



# COVID-19 and smoking: is nicotine the hidden link?

To the Editor:

LEUNG *et al.* [1] have recently published, in the *European Respiratory Journal*, a paper on the expression of angiotensin-converting enzyme II (ACE-2) in the small airway epithelia of smokers and COPD patients, discussing its effects on the risk of severe coronavirus disease 2019 (COVID-19). The authors found an increased expression of the ACE-2 gene in the airways of subjects with COPD and in current smokers. Indeed, a recent systematic review reporting data on the smoking habits of patients infected with severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), concluded that smoking may be associated with a negative progression of the disease and with the adverse outcome [2]. These conclusions were challenged in a correspondence by CAI [3] on the basis that a reliable mechanism explaining this association was missing. The need for these results to be supported by additional studies is quite clear, but we believe that a robust mechanistic explanation exists. Nicotine has a known influence on the homeostasis of the renin-angiotensin system (RAS) up-regulating the angiotensin-converting enzyme (ACE)/angiotensin (ANG)-II/ANG II type 1 receptor axis, and down-regulating the compensatory ACE-2/ANG-(1-7)/Mas receptor axis, contributing in turn to the development of cardiovascular and pulmonary diseases [4]. Different airway cells, such as bronchial epithelial cells, type II alveolar epithelial cells and interstitial lung fibroblasts, express nicotinic acetylcholine receptors (nAChR), specifically the  $\alpha 7$  subtype [5]. All these cells express components of the RAS [4]. In addition, nicotine increases the expression and/or activity of ACE in the lung [4], an increase which has been found also in the serum of smokers, and that required at least 20 min to return to control level [4]. ACE-2 serves as a physiologically relevant cellular entry receptor for SARS-CoV, for the human respiratory coronavirus NL63, and probably for SARS-CoV-2 [6]. ACE binds the SARS-CoV-2 S protein, and through its tissutal expression mediates the localisation and the efficiency of the infection [6]. Moreover, nicotine induces the epithelial-mesenchymal transition (EMT) [5, 7], a mechanism sufficient to allow “normal” differentiated cells to acquire the stem cell-like characteristics and properties. We planned experiments on human bronchial epithelial cells (HBEpC), obtained from Cell Applications Inc. ([www.cellapplications.com/product](http://www.cellapplications.com/product) number 502K-05a). Cells were maintained as adherent monolayer in complete bronchial/tracheal epithelial cell growth medium ([www.cellapplications.com/product](http://www.cellapplications.com/product)) at 37°C in 95% air/5% CO<sub>2</sub>, seeded at an initial density of  $7.5 \times 10^4$  cells·cm<sup>-2</sup>, and sub-cultured with a 0.25% trypsin-1 mM EDTA solution (Sigma-Aldrich, Milan, Italy) when cultures reached 80% confluence. HBEpC are derived from the surface epithelium of normal human bronchi non-diseased (*i.e.* asthma, COPD or type 2 diabetes). The morphology is consistent with epithelial origin, and is positive for epithelial cell marker cytokeratin 18. Semi-confluent HBEpC at fourth passage ( $7.5 \times 10^4$  cells·cm<sup>-2</sup>) were treated: 1) for 1 h with zero or  $1.0 \times 10^{-7}$  M nicotine (Sigma-Aldrich, Milan, Italy) dissolved in saline in complete medium; 2) with  $1.0 \times 10^{-6}$  M  $\alpha$ -Bungarotoxin ( $\alpha$ -BTX; Sigma-Aldrich, Milan, Italy) dissolved in saline, in the continued presence of nicotine at zero or  $1.0 \times 10^{-7}$  M for 1 h; 3) treated continuously with nicotine for additional passages, 1 passage every 48 h for a total of 16 passages. We showed, for the first time, that nicotine at  $1 \times 10^{-7}$  M (the concentration present on the alveolar lining fluids after one cigarette is in the range  $6 \times 10^{-6}$  to  $6 \times 10^{-5}$  M [5]) is able to increase ACE-2 (figure 1a) in HBEpC. Treatment with nicotine induces phospho-S6 ribosomal protein (Ser235/236), Akt1, phospho-Akt (Ser473), phospho-Akt (Thr308) and phospho-p44/42 MAPK (Thr202/Tyr204) (figure 1b). To verify the hypothesis that ACE-2 is induced by nicotine through  $\alpha 7$ -nAChR, HBEpC, at fourth passage, in the exponential growth phase, plated at a density of  $1 \times 10^6$  cells·mL<sup>-1</sup>, were incubated with  $\alpha 7$ -nAChR siRNA (0.1  $\mu$ g) diluted in 100  $\mu$ L of siRNA transfection medium. Transfection was performed as described by LI *et al.* [8], who transfected, successfully, HBE16 human airway epithelial cell line (unaffected cells). A clone of transfected HBEpC that did not express  $\alpha 7$ -nAChR proteins, also after



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**Nicotine via alpha7-nicotinic receptor induces ACE-2 overexpression in human bronchial epithelial cells (HBEpC)** <https://bit.ly/3eJ5b35>

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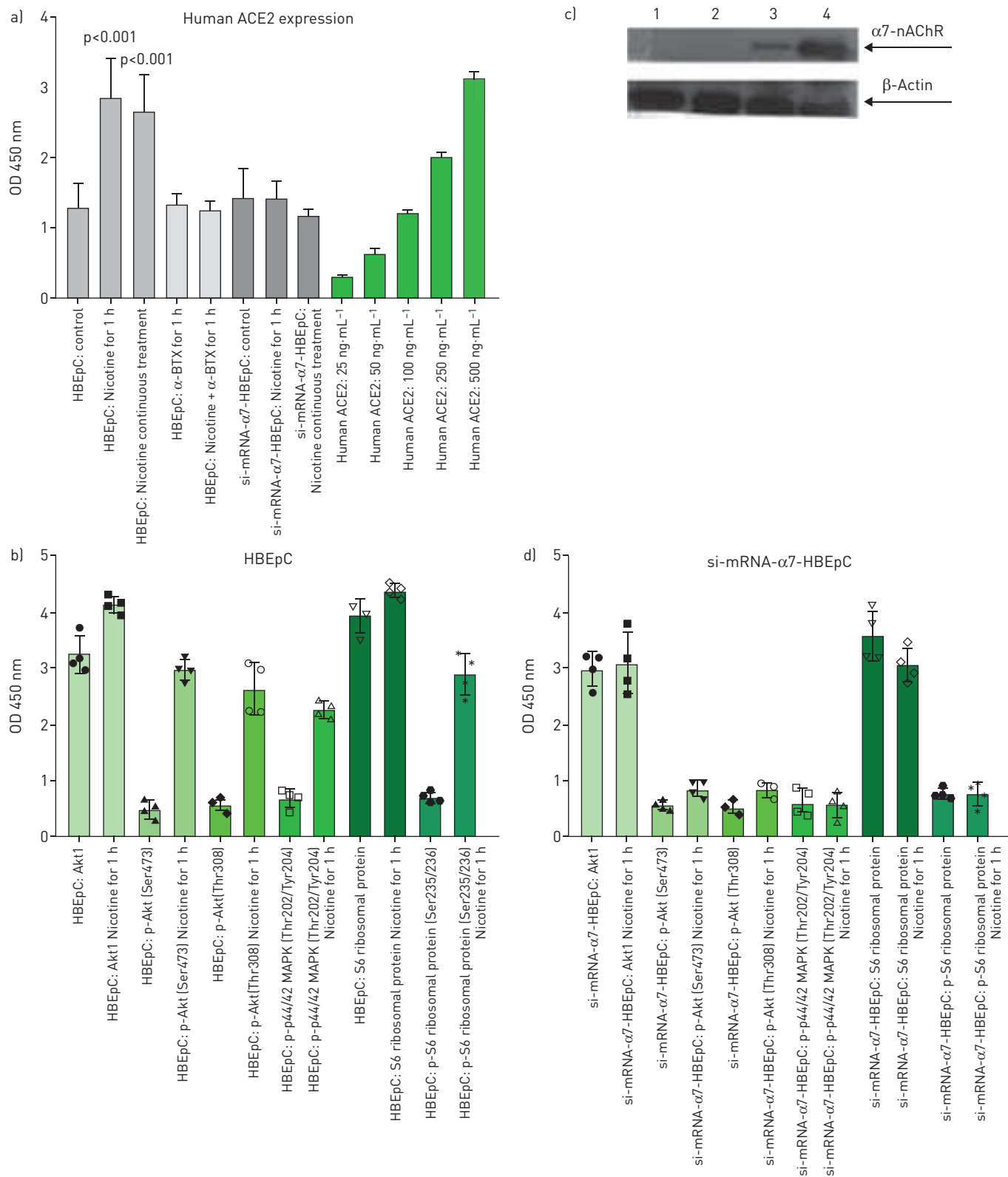


FIGURE 1 Effect of nicotine on human bronchial epithelial cells (HBEpC) or si-mRNA- $\alpha$ 7-HBEpC. a) Angiotensin-converting enzyme II (ACE-2) detection. ACE-2 was measured with human ACE-2 ELISA kit ab235649 ([www.abcam.com/human-ace2-elisa-kit-ab235649.html](http://www.abcam.com/human-ace2-elisa-kit-ab235649.html)) according to the manufacturer's instructions. Data are mean $\pm$ SEM; p-value was evaluated using t-test. The green plots are the human ACE-2 standard. Experiments were performed twice in triplicate. b) Induction of phospho-S6 ribosomal protein (Ser235/236), Akt1, phospho-Akt (Ser473), phospho-Akt (Thr308)

and phospho-p44/42 MAPK (Thr202/Tyr204) in HBEpC. Data were obtained using PathScan® cell growth Multi-target Sandwich ELISA kit n.7239 (Cell Signaling) PathScan® Cell Growth Multi-Target Sandwich ELISA Kit is a solid phase sandwich ELISA that combines the reagents necessary to detect endogenous levels of S6 ribosomal protein, phospho-S6 ribosomal protein [Ser235/236], Akt1, phospho-Akt [Ser473], phospho-Akt [Thr308] and phospho-p44/42 MAPK (Thr202/Tyr204). Data are mean $\pm$ SEM; p-value was evaluated using t-test. Experiments were performed twice in duplicate. c)  $\alpha$ 7-nAChR protein detection. Western blotting was performed as described previously [11]. Human  $\alpha$ 7-nAChR antibody NBP1-49348 was purchased from Novus Biologicals (www.novusbio.com). 1–2 si-mRNA- $\alpha$ 7-HBEpC treated with zero (lane 1) or  $1.0 \times 10^{-7}$  M nicotine (lane 2) for 1 h. 3–4 HBEpC treated with zero (lane 3) or  $1.0 \times 10^{-7}$  M nicotine (lane 4) for 1 h. Experiments were performed twice. d) as in panel b, but treated cells are si-mRNA- $\alpha$ 7-HBEpC.

treatment with nicotine (figure 1c), and is not able to induce phospho-S6 ribosomal protein (Ser235/236), Akt1, phospho-Akt (Ser473), phospho-Akt (Thr308) and phospho-p44/42 MAPK (Thr202/Tyr204) after nicotine treatment (figure 1d), was selected for further experiments. Nicotine did not induce ACE-2 in this clone (si-mRNA- $\alpha$ 7-HBEpC) (figure 1a). This observation supports the hypothesis that ACE-2 increase is specifically mediated by  $\alpha$ 7-nAChR. Moreover, when HBEpC were incubated simultaneously with nicotine and  $\alpha$ -BTX, an  $\alpha$ 7 nicotine antagonist [9], no induction of ACE-2 was observed (figure 1d). Importantly, treatment with nicotine,  $\alpha$ -BTX or with the combination is not cytotoxic (data not shown). On these bases, we suggest that smoking may promote cellular uptake mechanisms of SARS-CoV-2 through  $\alpha$ 7-nAChR signalling. A possible  $\alpha$ 7-nAChR down-stream mechanism may be the induction of phospho-Akt and phospho-p44/42 MAPK. This mechanism was hypothesised, partially, by OLDS and KABBANI [10] on their schematic model explaining how nicotine exposure increases the risk of SARS-CoV-2 entry into lung cells.  $\alpha$ 7-nAChR is present both in neuronal and non-neuronal cells (*i.e.* lung, endothelial, lymphocyte); consequently, smoking may impact COVID-19 pathophysiology and clinical outcome in several organ systems, including the brain.

**Patrizia Russo<sup>1,2</sup>, Stefano Bonassi<sup>1,2</sup>, Robertina Giacconi<sup>3</sup>, Marco Malavolta<sup>3</sup>, Carlo Tomino<sup>4</sup> and Fabrizio Maggi<sup>5,6</sup>**

<sup>1</sup>Clinical and Molecular Epidemiology, IRCSS San Raffaele Pisana, Rome, Italy. <sup>2</sup>Dept of Human Sciences and Quality of Life Promotion, San Raffaele University, Rome, Italy. <sup>3</sup>Advanced Technology Center for Aging Research, Scientific Technological Area, Italian National Institute of Health and Science on Aging (INRCA), Ancona, Italy. <sup>4</sup>Scientific Direction, IRCSS San Raffaele Pisana, Rome, Italy. <sup>5</sup>Dept of Translational Research, University of Pisa, Pisa, Italy. <sup>6</sup>Virology Division, Pisa University Hospital, Pisa, Italy.

Correspondence: Patrizia Russo, Clinical and Molecular Epidemiology, IRCSS San Raffaele Pisana, Via di Val Cannuta, 247, I-00166 Rome, Italy. E-mail: patrizia\_russo@hotmail.it

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# COVID-19 and nicotine as a mediator of ACE-2

From the authors:

We recently reported that current smokers and those with COPD had higher airway epithelial cell expression of the angiotensin-converting enzyme II (ACE-2) viral entry receptor [1]. We thus read with great interest the work of P. Russo and co-workers, which proposes a mechanism for this finding, namely that this upregulation is mediated by nicotine exposure specifically through the  $\alpha 7$  subtype of nicotine acetylcholine receptors ( $\alpha 7$ -nAChR). While exposure to increasing concentrations of nicotine caused epithelial cells to increase ACE-2 levels, subsequent gene silencing of  $\alpha 7$ -nAChR appeared to significantly dampen this response. A secondary transcriptome sequencing analysis of our cohort (consisting of 42 subjects who underwent bronchoscopy for epithelial cell brushings [1]) reveals evidence in support of this hypothesis. We found that airway epithelial cell expression of *CHRNA7*, encoding  $\alpha 7$ -nAChR, was significantly correlated with the expression of *ACE2* (Pearson  $r=0.54$ ,  $p=2.31\times 10^{-8}$ ) (figure 1). There was significantly higher *CHRNA7* expression in those with COPD ( $2.75\pm 0.73$  versus  $2.14\pm 0.43$  in those without COPD;  $p=1.47\times 10^{-4}$ ), with a trend towards higher expression in current smokers compared to former and never smokers ( $2.86\pm 0.92$  in current smokers,  $2.35\pm 0.57$  in former smokers, and  $2.27\pm 0.45$  in never smokers;  $p=6.16\times 10^{-2}$ ). *CHRNA7* was also negatively correlated with forced expiratory volume in 1 s percent predicted (Pearson  $r=-0.37$ ,  $p=2.83\times 10^{-4}$ ). Interestingly, *CHRNA7* was positively if weakly correlated with body mass index (Pearson  $r=0.14$ ,  $p=6.31\times 10^{-3}$ ), raising the intriguing possibility that nicotine receptor mediation of ACE-2 may also be related to why obese individuals have made up a considerable proportion of coronavirus disease 2019 (COVID-19) cases [2].

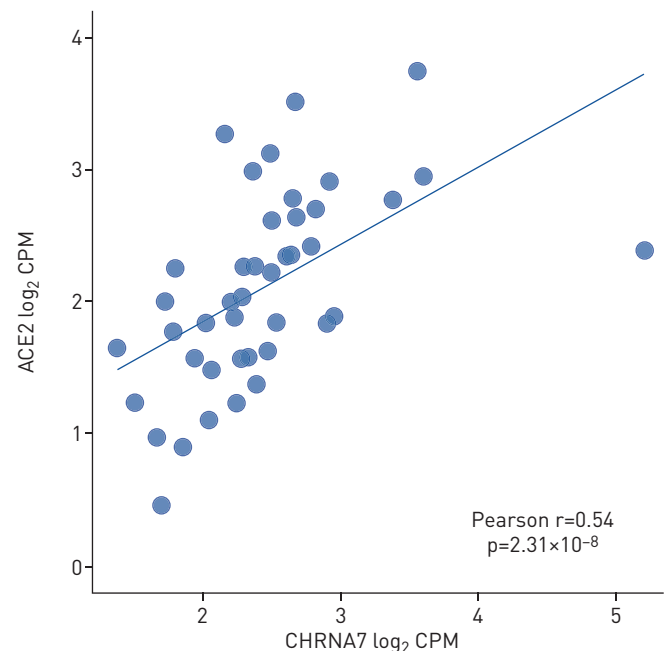


FIGURE 1 Transcriptome profiles generated through RNA-Seq of airway epithelial cells demonstrated a significant positive correlation between *ACE2* and *CHRNA7* expression.

Together, these data further help to characterise the connections between airway epithelial ACE-2, and  $\alpha 7$ -nAChR, and the unique vulnerability of patients with COPD to severe COVID-19.  $\alpha 7$ -nAChR's

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 **$\alpha 7$ -nAChR may upregulate ACE-2** <https://bit.ly/2xS0cfT>

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widespread abundance in the human body, from neuronal tissue to immune cells to the lung and digestive tract, and its various roles in diseases such as schizophrenia [3], Alzheimer's disease [4] and Parkinson's disease [5] has meant that considerable work has already been done to target  $\alpha 7$ -nAChR as a therapeutic modality. As an example,  $\alpha 7$ -nAChR antagonists for the purpose of smoking cessation have long been proposed [6] and the idea of potentially repurposing these compounds for a pandemic with few therapeutic options currently available is certainly appealing. Whether  $\alpha 7$ -nAChR-selective antagonists, such as methyllycaconitine [7] and  $\alpha$ -conotoxin [8], can meaningfully alter ACE-2 expression to prevent severe acute respiratory syndrome coronavirus 2 entry into the airway epithelium seems the next logical investigation in our furious pursuit for better therapeutics.

**Janice M. Leung<sup>1,2</sup>, Chen Xi Yang<sup>1</sup> and Don D. Sin<sup>1,2</sup>**

<sup>1</sup>Centre for Heart Lung Innovation, University of British Columbia, Vancouver, BC, Canada. <sup>2</sup>Division of Respiratory Medicine, Dept of Medicine, St. Paul's Hospital, Vancouver, BC, Canada.

Correspondence: Don D. Sin, Division of Respiratory Medicine, Dept of Medicine, St. Paul's Hospital, 1081 Burrard Street, Vancouver, BC V6Z1Y6, Canada. E-mail: Don.Sin@hli.ubc.ca

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