouse NK₁R mRNAs in the spinal cord of mice. Probes were designed to specifically target both human NK₁R-407 and NK₁R-311 or mouse NK₁R. Neurons were identified by Nissl staining. Corroborating with the qRT-PCR results, a probe common to human NK₁R-407 and NK₁R-311 hybridized to spinal neurons of hNK₁R-407 and hNK₁R-311 but not in wild-type

mice (Fig. 5D). Mouse NK₁R mRNA was detected in spinal neurons of wild-type mice (Fig. 5D).

SP-Evoked Nociception in hNK₁R-407 or hNK₁R-311 Mice. Since NK₁R-311 displays aberrant signaling by G proteins and βARRs and is resistant to SP-induced endocytosis (9, 27, 37), a comparison of the pronociceptive actions of SP between mice expressing full-length or truncated human NK₁R could provide insights into the contribution of NK₁R signaling and endocytosis to nociception. SP (0.1, 1, 3 μg/5 μL) was injected intrathecally to activate spinal neurons expressing the NK₁R in hNK₁R-407, hNK₁R-311, and wild-type mice. Paw withdrawal responses to stimulation with von Frey filaments were assessed over 1 h to evaluate mechanical allodynia. SP dose-dependently reduced withdrawal threshold

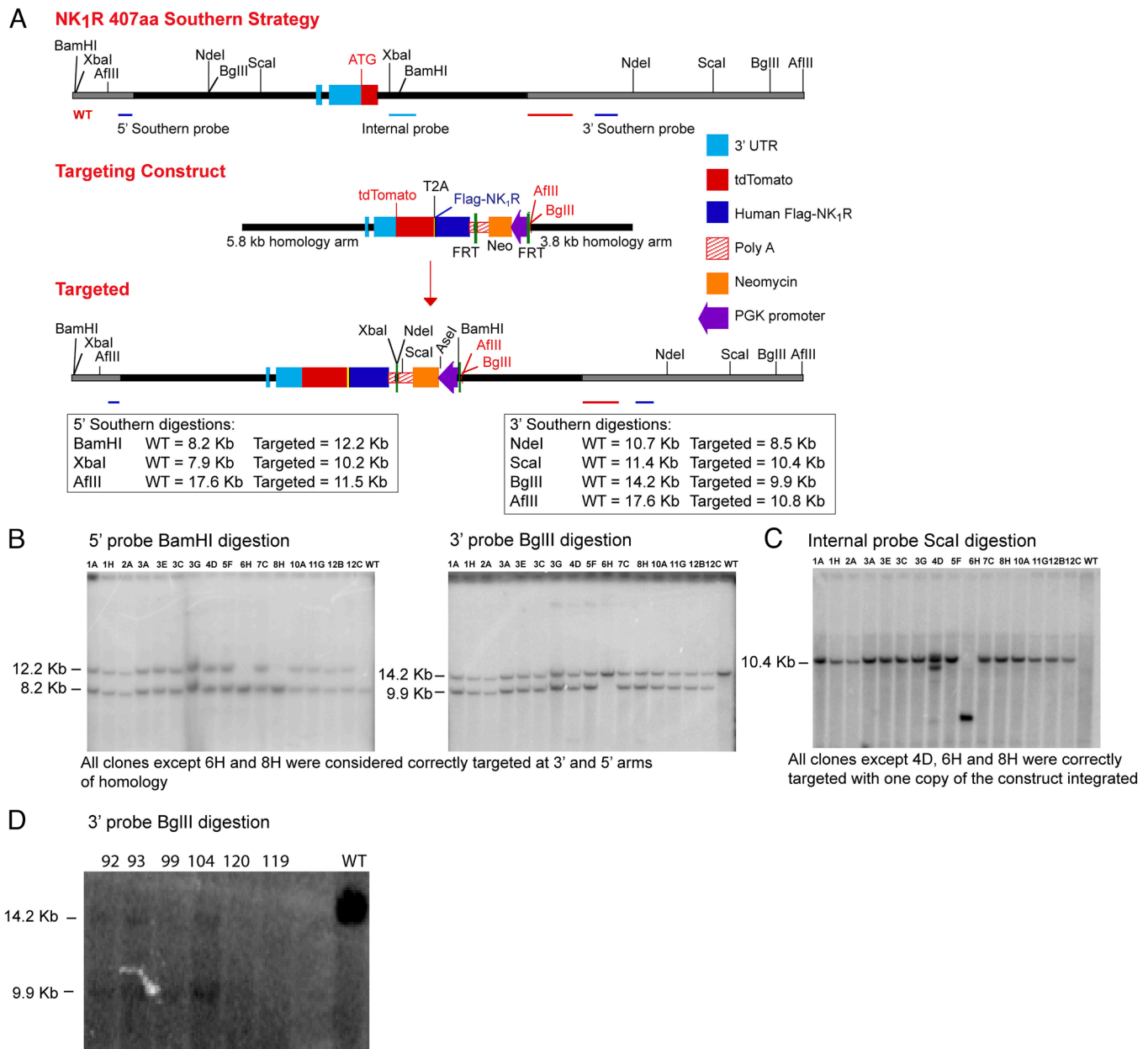


Fig. 4. Generation of knockin mice expressing hNK₁R-407. (A) NK₁R-407 Southern strategy and targeting construct comprising tdTomato, a T2A peptide cleavage site, Flag-hNK₁R-407, a downstream phosphoglycerine kinase neomycin cassette flanked by FRT sites, and downstream AflIII and BglIII sites. (B and C) Southern blots of selected clones with 5', 3', and internal probes confirming correct targeting. (D) Southern blot of F1 mice confirming NK₁R-407 expression. WT, wild-type.

to von Frey filaments in hNK₁R-407, hNK₁R-311, and wild-type mice, consistent with mechanical allodynia (Fig. 6 A–C). All doses of SP evoked mechanical allodynia, which was detected after 15 min, was maximal at 30 min, and returned to baseline at 60 min after injection. A comparison of mechanical allodynia between strains to the same dose of SP indicated identical mechanical allodynia in hNK₁R-407 and wild-type mice to all doses of SP (Fig. 6 D–I). Mechanical allodynia in hNK₁R-311 mice was less than that in NK₁R-407 and wild-type mice at all doses of SP, and this difference was significant at 1 μg SP (Fig. 6 E and H).

Although SP is the primary ligand for the NK₁R, SP can also activate the NK₂R and NK₃R, albeit with low affinity (25). To confirm that the SP nociceptive response was dependent on the NK₁R, AP (100 nM/5 μL) was coadministered with SP (1 μg/5 μL) by intrathecal injection. AP abolished SP-evoked mechanical

allodynia in hNK₁R-407, hNK₁R-311, and wild-type mice (Fig. 6 J and K).

SP activates the NK₁R on spinal neurons to evoke receptor endocytosis and persistent neuronal excitation. Inhibitors of clathrin-mediated endocytosis, and lipidated or nanoparticle-encapsulated NK₁R antagonists that target endosomes, prevent sustained neuronal excitation, which thus requires NK₁R signaling in endosomes (9, 17). To obtain further evidence for the contribution of NK₁R endocytosis to nociception, spinal cord slices from NK₁R-407 and NK₁R-311 mice were superfused with SP (10 nM, 2 min), and action potential firing was assessed by cell-attached patch-clamp recording. SP induced a rapid-onset burst of action potential discharge that was similar in spinal neurons of NK₁R-407 and NK₁R-311 (Fig. 6 L–N). In NK₁R-407 mice, neuronal excitation persisted for 12 min after SP washout. In contrast, in NK₁R-311 mice, neuronal excitation returned to baseline by 4 min after SP

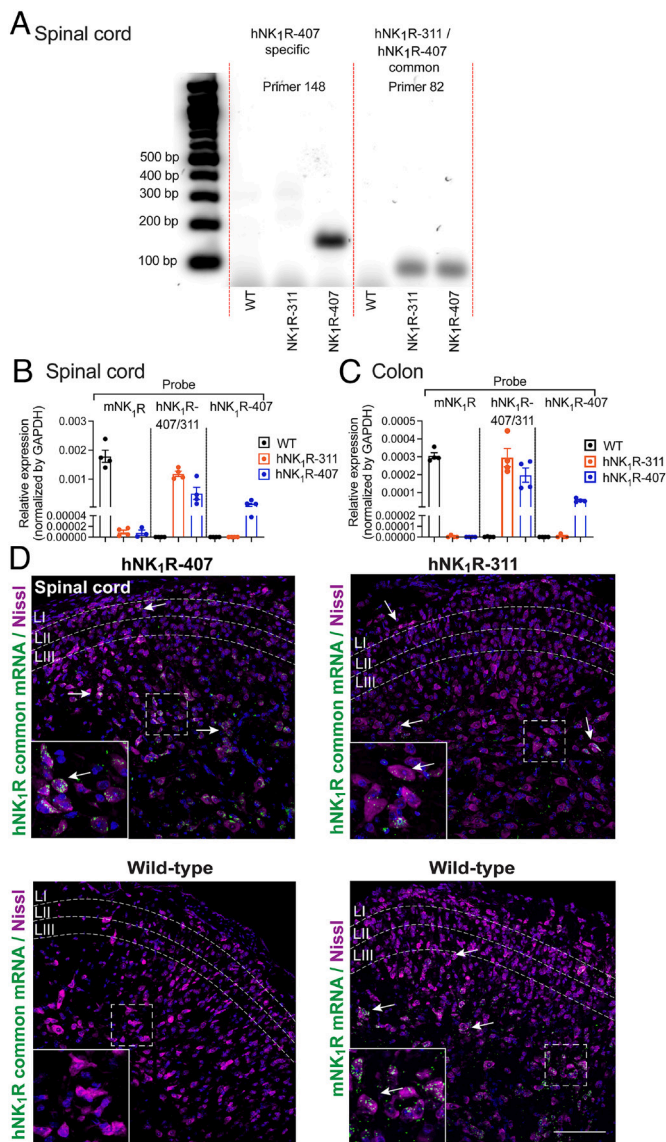


Fig. 5. Characterization of mice expressing hNK₁R-407 or hNK₁R-311. (A) PCR gels of the spinal cord from NK₁R-407, NK₁R-311, and WT mice probed with a primer (#148) specific for human NK₁R-407 or a primer (#82) that would detect both human NK₁R-407 and NK₁R-311. (B and C) qRT-PCR analysis of extracts of the spinal cord (B) and colon (C) probed with primers to mouse NK₁R, both human NK₁R-407 and NK₁R-311, or hNK₁R-407 alone. N = 4 mice. (D) RNAScope® in situ hybridization of sections of the spinal cord from NK₁R-407, NK₁R-311, or WT mice using probes to both human NK₁R-407 and NK₁R-311 or to mouse NK₁R, highlighting neurons with a Nissl stain (magenta). Representative images from n = 3 mice. Scale bar, 50 μ m, and in detail box, 20 μ m. L, lamina.

washout. AP prevented SP-induced excitation of spinal neurons in NK₁R-407 mice.

Attenuated SP-induced activation of spinal neurons and diminished nociception observed in mice expressing truncated NK₁R support the hypothesis that efficient coupling to G proteins, β ARRs, and the endocytic machinery are necessary for SP-evoked pain.

Antinociceptive Effects of NK₁R Antagonists in hNK₁R-407 Mice.

To assess whether effective antagonism of NK₁R signaling in endosomes correlates with antinociceptive activity, the effect of NK₁R antagonists was evaluated on capsaicin-induced mechanical allodynia in hNK₁R-407 mice. AP, NT, NT analogs (PS15, PS29, PS34, PS49) (10, 30, 100 or 300 nM, 5 μ L) or vehicle (control) was injected intrathecally immediately before intraplantar injection

of capsaicin. Withdrawal responses of the capsaicin-injected (right, ipsilateral) hindpaw to stimulation with von Frey filaments were assessed to evaluate mechanical allodynia. In vehicle-treated mice, capsaicin caused mechanical allodynia for 8 h (Fig. 7 A and B). Intrathecal AP (100 nM) inhibited capsaicin-evoked mechanical allodynia for 2 h. NT (100 nM) inhibited mechanical allodynia for 4 h. PS15, PS29, PS34, and PS49 (100 nM) had more efficacious and sustained antinociceptive effects. PS29 and PS34 completely prevented capsaicin-evoked mechanical allodynia for 4 h, with antinociceptive effects for at least 8 h. Higher doses of AP or NT (300 nM) did not further increase the magnitude or duration of the antinociception (Fig. 7 C, D, and F). In contrast, lower doses of PS34 (30 nM) still induced antinociception for 6 h. Thus, NT analogs that effectively inhibit endosomal signaling of the NK₁R also caused efficacious and persistent antinociception.

Discussion

The results support the hypothesis that SP stimulates recruitment of the NK₁R, mG α_{sq} , mG α_{si} , and β ARR2 to early endosomes, where an NK₁R, mG α , and β ARR signalosome may assemble and generate signals in subcellular compartments, including ERK activation in the nucleus. Although AP, NT, and NT analogs can disrupt complex formation and inhibit resultant signals, the actions of AP are rapidly reversed. NT analogs, in particular PS29 and PS34, demonstrate sustained antagonism of endosomal signaling and potent, efficacious, and long-lasting inhibition of nociception in mice expressing the human NK₁R. In mice expressing a truncated NK₁R variant with aberrant signaling and endosomal trafficking, SP is unable to cause sustained activation of spinal neurons and persistent nociception. The results identify criteria for the design of GPCR antagonists that are capable of penetrating endosomes and disrupting signalosomes and provide evidence for the contribution of endosomal NK₁R signaling in nociception.

NK₁R Signalosomes in Endosomes. BRET assays of NK₁R trafficking revealed that SP induces depletion of NK₁R from the plasma membrane and recruitment to early endosomes, recycling endosomes, and the Golgi apparatus. NK₁R trafficking was Dnm dependent because dominant-negative Dnm inhibited NK₁R endocytosis and accumulation in endosomes and the cis-Golgi apparatus. SP also stimulated the recruitment of mG α_{sq} , mG α_{si} , and β ARR2, but not mG α , to the plasma membrane and early endosomes. A nbBRET assay allowed simultaneous evaluation of the proximity of the NK₁R, an effector (mG α , β ARR2), and a plasma membrane or endosomal marker. These studies suggest that SP evokes assembly of NK₁R signalosomes with mG α_{sq} , mG α_{si} , or β ARR2 in early endosomes. Although BRET assays can be used to infer signalosome assembly, the approach evaluates protein proximity rather than interaction. However, structural studies support the concept that GPCRs could physically associate with G α and β ARRs in endosomes, forming “megaplexes” that continue to signal (12, 13). GPCR signalosomes in endosomes generate second messengers and activate enzymes in specific intracellular compartments, including nuclear ERK and cytosolic protein kinase C and cAMP in the case of the NK₁R (9). Compartmentalized signals allow GPCRs, which often couple to a common set of effectors, to regulate specific cellular functions, including activity of spinal neurons that is required for the central transmission of pain (9). Further studies are required to evaluate the physiological relevance of recruitment of the NK₁R and isoforms of G α and β ARRs to endosomes, including stimulation of cells that endogenously express these proteins with biologically relevant concentrations of SP.

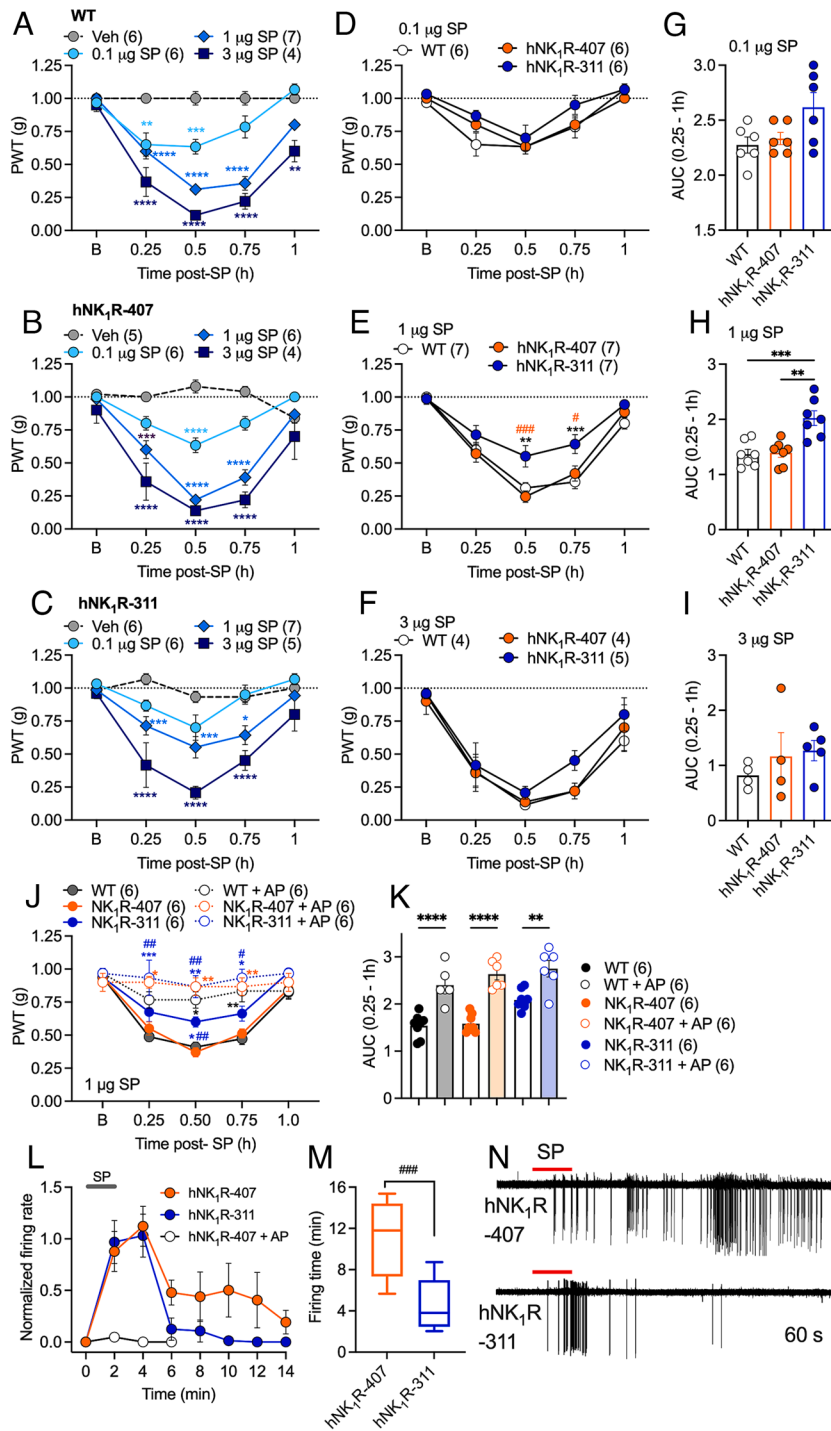


Fig. 6. SP-evoked nociception in hNK₁R-407, hNK₁R-311, and wild-type mice. (A–C) Substance P (SP; 0.1, 1, 3 µg/5 µL, intrathecal)-evoked mechanical allodynia in WT (A), NK₁R-407 (B), and NK₁R-311 (C) mice. Mechanical allodynia was measured 0.25 to 1 h after SP injection. (D–F) Mechanical allodynia to the same dose of SP in the three strains of mice. Paw withdrawal threshold (PWT) to stimulation with calibrated von Frey filaments was measured to assess mechanical allodynia. (G–I) AUC in the three strains of mice. (J and K) Effects of intrathecal injection of aprepitant (AP, 100 nM/5 µL) or vehicle on SP (1 µg/5 µL)-evoked mechanical allodynia in NK₁R-407, NK₁R-311, and WT mice. (L–N) Activation of spinal neurons in slice preparations of spinal cord from NK₁R-407 and NK₁R-311 mice after transient stimulation with SP (10 nM, 2 min). (L) Normalized firing rate. (M) Firing time. (N) Representative traces. Mean ± SEM; (N) denotes number of mice studied. **P* < 0.05, ***P* < 0.01, ****P* < 0.001, *****P* < 0.0001 vs. Veh, WT; #*P* < 0.05, ##*P* < 0.01, ###*P* < 0.001 vs. NK₁R-407. Two-way ANOVA, Sidák's multiple comparison test (A–F and J), one-way ANOVA, Tukey's multiple comparison test (G–I and K), or parametric unpaired two-tailed *t* test (M).

Antagonism of NK₁R Signaling in Endosomes. Whereas GPCR signaling at the plasma membrane is tightly regulated by GRK-mediated receptor phosphorylation and interaction with βARRs, GPCR signaling in endosomes is often sustained. The mechanisms that terminate endosomal signaling are not fully understood. However, degradation of SP in acidic endosomes by the transmembrane peptidase endothelin-converting enzyme 1

disrupts signalosomes and terminates endosomal signaling of the NK₁R (38, 39). The observations that inhibitors of endocytosis block compartmentalized signaling and nociception provide a link between endosomal signaling and pain and suggest that GPCRs in endosomes are an optimal target for the treatment of pain (9, 11, 16). GPCR antagonists conjugated to the transmembrane lipid cholesterol or encapsulated into nanoparticles accumulate in early

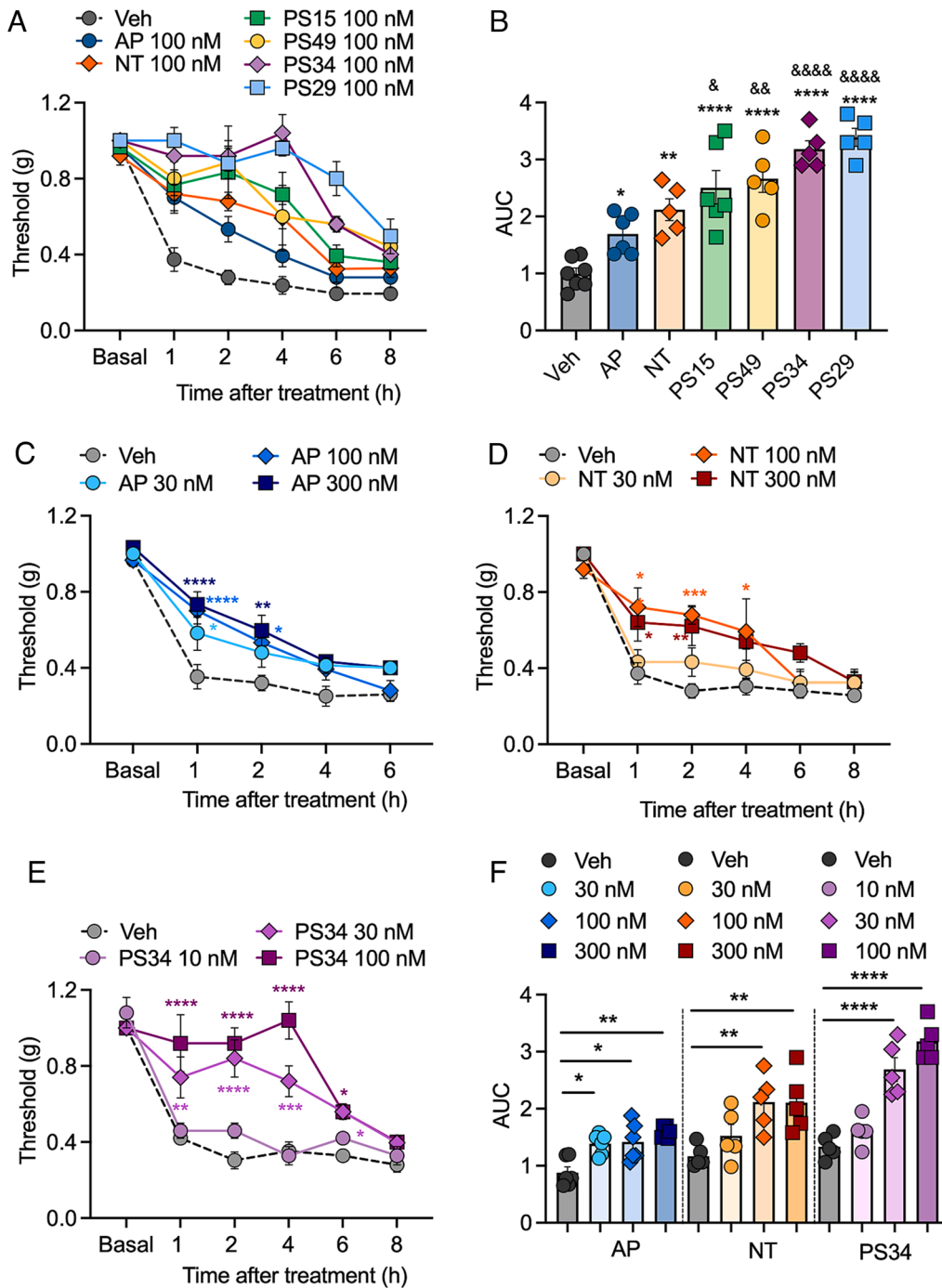


Fig. 7. Antagonism of capsaicin-evoked nociception in hNK₁R-407 mice. (A) Effects of intrathecal (i.t.) injection of aprepitant (AP), netupitant (NT), or NT analogs (PS15, PS49, PS34, PS29; 100 nM/5 μ L) on capsaicin-induced mechanical allodynia in hNK₁R-407 mice. (B) AUC in the same groups of mice. (C–E) Dose-response effects of AP (30, 100, 300 nM/5 μ L, i.t.), NT (30, 100, 300 nM/5 μ L, i.t.), or PS34 (10, 30, 100 nM/5 μ L, i.t.) on capsaicin-induced mechanical allodynia. (F) AUC in the same groups of mice. (N) denotes number of mice studied. * P < 0.05, ** P < 0.01, *** P < 0.001, **** P < 0.0001 vs. Veh; & P < 0.05, && P < 0.01, &&& P < 0.0001 vs. AP, one-way ANOVA, Tukey's multiple comparison test (B and F) or two-way ANOVA, Sidák's multiple comparison test (C, D, and E).

endosomes, block endosomal signaling, and provide more efficacious and long-lasting relief from pain than conventional antagonists (9, 11, 15, 17, 22, 40). However, lipidated and nanoparticle-encapsulated drugs are not suited to systemic administration, and pharmacokinetic considerations may limit their usefulness. Thus, the current investigation sought to identify criteria for the rational design of small-molecule antagonists of the NK₁R. The structure of the NK₁R complexed with NT was used to design derivatives of NT (35). NT analogs of variable lipophilicity (LogP, LogD) and acidity (pKa) were synthesized based on the presumption that antagonists with altered physicochemical behavior would penetrate plasma and endosomal membranes and compounds with ideal properties are retained in proximity to NK₁R signalosomes. A similar approach has been used to design fluorinated analogs of fentanyl with a low pKa. These efficiently activate μ -opioid receptors in the acidified

extracellular environment of damaged or diseased tissue, and thereby prevent pain without the detrimental on-target side effects usually evoked by engaging μ -opioid receptors in healthy tissues with normal extracellular pH (21, 41). A long residence time of antagonists in endosomes containing the NK₁R might be expected to abrogate endosomal signaling and prevent pain. NT analogs retained the ability to bind to the NK₁R in membrane preparations at extracellular (pH 7.4) and endosomal (pH 6.0) pH, and all potently inhibited SP-induced IP₁ formation. Lipophilic receptor antagonists may better diffuse into endosomes. Moreover, NK₁R conformation may change at a lower pH, conferring a strong and persistent receptor binding of suitable NT analogs. We suggest that PS29 and PS34 show favorable binding properties because such ligands are not charged at the lower pH in endosomes, leading to a more sustained inhibition of endosomal signaling. NT analogs

also potently inhibited SP-induced activation of nuclear ERK, which requires endosomal NK₁R signaling (9). However, whereas AP caused only a transient inhibition of SP-induced activation of nuclear ERK that was rapidly reversed after washout, the inhibition by NT and all NT analogs was maintained after washing cells to remove extracellular drug. Moreover, NT and NT analogs rapidly reversed SP-induced recruitment of mG α_{sq} to endosomes. Notably, the NT analogs PS34 and PS29 caused a considerably larger and more sustained inhibition of capsaicin-evoked allodynia than that of AP and NT, which had smaller and more transient antinociceptive effects. The LogD values of PS29 (LogD_{5,0} = 5.05, LogD_{7,0} = 4.99) and PS34 indicate high lipophilicity. Despite its acidic functional group, the LogD value of PS34 at pH 5 (LogD_{5,0} = 4.90, compared to LogD_{7,0} = 3.49) indicates that deprotonation of PS34 in endosomes is not crucial. Hence, PS34 remains neutral and lipophilic in endosomes. The administration of antagonists by injection into the intrathecal space where they would readily access dorsal horn spinal neurons expressing the NK₁R likely minimizes the possibility that pharmacokinetic differences account for the large and prolonged analgesic actions of PS34 and PS29. Future pharmacokinetic studies are required to fully exclude this possibility. Pharmacokinetic and pharmacodynamic studies are also necessary to determine whether parenterally administered antagonists such as PS34 and PS29 access the NK₁R in endosomes of spinal neurons and effectively alleviate inflammatory and neuropathic pain.

Kinetics and Physicochemical Properties of NK₁R Antagonists.

The most basic NT analogs, PS15 and PS49, show the fastest kinetics for the inhibition of endosomal signaling at high concentration. A rapid exchange between the protonated and nonprotonated forms under physiological conditions and pH-dependent protonation states, which are reflected by the experimental LogD values of PS15 (LogD_{7,0} = 5.06, LogD_{5,0} = 3.85), may promote their traffic to endosomes. Additionally, the exchange of the pyridine-core of NT to a more lipophilic benzene-ring (PS analogs) could accelerate the penetration of membranes.

SP-Induced Nociception in Mice Expressing Truncated NK₁R.

mRNA encoding NK₁R-311, which lacks most of the intracellular C-terminus, is highly expressed in the human brain (26) and the immune system (42). Lacking critical Ser and Thr residues that are potential sites of GRK phosphorylation and interaction with β ARRs, NK₁R-311 is unable to recruit β ARRs and does not undergo SP-induced endocytosis (9). The C-terminal domain is also necessary for efficient G α coupling, since NK₁R-311 responds only to high SP concentrations with reduced and delayed ERK signaling (27, 28). In knockin mice expressing human NK₁R-311, intrathecal injection of SP caused mechanical allodynia, although responses were smaller than those observed in mice expressing human NK₁R-407 or in wild-type mice, which were identical. Although SP caused an initial activation of spinal neurons from NK₁R-311 mice, the sustained response observed in neurons from NK₁R-407 mice was largely absent. Given that the sustained activation of spinal neurons is dependent on NK₁R endocytosis and consequent activation of ERK, the diminished response in neurons from NK₁R-311 mice may be attributable to diminished endocytosis and delayed ERK signaling. However, there are other possible explanations for the diminished nociception in NK₁R-311 mice. The inability of NK₁R-311 to respond to low concentrations of SP and efficiently couple to G proteins may also contribute to diminished nociception. Although altered levels of expression could explain abnormal nociception, the mRNA levels and distribution of the NK₁R were similar in NK₁R-311, NK₁R-407, and wild-type mice. It is also possible that SP may activate receptors other

than the NK₁R. SP at high concentrations can activate NK₂R and NK₃R (25), but AP, which is selective for the NK₁R over NK₂R and NK₃R, abolished SP-evoked nociception in NK₁R-311, NK₁R-407, and wild-type mice. It is not possible to exclude the possibility that SP activates other types of GPCRs, such as the mouse Mas-related GPCR MrgprB2, which responds to high concentrations of SP and is antagonized by high concentrations of AP (43). Although MrgprB2 appears to mediate the effects of SP on mouse mast cells, further work is required to determine whether MrgprB2 could be involved in the central actions of SP on nociception in mice.

In summary, lipophilic and acidic NK₁R antagonists cause sustained disruption of endosomal signaling and long-lasting antinociception. The results provide insights into strategies for antagonizing GPCRs in intracellular locations, with implications for designing improved treatments for disease.

Materials and Methods

SI Appendix includes detailed methods.

Design, Synthesis, and Analysis of NT Analogs. The NT-bound NK₁R structure (PDB: 6HLP) (35) was used for molecular docking (44). NT analogs were synthesized by the Suzuki reaction and analyzed by chromatography and mass spectrometry. pK_a was determined as described (45). TAMRA-NT binding to membranes from HEK293T cells expressing Nluc-NK₁R was measured using BRET (46). G-protein signaling was assessed by IP₁ assays (47, 48). TAMRA-NT uptake in cells expressing NK₁R-eGFP was studied by confocal microscopy.

BRET. NK₁R trafficking, mG α and β ARR2 recruitment, and G α /G $\beta\gamma$ dissociation were determined by BRET, ebBRET, and nbBRET (16).

FRET. Nuclear ERK activation was measured using nuc-EKAR FRET biosensor (9).

Generation and Characterization of Knockin Mice. Knockin mice were generated expressing full-length (407 aa) or truncated (311 aa) human NK₁R. PCR and RNAScope[®] probes to human NK₁R-407 and NK₁R-311, specific for human NK₁R-407, or specific for mouse NK₁R, were used to detect NK₁R mRNA. Nociception was studied after intrathecal injection of SP or NK₁R antagonists or intraplantar injection of capsaicin. Mechanical allodynia was assessed by measuring paw withdrawal response to von Frey filaments (9, 17, 49). Cell-attached patch-clamp recordings were made from spinal cord slices (9, 17, 49).

Statistics. Differences were assessed using Student's *t* test for two comparisons and 1- or 2-way ANOVA and Tukey's, Dunnett's, or Šidák's post-hoc test for multiple comparisons. *P* < 0.05 was considered significant at the 95% confidence level.

Data, Materials, and Software Availability. All study data are included in the article and/or *SI Appendix*.

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Author affiliations: ^aDepartment of Molecular Pathobiology, College of Dentistry, New York University, New York, NY 10010; ^bDepartment of Neuroscience and Physiology, Neuroscience Institute, New York University, New York, NY 10010; ^cPain Research Center, College of Dentistry, New York University, New York, NY 10010; ^dDepartment of Chemistry and Pharmacy, Friedrich-Alexander Universität Erlangen-Nürnberg, 91058, Erlangen, Germany; ^eGene Modification Platform, Monash University, Clayton, VIC 3168, Australia; ^fDrug Discovery Biology, Monash Institute of Pharmaceutical Sciences, Monash University, Parkville, VIC 3052, Australia; ^gNYU Dentistry Translational Research Center, College of Dentistry, New York University, New York, NY 10010; and ^hDepartment of Physiology and Monash Biomedicine Discovery Institute, Monash University, VIC 3800, Australia

Author contributions: A.H., C.J.P., R.T., P.S., S.T., R.L., H.H., D.W., J.R., N.A.V., D.P.P., D.D.J., B.L.S., W.L.I., P.G., and N.W.B. designed research; A.H., C.J.P., R.T., P.S., S.T., R.L., H.H., D.W., J.R., N.A.V., D.P.P., D.D.J., W.L.I., and P.G. performed research; A.R.B.T. contributed new reagents/analytic tools; A.H., C.J.P., R.T., P.S., S.T., R.L., H.H., D.W., J.R., N.A.V., D.P.P., D.D.J., W.L.I., P.G., and N.W.B. analyzed data; and A.H., C.J.P., R.T., P.S., B.L.S., W.L.I., P.G., and N.W.B. wrote the paper.

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