Themed Section: Nicotinic Acetylcholine Receptors

RESEARCH PAPER

Loops D, E and G in the Drosophila D α 1 subunit contribute to high neonicotinoid sensitivity of D α 1-chicken β 2 nicotinic acetylcholine receptor

Correspondence Kazuhiko Matsuda, Department of Applied Biological Chemistry, Faculty of Agriculture, Kindai University, 3327-204 Nakamachi, Nara 631-8505, Japan. E-mail: kmatsuda@nara.kindai.ac.jp

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Makoto Ihara¹, Mai Hikida¹, Hiroyuki Matsushita¹, Kyosuke Yamanaka¹, Yuya Kishimoto¹, Kazuki Kubo¹, Shun Watanabe¹, Mifumi Sakamoto¹, Koutaro Matsui¹, Akihiro Yamaguchi¹, Daiki Okuhara¹, Shogo Furutani¹, David B Sattelle² and Kazuhiko Matsuda¹

¹Department of Applied Biological Chemistry, Faculty of Agriculture, Kindai University, Nara, Japan, and ²Centre for Respiratory Biology, UCL Respiratory, University College London, London, UK

BACKGROUND AND PURPOSE

Neonicotinoid insecticides interact with the orthosteric site formed at subunit interfaces of insect nicotinic ACh (nACh) receptors. However, their interactions with the orthosteric sites at α -non α and α - α subunit interfaces remain poorly understood. The aim of this study was to elucidate the mechanism of neonicotinoid actions using the *Drosophila* D α 1-chicken β 2 hybrid nACh receptor.

EXPERIMENTAL APPROACH

Computer models of the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor in complex with imidacloprid and thiacloprid were generated. Amino acids in the D $\alpha 1$ subunit were mutated to corresponding amino acids in the human $\alpha 4$ subunit to examine their effects on the agonist actions of neonicotinoids on $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors expressed in *Xenopus laevis* oocytes using voltage-clamp electrophysiology.

KEY RESULTS

The $(D\alpha 1)_3(\beta 2)_2$ nACh receptor models indicated that amino acids in loops D, E and G probably determine the effects of neonicotinoids. The amino acid mutations tested had minimal effects on the EC₅₀ for ACh. However, the R57S mutation in loop G, although having minimal effect on imidacloprid's actions, reduced the affinity of thiacloprid for the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor, while scarcely affecting thiacloprid's action on the $(D\alpha 1)_2(\beta 2)_3$ nACh receptor. Both the K140T and the combined R57S;K140T mutations reduced neonicotinoid efficacy but only for the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor. Combining the E78K mutation with the R57S;K140T mutations resulted in a selective reduction of thiacloprid's affinity for the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor.

CONCLUSIONS AND IMPLICATIONS

These findings suggest that a triangle of residues from loops D, E and G contribute to the selective actions of neonicotinoids on insect-vertebrate hybrid nACh receptors.

LINKED ARTICLES

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Abbreviations

 $I_{\mbox{\scriptsize max}}$, normalized maximum response; nACh receptor, nicotinic ACh receptor



Introduction

Neonicotinoid insecticides such as imidacloprid and thiacloprid (Figure 1) have been used to control crop pests and pests of farm animals (Matsuda et al., 2001, 2005, 2009; Tomizawa and Casida, 2003, 2005; Jeschke et al., 2013; Casida and Durkin, 2017). Neonicotinoids interact with the orthosteric site of insect nicotinic ACh receptors (nACh receptors) mainly as partial and full agonists, although for certain compounds, super agonist and antagonist actions have also been described (Ihara et al., 2004, 2006; Brown et al., 2006; Tan et al., 2007; Matsuda et al., 2009). Unlike ACh, neonicotinoids are not hydrolyzed by acetycholinesterase and hence can persistently modulate nACh receptors (Matsuda et al., 2001). Neonicotinoids show selectivity for insect over vertebrate nACh receptors (Matsuda et al., 2001); they are broad spectrum insecticides taken up by roots and subsequently translocated into all plant tissues (Kagabu, 1997; Jeschke et al., 2013). Neonicotinoids currently make up >25% of world insecticide sales (Jeschke et al., 2013).

The reduced numbers of honey bees, bumble bees and other insect pollinators are a threat to the effective pollination of crop plants. Neonicotinoids have been suggested as one possible contributor to Colony Collapse Disorder, because they modulate the nACh receptors of bees as well as pest insect species (Gill *et al.*, 2012; Whitehorn *et al.*, 2012; Rundlof *et al.*, 2015). In view of the possible risk to bees, the EU restricted the use of three neonicotinoids (imidacloprid, clothianidin and thiamethoxam) in 2013 and continues to restrict their deployment subject to further assessment of the risk. Therefore, it is extremely important to understand the structural features that determine the selectivity and potency of neonicotinoids at nACh receptors.

The nACh receptors are membrane-spanning proteins containing an integral cation channel and play a crucial role in fast cholinergic neurotransmission in vertebrates and invertebrates (Changeux, 2012). Channel opening in response to ACh binding results in depolarization of nerve and muscle membranes (Miyazawa et al., 2003; Taly et al., 2005, 2006; Unwin, 2005; Taly, 2007; Unwin and Fujiyoshi, 2012; Nemecz et al., 2016). Although some nACh receptors are homomers or heteromers of α subunits (defined by a YXCC motif in loop C of the ACh binding site), most are heteromers of α and non- α subunits (Changeux, 2012). The N-terminal extracellular six loops (typically A, B, C from the α subunit and D, E, F from the non- α subunit) form the orthosteric site of such α /non- α heteromers (Corringer et al., 2000) to which ACh and neonicotinoids bind (Matsuda et al., 2005). Based on computational studies, we proposed that negatively charged nitro or cyano groups of neonicotinoids (Figure 1) interact with basic residues that are selectively present in insect nACh receptors (Matsuda et al., 2001, 2005, 2009). The resulting planar imidazolidine or related moieties facilitate interaction with π -electron-rich aromatic residues (Matsuda *et al.*, 2009). Mutations of Gln^{79} in chicken $\alpha 7$ (Shimomura *et al.*, 2002) and Thr^{77} in chicken $\beta 2$ subunit (Shimomura *et al.*, 2006), both of which are located in loop D. to basic residues. markedly enhanced imidacloprid sensitivity of nACh receptors containing these subunits, suggesting a possible interaction of the nitro group of imidacloprid with the added basic residues.

To demonstrate an interaction of loop D with neonicotinoids, we co-crystallized wild-type and the Q55R



Figure 1

Homology models of the ligand binding domain of fruit fly $D\alpha 1$ /chicken $\beta 2$ nACh receptor in complex with imidacloprid and thiacloprid. (A) Overall top view of the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor model generated from human $(\alpha 4)_2(\beta 2)_3$ nACh receptor docked with imidacloprid. The $D\alpha 1$ subunits at principal and complementary sides are coloured tan and yellow, respectively, whereas the $\beta 2$ subunits are coloured red. (B) Chemical structure of imidacloprid and close-up view of the imidacloprid binding site. (C) Chemical structure of thiacloprid and close-up view of the thiacloprid binding site. (C) Chemical structure of thiacloprid and close-up view of the thiacloprid binding site. Arg⁵⁷ and Lys¹⁴⁰ interacted electrostatically with the nitro group of imidacloprid (B) and the cyano group of thiacloprid (C). Glu⁷⁸ made salt bridges with Arg⁵⁷ and Lys¹⁴⁰ to form a 'loop D-E-G triangle' (B, C). In each panel, the main chains of the nACh receptors are drawn as cartoon, whereas Arg⁵⁷ (coloured orange), Lys¹⁴⁰ (coloured orange), Glu⁷⁸ (coloured blue) and the neonicotinoids are drawn as space filling models. For neonicotinoids, carbon-, nitrogen-, oxygen-, sulfur- and chlorine-atoms are coloured grey, blue, red, tan and green, respectively.

mutant of the ACh binding protein (AChBP) of *Lymnaea stagnalis*, which provides a surrogate of the nACh receptor orthosteric binding site, with neonicotinoids (imidacloprid, clothianidin, thiacloprid and the nitromethylene analogue of imidacloprid [CH-IMI]) (Ihara *et al.*, 2008, 2014a). The X-ray crystal structures revealed that basic residues in loop D interacted electrostatically with the nitro or cyano group of neonicotinoids. Also, we showed that desnitro-imidacloprid, an imidacloprid metabolite lacking the nitro group, bound to the AChBP, placing its guanidine tip in the opposite direction against loop D, supporting a role for loop D in the interactions of nACh receptors with neonicotinoids (Ihara *et al.*, 2014a).

We found that T77R;E79V mutations in loop D in the β 2 subunit of the avian $\alpha 4\beta 2$ nACh receptor resulted in enhanced neonicotinoid sensitivity, whereas they had no significant effect on the concentration–response curve for ACh, and hence, we predicted that inverse mutations in loop D in insects would lead to resistance (Shimomura *et al.*, 2006). In fact, an R81T mutation was later found in the β 1 subunit in a neonicotinoid-resistant field population of aphid *Myzus persicae*, supporting our proposal that neonicotinoids interact strongly with loop D (Bass *et al.*, 2011).

In the structural work, we unexpectedly discovered that CH-IMI, clothianidin and thiacloprid interacted with Lvs³⁴ on the β 1 strand (loop G) of the AChBP. Interestingly, insect nACh receptor α subunits possess basic residues in loop G, and thus, such residues could also underlie the potency and insect selectivity of neonicotinoids. Indeed, mutations of Ser⁵⁸ in the avian α 7 nACh receptor, which corresponds to Lys³⁴ in the Lymnaea AChBP, led to enhanced agonist actions of neonicotinoids, while reducing the actions of ACh, (-)nicotine and desnitro-imidacloprid (Ihara et al., 2014a). Since loop G is located in the complementary side of α subunits, we predicted that the α - α subunit interface may also contribute to interactions with neonicotinoids (Ihara et al., 2015). However, no evidence for this hypothesis has been provided using heteromers. In a relevant study, we showed that insect nACh receptor α subunits possess structural features that are favourable for interactions with neonicotinoids and identified loop C and the region upstream of loop B as potential contributors (Shimomura et al., 2005). However, it is unclear as to precisely which amino acids in the region upstream of loop B underpin the selective interactions with neonicotinoids.

To address these questions, we computationally modelled a full length fruit fly (Drosophila melanogaster) Da1-chicken (Gallus gallus) B2 nACh receptor in complex with imidacloprid, since this insect-vertebrate hybrid nACh receptor exhibits a higher neonicotinoid sensitivity than the chicken $\alpha 4\beta 2$ nACh receptor (Ihara *et al.*, 2003, 2014b). The model showed that not only Arg⁵⁷ in loop G, but also Lys¹⁴⁰ in loop E at the D α 1-D α 1 subunit interface interact with the nitro group of imidacloprid and Glu⁷⁸ supports this interaction by forming salt bridges. Thus, we mutated Arg⁵⁷, Lys¹⁴⁰ and Glu⁷⁸ in the *Drosophila* D α 1 subunit, as a representative of insect nACh receptor α subunits, to corresponding amino acids in the human $\alpha 4$ subunit, as a representative of mammalian nACh receptor a subunits, and investigated the effects of these mutations on the agonist actions of imidacloprid and thiacloprid on $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$

nACh receptors. By this means, we sought to address whether the amino acids in the $D\alpha 1-D\alpha 1$ subunit interface contribute to the selectivity of neonicotinoids for insect nACh receptors.

Methods

Xenopus laevis oocytes

Female Xenopus laevis were anaesthetized with benzocaine (ethyl 4-aminobenzoate) to reduce animal suffering as much as possible according to the UK Animals (Scientific Procedures) Act, 1986, and minimum amounts of oocytes were removed from anaesthetized frogs. Xenopus oocvtes were treated for 30 min with 2 mg·mL⁻¹ Type IA collagenase (Sigma Aldrich, St. Louis, MO, USA) in Ca²⁺-free standard oocyte saline Ca-free SOS: 100 mM NaCl, 2 mM KCl, 1 mM MgCl₂ and 5 mM HEPES; (pH 7.6 adjusted with NaOH) and transferred to SOS (100 mM NaCl, 2 mM KCl, 1.8 mM CaCl₂, 1 mM MgCl₂ and 5 mM HEPES; pH 7.6) (Ihara et al., 2003; Shimomura et al., 2006). The follicle cell layer was removed from oocytes manually with fine forceps, and defolliculated oocytes were injected with cRNAs. Animal studies are reported in compliance with the ARRIVE guidelines (Kilkenny et al., 2010; McGrath and Lilley, 2015).

cRNA preparation and injection into oocytes

The *Drosophila* $D\alpha 1$ subunit and the chicken $\beta 2$ subunit containing amino acid sequences in the Refseq database (Dα1: NP_524481; β2: NP_990144) were used as the wild-type subunits. Each cDNA was cloned into the pcDNA3.1 (+) vector (Thermo Fisher Scientific, Waltham, MA, USA). The nucleotide sequence of the Da1 subunit sequence was mutated by PCR. The cRNAs of wild-type and mutant Da1 subunits as well as of a wild-type chicken $\beta 2$ subunit were prepared by in vitro transcription from each respective cDNA, cloned in the pcDNA 3.1 (+) vector using the mMESSAGE mMACHINE T7 Ultra kit (Thermo Fisher Scientific) (Furutani et al., 2014). cRNAs were dissolved in RNase-free water at a concentration of 1 mg·mL⁻¹. Da1 and ß2 cRNA solutions were mixed at a ratio of 5:1 or 1:5 for reconstituting $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors, respectively, in Xenopus oocytes. To each oocyte, 50 nL of cRNA mixture solutions were injected, and the injected oocytes were incubated for 4-5 days prior to electrophysiology in SOS supplemented with penicillin (100 units ml^{-1}), streptomycin (100 μ g ml⁻¹), gentamycin (20 μ g ml⁻¹) and 2.5 mM sodium pyruvate (pH 7.6).

Electrophysiology

Two-electrode voltage-clamp electrophysiology was conducted using a GeneClamp 500B amplifier (Molecular Devices, Sunnyvale, CA, USA) at a holding potential of -100 mV (Matsuda *et al.*, 1998; Ihara *et al.*, 2003; Shimomura *et al.*, 2006). Oocytes were perfused extracellularly at a flow rate of 7–10 mL·min⁻¹ with SOS containing 0.5 μ M **atropine** to suppress the muscarinic receptor-mediated responses of oocytes. Neonicotinoids were dissolved in DMSO at 100 mM and diluted with SOS for preparing test solutions. DMSO concentrations in each test solution were 0.1% or



lower at which DMSO had no apparent effect on the nACh receptor response to ACh and neonicotinoids. ACh was directly dissolved in SOS immediately before the experiments. After successive applications of 10 μ M ACh to confirm reproducibility of the response, neonicotinoids were applied for 5 s, from lower to higher concentrations, at 3 min intervals. No irreversible nACh receptor desensitization was observed with treatments of 10 μ M or lower concentrations of ACh and 100 μ M or lower concentrations of neonicotinoids using this protocol.

Chemicals

Imidacloprid and thiacloprid were donated by Bayer Crop-Science. ACh was purchased from Sigma-Aldrich (St. Louis, MO, USA) and used without further purification.

Homology modelling of the fruit fly-chicken hybrid nACh receptors in complex with neonicotinoids

Homology modeling of $(D\alpha 1)_3(\beta 2)_2$ and docking with neonicotinoids were performed using Modeller version 9.17 (Webb and Sali, 2014). Amino acid sequences of the ligand binding domain of the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor were aligned to those of the human $(\alpha 4)_2(\beta 2)_3$ nACh receptor taken from the PDB file of 5KXI (Morales-Perez et al., 2016). Then, structural coordinates for corresponding amino acid regions were excised to make a structural template for the homology model. To generate an initial docking model, structural coordinates for imidacloprid (PDB: 2ZJU) and thiacloprid (3WTK) were manually transferred to that of 5KXI by using PyMOL (Schrödinger, New York, NY, USA), where only one neonicotinoid molecule was placed at the $D\alpha$ 1- $D\alpha$ 1 interface. The model was then refined by molecular dynamics combined with simulated annealing (Kirkpatrick et al., 1983).

Electrophysiology data analysis

The data and statistical analysis comply with the recommendations on experimental design and analysis in pharmacology (Curtis et al., 2015). Peak current amplitude of each inward current recorded from oocytes in response to bathapplied neonicotinoid and ACh was measured repeatedly. The peak amplitude of the response of oocytes expressing wild-type and mutant Da1B2 nACh receptors fluctuated considerably from several thousand nA to greater than 10 μ A and was therefore normalized to that induced by $10\ \mu M$ ACh. The concentration-normalized response relationships (ACh, n = 5; imidacloprid and thiacloprid, n = 6) were analysed by nonlinear regression using Prism 5 (GraphPad Software, La Jolla, CA, USA) to determine an EC₅₀ (M) and normalized maximum response (I_{max}) using oocytes from at least two female frogs according to the following equation:

 $Y = I_{max} \left(1 + 10^{\left(logEC_{50} - X\right) nH}\right)^{-1}$, where Y is normalized

response, X is log[agonist concentration (M)] and $n_{\rm H}$ is the Hill coefficient. The peak current amplitude of the response of the nACh receptors to 10 μ M ACh is presented as mean \pm SEM of 17 experiments. Differences between the values obtained for the wild-type and mutant nACh receptors were analysed by one-way ANOVA (Dunnett's test, P < 0.05).

Nomenclature of targets and ligands

Key protein targets and ligands in this article are hyperlinked to corresponding entries in http://www.guidetopharmacology.org, the common portal for data from the IUPHAR/ BPS Guide to PHARMACOLOGY (Southan *et al.*, 2016), and are permanently archived in the Concise Guide to PHARMA-COLOGY 2015/16 (Alexander *et al.*, 2015).

Results

Homology modelling of $(D\alpha 1)_3(\beta 2)_2$ nACh receptors in complex with neonicotinoids

To investigate a role for the D α 1-D α 1 orthosteric site of D α 1 β 2 hybrid nACh receptors in its interactions with neonicotinoids, the wild-type (D α 1)₃(β 2)₂ nACh receptors with imidacloprid and thiacloprid bound to them were modelled using the crystal structure of the human (α 4)₂(β 2)₃ nACh receptor (Figure 1A–C). In these nACh receptor neonicotinoid complexes, the two oxygens in the nitro group of imidacloprid formed hydrogen bonds with Arg⁵⁷ in loop G and Lys¹⁴⁰ in loop E (Figure 1B), whereas the cyano group of thiacloprid interacted mainly with Arg⁵⁷ (Figure 1C). In addition, Glu⁷⁸ in loop D was found to form salt bridges with Arg⁵⁷ and Lys¹⁴⁰, making a 'loop D-E-G triangle' (Figure 1B, C).

Effects of mutations on agonist actions of ACh and neonicotinoids on $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors

To explore the nACh receptor-neonicotinoid interactions observed in the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor models (Figure 1A, B), Arg⁵⁷ in loop G, Lys¹⁴⁰ in loop E and Glu⁷⁸ in loop D were mutated to serine, threonine and lysine, respectively, which are corresponding amino acids in the human $\alpha 4$ subunit (Figure 2), and the agonist actions of ACh, imidacloprid and thiacloprid on the wild-type and mutant $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors expressed in *Xenopus* oocytes were measured by voltage-clamp electrophysiology.

ACh showed agonist actions on the mutant as well as the wild-type $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors tested (Figure 3). For the wild-type nACh receptors, ACh showed slightly higher affinity, as measured by the EC₅₀ value, for the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor than the $(D\alpha 1)_2(\beta 2)_3$ nACh receptor (Table 1). The R57S, K140T, R57S;K140T and R57S; E78K;K140T mutations in the D α 1 subunit had a minimal effect on the EC₅₀ values of ACh, not only for the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor (Figure 4A) but also for the $(D\alpha 1)_2(\beta 2)_3$ nACh receptor (Figure 4B, Table 1). However, the E78K mutation slightly increased the ACh EC₅₀ value of the $(D\alpha 1)_2(\beta 2)_3$ nACh receptor (Figure 4B, Table 1).

When the peak current amplitude of the response to ACh was compared between the wild-type and mutant nACh receptors, the E78K and R57S;K140T mutations significantly reduced the peak current amplitude of the response to 10 μ M ACh of the (D α 1)₃(β 2)₂ and (D α 1)₂(β 2)₃ nACh receptors (Figures 3 and 4C, D, Table 1). In addition, the R57S and K140T mutations also reduced the peak current amplitude of the response to 10 μ M ACh in the (D α 1)₂(β 2)₃ nACh receptor.

MGSVLFAAVFIALHFATGGÜANPDAKRLYDDLLSNYNRL ÏRPVGNNSDR TVKMGLRLSQLIDVNLKNQ MTTNVWVEQË Signal peptide

WNDYKLKWNPDDYGGVDTLHVPSEHIWLPDIVLYNNADGNYEVTIMTKAILHHTGKVVWKPPAIYKSFCEIDVEYFPFDE

QTCFMKFGSWTYDGYMVDLRHLKQTADSDNIEVGIDLQDYYISVEWDIMRVPAVRNEKFYSCCEEPYLDIVFNLTLRRKT

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	Insect nAChR subunits		L	oot	G					-	.00	טנ				-	oop	E	
	Bombyx mori α1 (NP_001103388)	(53)	G	L	R	L	S	(74)	w	v	Е	Q	Е	(136)	v	w	Κ	Ρ	Ρ
	Drosophila melanogaster α1 (NP_524481)	(55)	G	L	R	L	S	(76)	w	۷	Е	Q	Е	(138)	۷	W	Κ	Ρ	Ρ
	Myzus persicae a1 (CAA57476)	(54)	G	L	R	L	S	(75)	w	L	Е	н	Е	(137)	М	w	Т	Ρ	Ρ
	Bombyx mori α2 (NP_001103397)	(52)	G	L	R	L	S	(73)	w	L	Е	н	Е	(135)	L	w	т	Ρ	Ρ
	Drosophila melanogaster α2 (NP_524482)	(75)	G	L	R	L	S	(96)	w	L	Е	н	Е	(158)	v	w	т	Ρ	Ρ
	Myzus persicae α2 (CAA57477)	(59)	G	L	Κ	L	S	(80)	w	۷	Е	Q	Е	(142)	v	w	Т	Ρ	Ρ
	Bombyx mori β1 (NP_001166819)	(66)	G	L	Α	F	۷	(87)	w	L	R	L	۷	(149)	L	w	۷	Ρ	Ρ
	Drosophila melanogaster β1 (NP_523927)	(58)	G	L	Α	F	۷	(79)	w	L	R	L	۷	(141)	L	w	۷	Ρ	Ρ
	Myzus persicae β1 (CAB87995)	(58)	G	L	Α	F	۷	(79)	w	L	R	L	۷	(141)	L	w	Т	Ρ	Ρ
	Drosophila melanogaster β2 (NP_524483)	(60)	G	L	К	L	S	(81)	w	v	К	Q	R	(143)	F	W	Е	Ρ	Ρ
	Vertebrate nAChR subunits																		
	Gallus gallus α3 (NP_989747)	(56)	Е	v	S	М	S	(77)	w	L	Κ	н	1	(139)	т	w	Т	Ρ	Ρ
	Homo sapiens α3 (NP_000734)	(65)	Е	v	S	М	S	(86)	w	L	Κ	Q	Т	(148)	т	w	Т	Ρ	Ρ
	Gallus gallus α4 (NP_990145)	(62)	G	L	S	Т	Α	(83)	w	۷	Κ	Q	Е	(145)	κ	w	М	Ρ	Ρ
	Homo sapiens α4 (NP_000735)	(67)	G	L	S	Т	Α	(88)	w	۷	Κ	Q	Е	(150)	Q	w	Т	Ρ	Ρ
	Gallus gallus α7 (NP_989512)	(56)	Т	L	S	L	М	(77)	w	L	Q	М	Υ	(139)	Q	Υ	L	Ρ	Ρ
	Homo sapiens α7 (NP_000737)	(56)	S	L	S	L	L	(77)	w	L	Q	М	S	(139)	Q	Υ	L	Ρ	Ρ
	Gallus gallus β2 (NP_990144)	(54)	М	v	S	L	Α	(75)	w	L	т	Q	Е	(137)	F	w	L	Ρ	Ρ
	Homo sapiens β2 (NP_000739)	(61)	М	v	S	L	Α	(82)	w	L	Т	Q	Е	(144)	F	w	L	Ρ	Ρ
	Gallus gallus β3 (NP_990143)	(56)	G	L	Κ	Т	S	(77)	w	L	Κ	Q	Е	(139)	т	w	М	Ρ	Ρ
	Homo sapiens β3 (NP_000740)	(59)	G	L	κ	I	S	(80)	w	L	Κ	Q	Е	(142)	v	w	т	Ρ	Ρ

Figure 2

A

Amino acid sequences of nACh receptors. (A) Amino acid sequences of the ligand binding domain of *Drosophila melanogaster* α 1 (D α 1) subunit (accession number: NP_524481). Mutated amino acids in loops D, E and G region are coloured cyan. (B) Multiple sequence alignments of loops D, E and G regions of insect, chicken and human nACh receptor subunits. Numbers in parentheses indicate the amino acid residue numbers taken from the sequence database. Basic- and acidic-residues are in blue and red boxes, respectively. Serine and threonine residues are coloured orange. Amino acid sequences of the *Drosophila* D α 1 and human α 4 subunits are boxed.

Effects of mutations in loops G and E of the $D\alpha 1$ subunit on agonist actions of neonicotinoids on $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors

Imidacloprid and thiacloprid were agonists (Figure 5) with a lower efficacy than ACh on the wild-type $D\alpha 1\beta 2$ nACh receptor (Table 2). The R57S mutation in the $D\alpha 1$ subunit had a minimal impact on the EC_{50} and I_{max} of imidacloprid (Figure 6), while reducing the p EC_{50} value of thiacloprid from 7.47 ($EC_{50} = 33.8$ nM) to 6.53 ($EC_{50} = 295$ nM) in the ($D\alpha 1$)₃($\beta 2$)₂ nACh receptor (Figure 7, Table 2). However, no such effect of the R57S mutation on the EC_{50} and I_{max} values of thiacloprid or imidacloprid was observed in the ($D\alpha 1$)₂($\beta 2$)₃ nACh receptor (Figures 6 and 7, Table 2).

In contrast to the R57S mutation, the K140T mutation in the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor reduced the efficacy of the neonicotinoids with a minimal shift of EC₅₀ (Table 2), having a greater effect on the actions of imidacloprid than those of thiacloprid (Figures 6 and 7). A more profound effect of

this mutation on the efficacy of imidacloprid was observed for the $(D\alpha 1)_3(\beta 2)_2$ compared to the $(D\alpha 1)_2(\beta 2)_3$ nACh receptors (Figure 6, Table 2). Also, similar effects on the agonist actions of neonicotinoids were observed in the R57S; K140T mutations with a higher impact on the I_{max} than the EC₅₀ (Figures 6 and 7, Table 2).

Effects of the E78K mutation in loop D of the $D\alpha 1$ subunit on agonist actions of neonicotinoids on $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors

In the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor model (Figure 1), Glu⁷⁸ forms salt bridges with Arg⁵⁷ and Lys¹⁴⁰. We examined the role of this interaction on the effects of the neonicotinoids; the E78K mutation was predicted to lead to an electrostatic repulsion by Arg⁵⁷ and Lys¹⁴⁰, resulting in an adverse effect on the orthosteric site. Indeed, ACh induced a much smaller amplitude current response than that seen in the wild-type nACh receptor (Figure 4C, D). Furthermore, imidacloprid



Figure 3

Currents recorded in response to ACh from wild-type and mutant $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors expressed in *X. laevis* oocytes.

BIP



Agonist actions of ACh on wild-type and mutant $D\alpha 1\beta 2$ nicotinic ACh receptors expressed in Xenopus oocytes^a

	(Dα1) ₃ (β2) ₂		(Dα1) ₂ (β2) ₃	
nACh receptors	pEC ₅₀	I _{max}	pEC ₅₀	I _{max}
Wild type	7.05 ± 0.06	0.981 ± 0.031	6.74 ± 0.05	1.026 ± 0.031
R57S	6.97 ± 0.06	0.985 ± 0.031	6.67 ± 0.05	1.035 ± 0.031
K140T	6.89 ± 0.05	0.978 ± 0.026	6.66 ± 0.04	1.009 ± 0.024
R57S;K140T	6.90 ± 0.05	1.006 ± 0.030	6.81 ± 0.07	1.071 ± 0.038
E78K	7.04 ± 0.04	0.979 ± 0.021	7.02 ± 0.05*	1.036 ± 0.023
R57S;E78K;K140T	7.00 ± 0.04	0.977 ± 0.024	6.83 ± 0.08	0.995 ± 0.043

^aData are presented as mean \pm SEM of repeated experiments (n = 5).

*The pEC₅₀ (-log EC₅₀) value differed significantly from that determined in the wild type (P < 0.05).



Figure 4

Concentration–response curves for ACh observed in $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ hybrid nACh receptors expressed in *X. laevis* oocytes and peak current amplitude of the ACh-induced responses. (A) Concentration–response curve for the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor; (B) concentration–response curve for the $(D\alpha 1)_2(\beta 2)_3$ nACh receptor. (C) Peak current amplitude of the response of $(D\alpha 1)_3(\beta 2)_2$ nACh receptor to 10 μ M ACh. (D) Peak current amplitude of the response of $(D\alpha 1)_2(\beta 2)_3$ nACh receptor to 10 μ M ACh. Each data point in panels (A, B) represents mean ± SEM (n = 5), whereas each bar graph in panels (C, D) represents mean ± SEM (n = 17). In panels (C) and (D), significant differences of the peak current amplitude of the response induced by 10 μ M ACh between the wild-type and mutant nACh receptors are indicated by an asterisks (* P < 0.05).

and thiacloprid were inefficient in activating the E78K mutant of $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors even at 100 μ M (Figure 5), hence the EC₅₀ could not be determined.

However, when the E78K mutation was combined with the R57S;K140T double mutations, the ability of the hybrid nACh receptor to respond to ACh (Figures 3 and 4) and

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Figure 5

Currents recorded in response to imidacloprid (IMI) and thiacloprid (THI) from wild-type and mutant $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors expressed in *X. laevis* oocytes.

neonicotinoids was restored (Figures 5–7). The triple mutations significantly reduced the pEC₅₀ value of thiacloprid from 7.47 to 6.37 (EC₅₀ = 424 nM) and the I_{max} value of imidacloprid from 0.165 to 0.119 for the $(D\alpha 1)_3(\beta 2)_2$ nACh

receptor (Figures 6 and 7, Table 2). In contrast, they had a minimal effect on the pEC_{50} and I_{max} values of the neonicotinoids for the $(D\alpha 1)_2(\beta 2)_3$ nACh receptor (Figures 6 and 7, Table 2).

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	(D α 1) ₃ (β 2) ₂				(Dα1) ₂ (β2) ₃			
	Imidacloprid		Thiacloprid		Imidacloprid		Thiacloprid	
nACh receptors	pEC ₅₀	lmax	pEC ₅₀	lmax	pEC ₅₀	lmax	pEC ₅₀	lmax
Wild type	7.20 ± 0.14	0.165 ± 0.009	7.47 ± 0.21	0.054 ± 0.005	7.24 ± 0.14	0.089 ± 0.005	7.70 ± 0.39	0.039 ± 0.003
R57S	6.99 ± 0.22	0.185 ± 0.015	$6.53 \pm 0.28^*$	0.064 ± 0.007	7.32 ± 0.15	$0.058 \pm 0.003^{*}$	7.53 ± 0.27	0.038 ± 0.004
K140T	7.29 ± 0.17	$0.032 \pm 0.002^*$	7.60 ± 0.26	$0.018 \pm 0.002^{*}$	7.05 ± 0.20	$0.048 \pm 0.003^{*}$	7.92 ± 0.15	$0.019 \pm 0.001^*$
R57S;K140T	7.08 ± 0.17	$0.075 \pm 0.004^*$	7.71 ± 0.14	$0.017 \pm 0.001^*$	7.07 ± 0.17	$0.115 \pm 0.006^{*}$	7.64 ± 0.21	0.045 ± 0.004
E78K	۹DN	QN	ND	DN	QN	ND	QN	ND
R57S;E78K;K140T	6.71 ± 0.32	$0.119 \pm 0.014^{*}$	$6.37 \pm 0.29^*$	0.043 ± 0.005	6.92 ± 0.25	0.083 ± 0.008	7.36 ± 0.17	0.031 ± 0.002
^a Data are presented as	mean ± SEM of rep€	sated experiments ($n = 6$	6).					

^bND: could not be determined because no neonicotinoid-induced currents were observed in Xenopus oocytes expressing the E78K mutant nACh receptor *The pEC₅₀ (-log EC₅₀) and I_{max} values differed significantly from those determined in the wild type (P < 0.05) BIP

Discussion

In this study, we showed for the first time that mutations of Arg⁵⁷ in loop G, Lys¹⁴⁰ in loop E and Glu⁷⁸ in loop D in the complementary (-) side of the D α 1 subunit, which are all located in the region upstream of loop B, are involved in determining the actions of neonicotinoids on the $D\alpha 1\beta 2$ hybrid nACh receptors heterologously expressed in X. laevis oocytes. The stoichiometry of the $D\alpha 1$ and $\beta 2$ subunits as well as the mutations in the Da1 subunit examined had minimal effects on the EC₅₀ value of ACh, pointing to selective interactions of ACh with the $D\alpha 1$ - $\beta 2$ orthosteric site rather than the $D\alpha 1$ -Da1 orthosteric site. The result is attributable to an electrostatic repulsion between ACh, which contains a quaternary ammonium, and the basic residues Arg⁵⁷ and Lys¹⁴⁰ that provide a positive charge at the $D\alpha 1$ - $D\alpha 1$ site.

Although the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor model (Figure 1A) indicates the proximity of the Arg⁵⁷ to the bound imidacloprid, the effect of the R57S mutation on the EC₅₀ was small. This does not contradict the model because the serine residue can form a hydrogen bond with the nitro group oxygen, thus permitting the access of imidacloprid. In contrast, the R57S mutation significantly shifted the EC50 of thiacloprid to a higher concentration (Table 2). An interpretation of this result is that the cyano group of thiacloprid more selectively interacts with the Arg⁵⁷ than the nitro group of imidacloprid, as shown by the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor model (Figure 1B, C). This provides one explanation for the R57S mutation having a more profound effect on the agonist action of thiacloprid than that of imidacloprid. Alternatively, the ability of the cyano group to form a hydrogen bond with the added serine residue is weaker than that of the nitro group, and thus, the R57S mutation may selectively reduce the action of thiacloprid.

We previously showed that the L118K mutation in loop E markedly enhanced the efficacy of imidaloprid, while reducing that of ACh for the chicken α7 nACh receptor. Therefore, we predicted that the lysine corresponding to Leu¹¹⁸ in insect nACh receptor α subunits (Amiri *et al.*, 2008) may partly account for the selective action of neonicotinoids. Indeed, the reduction in efficacy of imidacloprid and thiacloprid resulting from the K140T mutation was more profound in the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor than in the $(D\alpha 1)_2(\beta 2)_3$ nACh receptor (Figures 6 and 7, Table 2), supporting an interaction with the $D\alpha$ 1- $D\alpha$ 1 interface. If efficacy represents interaction with the activated state of the nACh receptor, the results suggest that Lys¹⁴⁰ interacts more strongly with imidacloprid than with thiacloprid in the activated state of the hybrid nACh receptor, as indicated by the homology models (Figure 1).

The nACh receptor model (Figure 1) predicts that the E78K mutation leads to repulsion in the triangle of Lys⁷⁸, Arg⁵⁷ and Lys¹⁴⁰, adversely altering the function of the nACh receptor. Indeed, the E78K mutation markedly reduced the capacity of the nACh receptor to respond to agonists (Figures 3-5, Tables 1 and 2), validating the model. The model also suggests that combining the E78K mutation with the R57S;K140T mutations will have a much smaller effect on the function of the nACh receptor compared with the single E78K mutation, since no electrostatic repulsion occurs within the triangle; this accords with the minimal impact these mutations then have on the agonist action of ACh in terms of affinity as well as efficacy (Figures 3 and 4, Table 1).





Figure 6

Concentration–response relationships for imidacloprid observed in wild-type and mutant $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors expressed in *X. laevis* oocytes. Each data point represents the mean ± SEM (n = 6).



Figure 7

Concentration–response relationships for thiacloprid observed in wild-type and mutant $(D\alpha 1)_3(\beta 2)_2$ and $(D\alpha 1)_2(\beta 2)_3$ nACh receptors expressed in *X. laevis* oocytes. Each data point represents the mean \pm SEM (n = 6).

The increased effect of the triple mutations in the Da1 subunit on the pEC₅₀ value of thiacloprid when compared with imidacloprid for the $(Da1)_3(\beta 2)_2$ nACh receptor (Figures 6 and 7, Table 2) may arise from lower capacity of the cyano group compared with the nitro group to form hydrogen bonds with the added serine and theronine residues.

We previously showed that the region upstream of loop B in the *Drosophila* $D\alpha 2$ subunit underlies the high neonicotinoid

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sensitivity of the D α 2 β 2 hybrid nACh receptor (Shimomura *et al.*, 2005). In the D α 1 subunit, the amino acids in the loop D-E-G triangle located upstream of loop B accounts, at least in part, for the higher sensitivity of the insect-avian hybrid nACh receptors to neonicotinoids than the avian α 4 β 2 nACh receptor (Matsuda *et al.*, 1998; Ihara *et al.*, 2003).

The R57S mutation significantly reduced the affinity (the pEC₅₀ value) of thiacloprid for the $(D\alpha 1)_3(\beta 2)_2$ nACh receptor (Figure 7, Table 2). Hence, it is predicted that mutations of corresponding basic residues in nACh receptors of pest insect species may result in resistance to this neonicotinoid as it had a minimal effect on the ACh concentration–response curve. However, we did not employ expressed nACh receptors composed only of subunits from insects due to the difficulty of robust functional nACh receptor expression, not only in *Xenopus* oocytes, but also in various cell lines (Lansdell *et al.*, 1997). A solution to this problem, exploring resistant field population for this mutation in pests controlled solely with neonicotinoids, is urgently needed.

It has been shown that the use of nACh receptor concatemers allows nACh receptor-ligand interactions to be analysed at specified orthosteric sites (Carbone et al., 2009; Mazzaferro et al., 2011, 2014; Benallegue et al., 2013). In this study, we did not define the sequence of the $D\alpha 1$ and $\beta 2$ subunit in the hybrid nACh receptors expressed in oocytes using a concatemer and, therefore, we cannot demonstrate unequivocally that the reduced neonicotinoid sensitivity resulting from the mutations tested can be attributed to the interactions with the $D\alpha$ 1- $D\alpha$ 1 interface. It will be necessary in the future to employ insect nACh receptor concatamers to study the mode of action of neonicotinoids. Nevertheless, it is reasonable to conclude that the structural changes in the complementary side of the Da1 subunit selectively affect neonicotinoid interactions with the hybrid nACh receptors and thus are likely to play an important role in determining the selective neonicotinoid actions on insect nACh receptors.

In summary, we have shown for the first time that triple rather than single mutations of amino acids in the loop D-E-G triangle of the *Drosophila* Da1 subunit to corresponding amino acids in the human a4 subunit significantly and selectively reduce the agonist actions of imidaloprid and thiacloprid on the $(Da1)_3(\beta 2)_2$ nACh receptor. The results add to our understanding of the mechanism of selectivity of neonicotinoids and provide new insights into the interactions of neonicotinoids with hybrid nACh receptors.

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Author contributions

M.I., M.H., H.M., K.Y., Y.K., K.K., S.W., M.S., K.M. (Matsui and Matsuda), A.Y., D.O. and S.F. conducted the

experiments; M.I., D.B.S. and K.M. designed the experiments and wrote the manuscript.

Conflict of interest

The authors declare no conflicts of interest.

Declaration of transparency and scientific rigour

This Declaration acknowledges that this paper adheres to the principles for transparent reporting and scientific rigour of preclinical research recommended by funding agencies, publishers and other organisations engaged with supporting research.

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