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Genetic Matters: Thirty years of progress using mouse models in nicotinic research

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Abstract

This Research Update summarizes thirty years of studies on genetic influences on responses to the acute or chronic administration of nicotine. Early studies established that various inbred mice are differentially sensitive to the effects of the drug. Classical genetic analyses confirmed that nicotine effects on locomotion, body temperature and seizures are heritable. A significant inverse correlation between the locomotor and hypothermic effects and the density of nicotine binding sites suggested that differential expression α4β2-neuronal nicotinic acetylcholine receptor (nAChR) mediated some of this genetic variability. Subsequent studies with α 4 and β 2 nAChR null (decreased sensitivity) and gain of function mutants (increased sensitivity) supports the role of the α4β2*nAChR subtype. However, null mutant mice still respond to nicotine, indicating that other nAChR subtypes also mediate these responses. Mice differing in initial sensitivity to nicotine also differ in tolerance development following chronic treatment: Those mice that are initially more sensitive to nicotine develop tolerance at lower treatment doses than less sensitive mice, indicating that tolerance is an adaptive response to the effects of nicotine.. In contrast, the sensitivity of mice to pre-pulse inhibition of acoustic startle response is correlated with the expression of α7-nAChR. While genetic variability in nAChR expression and function is an important factor contributing to differences in response to nicotine, the observations that altered activity of opioid, glutamate, and cannabinoid receptors among others also change nicotine sensitivity reinforces the proposal that the genetics of nicotine response is more complex than differences in nAChRs.

1. Introduction

Evidence for the importance of genetic factors in mediating tobacco use in humans was first provided by the R.A. Fisher in 1958 [1]. Since then many different approaches, including twin studies and more recently genome wide association studies have firmly established that genetic factors are important components in tobacco use in humans (see reviews [2–6]).

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Our research has used mouse models to investigate the role of genetics in mediating responses to nicotine. A useful initial step to assess the role of genetic factors on any response is the characterization of variability among defined genetic populations. The laboratory mouse is an excellent resource with which to begin the evaluation of genetic factors because of the availability of a large number of inbred strains. More recently, the mouse has been the species widely used to generate genetically modified lines, mostly gene knockout and knockin lines. Tests of the roles of specific genes on responses of interest are now possible.

2. Locomotor Activity and Body Temperature

2a. Inbred Mouse Strains and Classical Genetic Analysis

We initiated our studies on the role of genetic factors in mediating responses to nicotine using available inbred mouse strains. An early study examined the effect of an acute administration of nicotine by constructing full dose-response curves for several behavioral and physiological responses in four common inbred strains (BALB, C57BL/6, DBA/2 and C3H/IBG) [7]. Even with this fairly modest number of strains both quantitative differences (approximately a 4-fold difference in ED_{50} values for nicotine-induced hypothermia) and qualitative differences (locomotor depression in three of the inbred strains but locomotor activation in C3H mice in the open field arena) were observed. Certainly, genotype influenced response to nicotine in the mouse. However, with this limited number of mice a relationship between behavioral response and nicotinic receptor expression (measured with nicotine and α-bungarotoxin binding in tissue homogenates) could not be determined.

The observation of substantial strain differences in response to nicotine prompted two studies examining the heritability of these responses using a diallel cross. The parental strains for this analysis were the four strains screened initially (BALB, C57BL/6, DBA/2 and C3H/IBG) and A. All possible F1 hybrids were generated and tested for the effect of nicotine on hypothermia [8] and open-field activity [9]. Both analyses confirmed that strain differences exist and also demonstrated heritability of the nicotine-induced responses consistent with an additive/dominance model. A significant directional dominance toward increased sensitivity to nicotine that was particularly pronounced for the locomotor response was observed. That is, the hybrid mice were more sensitive to nicotine than predicted by the parental responses. This directional dominance was interpreted from an evolutionary point of view to be indicative of a selective advantage where increased sensitivity could protect against ingestion of toxic levels of nicotine.

The screen of inbred mice was subsequently expanded to include 19 strains [10]. A multicomponent test battery was designed to allow the measurement of several different responses to nicotine in an individual mouse. The battery consisted of measurements of the effects of nicotine on respiratory rate, acoustic startle response, crosses and rears in the Ymaze, heart rate and body temperature. The efficiency of the test battery allowed construction of full nicotine dose-response curves for each strain. Substantial differences in ED_{50} values (4–5 fold for most tests) were observed among the strains, further establishing the importance of genetic factors in mediating nicotine-induced responses. Correlational analysis of the results revealed that the effects of nicotine on the activity measures and body

temperature were very similar, a result confirmed by factor analysis. Overall these analyses indicated the existence of four groups of mice ranging from those that are very sensitive to nicotine (C57BL/10, C57BL/6 and A) to those that are very resistant (BUB and C58). Two additional subsets were also identified, one that is moderately sensitive (including DBA/1 and DBA/2) and a second that is moderately resistant (including C3H and CBA).

In order to investigate whether the variation in acute response to nicotine is a consequence of variability in expression of nicotinic receptors, the binding of nicotine and αbungarotoxin was measured in homogenates prepared from eight different brain regions. It is now well established that nicotine labels $α4β2*-nAChR$ sites [11–12] (the * represents the potential for additional subunits[13]) and α -bungarotoxin labels α 7-nAChR sites [14]. A significant overall negative correlation between the density of nicotine binding sites and ED50 values for nicotine effects on activity and body temperature was observed [15]. The correlation between α -bungarotoxin binding and these ED₅₀ values was not statistically significant. This result indicated that the density of α4β2-nAChR was inversely correlated with sensitivity to locomotor and hypothermic effects of nicotine: the higher α4β2*-nAChR expression, the lower the dose of nicotine necessary to elicit a response. However, these results should be and have been regarded as merely suggestive.

2b. nAChR Null, Gain of Function Mutants and Natural Variants

With the development of genetically modified mice the nAChR can either be deleted (null mutants) or mutated to enhance agonist sensitivity (gain of function) (see [16] for review). Both types of mutants have been generated for the Chrna4 and Chrnb2 genes, which encode the α4 and β2 nAChR subunits, respectively. The availability of these genetically modified mice allows the direct test of the effects of altered α4β2-nAChR expression on response to acute nicotine administration. The results presented in Figure 1 demonstrate the effect of deletion of either the α4 or β2 nAChR subunit gene or insertion of hyperactive α4 (L9'A) or β2 (V22'L) nAChR subunit gene on nicotine effects on Y-maze crosses or body temperature. Significant changes in sensitivity to acute nicotine injection were noted for mice from which wild-type versions of either the α 4 or β 2 nAChR subunits were deleted or replaced with a mutant hyperactive receptor subunit. Deletion of either α4 (new data) or β2 [17] resulted in a gene-dose dependent decrease in sensitivity to acute nicotine. In contrast, insertion of a hypersensitive version of either $α$ 4 (new data) or $β$ 2 [18] resulted in a genedose dependent increase in sensitivity to nicotine. It should be noted, that the null mutant mice still responded to acute nicotine administration illustrating that the α4β2-nAChR was not the only receptor subtype that regulates nicotine-induced hypomotility or hypothermia.

The studies described above concentrated on the relationship between the density of α4β2* nAChR expression and response to nicotine. However, a polymorphism representing a single point mutation in the Chrna4 gene changed in the primary sequence of the subunit (ala/thr difference at position 529) and an alteration of receptor function [19]. This mutation was originally identified in the long sleep (LS) and short sleep (SS) mice that were selected for their differential sensitivity to ethanol and also differ in response to nicotine [20]. Recombinant inbred (RI) strains generated by inbreeding mice isolated from a F2 cross of these mice were tested for their responses to acute nicotine administration. RI mice differing

in the Chrna4 polymorphism were also differentially sensitive to nicotine-induced hypomotility and hypothermia [21]. In heterologous expression systems, α4β2-nAChR can assemble with two alternative stoichiometries $[(\alpha 4)_2(\beta 2)_3]$ with high agonist sensitivity to agonists (HS form) and $\left(\alpha 4\right)_3$ $\left(\beta 2\right)_2$ with lower agonist sensitivity (LS)] [22–25]. These alternate stoichiometries are also found in mouse brain [26–27]. The observation that the A529T polymorphism affects the relative expression of the two alternative stoichiometric forms of the α4β2-nAChR receptor with intrinsic differences in sensitivity to activation by nicotinic agonists [28] suggests that this and perhaps other point mutations in a receptor subunit can alter nicotine responses by changing the ratio of HS to LS α4β2-nAChR. In addition, the relative expression of HS and LS forms of α4β2-nAChR can also be influenced by the 3' untranslated region of mRNA encoding the α4 nAChR subunit protein [29]. A mechanism such as this could contribute to the differential expression of HS and LS forms of the receptor throughout the brain and to alter relative sensitivity to of α4β2-nAChR to agonists, including nicotine, and change response to the drug. However, it is unknown whether alterations in ratio of HS and LS forms contribute to genetically determined differences in response to nicotine.

3. Antinociception

Investigating the differential responses of inbred strains has also been extended to an examination of the role of nicotine as an anti-nociceptive [30]. Significant differences in the potency of nicotine as an antinociceptive were noted among the seven inbred strains using two tests for thermal pain. The ED_{50} values for the tail-flick and hot plate tests were significantly positively correlated ($r = 0.89$). In addition, the ED₅₀ values for these responses showed a significant negative correlation to the density of α4β2*-nAChR binding sites in the hindbrain. These inverse correlations are reminiscent of those observed for the analysis of nicotine effects on locomotor depression and hypothermia [15] and indicate the importance of α4β2*-nAChR in mediating these anti-nociceptive effects of nicotine. This assignment is consistent with results obtained with α4 and β2 null mutant mice each of which required higher nicotine doses to block the thermal pain than those required for wild-type mice [12] However, testing of two F1 hybrids (C57BL/6 \times CBA and C57BL/6 \times DBA) indicated that the genetic architecture of the antinociceptive action of nicotine may not be simple: Overdominance toward increased sensitivity of the C57BL/6 \times DBA F1 hybrid was similar to the result from the diallel crosses for locomotor activity [9] and hypothermia [8]. However, over-dominance toward decreased sensitivity was observed for the C57BL/6 \times CBA F1 hybrid. It should also be noted that deletion of the α4β2*-nAChR subtype had a greater effect on the ED_{50} or maximal response for the hot-plate test than for the tail flick test. The α4 L9'S heterozygotes, which express a hypersensitive α4 subunit, showed enhanced sensitivity on the hot plate test with less effect for tail flick [30], further indicating a more important role for the α4β2*-nAChR subtype in mediating the response to nicotine for the hot plate.

The α5 nAChR subunit is an auxiliary subunit that does not function as a classical α nAChR subunit, but can co-assemble with $\alpha\beta$ nAChR subunit pairs to occupy the fifth position in the receptor pentamer. Incorporation of the α5 subunit in either an α4β2α5-nAChR or an α3β4α5-nAChR markedly affects the physiological and pharmacological properties of the

receptors without the α 5 subunit [31–33]. Consistent with altered pharmacology, α 5 null mutant mice are significantly less sensitive to the effects of an acute nicotine administration on locomotor activity and body temperature, as well as nicotine action as an antinociceptive [34]. Inasmuch as the α5 subunit can coassemble with both α4β2- and α3β4-nAChR subtypes, the effects of α5 subunit deletion cannot be unambiguously ascribed to either major subtype. However, the observation that deletion of either the α 4 or β 2 subunit decreases the hypomotive, hypothermic and antinociceptive [17] [35] [12] properties of nicotine suggests that the α4β2α5-nAChR mediates at least some of these responses.

4. Voluntary Oral Nicotine Consumption

The propensity of mice to self-administer nicotine is likely to be an important indicator of potential for nicotine abuse. Establishing models for intravenous nicotine self-administration in the mouse has, until recently, be extremely challenging [36]. We used oral self-selection of nicotine using the two-bottle choice paradigm to examine genetic influences on nicotine intake [37]. The six inbred mouse strains used in this study (A, BUB,C57BL/6, C3H, DBA/2, and ST/b) differ significantly in their responses to acute nicotine administration [10, 38]. These six strains also differed markedly in oral nicotine intake from water or 0.2% saccharin; total fluid consumption for saccharin solutions tended to be higher. The vehicle made little difference in the pattern of oral intake of nicotine for all strains other than ST/b mice. C57BL/6 mice consumed the most nicotine, while A, C3H and ST/b mice had low nicotine intake.

5. Conditioned Place Preference

Conditioned place preference has been successfully used to evaluatethe condition reinforcing effects of environmental stimuli associated with drug administration. Unlike the genetic analyses of the effects of nicotine on many other responses, much of the research on conditioned place preference has been conducted using genetically modified mice, rather than inbred strains. However, one study did note that C57BL/6 mice acquired conditioned place preference while DBA/2 mice did not [39]. In contrast to the relative paucity of data from strain comparisons, studies with knockout and knockin mice have helped to define the nAChR subtypes contributing to the development of conditioned place preference. Deletion of the β2 subunit eliminated conditioned place preference, while deletion of the α7 subunit did not, supporting the role of β2*-nAChR [40]. Subsequently, targeted deletion of the α4 subunit in dopaminergic neurons was found to eliminate conditioned place preference, supporting the role of α 4 β 2*-nAChR on the reward pathway [41]. However, in a different study little effect of a global deletion of the α4 subunit was noted, but mice expressing the α4 S6'F gain of function mutation achieved conditioned place preference at lower nicotine doses than did wild-type mice [42]. Mice differing in the α4 A529T polymorphism differed significantly in nicotine conditioned place preference; mice expressing the T variant developed conditioned place preference, while those with the A variant did not [43]. Interestingly, DBA mice express the A variant and C57BL/6 express the T variant a result consistent with the difference in conditioned place preference reported for these inbred strains [39]. The α5 subunit also contributes to nicotine-induced conditioned place preference [34]. Wild-type mice and α5 knockout mice showed similar dose-response

curves for conditioned place preference in the lower range of nicotine doses. However, after conditioning with higher nicotine doses, α5 knockouts continued to exhibit conditioned place preference while wild-type mice did not. This pattern of response is reminiscent of the effect of the deletion of the α5 subunit on intravenous nicotine self-administration where α5 knockout mice self-administer significantly more nicotine than wild-type mice [44]. Pharmacological studies have implicated the α6 subunit in conditioned place preference, as well [45]. Treatment of mice with the selective α6β2*-nAChR antagonist α-conotoxin MII (H9A, L15A) resulted in a dose dependent inhibition of nicotine-induced conditioned place preference. Overall, several different nAChR subtypes participate in nicotine-induced conditioned place preference and appear to participate both in the rewarding aspects (α4, α6 and β 2) as well as the adversive aspects (α 5) of this complex behavioral response.

6. Seizure Activity

Administration of relatively high doses of nicotine causes convulsions. Sensitivity to nicotine-induced seizures varies markedly among inbred strains administered nicotine either intraperitoneally or intravenously [38]. A greater than 2.5-fold difference in both ED_{50} and seizure latency were determined with ST/b mice being the most sensitive for both measures and DBA/2 among the most resistant. Subsequent comparison of seizure sensitivity to the density of α-bungarotoxin binding sites (now known to measure α7-nAChR) revealed a significant inverse correlation. This result suggested that mice expressing higher levels of α7-nAChR were more prone to nicotine –induced seizures. This potential relationship between seizure sensitivity and α-bungarotoxin binding was consistent with the results of previous studies in which the inheritance of these measures was examined with a classical genetic cross between C3H and DBA/2 mice that are relatively sensitive or resistant to nicotine-induced seizures, respectively [46–47]. These studies revealed that resistance to seizures was dominant as was expression of lower levels of α-bungarotoxin binding sites. These results are also consistent with the role for α 7-nAChR in mediating nicotine-induced clonic seizures. However, in the segregating F2 population the relationship between seizure sensitivity and α-bungarotoxin binding sites was not observed. No significant correlation was also reported when seizure sensitivity was compared to α-bungarotoxin binding as well as polymorphisms in the Chrna5 and Chrna7 genes [48]. A relatively small proportion of the F2 generation from an independent DBA by C3H cross mice was seizure sensitive. This result is consistent with the previous observation of dominance toward seizure resistance for these F2 mice. Although higher seizure sensitivity was noted in mice expressing the C3H polymorphism in the Chrna7 gene, the effect was more pronounced for the Chrna5 polymorphism, suggesting a possible role for $a5*-nAChR$ in modulating nicotine-induced seizures.

The contribution of α 7-nAChR to nicotine-induced seizures was subsequently examined using null mutant mice. Deletion of the α7 subunit did not alter the seizures elicited by nicotine, strongly indicating that the wild-type α7-nAChR did not mediate this response [49]. However, heterozygotic mice harboring a hyperactive α7-nAChR are indeed seizure sensitive, suggesting that stimulating a hyperactive α7-nAChR can indeed elicit seizure activity [50]. The appearance of nicotine-induced seizures in mice expressing hyperactive receptors has also been observed for the $α4$ subunit [51–52] and the $β2$ subunit [53]. This

general result suggests that gain of function mutations can recruit nAChR subtypes to mediate nicotine-induced seizures, subtypes that may not mediate this response without altered receptor sensitivity.

As mentioned above, a polymorphism in the Chrna5 gene was implicated in mediating nicotine-induced seizures in the C3H×DBA F2 population [48]. The observation that deletion of the α5 nAChR subunit substantially reduced sensitivity to nicotine-induced seizures confirmed this prediction [54]. Furthermore, demonstration that deletion of the β4 subunit or reduced expression of the α3 subunit both significantly reduced seizures elicited by nicotine [55]strongly argues that an α 3 β 4 α 5-nAChR subtype mediates this response.

6. Auditory Gating

Although the α7-nAChR may not be the primary mediator of nicotine-induced seizures, pharmacological studies have implicated this subtype in auditory gating [56]. A subsequent study compared auditory gating and α-bungarotoxin binding in hippocampus of eight different mouse strains. A significant inverse correlation was seen between the degree of auditory gating and the density of hippocampal α -bungarotoxin sites (α 7-nAChR) but not nicotine sites (α4β2*-nAChR) [57], indicating that mice with relatively low α7-nAChR displayed poor auditory gating. In order to test this relationship, gating was investigated using C3H wild-type and α7 heterozygotes, which express significantly fewer αbungarotoxin binding sites [58]. Indeed, the C3H α 7^{+/-} mice displayed much less auditory gating than did C3H^{+/+} mice, which except for the difference in α 7 expression have virtually the same genetic background. Additional support for the role of α7-nAChR in modulating auditory gating is supplied by pharmacological studies demonstrating improvement in gating in DBA/2 mice (a poor gaiter) following administration of α 7-nAChR selective nicotinic drugs [59].

7. Chronic Nicotine Treatment

Chronic exposure to nicotine results in the development of tolerance to the effects of nicotine and changes in the expression of nAChR. The increase in nAChR with high affinity for agonists was initially observed in rats [60] and mice [61]. Importantly, the nicotineinduced increase in these receptors also occurs in human tobacco users [62–64]. We have investigated the effects of genetic factors on development of tolerance to and regulation of nAChR expression following chronic nicotine treatment in mice.

Initially the four strains that had been tested for differences in response to acute nicotine treatment (BALB, C3H, C57BL/6 and C3H) were chronically treated with a single dose of nicotine (3 mg/kg/hr for 10 days) [65]. Tolerance to nicotine effects on locomotor activity and body temperature was noted in three of the strains, but C3H mice developed little tolerance following this treatment. A follow-up study in which DBA and C3H mice were treated with one of four nicotine doses (0, 2, 4 or 6 mg/kg/hr) confirmed that DBA mice develop increasing tolerance with increasing chronic treatment dose, while C3H mice develop very little tolerance [66].

These studies revealed significant genetic influences on the development of nicotine tolerance and suggested that mice that are initially less sensitive to the effects of nicotine (C3H) developed less tolerance following chronic nicotine treatment than mice that exhibited greater acute effects. The screen of 19 inbred strains [10] identified additional strains that were very sensitive (A) and very resistant (BUB) to acute nicotine administration. Mice from five inbred strains (most sensitive to least sensitive: $C57BL/6 > A$ $>$ DBA $>$ C3H $>$ BUB) chronically treated with nicotine (0, 0.5, 1, 2, 4 or 6 mg/kg/hr) were tested for tolerance in order to further examine genetic influences on tolerance development. As was the case with the previous studies, mice that were more sensitive to the acute effects of nicotine developed tolerance after treatment with lower nicotine doses than mice that were less sensitive to the acute effects $(C57BL/6 > A > DBA > C3H > BUB)$.

Increases in nicotine binding were noted in the six brain regions assayed. The extent of the increase varied among the regions, but the responses were similar among the strains [65– 67]. Chronic nicotine treatment did not affect the K_D for nicotine. The generally similar response of binding sites measuring α4β2*-nAChR among the strains that differ markedly in tolerance development indicates that changes in the expression of these receptors do not adequately explain the differences in the development of tolerance following chronic nicotine treatment among these strains. Alternative or additional mechanisms are required to define the strain differences.

Chrna and Chrnb knockout and knock-in are being used to investigate various aspects of nicotine dependence including their roles in reinforcement and withdrawal. is The results of these studies demonstrate important roles for specific subunits including α 4, α 5, α 6, β 2 and β4 nAChR subunits and are the subject to several recent reviews and will not be discussed here [68–70].

8. Comparison of Patterns of Response Across Genotypes

The scattergrams shown in Figure 2 compare the responses to nicotine measured during the screens for the inbred mouse strains. All responses were not measured in every strain, so the number of points on the scattergrams varies among the tests. In general, correlations among the various tests were not significant, indicating either that these responses either do not have a common genetic basis or too few strains have been tested. However, several responses for which regression lines are included in the panels were correlated: ED_{50} for antinociception measured by tail flick to that measured by hotplate ($r = 0.89$), ED_{50} for hypothermia and ED_{50} for antinociception measured with the hot plate (r = 0.82), ED_{50} for hypothermia and threshold tolerance dose for hypothermia ($r = 0.93$), and ED₅₀ for seizures and maximum oral nicotine intake $(r = 0.89)$. These significant correlations may indicate some shared genetic factors mediate the correlated behaviors.

The correlation between the two measures of thermal pain may not be surprising since both of these responses are significantly reduced in both α4 and β2 knockout mice [12], indicating the involvement of α4β2*-nAChR in mediating both responses. However, the effect of these gene deletions is not identical. A larger effect on the hot plate test was noted in both the α4 and β2 knockout mice as well as for the α4 L9'S knock-in heterozygote.

The positive correlation between ED_{50} values for hot plate antinociception and hypothermia was not quite as robust has that for the two measures of thermal pain. However, the fact that deletion of the α4 and β2 subunits significantly reduced sensitivity to nicotine for both of these measures, albeit to a different degree, is consistent with some shared mechanism of action.

The relationship between sensitivity to acute responsiveness to nicotine and the chronic treatment dose that elicits tolerance has been noted previously [67]. This observation indicates that tolerance develops to the effects of nicotine. That is mice that are initially more sensitive to the effects of the drug (such as C57BL/6) develop tolerance after exposure to lower nicotine doses than mice that are less responsive to an acute dose of nicotine (such as BUB). This differential tolerance development is not directly related to the up-regulation of nicotine binding sites since the pattern of change in these binding sites is generally similar for each of the five strains tested.

The significant correlation between oral nicotine intake and ED_{50} values for nicotineinduced seizures has also been noted previously [37]. Mice that are more sensitive to this adverse effect of nicotine consumed less nicotine. This result suggests that the perception of an unpleasant effect of nicotine, perhaps indicated by sensitivity to nicotine-induced seizures, limits voluntary oral intake of nicotine. Adverse effects of the drug may be the limiting factor in oral nicotine consumption. The observation that α5 knockout mice, which are resistant to nicotine-induced seizures [54], self-administer significantly more nicotine that wild-type mice [44] also indicates that reduction in adverse responses to nicotine facilitates drug intake.

9. Summary and Discussion

Genetic factors clearly influence several different physiological and behavioral responses to nicotine administered either acutely or chronically. Investigations using genetically modified mice have identified the importance of several different nAChR subunits, and consequently different nAChR subtypes, in mediating many of the responses in mice. With the advent of genome wide association studies, it has been demonstrated that variation among CHRN genes in either translated or untranslated regions (particularly CHRNA5, which encodes the α5 nAChR subunit) contribute to several different aspects of human tobacco use [71–73]. However, these variations in CHRN genes account for a relatively small amount of the genetic variance clearly indicating that other factors exist. One of these factors is differences in nicotine metabolism [74]. Several genetic factors are now known to influence nAChR function and/or expression. Probably most obvious changes are mutations leading to changes in primary sequence of a nAChR subunit and subsequently to functional change. Single point mutations, frequently leading to a gain of function of the mutant nAChR, contribute to the relatively rare syndrome, autosomal dominant frontal lobe epilepsy (reviewed in [75–77]). Animal models that express the mutant subunits have been developed [78]. Studies with mice engineered to express these mutations demonstrate that the gain of function mutations increase sensitivity to nicotine for locomotor activity and body temperature [18, 79]. In addition, gain of function mutations also alter sensitivity to nicotine-induced convulsions such that activation of nAChR subtypes containing α 4, α 7 or

β2 subunits that do not normally mediate this response elicit seizure activity [50–51, 53, 80– 81]. In addition to the profound effect of the hypersensitive nAChR subunits, more subtle changes in expression can alter drug responses. A naturally occurring mutation of the α4 subunit of mice also seems to elicit changes in stoichiometry and modifies responses to nicotine for several behaviors including hypothermia and open field activity [43]. Changes in the expression of α 4 and β 2 nAChR subunits alter the expression of two stoichiometric forms of α4β2-nAChR that are differentially sensitive to activation by agonists, including nicotine [27]. Stoichiometric changes were also observed for α4 transcripts lacking a 3' untranslated region [29]. Relatively subtle structural changes can also markedly affect receptor function. Inclusion of the α5 subunit in α4β2α5-nAChR markedly alters physiological activity [32] and deletion of the α5 subunit dramatically reduces sensitivity to nicotine *in vivo* [34, 54], illustrating an important role for this auxiliary subunit consistent with the well established role of the α 5 subunit in human smoking [82–83]. These examples illustrate that differences in nAChR expression and function are important factors in mediating response to nicotine.

Nevertheless, the importance of genetic factors, in addition to variation in nAChR subtype [34, 40, 45], distribution [41], and primary sequence [43] that also influence response to nicotine can be illustrated for conditioned place preference. Nicotine conditioned place preference is modified by changes in the expression of cannabinoid receptors [84–85], NMDA receptors [86], μ-opioid receptors [87], δ-opioid receptors [88], galanin [89], protein kinase C ε activity [90], and CREB expression [91]. Furthermore, nAChR are known to mediate the release of neurotransmitters dopamine, GABA, glutamate, and serotonin [92– 93] and hormones such as corticosterone [94]. Thus, genetic variation in any of these processes can affect responses to nicotine. Consequently, while the genetic influences on nicotine mediated behaviors are surely affected by the complex array of and variations in nAChR, themselves, genetic diversity of the myriad processes that are mediated by or interact with nAChR no doubt contribute significantly to the complex phenotypes observed in response to nicotine exposure. The complexity of the genetic influences on nicotine response is illustrated to some extent by the observation that the inbred BUB/Bn mouse strain, which expresses relatively normal levels of α 4 β 2*nAChR, [10] is less affected by acute nicotine administration than the β2 null mutant [17].

These genetic studies illustrate the complexity of the physiological, biochemical and behavioral responses observed following exposure to nicotine. Indeed, 30 years of research has progressed from the time before central nervous system nAChR were recognized as real until a diverse array of differentially distributed subtypes have been identified [95–100]. Further study of the role of these diverse nAChR and their interactions with the myriad of neuronal pathways will no doubt demonstrate additional levels of complexity underlying responses to acute and chronic nicotine and may reveal the basis for the persistence of tobacco use.

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Figure 1.

Effects of α4 or β2 gene deletion or gain of function mutations on acute responses to nicotine. Mice differing in either α4 genotype (α4 wild-type, heterozygotes or null mutants or wild-type, α4 L9'A homozygotes or hemizygotes) or β2 genotype (β2 wild-type, heterozygotes or null mutants or wild-type, \β2 V289L homozygotes or hemizygotes) were injected intraperitoneally with the indicated doses of nicotine (free base). Activity in the Ymaze was measured for 3 min beginning 3 min after injection and rectal body temperature was measured 15 min after injection. Each point on the dose-response curves represents data from 5–16 mice at each dose. The different α4 genotypes are represented by red symbols (+ +, wild-type, red; +−, heterozygote, pink; −−, null mutant, white; +M, α4L9'A hemizygote, dark red; MM, α4 9'AA homozygote, darkest red). The different β2 genotypes are represented by blue symbols (++, wild-type, blue; +−, heterozygote, light blue; −−, null mutant, white; +M, β2VL hemizygote, darker blue and MM β2 LL homozygote, darkest blue). The lines for each genotype represent the least squares curved fit of the data to modified Hill equations. The effect of variation of genotype on the ED_{50} \pm SEM values (mg/kg) calculated from the Hill fits are shown in the lower panels. Note that ED_{50} values increase with gene deletion and decrease with introduction of a gain of function mutation.

(mg/kg/hr)

Figure 2.

Correlations of relative sensitivity of inbred strains for seven measures of nicotine-induced changes. Scattergrams were constructed to compare measures of relative sensitivity of inbred mouse strains to each of seven independently measured responses to nicotine. Each individual panel presents the correlations between the measures of sensitivity to nicotine \pm S.E.M for two tests $(ED_{50}$ or similar value) for the mouse strains for which each measurement was conducted. Note that the numbers of points on each scattergram differ owing to variation in the number of strains measured for each test. The ordinate and abscissa for the responses on the diagonal in the figure change for the various comparisons. The axis

label for each measurement is provided near the vertex of the panels in the horizontal and vertical directions and is read upwards as the abscissa for figures above the label and to the right as the ordinate for figures to the right of the label. For responses that were significantly correlated, regression lines passing through the origin are shown. Correlation coefficients for these measures were: ED_{50} hypothermia vs. ED_{50} hot plate, $r = 0.82$; ED_{50} , hot plate vs. ED_{50} tail flick, $r = 0.89$; ED_{50} hypothermia vs. minimal tolerance dose, $r = 0.93$; ED_{50} clonic seizures vs. Maximal oral nicotine intake, $r = 0.89$. All other correlation coefficients were less than 0.55. Note that relatively few points are included for several pairs of tests reflecting limited number of strains tested for both measures.