Title: SARS-CoV-2 Spike protein hijacks VEGF-A/Neuropilin-1 receptor signaling to induce analgesia

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Abstract: Global spread of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) continues unabated. Binding of SARS-CoV-2's Spike protein to host angiotensin converting enzyme 2 triggers viral entry, but other proteins may participate, including neuropilin-1 receptor (NRP-1). As both Spike protein and vascular endothelial growth factor-A (VEGF-A) – a pro-nociceptive and angiogenic factor, bind NRP-1, we tested if Spike could block VEGF-A/NRP-1 signaling. VEGF-A–triggered sensory neuronal firing was blocked by Spike protein and NRP-1 inhibitor EG00229. Pro-nociceptive behaviors of VEGF-A were similarly blocked via suppression of spontaneous spinal synaptic activity and reduction of electrogenic currents in sensory neurons. Remarkably, preventing VEGF-A/NRP-1 signaling was antiallodynic in a neuropathic pain model. A 'silencing' of pain via subversion of VEGF-A/NRP-1 signaling may underlie increased disease transmission in asymptomatic individuals.

One Sentence Summary: SARS-CoV-2's Spike protein promotes analgesia by interfering with VEGF-A/NRP1 pathway, which may affect disease transmission dynamics.

Main Text: Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is the causative agent of COVID-19, a coronavirus disease that, as of July 17, has infected more than 13.8 million people and caused nearly 600,000 deaths worldwide (*1*). Most patients infected with SARS-CoV-2 report mild to severe respiratory illness with symptoms such as fever, cough and shortness of breath (*2*). On the other hand, a subset of patients who are diagnosed by a positive nuclei acids test but are either asymptomatic or minimally symptomatic (*2*). Increasing evidence shows that asymptomatic individuals can spread the virus efficiently, and the emergence of these silent spreaders of SARS-CoV-2 has limited control of the pandemic (*3*). Pain is a rising concern in symptomatic patients, likely emanating from a direct attack of SARS-CoV-2 on cells and the "cytokine storm" unleashed by affected cells (*4, 5*). Whether asymptomatic or minimally

symptomatic individuals have reduced pain thresholds, or whether their pain is silenced is unknown, but either could contribute to increased disease transmission dynamics.

The surface expressed angiotensin converting enzyme 2 (ACE2) has been lionized as the main receptor for uptake of SARS-CoV-2 (6-8). Emerging evidence points to a subset of ACE2 expressing sensory neurons (9) that synapse with spinal and brainstem CNS neurons to produce neurological effects, including headache and nerve pain (10, 11). Curiously, ACE2 is not present in most neurons (9), despite increasing reports of neurological symptoms being common in COVID-19 patients (10). Paradoxically, though the levels of ACE2 expression decline in aging (12), increased COVID-19 severity was noted in older patient populations, such as that of Italy's (13), supporting the contention that ACE2 is not the sole gateway for entry of SARS-CoV-2 (14).

Two recent reports demonstrated that the SARS-CoV-2 Spike protein can bind to the b1b2 domain of the neuropilin-1 receptor (NRP-1). This interaction occurs through a polybasic amino acid sequence ([682]RRAR[685]), termed the 'C-end rule' (CendR) motif, which significantly potentiates its entry into cells (15, 16). Importantly, 'omic' analyses revealed a significant upregulation of *NRP-1* in biological samples from COVID-19 patients compared to healthy controls (15). Using vascular endothelial growth factor-A (VEGF-A), a physiological ligand for the b1b2 pocket in NRP-1, we interrogated whether the Spike protein, the major surface antigen of SARS-CoV-2, could block VEGF-A/NRP-1 signaling to affect pain behaviors. Given parallels between the pro-nociceptive effects of VEGF-A in rodents (17, 18) and humans (18, 19) and clinical findings demonstrating increased VEGF-A levels in bronchial alveolar lavage fluid from COVID-19 patients (20) coupled with substantially lower levels in the sera of asymptomatic individuals compared to symptomatic patients (2), a secondary question was to test whether Spike protein could confer analgesia. We found that VEGF-A sensitizes nociceptor activity, a hallmark of neuropathic pain (21), which was blocked by the Spike protein and NRP-1 inhibitor EG00229 (22). Furthermore, we identify a novel analgesic role for Spike protein, which is mirrored by NRP-1 inhibition.

Results

Ligand specific engagement of NRP-1 signaling induces nociceptor activity and pain.

Initially, we assessed the involvement of Spike and NRP-1 in the VEGF-A/NRP-1 pathway. We plated sensory neurons on multiwell microelectrode arrays (MEAs), an approach enabling multiplexed measurements of spontaneous, as well as stimulus-evoked extracellular action potentials from large populations of cells (23). VEGF-A increased spontaneous firing of dorsal root ganglion (DRG) neurons, which was blocked by the receptor binding domain (RBD) of the Spike protein and by the NRP-1 inhibitor EG00229 (Fig. 1A). In contrast, ligands VEGF-B (ligand for VEGFR1 – a co-receptor for NRP-1 (24)) and semaphorin 3A (Sema3A, ligand for plexin receptor – also a co-receptor for NRP-1) (25, 26)) did not affect the spontaneous firing of nociceptors (Fig. 1A). The lack of effect of VEGF-B and Sema3A rule out a role for VEGF-R1 and plexin, respectively, thus implicating a novel ligand-, VEGF-A, and receptor-, NRP-1, specific pathway driving nociceptor firing.

As both VEGF-A and Spike protein share a common binding pocket on NRP-1 (Fig. 1B) (15, 16, 27), we asked if the Spike protein could block VEGF-A/NRP-1 signaling to affect pain behaviors. Consistent with previous reports (17, 18), we confirmed that VEGF-A is pro-nociceptive as intra-plantar injection of VEGF-A decreased both paw withdrawal thresholds

(Fig. 1C, D and Table S1) and latencies to a thermal stimulus (Fig. 1E, F and Table S1). Preventing VEGF-A from binding to NRP-1 with the NRP-1 inhibitor EG00229, or Spike from activating VEGF-A/NRP-1 signaling, blunted the mechanical allodynia and thermal hyperalgesia induced by VEGF-A alone (Fig. 1C-F and Table S1). Neither Spike nor EG00229 alone had any effect on these behaviors (Fig. 1C-F and Table S1). Together, these data provide functional evidence that VEGF-A/NRP-1 signaling promotes a pain-like phenotype by sensitizing nociceptor activity (Fig. 1G).

VEGF-A–mediated increases in DRG ion channel currents are normalized by disruption of VEGF-A/NRP-1 signaling.

To gain insight into the mechanism by which VEGF-A contributed to increased nociceptor activity, we postulated that ion channels in DRGs may be affected, as these contribute to nociceptive plasticity (*28*). Typical families of Na^+ currents from small diameter DRG neurons are shown in Figure 2A. VEGF-A facilitated a 1.9–fold increase in total Na^+ currents compared to vehicle (PBS)-treated DRGs, which was completely blocked by Spike protein (Fig. 2B, C). Spike protein alone did not affect Na^+ currents (Fig. 2B, C and Table S1). Since this decreased current could arise from changes in channel gating, we determined if activation and inactivation kinetics of DRG Na^+ currents were affected. Half-maximal activation and inactivation ($V_{1/2}$), as well as slope values (*k*) for activation and inactivation, were no different between the conditions tested (Fig. 2D, E and Tables S1, S2), except for an ~8 mV hyperpolarizing shift in sodium channel inactivation induced by co-treatment of VEGF-A and EG00229 (Table S2). Similar results were obtained for the NRP-1 inhibitor EG00229, which also inhibited the VEGF-A mediated increase in total Na^+ currents (Fig. 2F-H and Table S1) but had no effect on the biophysical properties (Fig. 2I, J and Tables S1, S2).

As calcium channels play multiple critical roles in the transmission and processing of pain-related information within the primary afferent pain pathway (*28*), we evaluated if they were affected. We focused on N-type (CaV2.2) channels as these mediate neurotransmitter release at afferent fiber synapses in the dorsal horn and are critical in the pain matrix (*29*). VEGF-A facilitated a 1.8–fold increase in total Ca^{2+} currents compared to vehicle (PBS)-treated DRGs, which was completely blocked by Spike protein (Fig. 2K-M and Table S1). Spike protein alone did not affect Ca^{2+} currents (Fig. 2B, C). Additionally, we did not observe any changes in activation and inactivation kinetics between the conditions tested (Fig. 2N, O and Tables S1, S2). Similar results were obtained for the NRP-1 inhibitor EG00229, which inhibited the VEGF-A mediated increase in N-type Ca^{2+} currents (Fig. 2P-R and Table S1) but had no effect on the biophysical properties (Fig. 2S, T and Tables S1, S2). These data implicate Spike protein and NRP-1 in Na^+ and Ca^{2+} (CaV2.2) channels in VEGF-A/NRP-1 signaling.

VEGF-A enhances synaptic activity in the lumbar dorsal horn that is that normalized by inhibition of NRP-1 signaling and Spike protein.

The spinal cord is an integrator of sensory transmission where incoming nociceptive signals undergo convergence and modulation (*30*). Spinal presynaptic neurotransmission relies on DRG neuron action potential firing and neurotransmitter release. From these fundamental physiological principles, as well as the results described above, we were prompted to evaluate whether synaptic activity was affected in the lumbar dorsal horn. The amplitudes of spontaneous excitatory postsynaptic currents (sEPSCs) of neurons in the *substantia gelatinosa* region of the lumbar dorsal horn were not affected by VEGF-A (Fig. 3A, B and Table S1). In contrast, VEGF-

A application increased sEPSC frequency by ~3.6–fold, which was reduced by ~57% by inhibition of NRP-1 with EG00229 and ~50% by Spike protein (Fig. 3A, C and Table S1). Amplitude and inter-event interval cumulative distribution curves for sEPSCs are shown in Figure 3D, E. When compared to vehicle controls, VEGF-A, with or without NRP-1 inhibitor or Spike protein, had no effect on the cumulative amplitude distribution of the spontaneous EPSCs (Fig. 3D and Table S1) but changed the cumulative frequency distribution of spontaneous EPSCs with significantly longer inter-event intervals (Fig. 3E and Table S1). Together, these data suggest a presynaptic mechanism of action of Spike protein and NRP-1.

Spike protein and inhibition of NRP-1 confer anti-nociception in the spared nerve injury model (SNI) of chronic neuropathic pain.

We used the spared nerve injury (SNI) model of neuropathic pain, chosen because it produces a reliable and consistent increase in pain sensitivity (*31*), to evaluate the potential of disruption of the VEGF-A/NRP-1 pathway to reverse nociception. In rats with SNI, intrathecal application of spike, decreased the phosphorylation of VEGFR2 (Y1175) on both the contralateral (non-injured) and ipsilateral side (injured) (Fig. 4A, B). This shows that Spike can inhibit VEGF-A signaling in a rat model of chronic neuropathic pain. SNI injury efficiently reduced paw withdrawal thresholds (PWTs) (mechanical allodynia, Fig. 4C and Table S1) 10 days post injury. Spinal administration of Spike protein significantly increased PWTs (Fig. 4C and Table S1) for 5 hours. Analysis of the area under the curve (AUC) confirmed the reversal of mechanical allodynia (Fig. 4D and Table S1) compared to vehicle-treated injured animals. Similar results were obtained with inhibition of NRP-1 signaling with EG00229 (Fig. 4E, F and Table S1).

Discussion

Our data show that SARS-CoV-2 Spike protein subverts VEGF-A/NRP-1 pro-nociceptive signaling. Relevant to chronic neuropathic pain, Spike negated VEGF-A–mediated increases in: (*i*) voltage-gated sodium and N-type calcium current densities in DRG neurons; (*ii*) spontaneous firing in DRG neurons; (*iii*) spinal neurotransmission; and (*iv*) mechanical allodynia and thermal hyperalgesia. Consequently, Spike protein was analgesic in a nerve injury rat model. Based on the reported increase in VEGF-A levels in COVID-19 patients (*20*), one would expect to observe increased pain-related symptoms. However, our data suggest that the SARS-CoV-2 Spike protein hijacks NRP-1 signaling to ameliorate VEGF-A mediated pain. This raises the possibility that pain, as an early symptom of COVID-19, may be directly dampened by the SARS-CoV-2 Spike protein. Leveraging this atypical function of SARS-CoV-2 Spike protein may yield a novel class of therapeutics for pain.

Clinical findings that VEGF-A contributes to pain are supported by observations that in osteoarthritis increased VEGF expression in synovial fluids has been associated with higher pain scores (*32*). VEGF-A has been reported to enhance pain behaviors in normal, nerve-injured and diabetic animals (*18, 19*). Our data shows that VEGF-A elicits long-lasting (up to 24 hours) mechanical allodynia and thermal hyperalgesia in naïve rats, thus supporting the premise that VEGF-A is pro-nociceptive. VEGF is augmented in serum of rheumatoid arthritis patients (*17, 33*). VEGF-A was increased in the bronchial alveolar lavage fluid from COVID-19 patients (*20*). The levels of VEGF-A were substantially lower in the sera of asymptomatic individuals compared to symptomatic individuals and matched those found in healthy controls (*2*). Conversely, transcript levels of the VEGF-A co-receptor NRP-1 were increased in COVID-19

patients compared to healthy controls (*15*). NRP-1 is a dimeric transmembrane receptor that regulates pleiotropic biological processes, including axon guidance, angiogenesis and vascular permeability (*34-36*). In chronic neuropathic pain, a concomitant increase of NRP-1 and VEGF-A have been reported in DRG neurons (*37*). We observed that pharmacological antagonism of the VEGF-A binding b1b2 domain of NRP-1 using EG00229 resulted in decreased mechanical allodynia in SNI. A corollary to this, VEGF-A–induced DRG sensitization and the related allodynia/hyperalgesia in naïve rats was annulled by NRP-1 antagonism. This work identifies a heretofore unknown role of VEGF-A/NRP-1 signaling in pain. Our working model shows that VEGF-A engagement of NRP-1 is blocked by Spike protein consequently decreasing activities of two key nociceptive voltage-gated sodium, likely NaV1.7, and calcium (CaV2.2) channels (Fig. 1G). The resulting decrease in spontaneous DRG neuronal firing by Spike protein translates into a reduction in pain (Fig. 1G). Characterization of the molecular cascade downstream of VEGF-A/NRP-1 signaling awaits further work.

Altogether, our data suggest that interfering with VEGF-A/NRP-1 using SARS-CoV-2 Spike or the NRP-1 inhibitor EG00229 is analgesic. For cancer, this pathway has been extensively targeted for anti-angiogenesis. A monoclonal antibody targeting VEGF-A (bevacizumab, Avastin®) has been used as a cancer treatment (*38*). A phase 1a clinical trial of NRP-1 antibody MNRP1685A (Vesencumab®), targeting the against VEGF-binding site of NRP-1, was well-tolerated in cancer patients (*39*). While neither antibody has been evaluated in neuropathic pain, our preclinical work provides a rationale for targeting the VEGF-A/NRP-1 pro-nociceptive signaling axis in future clinical trials.

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Supplementary Materials:

Materials and Methods

Tables S1-S2

References (*41-51*)

Fig. 1. Ligand specific engagement of NRP-1 signaling induces nociceptor activity, promoting a pain-like phenotype. (**A**) Mean action potential firing rates (Hz, event per second) of cultured DRG sensory neurons incubated for 30 min with VEGF-B (3 nM), Sema3A (100 ng), VEGF-A (1 nM), VEGF-A plus Spike (100 nM) or VEGF-A plus NRP-1 inhibitor EG00229 (30 µM) (*22*). Of the ligands tested, only VEGF-A, acting on VEGFR2, is a ligand for NRP-1 that triggers an increase in spontaneous firing of nociceptors. Data is shown as mean ± s.e.m. and was analyzed by non-parametric two-way analysis of variance (post hoc: Sidak). P values, versus control (PBS) or VEGF-A, are indicated. EG00229 is an NRP-1 inhibitor (PDB 3i97). (**B**) *Top*: VEGF-A heparin binding domain (gray cartoon with R323 in sticks) in complex with the NRP-1 b1 domain (white surface with binding site in red; PDB 4deq (*27*)). *Middle*: Peptide from C-terminus of furin cleaved SARS-CoV-2 Spike protein 681-PRRAR-685 (blue sticks) docked to NRP1-b1 domain (white surface with binding site in red; PDB 6fmc (*40*)) using Glide (Schrödinger). Additional Spike residues 678-TNS-680 modeled for illustration purposes only (blue cartoon). *Bottom*: Compound EG00229 (cyan sticks) in complex with NRP-1 b1 domain ((white surface with binding site in red; PDB 3i97 (*22*)). Paw withdrawal thresholds (**C, D**) or latencies (**E, F**) for male naïve rats injected in the paw with VEGF-A (10 nM), Spike (100 nM), EG00229 (300 µM) or PBS (vehicle), alone or in combination (50 µl/rat; = 6-12). For clarity, statistical significance is not presented in the time course graphs, instead it is presented in Table S1. Panels **D** and **F** are the area under the curve for 0-24 hours. Data is shown as mean ± s.e.m. and was analyzed by non-parametric two-way analysis of variance where time was the within subject factor and treatment was the between subject factor (post hoc: Sidak), *p<0.05. Areas under the curve were compared by Mann-Whitney test. The experiments were analyzed by an investigator blinded to the treatment. For full statistical analyses see Table S1. (**G**) Schematic illustration of the hypothesis that SARS-CoV-2 Spike protein binding to NRP-1 b1b2 domain triggers an intracellular cascade that increases sodium and calcium channel activity to increase nociceptor activity culminating in enhanced pain.

Fig. 2. VEGF-A–mediated increase in sodium and calcium currents is normalized by Spike protein or NRP-1 inhibition in DRG neurons. Representative sodium current traces (**A, F**) recorded from small-sized DRGs neurons, incubated for 30 min with the indicated treatments, in response to depolarization steps from −70 to +60 mV from a holding potential of −60 mV. Summary of current-voltage curves (**B, G**) and normalized peak (**C, H**) currents (pA/pF) from DRG neurons as indicated. Boltzmann fits for normalized conductance G/G$_{max}$ voltage relationship for voltage dependent activation (**D, I**) and inactivation (**E, J**) of the sensory neurons as indicated. Representative calcium current (via N-type channels) traces (**K, P**) recorded from small-sized DRGs neurons, incubated for 30 min with the indicated treatments, in response to holding voltage of −60 mV with 200-ms voltage steps applied at 5-s intervals in +10 mV increments from −70 to +60 mV. Pharmacological isolation of N-type (CaV2.2) current was achieved with a cocktail of toxins/small molecules. Summary of current-voltage curves (**L, Q**) and normalized peak (**M, R**) currents (pA/pF) from DRG neurons as indicated. Boltzmann fits for normalized conductance G/Gmax voltage relations for voltage dependent activation (**N, S**)

and inactivation (**O**, **T**) of the sensory neurons as indicated. Error bars indicate mean ± s.e.m. Half-maximal activation and inactivation ($V_{1/2}$) and slope values (k) for activation and inactivation were not different between any of the conditions (p >0.9999, Kruskal-Wallis test with Dunn's post hoc); values presented in Table S2. P values of comparisons between treatments are as indicated; for full statistical analyses see Table S1.

Fig. 3. sEPSC frequency is reduced by pharmacological antagonism of NRP-1 by EG00229. (**A**) Representative traces of spontaneous excitatory postsynaptic currents (sEPSC) from neurons from the substantia gelatinosa in the superficial dorsal horn (lamina I/II) treated for at least 30 min with the indicated conditions. Summary of amplitudes (**B**) and frequencies (**C**) of sEPSCs for all groups are shown. Cumulative distribution of the sEPSCs amplitude (**D**) and the inter-event interval (**E**) recorded from cells as indicated. Perfusion of 30 μM EG00229 decreased spontaneous excitatory synaptic transmission (**A-E**) in lumbar dorsal horn neurons. P values of comparisons between treatments are as indicated; for full statistical analyses see Table S1.

Fig. 4. SARS-CoV-2's Spike protein and NRP-1 antagonism reverses SNI-induced mechanical allodynia. Spared nerve injury (SNI) elicited mechanical allodynia 10 days after surgery. (**A**) Representative immunoblots of NRP-1, total and pY1175 VEGF-R2 levels at pre-synaptic sites in rat spinal dorsal horn after SNI. (**B**) Bar graph with scatter plots showing the quantification of n= 4 samples as in **A** (*p<0.05, Kruskal-Wallis test). Paw withdrawal thresholds for SNI rats (male) administered saline (vehicle) or receptor binding domain of the Spike protein (2.14 μg/5μl) intrathecally (i.t.); n = 9-12) (**C**) or the NRP-1 inhibitor EG00229 (2.14 μg/5μl; n = 6) (**E**). (**D**, **F**) Summary of data shown in panels **C** and **E** plotted as area under the curve (AUC) for 0-5 hours. P values, versus control, are indicated. Data is shown as mean ± s.e.m. and was analyzed by non-parametric two-way analysis of variance where time was the within subject factor and treatment was the between subject factor (post hoc: Sidak). AUCs were compared by Mann-Whitney test. The experiments were analyzed by an investigator blinded to the treatment. P values of comparisons between treatments are as indicated; for full statistical analyses see Table S1.

A

Contra | Ipsi

PBS | Spike | PBS | Spike

VEGFR2
- pY1175
- Total

NRP1

Flotilin

Pre-synaptic fraction

B

○ PBS ○ Spike ● PBS ● Spike

normalized expression

pY1175 | Total | NRP1
VEGFR2

C

● PBS vehicle ● Spike 2.14 μg in 5 μl (i.t.)

Paw withdrawal threshold (g)

a
p=0.0002
a
a
a
p=0.0063

a = p<0.0001

Pre-SNI 0 1 2 3 4 5
Post-i.t. injection (hours)

D

p=0.0002

Area under the curve

(12) (9)

E

● PBS vehicle ● EG00229 2 μg in 5 μl (i.t.)

Paw withdrawal threshold (g)

a a p=0.0006 a
a

a = p<0.0001

Pre-SNI 0 1 2 3 4 5
Post-i.t. injection (hours)

F

p=0.0087

Area under the curve

(6) (5)