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## Nicotinic $\alpha 7$ Acetylcholine Receptor ( $\alpha 7$ nAChR) in Human Airway Smooth Muscle

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### Abstract

Diseases such as asthma are exacerbated by inflammation, cigarette smoke and even nicotine delivery devices such as e-cigarettes. However, there is currently little information on how nicotine affects airways, particularly in humans, and changes in the context of inflammation or asthma. Here, a longstanding assumption is that airway smooth muscle (ASM) that is key to bronchoconstriction has muscarinic receptors while nicotinic receptors (nAChRs) are only on airway neurons. In this study, we tested the hypothesis that human ASM expresses  $\alpha 7$ nAChR and explored its profile in inflammation and asthma using ASM of non-asthmatics vs. mild-moderate asthmatics. mRNA and western analysis showed the  $\alpha 7$  subunit is most expressed in ASM cells and further increased in asthmatics and smokers, or by exposure to nicotine, cigarette smoke or pro-inflammatory cytokines TNF $\alpha$  and IL-13. In these effects, signaling pathways relevant to asthma such as NF $\kappa$ B, AP-1 and CREB are involved. These novel data demonstrate the expression of  $\alpha 7$ nAChR in human ASM and suggest their potential role in asthma pathophysiology in the context of nicotine exposure.

### Keywords

asthma; inflammation; cigarette smoke; nicotinic receptors

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## Introduction

Airway smooth muscle (ASM) cells play a critical role in bronchial structure and function via their role in regulation of contractility and remodeling including proliferation and extracellular matrix production in the context of diseases such as asthma and COPD (Amrani et al., 2020; Black, Panettieri, Banerjee, & Berger, 2012; M. Lam, Lamanna, & Bourke, 2019; Lo, Kennedy, Kurten, Panettieri, & Koziol-White, 2018; Prakash, 2013, 2016). Factors such as cigarette smoke are known to exacerbate asthma and contribute to COPD (Beasley, Semprini, & Mitchell, 2015; Burney, Jarvis, & Perez-Padilla, 2015; Ellis & Soliman, 2015; Gibson-Young, Martinasek, Tamulevicius, Fortner, & Alanazi, 2020; Lundback, Backman, Lotvall, & Ronmark, 2016; May & Li, 2015; Postma, Bush, & van den Berge, 2015; Sheehan & Phipatanakul, 2015; St Claire et al., 2020; Traboulsi et al., 2020; Viegi, Maio, Pistelli, Baldacci, & Carrozzi, 2006; Wurst, Kelly-Reif, Bushnell, Pascoe, & Barnes, 2016). However, the mechanisms by which CS affects ASM *per se* in airway hyperresponsiveness are still under investigation (Aravamudan et al., 2014; Sathish et al., 2015; Vogel et al., 2014; Wylam et al., 2015; Zuo et al., 2018).

There is increasing recognition that nicotine, the major active element of CS, contributes to increased airway hyperresponsiveness (Chun, Moazed, Calfee, Matthay, & Gotts, 2017; Clapp & Jaspers, 2017; Gibson-Young et al., 2020; Jankowski, Brozek, Lawson, Skoczynski, & Zejda, 2017; St Claire et al., 2020; Traboulsi et al., 2020). Such effects of nicotine are relevant to both smoking and nicotine delivery devices such as e-cigarettes. However, the mechanisms by which nicotine affects ASM or how such effects change in asthma or inflammation, are not well-understood. Previous studies in rat ASM have found nicotine elevates intracellular  $Ca^{2+}$  concentration ( $[Ca^{2+}]_i$ ) (Hong & Lee, 1996; Jiang et al., 2014). Classically, nicotine acts through nicotinic acetylcholine receptors (nAChRs). However, in ASM, the common assumption is that contractility involves muscarinic receptors. Less is known regarding expression or functionality of nAChR in ASM. Limited data show  $\alpha 7$ nAChR is expressed in rat ASM (He et al., 2014; Hong & Lee, 1996; Jiang et al., 2014). Whether human ASM expresses nAChR subunits is not known, nor is the effect of inflammation, asthma or CS on nAChR portfolios that could contribute to observed effects of nicotine or CS on the airway. Therefore, the present study was designed to explore nAChR in human ASM with the hypothesis that human ASM expresses  $\alpha 7$ nAChR and its expression is increased in inflammation and asthma, providing mechanistic basis for nicotine effects in the airway.

## Materials and Methods

### Reagents and supplies:

Cell culture reagents and supplies including Dulbecco's Modified Eagle's Medium/F12 (DMEM/F12), trypsin and Antibiotic-Antimycotic (AbAm) were purchased from Invitrogen (Carlsbad, CA). Charcoal stripped fetal bovine serum (FBS) was from Sigma-Aldrich (St. Louis, MO).  $\alpha 7$ nAChR antibody was from Novus Biologicals (Littleton, CO). Other antibodies for  $\alpha 4$ nAChR,  $\beta 2$ nAChR, caldesmon,  $\alpha$ -SMA, calponin-1 and transgelin/SM-22 were obtained from Abcam (Cambridge, UK).  $\beta$ -actin antibody was from ABM Biological Materials (Richmond, Canada). Pro-inflammatory cytokines, tumor necrosis factor alpha

(TNF $\alpha$ ) and interleukin-13 (IL-13) were from R&D Systems (Minneapolis, MN). RIPA cell lysis/extraction buffer and A/G PLUS Agarose beads for co-immunoprecipitation techniques were obtained from Thermo Fisher Scientific (Waltham, MA) and SantaCruz Biotechnology (Dallas, TX) respectively. Pharmacological inhibitors for signaling pathways NF $\kappa$ B (SN-50) and STAT6 (AS1517499) were from SantaCruz Biotechnology (Dallas, TX) and Millipore Sigma (Burlington, MA) respectively. STAT3 (SD 1008), AP1 (SR 11302), CREB (666-15) and MAPK (PD 98059) were from Tocris Bioscience (Bristol, UK). Fluorescent secondary antibodies were from Li-Cor Biosciences (Lincoln, NE). Other chemicals/drugs/antibodies were from Sigma-Aldrich unless otherwise specified.

#### **Primary human ASM cells:**

A previously described technique (Aravamudan et al., 2014; Sathish et al., 2014) was followed to isolate human ASM cells from lung samples incidental to patient thoracic surgery at Mayo Clinic, Rochester, MN (approved by the Mayo Clinic Institutional Review Board and considered minimal risk). Briefly, third to sixth generation bronchi were isolated from lung resections for non-infectious indications including focal cancers (bronchoalveolar carcinoma was excluded). Asthma diagnoses and severity was based on standard clinical criteria including pulmonary function testing, as noted in patients' clinical records. Non-asthmatics were defined as patients with no noted history of lung disease and were clinically deemed to have normal lung function. Samples were from both male and female adults ranging from 21 to 80 years of age. ASM was dissected from healthy appearing lung areas identified with the help of a pathologist. Samples were enzymatically dissociated as previously described (Aravamudan et al., 2014; Sathish et al., 2014) to isolate ASM cells, which were placed in cell culture under standard conditions of 37°C (5% CO<sub>2</sub>, 95% air) using DMEM/F12 supplemented with 10% FBS and 1% AbAm. Cells <5 passages of subculture were serum-deprived for a minimum of 24 hours prior to all experimental procedures. Furthermore, human ASM cell phenotype validations were also performed periodically (Supplement Figure 1).

#### **Cell Treatment:**

Following serum starvation, cells were treated with TNF $\alpha$  (20ng/ml) or IL-13 (50ng/ml) in the presence/absence of inhibitors for 24 h. For signaling studies, non-asthmatic ASM cells were treated with pharmacological inhibitors 2h prior to addition of TNF $\alpha$  or IL-13 for another 24h. Expression was assessed at protein and mRNA levels.

#### **qRT-PCR:**

Cells were washed with RNA-grade DPBS, trypsinized and centrifuged. Total RNA was extracted from cells using Quick-RNA™ MiniPrep kit (Zymo Research, Irvine, CA) following manufacturer's protocol and complementary DNA was synthesized using OneScript cDNA Synthesis Kit (Richmond, BC, Canada). Standard qPCR techniques were followed (optimized for Roche LC480 Light Cycler) using QuantStudio 3 RT-PCR system as per the manufacturer's instructions. The following primers obtained from Integrated DNA Technologies (Coralville, IA) were used for qRT-PCR analysis:  $\alpha$ 7nAChR (forward 5'-CAA TGG AGA ATG GGA CCT AGT G-3', reverse 5'-GCA GCA TGA AGA CGG TAA GA-3'), s16 (forward 5'-CAA TGG TCT CAT CAA GGT GAA CGG-3', reverse 5'-CTG

GAT AGC ATA AAT CTG GGC-3'). Fold changes in mRNA expression were calculated by normalization of cycle threshold [C(t)] value of target genes to reference gene S16 using the Ct method.

### **Immunoblotting:**

ASM cell lysates were prepared using cell lysis buffer (Cell Signaling Technologies, Beverly, MA) containing protease and phosphatase inhibitors using previously described methods (Aravamudan et al., 2014; Sathish et al., 2014). Resultant supernatants were assayed for total protein content using the DC protein Assay kit (BioRad, Hercules, CA, USA). Approximately 30 µg of each lysate were loaded on 10% SDS-page and transferred onto 0.22 µm PVDF membranes. Non-specific binding was blocked using 5.0% bovine serum albumin (BSA) and membranes probed overnight at 4°C with antibodies of interest. Blots were then incubated with Li-Cor secondary antibodies. Protein expression detection and densitometry were performed on a Li-Cor Odyssey IR scanning system (Lincoln, NE). Band intensities were normalized against β-actin.

### **Co-immunoprecipitation (Co-IP) of α4nAChR, α7nAChR and β2nAChR:**

Co-IP assays were performed according to previously published protocols (Sathish, Xu, Karmazyn, Sims, & Narayanan, 2006; Townsend, Sathish, Thompson, Pabelick, & Prakash, 2012). Briefly, fully confluent human ASM cells were washed twice in ice-cold DPBS. After washing, cells were scraped off directly with a cell scraper using 1X RIPA Lysis and Extraction Buffer. Cells were then centrifuged at 14,000 rpm for 10 min, and the supernatant was collected followed by protein estimation. For α4nAChR pull-down assays, A/G PLUS–agarose beads (Santa Cruz Biotechnology) were incubated with 2% BSA in DPBS for 2 h at room temperature. Post incubation with BSA, A/G PLUS agarose beads were vortexed for 30 sec and centrifuged in a Beckmann centrifuge for 10 min at 14,000 rpm to remove residual BSA. Then, BSA-treated A/G Plus agarose beads were incubated with 0.2 µg primary antibody (α4nAChR) by gentle rocking for 30 min at room temperature. Then agarose-bound α4nAChR was added to each protein sample (via RIPA extraction as above) at a concentration of 1mg/mL with gentle rocking overnight at 4°C. The sample mixtures containing protein sample, A/G PLUS agarose beads and α4nAChR antibody were vortexed for 30 sec and centrifuged for 30 min at 14,000 rpm. The mixtures were washed five times with cold washing buffer (20mM HEPES (pH 7.5), 150mM NaCl, 1mM EDTA and 0.5% Tween-20 in DPBS) to remove unbound protein. The sample mixtures were then denatured with 1X SDS sample buffer before loading in a 4-20% polyacrylamide gel. After blocking with non-fat milk suspension, the PVDF membranes were probed for α7nAChR or β2nAChR overnight at 4°C, followed by the Li-Cor Odyssey CLx Imaging detection using a Li-Cor IRDye secondary antibodies (Lincoln, NE).

### **Immunofluorescence:**

ASM cells were fixed in 4% paraformaldehyde for 15 min, washed in DPBS, blocked with 4% normal donkey serum, and exposed overnight to rabbit anti-α7nAChR antibody. Specificity of the antibody to α7nAChR was tested by pre-exposure of living ASM cells to alpha-bungarotoxin (αBTX; List Biologicals) prior to fixation. Cells were then washed with PBS and exposed to Cy3-conjugated donkey anti-rabbit secondary antibody. Nuclei were

counterstained with DAPI. Stained samples were imaged on a Nikon Ti-U inverted microscope with a 40X/1.3 objective lens and epifluorescence attachments.

### Statistical analysis:

All experiments were performed in human ASM cells from at least 3 patients (non-asthmatic/asthmatic) and performed in duplicate for each patient sample for qRT-PCR and western analyses. Statistical comparisons were made using either one-way ANOVA or two-way ANOVA as appropriate, followed by Bonferroni *post-hoc* multiple comparisons test using GraphPad Prism version 8.0.0 for Windows, GraphPad Software, San Diego, California USA, [www.graphpad.com](http://www.graphpad.com). Statistical significance was tested at a level of  $p < 0.05$ . All values are expressed as means  $\pm$  SEM. “N” values representing numbers of patients’ samples in the figure legends.

## Results

### $\alpha$ 4nAChR and $\alpha$ 7nAChR are expressed in human ASM

ASM cells from non-asthmatic patients were processed for mRNA-based portfolio of nAChR subunits known to be expressed in humans (Changeux & Taly, 2008). Interestingly, only  $\alpha$ 4nAChR and  $\alpha$ 7nAChR showed substantial expression with much smaller expression of the  $\beta$ 2 subunit (Fig. 1A). In comparison, other subunits were minimally expressed. Based on these data, we further focused only on the  $\alpha$ 4 and  $\alpha$ 7 subunits. Interestingly, asthmatic ASM showed significantly decreased expression of  $\alpha$ 4nAChR protein (Fig. 2A). Furthermore, in cells exposed to TNF $\alpha$  or IL-13, there was no significant change in  $\alpha$ 4nAChR levels, with a trend towards small increase with either cytokine (Fig 2B). In contrast, asthmatic ASM showed significantly increased expression of  $\alpha$ 7nAChR mRNA (Fig. 3A) and protein (Fig. 3B). Immunofluorescence staining of ASM cells showed plasma membrane localization of  $\alpha$ 7nAChR (Fig 1B) absent in cells pre-exposed to  $\alpha$ BTX. Furthermore, protein expression of  $\alpha$ 7nAChR was significantly elevated in patient samples from smokers compared to non-smokers (Fig. 4). Accordingly, subsequent studies focused on  $\alpha$ 7nAChR.

### Cytokine-induced changes in $\alpha$ 7nAChR in human ASM

We next examined if cytokines affected expression of  $\alpha$ 7nAChR, and differences between non-asthmatic vs. asthmatic ASM. Therefore, ASM cells were exposed to TNF $\alpha$  (20ng/ml) or IL-13 (50ng/ml) for 24h and harvested at 6 h, 12 h and 24 h time intervals. qRT-PCR showed that expression of  $\alpha$ 7nAChR was increased in the presence of either cytokine in non-asthmatics (Fig. 5) but significantly only at 24 h (Fig 5E). In comparison, there were significant increases in  $\alpha$ 7nAChR mRNA expression at all three time-points for asthmatic ASM (Fig. 5B, D, F). Interestingly, IL-13 appeared to have a more pronounced effect in comparison to TNF $\alpha$  particularly in asthmatics. This was not necessarily reflected by protein expression which nonetheless showed increased  $\alpha$ 7nAChR in non-asthmatic ASM with both cytokines compared to vehicle control (Fig 6A). Interestingly, 1% or 2% cigarette smoke extract (CSE) also resulted in increased  $\alpha$ 7nAChR (Fig 6B).

### **Mechanisms of $\alpha 7$ nAChR regulation in human ASM**

We examined the role of transcriptional regulation of  $\alpha 7$ nAChR, focusing on STAT6, NF $\kappa$ B, and AP-1 relevant to airway inflammation, using pharmacological inhibitors given 2 h before exposure to either TNF $\alpha$  or IL-13 for 24 h. Interestingly, STAT6 inhibition (20 nM AS151749) significantly reduced IL-13 effects on  $\alpha 7$ nAChR but not TNF $\alpha$  (Fig 7A). Pretreatment with an NF $\kappa$ B inhibitor (SN50, 20  $\mu$ M) or AP-1 inhibitor (SR1302; 1  $\mu$ M) significantly blunted  $\alpha 7$ nAChR enhancement by either cytokine (Fig 7B and Fig 5C).

To determine the role of major cytosolic signaling pathways, we focused on the ERK pathway using a MAPK inhibitor (PD98059; 2  $\mu$ M). Interestingly, we found no significant effects of this inhibitor on either TNF $\alpha$  or IL-13-induced increase in  $\alpha 7$ nAChR expression (Fig 8A). CREB is an important cytoplasmic-nuclear trafficking molecule in the transcriptional machinery of ASM (Prakash, 2016). Interestingly, only TNF $\alpha$ -induced increase in  $\alpha 7$ nAChR expression was blunted on pharmacological inhibition of CREB (Fig. 8B).

To investigate interactions amongst  $\alpha 4$ ,  $\alpha 7$  and  $\beta 2$  receptor subunits, we performed Co-IP experiments which showed that both  $\alpha 7$ nAChR and  $\beta 2$ nAChR interact with  $\alpha 4$ nAChR to some extent in human ASM cells. However, this colocalization was not significantly different between non-asthmatics and asthmatics (Fig. 9).

## **Discussion**

Inflammation in asthma enhances ASM Ca<sup>2+</sup> and contractility response to bronchoconstrictor agonists such as ACh (Guedes et al., 2015; Koopmans et al., 2014; Prakash, 2013, 2016; Wright et al., 2013). ACh effects on ASM are usually associated with muscarinic receptors, while nAChRs are generally considered to act as Ca<sup>2+</sup> channels in preganglionic neurons (Belmonte, 2005; Canning, 2006; Canning & Fischer, 2001; Pieper, 2012; Racke & Matthiesen, 2004; Rogers, 2002; van der Velden & Hulsmann, 1999). However, in the context of CS exposure as well as nicotine delivery devices such as e-cigarettes, it is important to consider whether and how inhaled nicotine influences airway cells such as epithelium and ASM. Clinical data shows that CS exacerbates asthma (Beasley et al., 2015; Burney et al., 2015; Ellis & Soliman, 2015; Lundback et al., 2016; May & Li, 2015; Postma et al., 2015; Sheehan & Phipatanakul, 2015; Wurst et al., 2016) and that e-cigarettes while seen as friendly for asthmatics (Lucchiari et al., 2016; Meo & Al Asiri, 2014; Oh & Kacker, 2014; Orellana-Barrios, Payne, Mulkey, & Nugent, 2015; Rowell & Tarran, 2015; Walton et al., 2015) may in fact not be as benign for asthmatics (Hsieh, 2016; Oh & Kacker, 2014) and are increasingly seen as toxic (Chun et al., 2017; Clapp & Jaspers, 2017; Jankowski et al., 2017; Javed, Kellesarian, Sundar, Romanos, & Rahman, 2017; G. Kaur, Muthumalage, & Rahman, 2018; McEvoy & Spindel, 2017). Nicotine also remains relevant as long as smoking exists. However, the mechanisms of nicotine action even in normal airways are surprisingly understudied, and barely understood in asthmatic airways. Furthermore, little is known about nAChR expression and roles in human ASM, inflammation or asthma. The present data show that human ASM in fact do express nAChR, largely  $\alpha 7$ nAChR. Furthermore,  $\alpha 7$ nAChR expression is increased with exposure to cigarette smoke or asthma-relevant cytokines such as TNF $\alpha$  or IL-13, and importantly in

asthmatics and smokers. Given the roles of ASM in airway hyperreactivity (Guedes et al., 2015; Koopmans et al., 2014; Prakash, 2013, 2016; Wright et al., 2013) and remodeling, (Berair & Brightling, 2014; Guedes et al., 2015; Hirota & Martin, 2013; Noble et al., 2014; Prakash, 2013, 2016) these findings on  $\alpha 7$ nAChR become important.

Nicotine acts on nAChRs of airway-related vagal preganglionic neurons in the central nervous system (Dehkordi, Kc, Balan, & Haxhiu, 2006; Q. Gu, Ni, & Lee, 2008; Krasteva et al., 2011; Lee et al., 2007; Lee & Gu, 2009). Cholinergic nerves enhance bronchoconstriction, mucus secretion, and bronchial vasodilation (Belmonte, 2005; Canning, 2006; Canning & Fischer, 2001; Racke & Matthiesen, 2004; Richardson & Peatfield, 1987), but abnormal cholinergic innervation in asthma has not been definitively proven. Nicotine absorption in blood (Schneider, Lunell, Olmstead, & Fagerstrom, 1996; Schneider, Olmstead, Franzon, & Lunell, 2001) could influence airway innervation, but inhaled nicotine certainly has the potential to directly target airway epithelium, ASM and fibroblasts: cell types that play critical roles in asthma (Davies, 2014; Loxham, Davies, & Blume, 2014; Pain et al., 2014; Parulekar, Atik, & Hanania, 2014; Prakash, 2013). Yet, effects of inhaled nicotine or even mechanisms of nicotine action via nAChRs on normal resident cells of the airway, particularly ASM are not well-studied (Hahn, Lang, Bleicher, Zwerenz, & Rausch, 1992; Hartiala, Mapp, Mitchell, & Gold, 1985; He et al., 2014; Hong & Lee, 1996; Jiang et al., 2014). Here, this likely reflects assumptions that nAChRs are involved in airway reflexes, while ASM have only mAChRs (Belmonte, 2005; Canning, 2006; Canning & Fischer, 2001; Pieper, 2012; Racke & Matthiesen, 2004; Rogers, 2002; van der Velden & Hulsmann, 1999). In this regard, there is also no information on how nicotine influences asthmatic airways, or whether/how inflammation alters nAChR expression and signaling in asthma. The present study provides initial evidence for nAChR expression in human ASM, showing increased expression in asthmatics and smokers, as well as with pro-inflammatory cytokines that could contribute to increased cholinergic responses directly at the level of the ASM.

There are 16 nAChR subunits ( $\alpha 1-7$ ,  $\alpha 9-10$ ,  $\beta 1-4$ ,  $\delta$ ,  $\epsilon$ ,  $\gamma$ ) with different pentameric configurations (Changeux & Taly, 2008; Fasoli & Gotti, 2015; Leonard & Bertrand, 2001; Zoli, Pistillo, & Gotti, 2015), but “muscle-type” subunits ( $\alpha 1$ ,  $\beta 1$ ,  $\delta$ ,  $\epsilon$ ,  $\gamma$ ) known in skeletal muscle are unlikely to be involved in the airway. Indeed, our data in human ASM show no mRNA expression for these subunits. Among “neuronal” subunits,  $\alpha 7$  and  $\alpha 9$  form homopentameric nAChRs, and importantly,  $\alpha 7$ nAChRs show high  $\text{Ca}^{2+}$  permeability (Leonard & Bertrand, 2001), and are thus appealing in the context of airway reactivity.  $\alpha 7$ nAChR is expressed in neurons (Albuquerque et al., 2000), epithelium (Y. Wang et al., 2001), endothelium (Y. Wang et al., 2001), macrophages (H. Wang et al., 2003), and arterial smooth muscle (Bruggmann, Lips, Pfeil, Haberberger, & Kummer, 2002). In epithelium,  $\alpha 4$ ,  $\alpha 5$  and  $\alpha 7$  levels correlate with lung function (Prakash, 2016), but nicotine enhances only  $\alpha 7$ . There are limited data on nAChR profiles in ASM with rat ASM showing high  $\alpha 7$ nAChR (Jiang et al., 2014). In this regard, our data showing largely  $\alpha 7$ nAChR in the human ASM is consistent. Here, we have used highly specific  $\alpha$ BTX for detection of surface  $\alpha 7$ nAChR in human ASM cells. We further show  $\alpha 4$ nAChR expression, albeit much lower compared to  $\alpha 7$ nAChR. Thus, in terms of human ASM, it is likely that  $\alpha 7$ nAChR and perhaps to a lesser extent  $\alpha 4$ nAChR are the key subunits that may be important towards

functional cholinergic or nicotinic effects. Our Co-IP findings suggest that there is an interaction of  $\alpha 4$  with  $\alpha 7$  and  $\beta 2$  subunits. These interactions may be functionally important for  $\alpha 7$ -mediated effects of nicotine in asthma. However, such functional effects remain to be investigated.

We find that  $\alpha 7$ nAChR is substantially increased by TNF $\alpha$  and IL-13: cytokines relevant to asthma (Lerner et al., 2015; Lin et al., 2010; Rincon & Irvin, 2012; Svenningsen & Nair, 2017), and known to affect ASM (Ammit et al., 2007; De, Zelazny, Souhrada, & Souhrada, 1995; Delmotte & Sieck, 2015; Guedes et al., 2015; Jia et al., 2013; D. Kaur et al., 2015; Matsumoto et al., 2012; Robinson et al., 2015; Sathish et al., 2014). We further find that  $\alpha 7$ nAChR is increased in ASM from both asthmatics and smokers. Thus, regardless of what function  $\alpha 7$ nAChR may have in ASM, such effects are likely to be exacerbated in inflammation relevant to both asthma and cigarette smoke exposure. In contrast,  $\alpha 4$ nAChR expression is not substantially altered in asthmatics and only slightly enhanced by cytokines. Whether this is functionally important remains to be determined. Thus, at least on the human ASM cellular level,  $\alpha 7$ nAChR may be particularly important.

How upstream regulation of ASM  $\alpha 7$ nAChR occurs with inflammation or in asthma has not been previously explored. This is an important aspect since intracellular  $\alpha 7$ nAChR is non-responsive to nicotine and presumably ACh unless nAChR- and  $\alpha 7$ -specific ER chaperones “resistance to inhibitor of cholinesterase” (RIC-3) (Ben-David et al., 2016; Ben-David & Treinin, 2017; Cohen Ben-Ami et al., 2009; Millar, 2008) and NACHO (S. Gu et al., 2016; Matta et al., 2017; Rex, Shukla, Gu, Brecht, & DiSepio, 2017) are expressed, and increase plasma membrane levels of the receptor. Ongoing studies in human ASM do show RIC-3 and NACHO (Borkar, Sathish, Pabelick, unpublished observations). Nonetheless, our immunofluorescence studies do show plasma membrane expression of  $\alpha 7$ nAChR suggesting functional receptor levels (although we recognize there is likely substantial intracellular expression as well). Whether such plasma membrane  $\alpha 7$ nAChR or the associated chaperones are increased in inflammation or asthma remains to be determined. Regardless, our studies do find that cytokines can regulate  $\alpha 7$ nAChR expression via multiple signaling pathways all relevant to airway inflammation (Berair & Brightling, 2014; Berair, Saunders, & Brightling, 2013; Hirota & Martin, 2013; Manuyakorn, Howarth, & Holgate, 2013; Noble et al., 2014; Prakash, 2013, 2016). Here, pathways such as NF $\kappa$ B and AP-1 (relevant to TNF), STAT6 (relevant to IL-13) and CREB appear to be important. Interestingly these pathways are also downstream effectors of nicotine at least in immune cells (Bregeon et al., 2011; Gahring, Myers, Dunn, Weiss, & Rogers, 2017; D. C. Lam et al., 2016; Narumoto et al., 2010; Narumoto et al., 2013; Sitapara et al., 2014; Su, Matthay, & Malik, 2010). Thus, it is possible that  $\alpha 7$ nAChR activation in airways promotes inflammatory signaling. Indeed, nicotine has been shown to have pro-proliferative, pro-inflammatory and anti-apoptotic activity in rat ASM (He et al., 2014).

In summary, the present study shows that  $\alpha 7$ nAChR is the major nicotinic receptor expressed in human ASM, with its expression significantly increased in asthmatics and smokers, as well as with inflammation. Thus,  $\alpha 7$ nAChR may play a key role in ASM hyperresponsiveness and other functional changes that occur in asthma and with nicotine or cigarette smoke exposure.



## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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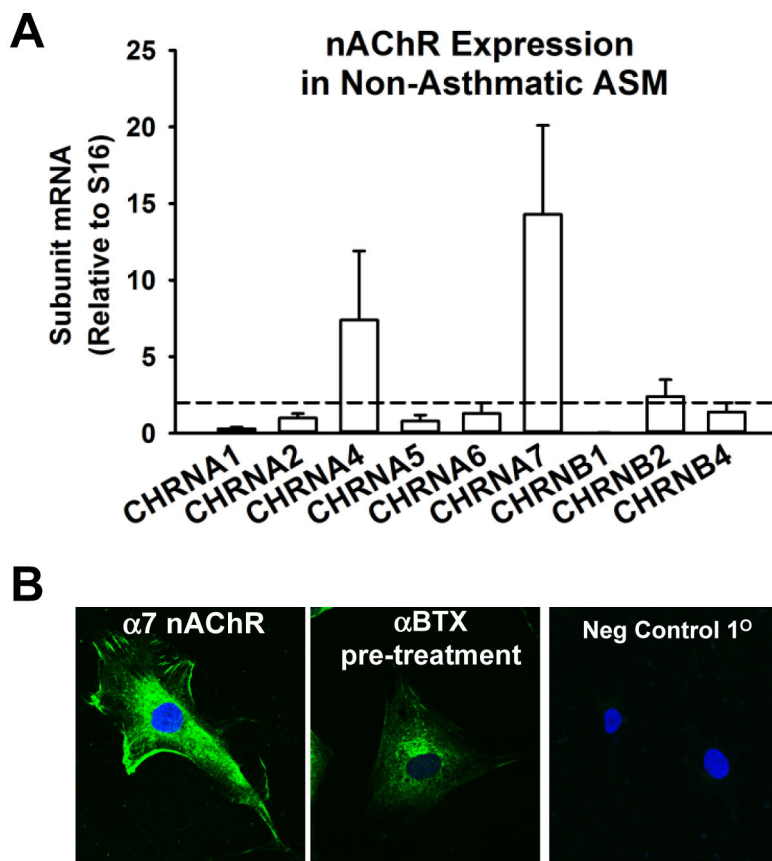
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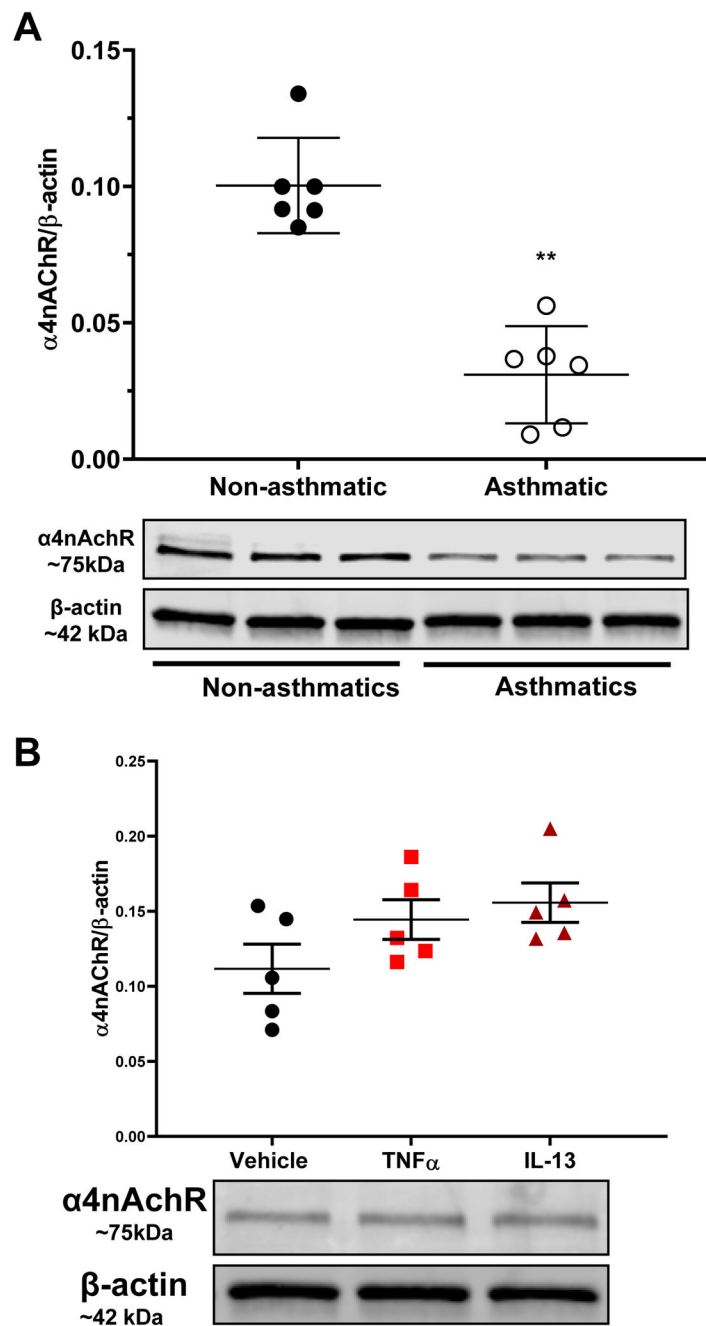
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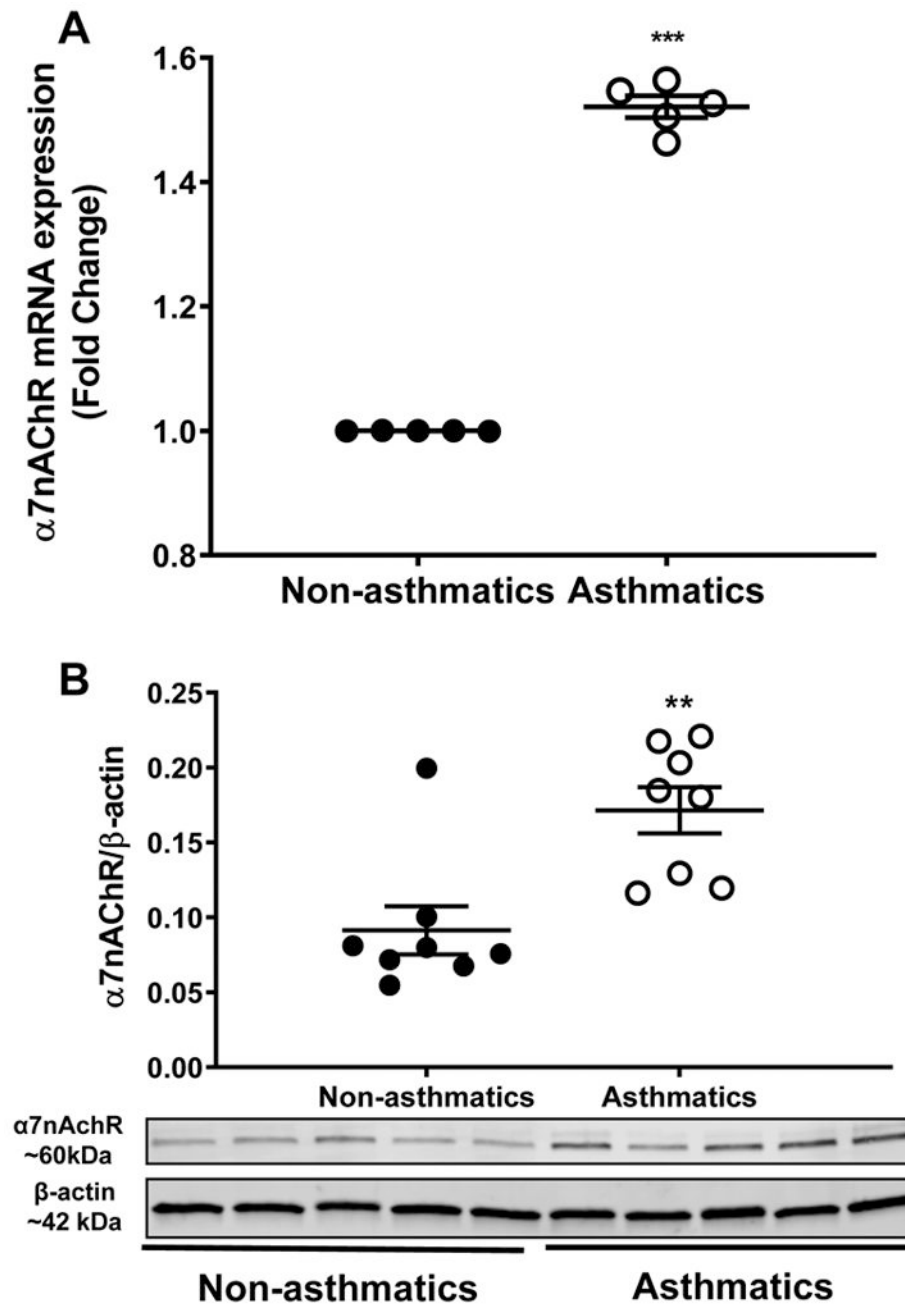
**Figure 1:**

Expression of nicotinic cholinergic receptor (nAChR) subunits in human airway smooth muscle (ASM) cells. mRNA studies in non-asthmatic human ASM cells show significantly higher expression of the  $\alpha 4$  and particularly  $\alpha 7$  subunits with small expression of  $\beta 2$  subunit, but otherwise minimal to no expression of other subunits (A).  $\alpha 7$ nAChR is expressed on ASM cell plasma membrane, evidenced by immunofluorescence (green) that is prevented by alpha-bungarotoxin ( $\alpha$ BTX) binding of the receptor (B). Data represented as mean  $\pm$  SEM (N=4).

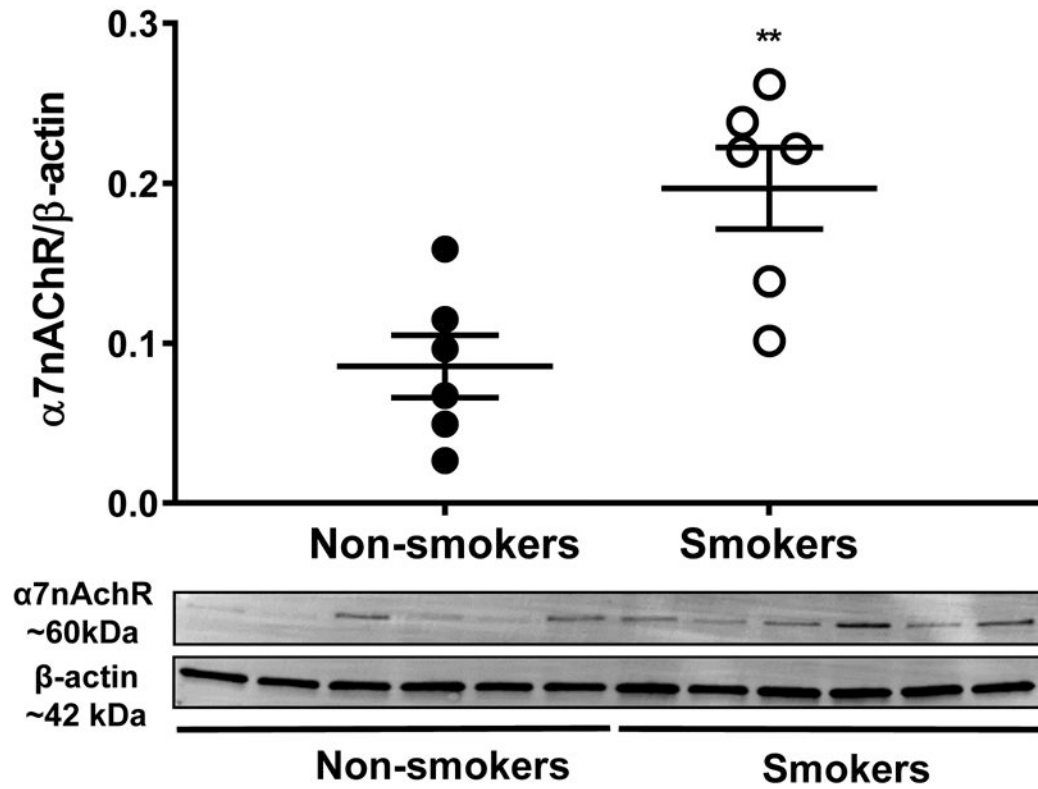




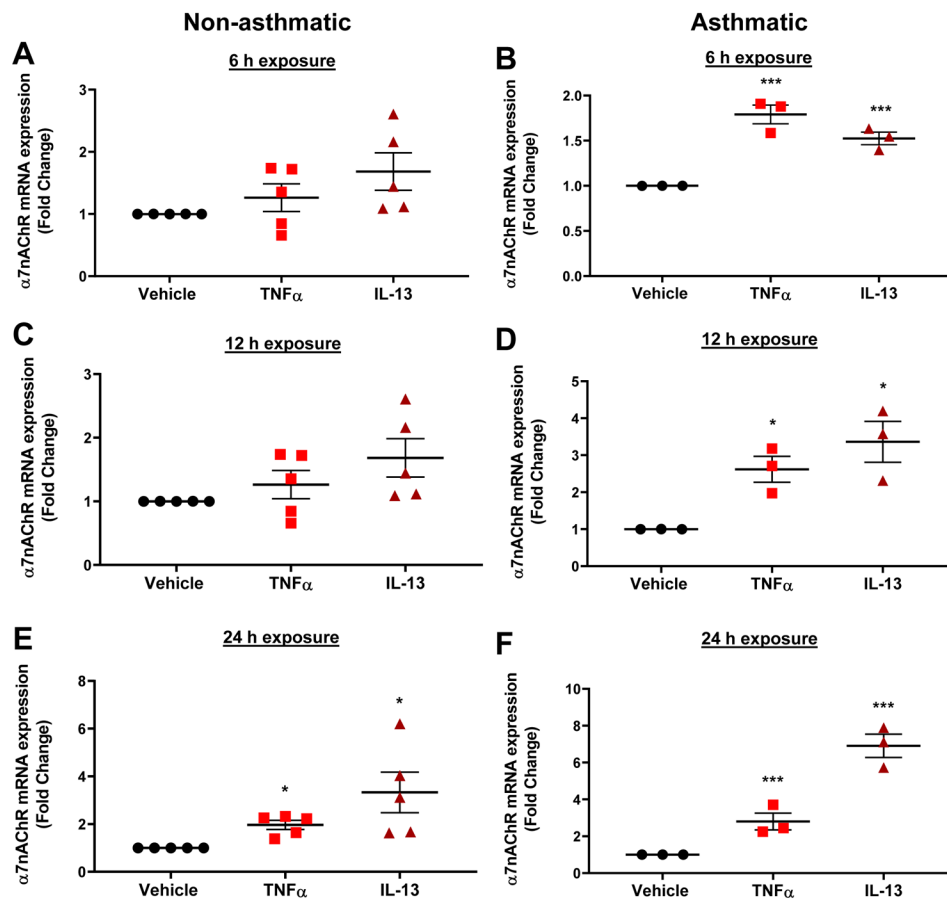
**Figure 2:**  $\alpha 4nAChR$  in human ASM cells. Western blotting showed that expression of  $\alpha 4nAChR$  is lower in asthmatic ASM compared to non-asthmatics (A). Exposure of non-asthmatic ASM cells to pro-inflammatory cytokines (20ng/ml  $TNF\alpha$  or 50ng/ml IL-13) resulted in only slight increases in  $\alpha 4nAChR$  compared to vehicle (B). \*\* $p < 0.05$  vs. non-asthmatics. Data represented as mean  $\pm$  SEM (N=5-6).



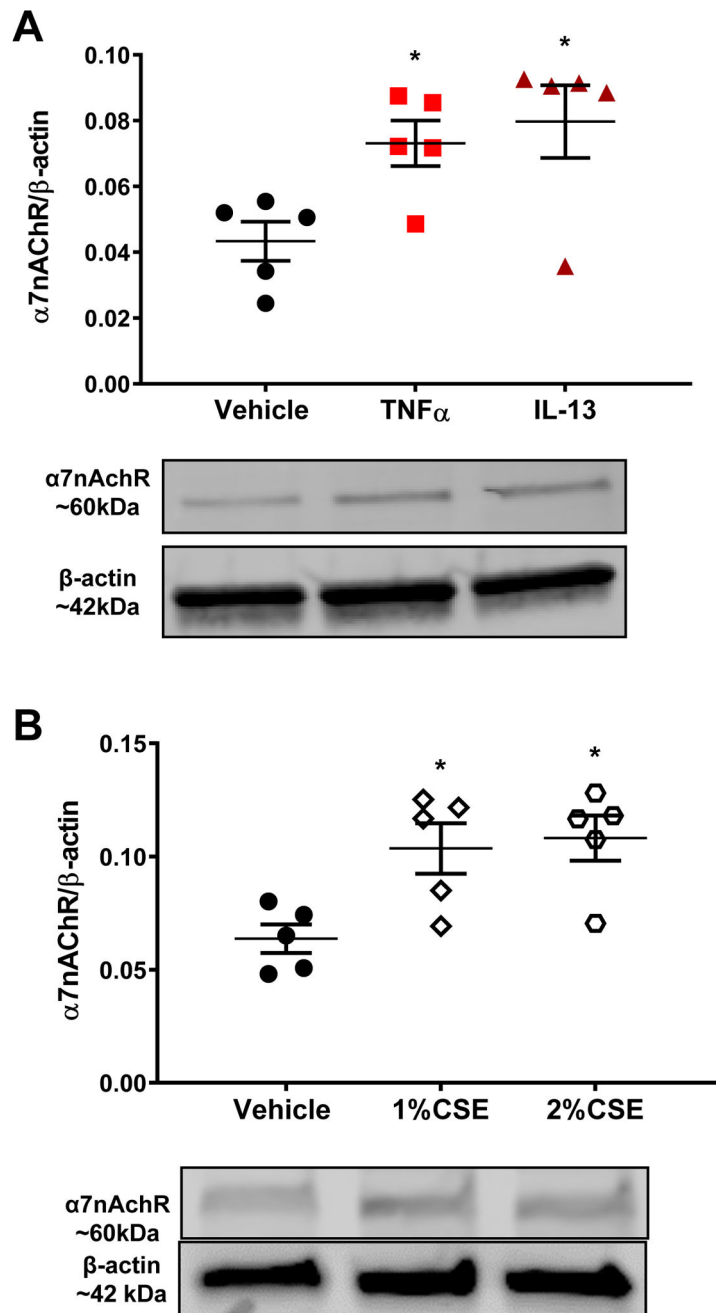
**Figure 3:** α7nAChR in human ASM cells. qRT-PCR (A) and western blotting (B) of ASM from non-asthmatics vs. asthmatics showed higher expression of α7nAChR in asthmatics. \*\* $p < 0.05$ , \*\*\* $p < 0.001$  vs. non-asthmatics. Data represented as mean ± SEM (N=5-8).



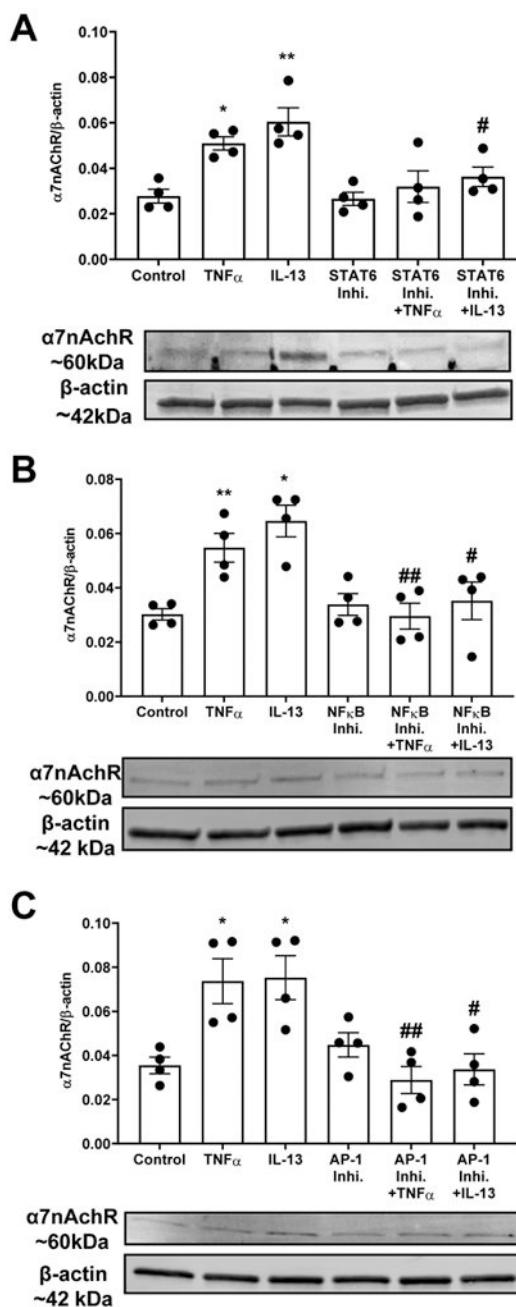
**Figure 4:**  $\alpha 7nAChR$  in ASM of smokers. Western analysis showed higher expression of  $\alpha 7nAChR$  in human ASM cells of smokers compared to non-smokers. \*\* $p < 0.01$  vs. Non-smokers. Data represented as mean  $\pm$  SEM (N=6).



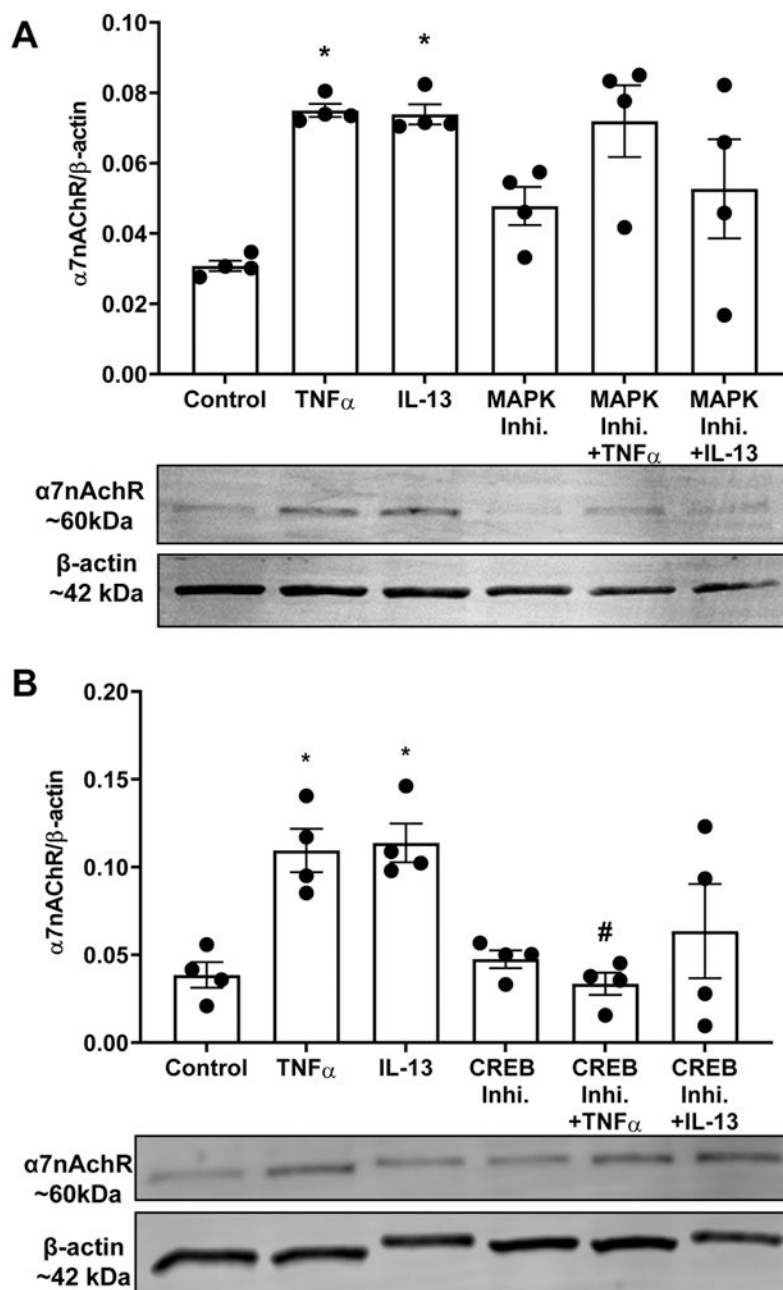
**Figure 5:** Effect of inflammatory cytokines on  $\alpha 7$ nAChR in human ASM cells. qRT-PCR analysis showed that in non-asthmatic ASM, 24h of TNF $\alpha$  or IL-13 exposure significantly increased  $\alpha 7$ nAChR expression (E) while 6 h or 12 h only trended towards an increase for either cytokine (A and C). On the other hand, expression of  $\alpha 7$ nAChR was significantly increased for all time points for asthmatic ASM cells (B, D, F). \* $p < 0.05$ , \*\*\* $p < 0.001$  vs. vehicle. Data represented as mean  $\pm$  SEM (N=3-5).



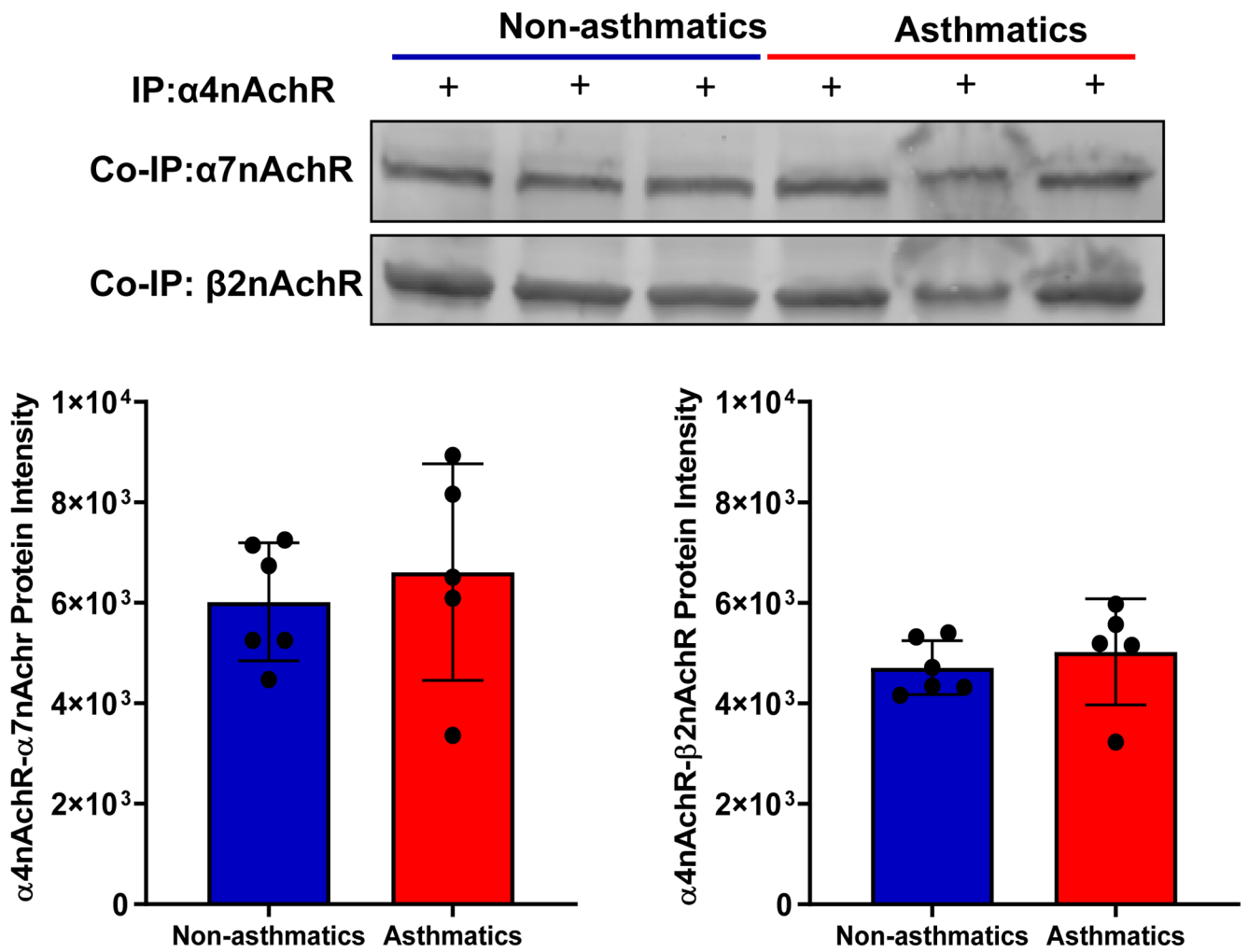
**Figure 6:** Effect of inflammatory cytokines and cigarette smoke on  $\alpha 7$ nAChR in human ASM. The effect of pro-inflammatory cytokines (TNF $\alpha$  and IL-13) and cigarette smoke extract (CSE; 1% and 2%) on  $\alpha 7$ nAChR was explored by western blotting. All these insults resulted in significant increase in  $\alpha 7$ nAChR expression. \* $p < 0.05$  vs. vehicle. Data represented as mean  $\pm$  SEM (N=5).



**Figure 7:** Inflammation-associated nuclear signals regulate  $\alpha 7nAChR$  in human ASM. In non-asthmatic human ASM cells, pharmacological inhibition of STAT6 (20 nM AS151749; A), NF $\kappa$ B (20  $\mu$ M SN50; B) or AP-1 inhibitor (1  $\mu$ M SR1302; C) significantly blunted the enhancing effects of TNF $\alpha$  or IL-13 on  $\alpha 7nAChR$ , with STAT6 inhibition showing specificity for IL-13. \* $p$ <0.05, \*\* $p$ <0.01 vs. vehicle, # $p$ <0.05, ## $p$ <0.01 vs. respective inflammatory cytokine. Data represented as mean  $\pm$  SEM (N=4).



**Figure 8:** Inflammation-associated cytosolic signals regulate  $\alpha 7nAChR$  in human ASM. In non-asthmatic human ASM cells, pharmacological inhibition of p42/44 MAPK (2  $\mu$ M PD98059; A) or CREB (5  $\mu$ M 666-15; B) significantly blunted the enhancing effects of TNF $\alpha$  or IL-13 on  $\alpha 7nAChR$ . CREB inhibition showed specificity for TNF $\alpha$ . \* $p$ <0.05 vs. Vehicle, # $p$ <0.05 vs. respective inflammatory cytokine. Data represented as mean  $\pm$  SEM (N=4).



**Figure 9:**

$\alpha$ 4nAChR interacts with  $\alpha$ 7nAChR and  $\beta$ 2nAChR in human ASM. Immunoprecipitation with  $\alpha$ 4nAChR and subsequent co-immunoprecipitation of  $\alpha$ 7nAChR and  $\beta$ 2nAChR using western blot analyses revealed that  $\alpha$ 7nAChR and  $\beta$ 2nAChR interacts with  $\alpha$ 4nAChR in both non-asthmatic and asthmatic ASM cells with no significant difference observed between the groups. Data represented as mean  $\pm$  SEM (N=5-6).