



# The $\alpha 9\alpha 10$ acetylcholine receptor: A non-neuronal nicotinic receptor<sup>☆</sup>

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## ABSTRACT

Within the superfamily of pentameric ligand-gated ion channels, cholinergic nicotinic receptors (nAChRs) were classically identified to mediate synaptic transmission in the nervous system and the neuromuscular junction. The  $\alpha 9$  and  $\alpha 10$  nAChR subunits were the last ones to be identified. Surprisingly, they do not fall into the dichotomic neuronal/muscle classification of nAChRs. They assemble into heteropentamers with a well-established function as canonical ion channels in inner ear hair cells, where they mediate central nervous system control of auditory and vestibular sensory processing. The present review includes expression, pharmacological, structure-function, molecular evolution and pathophysiological studies, that define receptors composed from  $\alpha 9$  and  $\alpha 10$  subunits as distant and distinct members within the nAChR family. Thus, although  $\alpha 9$  and  $\alpha 10$  were initially included within the neuronal subdivision of nAChR subunits, they form a distinct clade within the phylogeny of nAChRs. Following the classification of nAChR subunits based on their main synaptic site of action,  $\alpha 9$  and  $\alpha 10$  should receive a name in their own right.

## 1. Introduction

A wide spectrum of ion channel families has evolved over billions of years and plays fundamental roles in all domains of life. Included in this wide variety, the superfamily of pentameric ligand-gated ion channels (pLGICs) is the largest and most functionally diversified within those ion channels which are gated in response to ligand binding [1,2]. The discovery of pLGICs in bacterial species and Archaea indicates an ancient origin for this receptor superfamily [1–4]. In vertebrates, pLGICs are named the Cys-loop family, since all member subunits contain a disulfide cysteine bridge in the extracellular domain which closes a loop comprising 13 amino acids [1,5–7]. This Cys-loop family includes the nicotinic acetylcholine receptors (nAChRs), serotonin type 3 receptors (5-HT<sub>3</sub>), gamma aminobutyric acid type A receptors (GABA<sub>A</sub>) and glycine receptors [1,2,8].

Vertebrate nAChRs are non-selective cation channels. To date, 19 different nAChR subunits have been described:  $\alpha 1$ - $\alpha 10$ ,  $\beta 1$ - $\beta 4$ ,  $\gamma$ ,  $\delta$  and  $\epsilon$  [1,9,10], including  $\alpha 11$  and  $\beta 5$  only identified in some fish species [11].

The vertebrate nAChRs have been classically divided into two subgroups, following their main expression pattern: neuronal and muscle nAChRs. The last nAChR subunits to be identified were  $\alpha 9$  and  $\alpha 10$ , whose main function has been deciphered in inner ear mechanosensory hair cells and lateral line systems [12–18]. The general structure of nAChRs is described in other manuscripts within this special issue and will not be covered here. The present review includes expression, pharmacological, structure-function, molecular evolution and pathophysiological studies, that define receptors composed from  $\alpha 9$  and  $\alpha 10$  subunits as distant and distinct members within the nAChR family.

## 2. Expression of $\alpha 9$ and $\alpha 10$ nAChR subunits

The  $\alpha 9$  subunit was first isolated from a rat olfactory epithelium cDNA library [17]. Early on, in situ hybridization analysis localized  $\alpha 9$  to rat cochlear [17,19] and vestibular hair cells [20–22], the nasal (non-neuronal) epithelium, the pars tuberalis of the pituitary [17], and bone marrow [23]. The  $\alpha 10$  subunit was first isolated from a rat cochlear

**Abbreviations:** ACh, acetylcholine; pLGICs, pentameric ligand-gated ion channels; nAChRs, nicotinic cholinergic receptors; 5-HT<sub>3</sub>, serotonin type 3 receptors, GABA<sub>A</sub>, gamma aminobutyric acid type A receptors; MOC, medial olivocochlear; OHCs, outer hair cells; ERBB2, receptor tyrosine-protein kinase erbB-2; TMIE, transmembrane inner ear; IL-1 $\beta$ , interleukine 1 $\beta$ ; PAMs, positive allosteric modulators, TM, transmembrane region.

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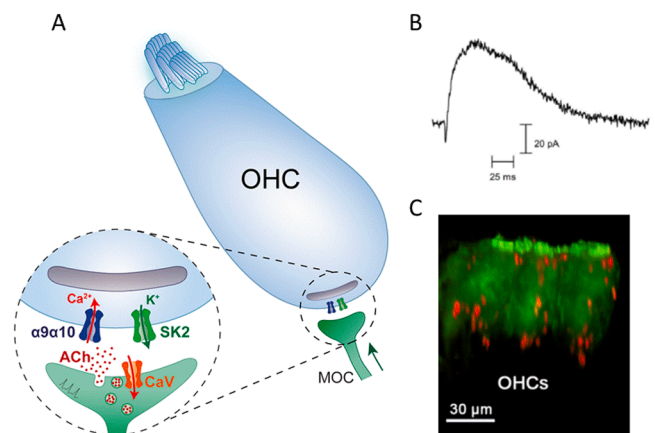
[16] and subsequently from a human cochlear cDNA library [18]. Similar to the  $\alpha 9$  subunit, early studies localized this subunit to cochlear and vestibular hair cells [16,24,25], pars tuberalis of the pituitary and keratinocytes [18]. Further studies have described the expression of  $\alpha 9$  and/or  $\alpha 10$  subunits in several non-neuronal, mostly epithelial cells, such as the adrenal medulla [26], immune cells [27–33], keratinocytes [34], urothelium [35], placenta [36], bronchial epithelium [37], to name a few expression sites. Several studies have reported the expression of  $\alpha 9$  subunits in the retina [38–40]. Since retinal vascular endothelial cells express the  $\alpha 9$  subunit mRNA [40], caution should be taken when arriving at conclusions of neuronal expression when performing RT-PCR experiments from retinal tissue. Further experiments are needed in order to determine a putative neuronal function of  $\alpha 9^*$  in the retina. Although some studies have claimed the presence of  $\alpha 9$  and  $\alpha 10$  subunits in dorsal root ganglia [41], these results have not been replicated. Thus, in a meta-analysis of single-cell gene expression data from mouse dorsal root ganglia neurons,  $\alpha 9$  expression was not detected and  $\alpha 10$  only at very low levels in a few cell subtypes or not found at all [42–44], an observation in line with that reported by quantitative PCR and functional assays [45]. This has been further confirmed with RNAscope in situ hybridization [46]. However, this same study has shown the presence of  $\alpha 9$  ( $\alpha 10$  at very low levels) mRNA in 25% of human dorsal root ganglia neurons [46]. Whether this peculiar differential expression of  $\alpha 9$  subunits in humans is extended to other sites, is unknown at present. In addition, although  $\alpha 9$  and  $\alpha 10$  neuronal and brain expression has also been proposed [47–49], a wealth of experimental data indicates that these nAChR subunits are not expressed in the central nervous system (for review see [50]). Thus, taken into account their known main expression pattern,  $\alpha 9$  and  $\alpha 10$  subunits should not be included within the neuronal subdivision of nAChRs.

### 3. Physiology of $\alpha 9^*$ and $\alpha 10^*$ nAChRs

#### 3.1. The inner ear hair cell nicotinic receptor

The first characterized, and best known function of  $\alpha 9\alpha 10$  nAChRs, is that in inner ear mechanosensory hair cells [16,17]. Moreover, it is only in hair cells that the  $\alpha 9\alpha 10$  nAChR has been unequivocally described to serve a canonical ion channel function in vivo. Thus, the proposal to name  $\alpha 9$  and  $\alpha 10$  as the hair cell subunit branch of nAChRs [42]. An ancestral feature of vertebrate inner ear hair cells is the presence of efferent innervation from (mainly) cholinergic neurons located in the ventral hindbrain that project all the way to the peripheral sensory system [51]. In adult mammals medial olivocochlear (MOC) neurons make direct synaptic contacts with the base of outer hair cells (OHCs), the cochlear amplifier, and inhibit somatic electromotility and thus sound amplification [51]. During a critical pre-hearing period, MOC neurons transiently contact inner hair cell, eliciting  $\alpha 9\alpha 10$  mediated responses [52,53]. Inhibition is brought about by the release of acetylcholine (ACh) from efferent terminals, the activation of very high calcium permeable  $\alpha 9\alpha 10$  nAChRs [54–56] of OHCs, coupled to the subsequent activation of  $\text{Ca}^{2+}$  activated SK2  $\text{K}^+$  channels and hair cell hyperpolarization [57,58] (Fig. 1). This gives rise to the biphasic nature of postsynaptic responses to ACh and is the only example of a nicotinic receptor serving an overall inhibitory synaptic transmission [53,59–62]. A similar biphasic response mediated by  $\alpha 9^*$  receptors coupled to  $\text{K}^+$  channels mediates efferent inhibition of Type II vestibular hair cells [63]. While  $\alpha 9$  subunits can assemble into functional homomeric receptors when expressed in heterologous systems [16,18], these do not play a major role in inner ear hair cells in vivo, as seen in *Chrna10* gene knockout mice [14].

A detailed analysis of the synaptic properties of the MOC-OHC synapse performed in cochlear explants, has shown that a correct operation of this feedback requires matching of acoustic input with the strength of cochlear inhibition [64–66]. This is driven by the rate of MOC activity, by short-term facilitation mechanisms present at the MOC-OHC synapse



**Fig. 1.**  $\alpha 9\alpha 10$  nAChRs mediate transmission at the MOC efferent-hair cells synapses of the cochlea. A. Schematic diagram of the efferent synapse between a medial olivocochlear fiber and an outer hair cell, highlighting the entry of calcium through the  $\alpha 9\alpha 10$  nAChR and the coupling the SK2 channel. B. Representative spontaneous synaptic current recorded in a P10 OHC from a mouse apical cochlear coil, voltage-clamped at  $-60\text{mV}$ . A rapid inward current, mediated by the  $\alpha 9\alpha 10$  nAChR is followed by a larger and longer-lasting outward current, mediated by the SK2 potassium channel. C. Cy3-RgIA-5727 labelling (red) of  $\alpha 9\alpha 10$  nAChRs in OHCs.

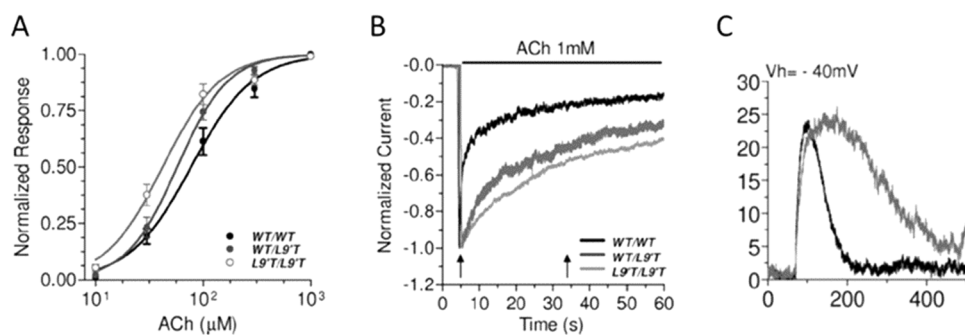
A. (Reproduced from Lipovsek et al., 2021, under the Creative Commons license (<http://creativecommons.org/licenses/by/4.0/>)). B. (Reproduced from Katz and Elgoyhen, 2014, under the Creative Commons license (<http://creativecommons.org/licenses/by/4.0/>)). C. (Reproduced from Fisher et al., 2021, under the Creative Commons license (<http://creativecommons.org/licenses/by/4.0/>)).

[61,67], and by a tight compartmentalization of calcium ions at the base of hair cells [68,69]. In this regard,  $\alpha 9$  and  $\alpha 10$  knockout mice lack OHCs ACh-evoked currents, and therefore efferent inhibition of OHC function [14,15]. On the other hand, mice carrying a threonine for leucine substitution at position 9' (L9 T) of transmembrane region (TM) 2 of the  $\alpha 9$  subunit, lead to hair cell ACh-evoked responses with increased agonist apparent affinity and decreased desensitization kinetics [70] (Fig. 2). This greatly elongates hair cell-evoked inhibitory post-synaptic currents and  $\text{Ca}^{2+}$  signals, leading to enhanced and sustained synaptic responses (Fig. 2) and hair cell hyperpolarization upon high-frequency stimulation of MOC terminals and a longer time course of efferent MOC suppression [70,71]. Since the kinetics of this synapse sets the function of the efferent MOC system, the development of positive allosteric modulators (PAMs) of  $\alpha 9\alpha 10$  nAChRs have been suggested as possible candidates to treat inner ear disorders [72,73].

The generation of  $\alpha 9$  and  $\alpha 10$  knockout mice [14,15,74] and the L9 T knockin [70] have helped elucidate the function of the efferent system in inner ear physiology. Thus, experiments performed with these genetically modified mice have supported the participation of the efferent system in the development of the entire auditory circuitry [71,75–80], in sound localization [81], in the suppression of auditory distractors during selective attention to visual stimuli [82,83], in the protection of the cochlea from the damage produced by the exposure to loud sounds [70,84], and in age-related hearing loss [85]. The use of these genetically modified mice are beginning to disentangle the effect of the efferent innervation to the vestibular system [86].

#### 3.2. Non-auditory functions of $\alpha 9^*$ and $\alpha 10^*$ nAChRs

The non-auditory functions of  $\alpha 9^*$  and/or  $\alpha 10^*$  nAChRs are less clear and are mainly the result of the expression of  $\alpha 9$  and  $\alpha 10$  subunits in epithelial cells outside the inner ear. Of particular importance is the role of ACh as a regulator of the immune system with pro- and anti-inflammatory potential [87,88]. In this regard, the presence of  $\alpha 9$  and



**Fig. 2.** Responses in  $\alpha 9$  L9 T knockin mice. A. Concentration-response curves to ACh performed in P9–P10 IHCs.  $\alpha 9$  L9 T hair cells are hypersensitive to ACh. B. Wave-form of responses to 1 mM ACh during 1 min ACh application in P6–P10 IHCs voltage-clamped at  $-90$  mV.  $\alpha 9$ (L9 T) $\alpha 10$  receptors exhibit slower desensitization kinetics. C. Spontaneous synaptic currents in inner hair cells voltage-clamped at  $-40$  mV, showing prolonged responses in  $\alpha 9$  L9 T knockin mice. Modified from Taranda et al., 2009, under the Creative Commons license (<http://creativecommons.org/licenses/by/4.0/>).

$\alpha 10$  nAChRs in immune cells leads to their participation in several immune and inflammatory responses. For example, brain activity directly enhances the adaptive immune responses in lymphoid organs in a manner that requires B-cell responsiveness to ACh mediated by  $\alpha 9^*$  nAChRs [32]. In addition,  $\alpha 9^*$  nAChRs are involved in endogenous pro-inflammatory mechanisms required for disease initiation and evolution of experimental autoimmune encephalomyelitis, a mouse model for multiple sclerosis, likely via the endogenous agonist acetylcholine [89–91]. Moreover, several antagonists of  $\alpha 9^*$  nAChR have been shown to be analgesic and anti-inflammatory in animal models of neuropathic pain [33,92]. In contrast,  $\alpha 9^*$  nAChRs mediate anti-inflammatory effects during the acute phase response of systemic inflammation caused by trauma or infection [93].

Signaling through  $\alpha 9^*$  nAChR is also important for keratinocyte adhesion and completion of the very early stages of epithelialization [94,95] and form part of the skin non-neuronal cholinergic system which comprises several nicotinic and muscarinic receptors [96]. Moreover, the human  $\alpha 9^*$  nAChR is targeted in Pemphigus vulgaris autoimmunity, suggesting its participation in skin pathology [34]. In addition, a role for  $\alpha 9^*$  nAChRs in stress responses has been suggested by comparing the stress- and affect-related phenotypes of wild-type and  $\alpha 9$  knockout mice [97]. Thus,  $\alpha 9$  knockout mice show decreased stress-induced arousal and increased anxiety-like behavior after a sub-chronic restraint stress. This might be related to the expression of  $\alpha 9$  nAChR subunits in the adrenal medulla [26]. Finally,  $\alpha 9^*$  nAChR have been increasingly involved in the formation, progression, and metastatic spread of various tumors, including lung, breast, glioblastoma, and melanoma cancers (for review see [98,99]).

Recent evidence supports the existence of non-canonical signaling pathways used by ligand-gated ion channels [100]. In particular, nAChRs expressed in numerous non-neuronal cell types serve functions that do not require canonical ion channel function, but rather affect intracellular signal transduction pathways and regulate gene expression [101]. The pharmacology for such non-canonical signaling through nAChRs is different from channel activation [101]. The metabotropic mode of action of nAChRs was first shown for the  $\alpha 7^*$  nAChR in leukocytes [102]. In T cells, activation of  $\alpha 7^*$  nAChRs induces metabotropic signaling that results in an increase of intracellular  $\text{Ca}^{2+}$  concentrations, independent of obvious ionotropic receptor functions [102]. It seems likely that the effects of nAChRs in immune cells is always mediated by non-canonical metabotropic pathways, since no ion-currents have been detected in response to nAChR agonists [28, 102–105].

In human and murine monocytes, ACh acting as a paracrine or autocrine regulator elicits a metabotropic response to reduce ionotropic  $\text{P}2\times 7$  mediated ATP-induced intracellular calcium signals that lead to inflammasome activation, by limiting the release of the pro-inflammatory interleukine 1 $\beta$  (IL-1 $\beta$ ) into the circulation. Moreover, phosphocholine modifies macromolecules produced by pathogens and canonical nicotinic agonists like nicotine and choline inhibit IL-1 $\beta$  release and trigger anti-inflammatory mechanisms via  $\alpha 7$  and  $\alpha 9\alpha 10$

nAChRs [104,105]. Since phosphocholine and nicotine do not evoke ion current responses in  $\alpha 9$  and  $\alpha 10$  cRNA injected *Xenopus laevis* oocytes,  $\alpha 9\alpha 10$  nAChR-mediated immuno-modulatory effect is not due to a canonical ionotropic receptor function [105]. Moreover, choline-gated currents in *Xenopus* oocytes expressing human  $\alpha 9\alpha 10$  nAChR are reversibly inhibited by phosphocholine, resembling a silent agonist or antagonist function [105]. Furthermore, metabolites of phosphatidylcholines such as lysophosphatidylcholine and glycerophosphocholine also trigger  $\alpha 9\alpha 10$  nAChR metabotropic activity in monocytes, but do not elicit ion currents in  $\alpha 9\alpha 10$ -expressing oocytes [106]. The mechanism downstream of nAChR activation controlling monocyte ATP ion-channel function is for the most part unknown and currently under investigation [107].

Another example of non-canonical signaling of  $\alpha 9^*$  and  $\alpha 10^*$  nAChRs is that in cancer cells (for review see [99,108]). In a recent work, several  $\alpha 9$  nAChR interacting and downstream proteins were identified from a resource of cancer membrane protein-regulated networks (CaMPNets), containing 63,746 high confidence protein–protein interactions for 1962 membrane proteins, using expression profiles from 5922 tumors across 15 human cancers [98]. These were related to cell growth and communication in more than six cancer types, thus suggesting that  $\alpha 9$  plays a role in the formation, progression, and metastatic spread of various tumors [98]. These interactions were further verified by immunoprecipitation and FRET analysis and include proto-oncogene tyrosine-protein kinase, stratifin, receptor tyrosine-protein kinase erbB-2 (ERBB2), receptor tyrosine-protein kinase erbB-3, tyrosine 3-Monooxygenase/Tryptophan 5-Monooxygenase Activation Protein Gamm, and casein kinase I isoform delta. Moreover, this work shows dissociation of the  $\alpha 9$ /ERBB2 complex in the presence of nicotine, indicating that the  $\alpha 9$  interactome in cancer cells might be a pharmacotherapeutic target [98].

#### 4. Structure-function of $\alpha 9^*$ and $\alpha 10^*$ nAChRs

Early on classical biochemical and mutagenesis analysis [109,110], together with the low resolution of the Torpedo muscle-type nAChR [111,112], delineated the structure of nAChRs. These studies have been recently refined by the crystal structures of the molluscan ACh binding proteins (AChBPs) [113,114], the high resolution monomeric extracellular domains of the  $\alpha 1$ ,  $\alpha 9$  and  $\beta 3$  [115–117,118] and homopentameric  $\alpha 2$  nAChRs [119], the  $\alpha 7$ -AChBP chimera [119], the near-intact  $\alpha 4\beta 2$  [120], the cryoEM structures of  $\alpha 4\beta 2$  [121] and of the near-intact  $\alpha 7$  nAChR [122,123].

nAChRs are pentameric assemblies of homologous subunits. Each has a similar structure with a large extracellular N-terminal region, four transmembrane helices (TM1–TM4), and an intracellular domain between TM3 and TM4 [112,113,124]. The orthosteric agonist binding site lies at the interface of the extracellular domains of adjacent subunits and is formed by six noncontiguous regions (loops A–F). Each binding site is composed of a principal component or (+) face provided by loops A–C of one subunit and a complementary component (–) comprising loops D–F

of the adjacent subunit [112,113,124]. Moreover, nAChRs can be assembled from either homomeric or heteromeric subunit combinations [125]. In the case of homomeric receptors, such as  $\alpha 7$ , the ACh binding sites are composed from the identical principal and complementary components and are therefore equivalent. In contrast to homomeric nAChRs, heteromeric receptors can have nonequivalent ACh binding sites provided by different subunit interfaces due to the combination of  $\alpha$  and  $\beta$  subunits, in the case of neuronal receptors [125].

So far only functional heteromeric assemblies of  $\alpha 9$  and  $\alpha 10$  subunits have been reported, lacking convincing evidence of assembly with other nAChR subunits [17,42,126]. Taking into account its tissue distribution, sequence identity to other nAChR subunits and pharmacology and evolution,  $\alpha 9$  and  $\alpha 10$  subunits are atypical members of the nAChR family. Different to muscle and neuronal nAChRs, when combined into a heteromeric assembly they form a receptor composed only of  $\alpha$  subunits [16,18]. A  $(\alpha 9)_2(\alpha 10)_3$  stoichiometry has been determined for the rat recombinant receptor [127], although expression of a 10-fold excess of  $\alpha 9$  in *Xenopus* oocytes can lead to an additional receptor isoform with a  $(\alpha 9)_3(\alpha 10)_2$  stoichiometry [128]. The arrangement of subunits in the pentamer and the relative contribution of each subunit to the binding pocket of the heteromeric  $\alpha 9\alpha 10$  receptor is unknown, since only monomers of the  $\alpha 9$  human extracellular domain have been crystallized [115,116]. Since murine  $\alpha 9$  subunits can form functional homomeric  $\alpha 9$  receptors [17], the  $\alpha 9$  subunit is capable of providing principal and complementary components to functional agonist binding sites. In contrast, initial experiments indicated that rat and human  $\alpha 10$  subunits do not form functional channels when heterologously expressed in *Xenopus laevis* oocytes [16,18], leading to the proposal that  $\alpha 10$  might only provide complementary components to agonist binding and serve as a structural subunit, similar to the  $\beta$  subunits of heteromeric nAChRs [58]. However, this is not the case, since site-directed mutagenesis coupled to molecular docking analysis indicate that murine  $\alpha 10$  subunits contribute to the principal component of the binding site [129]. In addition, these studies have shown that the contribution of  $\alpha 9$  and  $\alpha 10$  to the complementary component of the binding site is nonequivalent, since mutagenesis studies have indicated that the binding site does not equally involve residue W55 of loop D (which is key for ACh binding and gating of nAChRs [10,130]) when either provided by  $\alpha 9$  or  $\alpha 10$  [129]. Moreover, docking analysis show that for rat  $\alpha 9\alpha 10$  nAChRs the most frequent conformations with ACh bound in the correct orientation are observed at the interface in which  $\alpha 10$  contributes to the principal and  $\alpha 9$  to the complementary component [129]. The non-equivalent contribution of murine  $\alpha 9$  and  $\alpha 10$  subunits to the complementary component of the binding site might be due to differences in the subunit amino acid sequence. Thus, residues on a radial distribution of 5 Å from the ACh binding site are the same in  $\alpha 9$  and  $\alpha 10$  for the principal components (Y93, S148, W149, Y190, C192, and Y197), whereas an excess of positively charged residues are present in the complementary component of  $\alpha 10$  compared to  $\alpha 9$  [124,129]. This might produce a repulsion of positively charged compounds and less favorable contribution of  $\alpha 10$  to the (-) face. In addition, recent experiments show that ACh-evoked currents can be obtained in *Xenopus* oocytes that express human  $\alpha 10$  homomers, by exposing the oocytes to the alkaloids strychnine, brucine, or methyllycaconitine, further indicating that this subunit can contribute to principal and complementary components of the binding site [131]. Finally, the crystal structure of the potent  $\alpha 9\alpha 10$  receptor antagonist  $\alpha$ -CTx RgIA bound to the monomeric extracellular domain of  $\alpha 9$ , followed by molecular dynamics experiments [115], in addition to mutagenesis analysis [132] have shown that this conotoxin interacts at the  $\alpha 9(+)\alpha 9(-)$ ,  $\alpha 10(+)\alpha 9(-)$  rather than the  $\alpha 9(+)\alpha 10(-)$  nAChR subunit interface, indicating the participation of  $\alpha 10$  subunits mainly to the principal component of the binding site.

One distinct feature of the recombinant and native hair cell  $\alpha 9\alpha 10$  nAChR is its pharmacological profile. Although the  $\alpha 9\alpha 10$  nAChR is activated by ACh, most classical nicotinic receptor agonists, including nicotine, do not elicit responses and block ACh-evoked currents [16,18,

133]. Based on the  $\alpha 9$  extracellular domain crystal structures [115,116] and molecular dynamics simulations [132], it has been suggested that the mode of engulfment of a ligand by loop C may differ significantly in  $\alpha 9\alpha 10$  nAChRs compared with other nAChR subtypes [124], due to an excess accumulation of positively charged residues in (-) sides of  $\alpha 9$  and  $\alpha 10$  subunits. This could explain the conversion of classical agonists to antagonists in the case of  $\alpha 9$ -containing nAChRs [124]. In addition, the canonical metabotropic muscarinic receptor agonist muscarine also blocks  $\alpha 9\alpha 10$  nAChRs [16]. Moreover, rat and chicken  $\alpha 9\alpha 10$  nAChRs are blocked by the classical nicotinic and muscarinic antagonists curare and atropine, respectively, thus exhibiting a mixed nicotinic and muscarinic pharmacological profile [16,134]. Most strikingly,  $\alpha 9\alpha 10$  nAChRs are blocked by the glycinergic antagonist strychnine, the GABAergic antagonist bicuculline and the serotonin type 3 receptor antagonists ICS 205–930 and ondansetron, thus sharing pharmacological profiles with other members of the Cys-loop family of pentameric ligand-gated ion channels [16,135,136]. Compared to muscle and  $\alpha 7$  which are irreversibly blocked by  $\alpha$ -bungarotoxin,  $\alpha 9\alpha 10$  nAChRs are reversibly blocked by this compound [16,18], further indicating differential structural determinants and kinetics of ligand binding. Several highly potent and selective peptides that block  $\alpha 9\alpha 10$  nAChR have been isolated from the venom of *Conus* marine snails [137–141]. These open up new opportunities for developing novel peptide therapeutics and molecular probes to study the structure and function of  $\alpha 9\alpha 10$  nAChR [92,138,142].

Recombinant  $\alpha 9\alpha 10$  nAChR were only classically expressed in *Xenopus* oocytes [16,18], due to their inefficient expression in cell lines [143], with the exception of mammalian GH4C1 cells [54]. However, using genome-wide complementary DNA screening to identify functional partners for  $\alpha 9\alpha 10$ , choline acetyltransferase and transmembrane inner ear (TMIE) were isolated and both elicit  $\alpha 9\alpha 10$  nAChR expression in HEK cells [144]. Since expression is also increased by pre-incubation of cells with either ACh or the potent  $\alpha 9\alpha 10$  antagonist methyllycaconitine, the isolation of choline acetyltransferase most likely indicates that expression is enhanced by ligand binding [144]. On the other hand, although co-immunoprecipitation experiments suggest that TMIE might be an auxiliary component of the  $\alpha 9\alpha 10$  receptor [144], its mechanism of action is less clear. TMIE is mainly implicated in mechanotransduction near the tips of hair cell stereocilia [145] and mutations in this gene leads to human deafness [146], a phenotype not seen in  $\alpha 9$  and  $\alpha 10$  knockout mice [14,15,80]. Moreover, TMIE knockout mice do not have a clear lack of efferent function phenotype [144]. Single channel analysis will be required in order to decipher the effect of TMIE in  $\alpha 9\alpha 10$  nAChR function.

## 5. Evolution of $\alpha 9$ and $\alpha 10$ nAChRs subunits

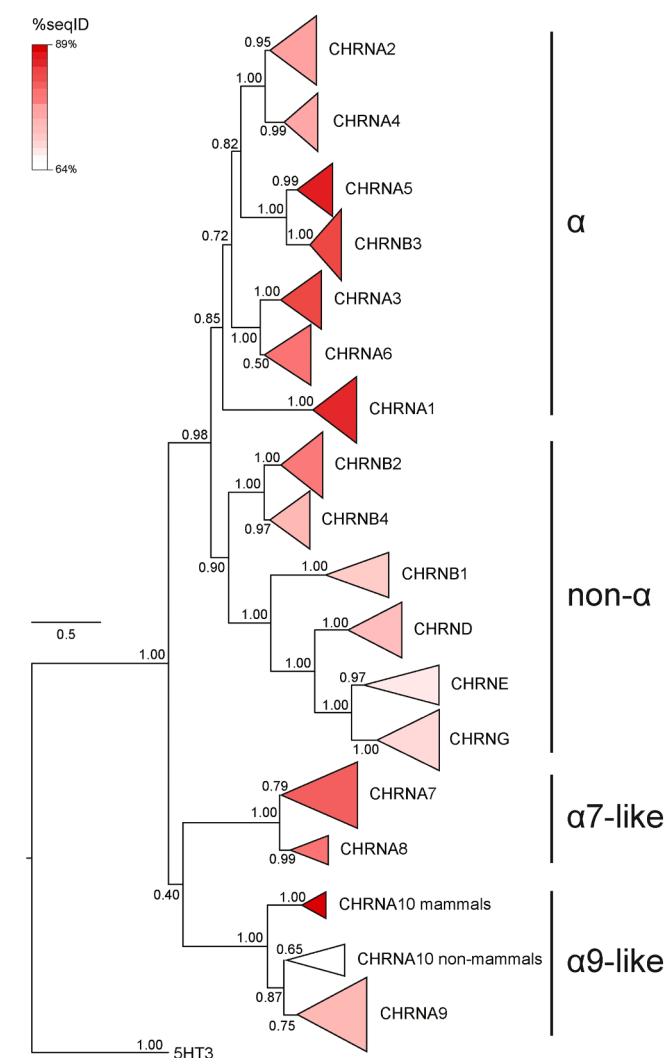
The cloning of the  $\alpha 9$  nAChR subunit and the initial comparison of its amino acid sequence with that of other members of the nAChR family, indicated that whereas neuronal nAChR  $\alpha$  subunits and the muscle  $\alpha 1$  subunit share amino acid sequence identities that range from 48% to 70%, the sequence identity between  $\alpha 9$  and all known nAChR subunits is less than 39%. This, taken together with the fact that the intron-exon structure of the gene encoding the  $\alpha 9$  subunit differs from that of other known nAChR subunit genes, indicate that  $\alpha 9$  is a distinct and distant member of the nAChR family [17]. This has been proved in recent years, with the evolutionary analysis of the increasing amount of known nAChR receptor subunit sequences from different species and *in silico* molecular evolutionary tools, coupled to mutagenesis analysis [9, 11,42,55,147].

Several phylogenetic analyses of the nAChR family of subunits have been reported in the literature [11,42,55,147–155]. Some of them pre-date the high throughput sequencing era, include few sequences and therefore are less informative. The ancestral subunit from which the entire family of extant nAChR subunits radiated, is unknown. The presence of both a neuronal/muscle-like subunit and a  $\alpha 7$ - like subunit



in the last common ancestor of Bilateria is supported by the identification of corresponding groups of subunits in primitive invertebrates by several groups [11,151,153,156–159]. In contrast, current evidence is weak in order to clearly establish the presence of an  $\alpha 9$ -like subunit in the last common ancestor of Bilateria. As sequences become available, further comprehensive analysis of Eumetazoan nAChR subunits will be required in order to claim an ancient and early split of the  $\alpha 9$  nAChR [9], as suggested early on [135,148–150,160] and from then onwards paraphrased in the literature.

The evolution of nAChR subunits in the vertebrate lineage has been extensively studied and the results obtained replicated and reliable. The last common ancestor of vertebrates had a repertoire of 10 genes coding for nAChR subunits [11]. The extant complement of vertebrate subunits most likely resulted from the first and second rounds of tetraploidization which took place in the stem vertebrate branch, gene losses and occasional de novo duplications and, in the teleost lineage, a third whole genome duplication. Thus, the different extant subunits are encoded by paralogous genes, most likely derived from five paralogs [11].



**Fig. 3.** Phylogenetic tree of vertebrate nicotinic subunits. The branches corresponding to the same subunits of different species were collapsed to their respective node. The length of each triangle indicates the sequence divergence from the corresponding node. Triangle colors show the average sequence identity within the branch. Bootstrap values are indicated by the numbers in each branch. The scale bar indicates the number of amino acid substitutions per site.

Modified from Marcovich et al., 2020 and Lipovsek et al., 2021, under the Creative Commons license (<http://creativecommons.org/licenses/by/4.0/>).

The latest phylogenetic study of the nAChR family includes 392 amino acid sequences from 17 nAChR subunits of 29 vertebrate species [42]. The family of nAChR subunits can be divided into four groups:  $\alpha$ , non- $\alpha$ ,  $\alpha 7$ -like, and  $\alpha 9$ -like (Fig. 3). Due to the lowest degree of sequence identity when compared against all other nAChR subunits,  $\alpha 9$  and  $\alpha 10$  subunits form their own group and the outermost branch on the tree. It should be once again noted that, since all phylogenetic trees have been built based only on sequence identity, this position of the  $\alpha 9/\alpha 10$  group cannot be used as evidence for an ancestral origin of these subunits [9].

One important and peculiar feature of the tree is that, due to their high degree of coding sequence conservation, all sequences for each one of the nAChR subunits fall within their own respective branch, except  $\alpha 10$ . These latter subunits are unique in presenting a segregated grouping of orthologs. Thus, non-mammalian  $\alpha 10$  subunits form a sister group to all  $\alpha 9$  subunits, while mammalian  $\alpha 10$  subunits are an outgroup to the  $\alpha 9$ /non-mammalian  $\alpha 10$  branch [42]. This differential grouping of the  $\alpha 9$ -like branch of subunits has been systematically reported in several studies, which have included different number of sequences from different species [55,147,153]. The grouping of  $\alpha 10$  orthologs into differential branches results from a low percentage of amino acid sequence identity between vertebrate  $\alpha 10$  subunits, together with a high conservation of  $\alpha 10$  subunit sequences within the mammalian lineage. The variability in primary sequence of non-mammalian  $\alpha 10$  subunits has resulted from a high rate of acquisition of non-synonymous amino acid substitutions in the coding region of the protein [11,42,55,147], that were subjected to Darwinian positive selection during the course of evolution. Thus, signatures of positive selection acting on CHRNA10 have been determined by several *in silico* molecular evolutionary analysis [55,147]. Moreover, the *in silico* search for site-specific shifts in the amino acid biochemical state between clades, indicates functionally significant amino acid changes when comparing  $\alpha 9$  and  $\alpha 10$  mammalian subunits with their sauropsid counterparts, and no changes in the case of neuronal nAChR subunits [42].

The distinct evolutionary trajectories of  $\alpha 9$  and  $\alpha 10$  compared to that of neuronal nAChR subunits has prompted functional and mutagenesis analysis in order to understand their physiological consequences. Thus, a comprehensive analysis of functional and biophysical properties (ACh sensitivity, calcium permeation and modulation, desensitization rate, voltage-current relationship) of  $\alpha 9/\alpha 10$  receptors from three representative tetrapod species (rat, chicken, and frog) [42] and zebrafish [13], have shown striking differences across them, following the functional divergence analysis at the amino acid sequence. This is in strike contrast to the high degree of functional conservation observed for rat, chicken and frog neuronal  $\alpha 4\beta 2$  and  $\alpha 7$  nAChRs [42].

Positively selected sites identified in the  $\alpha 10$  subunit reside mainly within the conserved ligand-binding and gating regions of the protein [55,147]. Thus, the asymmetric contribution of  $\alpha 9$  and  $\alpha 10$  subunits to the (-) face of the ACh binding site described above might result from the adaptive evolution that occurred in mammalian CHRNA10 genes. For example, residue 117 in Loop E of the complementary component of the binding site, is a positively charged arginine in mammalian  $\alpha 10$  subunits and a non-charged threonine or methionine in  $\alpha 9$  and non-mammalian  $\alpha 10$  subunits [147]. Homology models of the extracellular domain with ACh docked in the binding site show that the positively charged R117 in rat  $\alpha 10$  is located  $\sim 8-9$  Å from the ACh amino group and can thus produce repulsion of the agonist in mammalian, but not in non-mammalian subunits [129]. This seems to be the case, since contrary to that observed for rat subunits, in chicken  $\alpha 9/\alpha 10$  nAChRs both  $\alpha 9$  and  $\alpha 10$  equally contribute to the (-) face of the intersubunit interface when expressed in *Xenopus* oocytes [129]. Moreover, when rat  $\alpha 10$  R117 is replaced by a methionine, mimicking that seen in non-mammalian  $\alpha 10$  and all  $\alpha 9$  subunits, R117M rat mutant  $\alpha 10$  subunits can readily contribute to the complementary component of the binding site of  $\alpha 9/\alpha 10$  receptors [129]. In addition, the evolutionary changes acquired in the mammalian  $\alpha 10$  (-) face of the binding site rendered a rat  $\alpha 9/\alpha 10$  nAChR in which choline acts as a partial agonist, when compared

to its full efficacy in chicken receptors [161].

Clade-specific functionally significant amino acid changes also occurred during the evolution of the  $\alpha 9$  subunits, albeit with a lower prevalence when compared to  $\alpha 10$  [162]. Using molecular dynamics simulations and an evolutionary-based mutagenesis strategy that included ancestral sequence reconstructions, three specific amino acid substitutions in the  $\alpha 9$  subunit that rendered a high calcium permeable mammalian (but not chicken)  $\alpha 9\alpha 10$  nAChR were identified, stemming from a low calcium permeable amniote ancestor. The  $\alpha 9$  identified sites are located at the extracellular vestibule (D110 and S127) and at the exit of the channel pore (4'A) [162].

Taken together, the inter-clade sequence divergence and the positive selection of non-synonymous substitutions in mammalian  $\alpha 9$  and/or  $\alpha 10$  subunits, indicate that contrary to that observed for neuronal nAChR subunits, the evolution of the former subunits has been dominated by functionally significant changes in the coding sequences. This might relate to the fact that hair cells, where the  $\alpha 9\alpha 10$  nAChR plays a major sensory synaptic role as a canonical ion channel, do not express other nAChR subunits [19,20]. Thus, innovation of receptor function across species during evolution could only occur through changes in functional properties of the  $\alpha 9\alpha 10$ , based on changes at the protein coding sequence and not through toggling between alternative expressed subunits, as observed for neuronal nAChRs [9,42]. The evolutionary trajectory of the  $\alpha 9\alpha 10$  nAChR is in line with what has been described for an increasing number of inner ear expressed genes, pinpointing adaptive molecular evolution of protein sequences as a major player in the emergence of morphological and functional innovations in the mammalian inner ear [163,164].

## 6. Targeting $\alpha 9\alpha 10$ nAChRs in pharmacotherapeutics

### 6.1. Inner ear disorders

Hearing loss is one of the most prevalent sensory disabilities and a major public health problem. It impairs language development and education, communication, mental health, employment, with impact in interpersonal relationships and/or psychosocial well-being [165,166]. Moreover, recent studies indicate that hearing loss accelerates cognitive decline and dementia risk in older adults [167–169]. The World Report on Hearing from the World Health Organization [170] and the Global Burden of Disease Study 2019 [171], estimate that 1.5 billion people worldwide will have some hearing decline during their life and that at least 430 million will require intervention, and this will escalate in the years to come. Hearing loss is sometimes accompanied with phantom sound perception, also known as tinnitus, which can be a debilitating condition on its own [172–174]. Within the lifestyle and environmental factors that can be prevented and contribute the most to hearing loss and tinnitus is the exposure to elevated noise (occupational, recreational or environmental) [175–177]. Thus, hearing loss and tinnitus require an urgent call for global attention.

Although a wide variety of drugs have been investigated and/or used in the treatment of noise-induced hearing loss and tinnitus, there is no effective drug on the market [73,178–180]. The fact that the efferent system protects the inner ear from damage produced by exposure to overly loud sounds [70,84,181,182], the enhancement of the efferent feedback to the inner ear by activating  $\alpha 9\alpha 10$  nAChRs, appears as an attractive approach. The phenotype of  $\alpha 9$  L9T knockin mice, which are resistant to permanent noise-induced and hidden hearing loss and have a delayed presbycusis [70,84,85], resembles the effect of type II PAMs described for other nAChRs [183–185], since efferent synaptic responses are prolonged, ACh responses are potentiated and have a slower desensitization kinetics [70,71] (Fig. 2). PAMs which reinforce endogenous cholinergic neurotransmission without directly stimulating the target receptors, do not lead to receptor permanent desensitization, have less side effects and have attracted attention to target neuronal nAChRs in CNS disorders [186]. Thus, the development of PAMs that target

$\alpha 9\alpha 10$  nAChRs appear as attractive molecules to treat/prevent noise-induced hearing loss. The observation that the store active compound ryanodine [187] and the serotonin type 3 receptor agonist 1-(*m*-chlorophenyl)-biguanid [136], are not agonists but potentiate  $\alpha 9\alpha 10$ -mediated ACh-responses, indicates that as for other nAChRs, PAMs that target  $\alpha 9\alpha 10$  can be developed. The recent successful expression of  $\alpha 9\alpha 10$  receptors in HEK cells coupled to FLIPR calcium assay [144], opens the possibility of high throughput screening for  $\alpha 9\alpha 10$  nAChR selective lead compounds. In addition, the availability of the crystal structure of the  $\alpha 9$  subunit extracellular domain [116], coupled to radioligand binding assays of compounds to purified extracellular  $\alpha 9$  protein fragments and mutagenesis structure-function analysis, will aid to decipher the orthosteric or allosteric interaction of novel  $\alpha 9\alpha 10$  nAChR ligands [188].

### 6.2. Non-auditory $\alpha 9\alpha 10$ nAChR pharmacotherapeutics

The fact that the  $\alpha 9\alpha 10$  nAChR has a restricted expression pattern, has no CNS expression [17,19,50,189] and has a differential pharmacological profile, makes this receptor a suitable pharmacotherapeutic target for the design of drugs for inner ear disorders with less side effects, specially of central origin. However, the expression of  $\alpha 9$  and  $\alpha 10$  subunits in epithelial cells outside the inner ear serving non-canonical cholinergic effects should be considered, both as potential targets of side effects when using compounds as otoprotectants and as targets for pharmacotherapeutics. Side effects of drugs used for inner ear disorders can be circumvented by local delivery using gels and nanoparticles [190,191].

The pharmacotherapeutic activity of  $\alpha 9$  and/or  $\alpha 9\alpha 10$  nAChR antagonists like conotoxins have been investigated in several types of pain in animal models [92,192–194]. The analgesic effect is most likely due to the participation of  $\alpha 9^*$  nAChR which are present in immune cells involved in inflammatory processes (for an extended review see [33]). Moreover,  $\alpha 9$  and/or  $\alpha 10$  nAChR subunit blockers have been also suggested to be of use in animal models of immune diseases such as experimental autoimmune encephalomyelitis [89,90]. In addition, pCF3-N,N-diethyl-N'-phenyl-piperazine (pCF3-diEPP), in part due to its non-canonical effect on  $\alpha 9^*$  nAChRs, has been suggested as useful in the prevention of life-threatening trauma-induced inflammation, either as an emergency treatment for accident victims or in the context of major surgery [195]. The non-competitive  $\alpha 9\alpha 10$  nAChR antagonist alphaO-Conotoxin GeXIVA and the competitive antagonist RgIA4, result in antitumor effects [196–198]. Thus, the non-auditory functions of  $\alpha 9^*$  and  $\alpha 10^*$  nAChRs open a wide variety of uses of potential drugs that target the non-canonical metabotropic signalling of these receptors.

## 7. Conclusions and forward

Much has been advanced since the first cloning of  $\alpha 9$  [17] and  $\alpha 10$  [16], the last nAChR subunits to be identified. For the auditory field, the puzzle concerning the structure of the hair cell nAChR and the decades long lasting debate between a nicotinic versus a muscarinic ACh receptor [134,199–201], was solved. This enabled a precise characterization of the pharmacology, biophysical and structure function analysis of the  $\alpha 9\alpha 10$  nAChR (for review see [9,57,58,67]) and the crystallization of the  $\alpha 9$  monomeric extracellular domain [115,116]. Moreover, the kinetics of *ex vivo* hair cells  $\alpha 9\alpha 10$  nAChR-mediated currents and the properties of efferent synaptic transmission have been extensively analyzed and results replicated by several laboratories (for review see [67]). In addition, the generation of *Chrna9* and *Chrna10* genetically modified mice [14,15,70,74,202], has helped elucidate the function of the inner ear efferent system in audition. It should be noted that it is only in hair cells where a clear ionotropic function of  $\alpha 9\alpha 10$  nAChR has been reproducibly reported. Only one report has proposed that synaptic currents in chromaffin cells are due to  $\alpha 9^*$ , in addition to  $\alpha 7^*$  and  $\alpha 3^*$  nAChRs [26]. This is based on the observation that the frequency

and amplitude of evoked post-synaptic currents are blocked by the  $\alpha 9^*$  antagonist  $\alpha$ -RgIA and on the fact that chromaffin cells express the  $\alpha 9$  subunit [26]. These results should be expanded using additional compounds in order to perform an entire characterization of the pharmacological profile of responses, in addition to experiments in *Chrna9*, *Chrna10*, *Chrna3* and *Chrna7* knockout mice, to unequivocally ascertain the molecular nature of the receptors mediating ACh ion channel responses in chromaffin cells.

From the pharmacological point of view, the cloning of the  $\alpha 9$  and  $\alpha 10$  nAChRs added "odd cousins to an old family" of nicotinic pentameric ligand channels [9]. *In vivo*, ionotropic responses require the assembly of both subunits into pentameric receptors [14,16,18] (with the exception of the zebrafish lateral line hair cell receptor which appears to be a  $\alpha 9$  homomeric nAChR [13]). No clear evidence of co-assembly with other nAChRs has been reported [17,126].  $\alpha 9$  and  $\alpha 10$  form a distinct and distant clade within the nAChR phylogeny and have a distinct evolutionary trajectory when compared to neuronal and muscle nAChR subunits [42,55,147]. Moreover, they have a distinct, mainly non-neuronal expression pattern [9,17,19,42,50,203]. In addition,  $\alpha 9$  and  $\alpha 9\alpha 10$  nAChRs have peculiar pharmacological and biophysical properties [16–18,56,133,135,136,204,205]. Therefore, the  $\alpha 9$  and  $\alpha 10$  subunits are functionally isolated from the remainder nicotinic subunits. Although within the dichotomic subdivision of nAChR subunits into muscle and neuronal  $\alpha 9$  and  $\alpha 10$  have been classically and repeatedly included within the neuronal subgroup [10], based on their evolutionary trajectory, expression pattern, biophysical and pharmacological properties, they are clearly non-neuronal. Thus, as previously proposed [9],  $\alpha 9$  and  $\alpha 10$  subunits could be re-classified as "hair cell" nAChR subunits, based on their known canonical ion channel function mediating efferent olivocochlear inhibition of inner ear hair cells.

Finally, the increasing knowledge of the canonical and non-canonical functions of  $\alpha 9^*$  and  $\alpha 10^*$  nAChRs paves the way to new pharmacotherapeutical applications and urges the discovery of new selective and safe lead molecules that target these receptors.

#### CRedit author statement

The review was completely written by Ana Belén Elgoyhen.

#### Declarations of interest

none.

#### Data Availability

No data was used for the research described in the article.

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