

Modulation of the α 7 nicotinic acetylcholine receptor by the "neurotoxin-like region" of the SARS-CoV-2 spike protein is selective, allosteric, and concentrationspecific (a role in aggression and anxiety?) | 1

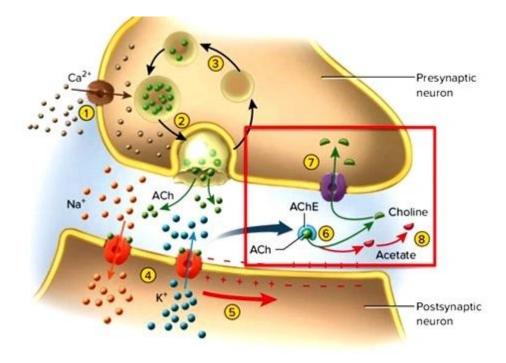
The SARS-CoV-2 spike (S) protein contains a neurotoxin-like region that has sequence similarities with the ectodomains of the rabies virus strains and snake neurotoxins (for example, α -bungarotoxin from snake *Bungarus* genera). In this work, the authors from the United States investigated whether the neurotoxin-like region of the SARS-CoV-2 S protein targets nicotinic acetylcholine receptors (nAChRs) and whether nicotine modifies this potential interaction.

AChRs are categorized as either metabotropic muscarinic (mAChRs) or ionotropic nicotinic acetylcholine receptors (nAChRs). Ionotropic neuromuscular and neuronal nAChRs are pentameric, cation-conducting channels that respond to the endogenous neurotransmitter acetylcholine (ACh) and are involved in signal transduction. Fast and non-selective opening of membrane-bound, excitatory cation channels results from their activation. The neuronal nAChRs are widely distributed throughout the central nervous system (CNS), particularly in the forebrain and brainstem. They have important physiological functions, including neurotransmitter release, mediation of cholinergic excitatory neurotransmission, synchronization of neuronal activity, and regulation of essential physiological functions such as cognition, arousal, sleep, fatigue, anxiety, nutrition, central processing of pain, attention, and behavior (aggression, mood, and impulsivity).

In addition, nAChRs are present in numerous immune cells, including macrophages, T cells, dendritic cells, and B-cells. ACh is involved in an anti-inflammatory response called the cholinergic anti-inflammatory pathway (CAP) that suppresses inflammation in the brain and peripheral tissues *via* the autonomic nerve fibers.



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Acetylcholine neurotransmission

The nAChRs are composed of different homomeric or heteromeric combinations of 16 subunits, designated as $\alpha 1 - \alpha 7$, $\alpha 9$, $\alpha 10$, $\beta 1 - \beta 4$, δ , ε , and γ , based on sequence homology. The subunits $\alpha 2 - \alpha 7$, $\alpha 9 - \alpha 10$, and $\beta 2 - \beta 4$ are expressed by neurons and are referred to as the neuronal nAChR subunits. The nAChRs localized at the mammalian neuromuscular junctions are composed of ($\alpha 1$)2 $\beta 1\delta \gamma$ subunits in developing muscle, with the subunit ε substituting the subunit γ in mature muscle. https://doi.org/10.1016/j.heliyon.2022.e10434

The pharmacological and biophysical properties of these receptors are determined by the composition of nAChR subunits. The most common nAChRs in the CNS, $\alpha 4\beta 2$ and the $\alpha 7$ subtypes are linked to hyperactivity, aggression, and anxiety. According to previous rodent and human studies, neuronal $\alpha 7$ nAChR plays a critical role in the modulation of aggression behavior. It was demonstrated that the $\alpha 7$ nAChR was necessary for the anti-aggressive or 'serenic' effects of systemic administration of nicotine. The hippocampal $\alpha 7$ nAChR seems to directly regulate aggression in mice, whereas the loss of its function increases aggression.

Changeux proposed initially that the neurotoxin-like region of the SARSCoV-2 S protein interacts with

nAChRs. https://comptes-rendus.academie-sciences.fr/biologies/item/CRBIOL_2020__343_1_ 33_0/ Farsalinos et al. identified a "toxin-like" amino-acid sequence in the receptor binding domain (RBD) of the SARS-CoV-2 S protein (amino-acid 375–390) which shows significant



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sequence homology with the neurotoxin homolog NL1, one of the many snake venom toxins interacting with nAChRs. They demonstrated that the main interaction occurs between the amino-acid sequence 381-386 of the RBD of the SARS-CoV-2 S protein and the amino-acid sequence *189–192* of the extracellular domain of the α9 nAChR, which is the core of nAChR "toxin-binding site". According to the authors, these findings strongly supported the hypothesis of the significant role of nicotinic cholinergic system dysregulation in the pathophysiology of COVID-19. https://www.mdpi.com/1422-0067/21/16/5807 Later in silico study predicted that the neurotoxin-like region of the SARS-CoV-2 RBD interacts with high variability with models of the α 7 and α 4 β 2 nAChR. Agonists of nAChR are thought to stabilize a compact C loop conformation, whereas antagonists prevent C loop closure. In the $\alpha 4\beta 2$ model, the neurotoxin-like region was unable to bind deeply into the orthostericbinding site, keeping the C loops of the receptor in an open conformation. Nevertheless, in the α7 nAChR model of nAChR, the neurotoxin-like region showed multiple modes of C loop, ranging from an open conformation to a semiclosed structure if it moved deeper into the binding pocket. https://www.sciencedirect.com/science/article/pii/S0006349521001466 Other studies have also discussed the role of cholinergic deficiency in the formation of various syndromes in COVID-19. A recent animal study has shown that inoculation of the S1 subunit of the SARS-CoV-2 S protein in the olfactory cavity resulted in increased apoptosis of the olfactory system, brain inflammation, and reduced ACh levels in the mouse brain. The administration of donepezil, a central cholinergic agent and acetylcholine esterase inhibitor, normalized inflammatory cytokines that were previously elevated and mitigated enhanced apoptosis in the olfactory bulb.

https://discovermednews.com/s1-protein-causes-brain-inflammation-and-decreases-the-acety lcholine-levels/

About the study

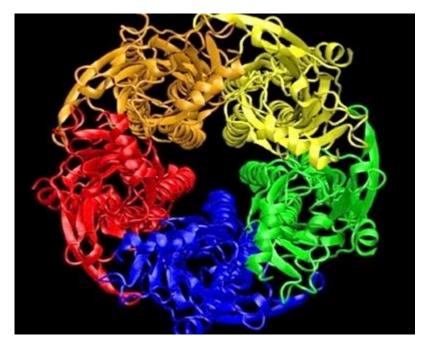
The authors employed two-electrode voltage clamp electrophysiology to investigate the interaction of subtypes α 7, α 4 β 2, α 3 β 4, and α 3 β 2 of nAChRs with SARS-CoV-2 glycoprotein peptide (ScoV2P), which is the neurotoxin-like region of SARS-CoV-2, and the SARSCoV-2 ectodomain (SCoV2ED). These nAChRs were chosen because they are expressed in the target tissues of SARS-CoV-2, such as the nose, lungs, CNS, and some immune cells. The researchers hypothesized that SCoV2P and SCoV2ED will antagonize the chosen subtypes of nAChR, and that nicotine will enhance this inhibition effect.

Live-cell confocal microscopy was used to confirm that SCoV2P interacts with nAChRs in



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transfected neuronal-like N2a and human embryonic kidney 293 (HEK293) cells.



 $\alpha 7 \; nAChR$

Results

High concentrations of ScoV2P, a neurotoxin-like region of SARS-CoV-2, moderately inhibited the $\alpha 4\beta 2$, $\alpha 3\beta 4$, and $\alpha 3\beta 2$ subtypes of nAChRs.

The ScoV2P exerted an opposing, bimodal, and concentration-specific effect on the α 7 nAChR. A high concentration of ScoV2P significantly inhibited α 7 nAChR and reduced ACh potency, whereas low concentrations potentiated ACh-induced currents. In addition, the results showed that SCoV2P can bind to α 7nAChRs in different orientations and exerts its dual action- potentiation or inhibition of the α 7 nAChR through an allosteric mechanism. SCoV2P did not have any similar allosteric actions on heteromeric nAChRs. At high concentrations, ScoV2P inhibited the ACh-induced currents of α 4 β 2, α 3 β 2, and α 3 β 4 nAChRs with some isoform-specific effects.

Confocal imaging confirmed that ScoV2P interacts with $\alpha 7$ nAChRs expressed on the cell surface.

These findings revealed an antagonizing selective effect of ScoV2P on nAChRs, with a high preference for the α 7 subtype. This confirms that the α 7nAChR is a target for the neurotoxin-like region of the SARS-CoV-2. These results also suggest that SCoV2P binding



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to $\alpha 7$ nAChRs can prime receptors for agonist activation.

Importantly, pretreatment with nicotine enhanced the modulation of α 7 nAChR responses by SCoV2P and ScoV2ED, resulting in increased potentiation *via* a mechanism that resensitizes nicotinic desensitized receptors. In the absence of nicotine, the α 4 β 2, α 3 β 2, and α 3 β 4 subtypes of nAChRs were minimally inhibited. Pretreatment with 200 nM of nicotine enhanced the inhibition of currents of (α 4 β 2)2 α 4 and (α 3 β 2)2 α 3 nAChRs by ScoV2P and made the inhibition of currents of (α 3 β 2)2 α 3 nAChR similar to the inhibition of α 7 nAChR (40%). According to the authors, nicotine pretreatment broadened the effects of SCoV2P by inhibiting not only the α 7 subtype of nAChRs, but also α 4 β 2 and α 3 β 2 subtypes. They also said that the ability of the SARSCoV-2 neurotoxin-like region to resensitize desensitized α 7 nAChRs in tobacco users could further activate their cholinergic anti-inflammatory response, allowing for higher levels of viral replication.

Conclusion

These results demonstrated that nAChR subtypes interact with the neurotoxin-like region of the SARS-CoV-2, confirming that the α 7 nAChR is a target for the SARS-CoV-2. Low concentrations of SCoV2P and SCoV2ED positively modulated ACh-mediated currents *via* the facilitation of α 7 nAChRs transition to the active conformation, whereas higher concentrations of both ScoV2P and ScoV2ED switched the modulation activity to inhibition.

The α 7 nAChRs are present in numerous immune cells, including macrophages, T cells, dendritic cells, and B-cells. The activation of α 7 nAChRs in the cholinergic antiinflammatory pathway suppresses the production of proinflammatory cytokines. The recombinant trimeric rabies virus glycoprotein binds to α 7 nAChRs expressed on monocyte-derived macrophages, activating the cholinergic anti-inflammatory pathway. This suppressed macrophages' function as T-cell activators, indicating that the rabies virus could evade the immune system by inducing an anti-inflammatory state in human macrophages through interactions with α 7 nAChR. In addition, the dorsal root ganglion, which is believed to be a crucial bridge allowing the rabies virus to pass from the periphery into the CNS expresses a variety of neuronal nAChR subtypes, including α 7. https://doi.org/10.1016/j.heliyon.2022.e104

The authors concluded that understanding how the SARS-CoV-2 neurotoxin-like region affects different nAChR subtypes and associated isoforms provides an understanding of COVID-19 pathophysiology, which very likely facilitates the development of targeted



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therapeutics.

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