



“Expression of endocannabinoid system components in human airway epithelial cells: impact of sex and chronic respiratory disease status”. Matthew F. Fantauzzi, Jennifer A. Aguiar, Benjamin J-M. Tremblay, Michael J. Mansfield, Toyoshi Yanagihara, Abiram Chandiramohan, Spencer Revill, Min Hyung Ryu, Chris Carlsten, Kjetil Ask, Martin Stämpfli, Andrew C. Doxey and Jeremy A. Hirota. *ERJ Open Res* 2020; 6: 00128-2020.

This article was originally published with an error in figure 3, in which part b was a duplication of part a. The corrected figure is shown below and has been corrected in the article itself.

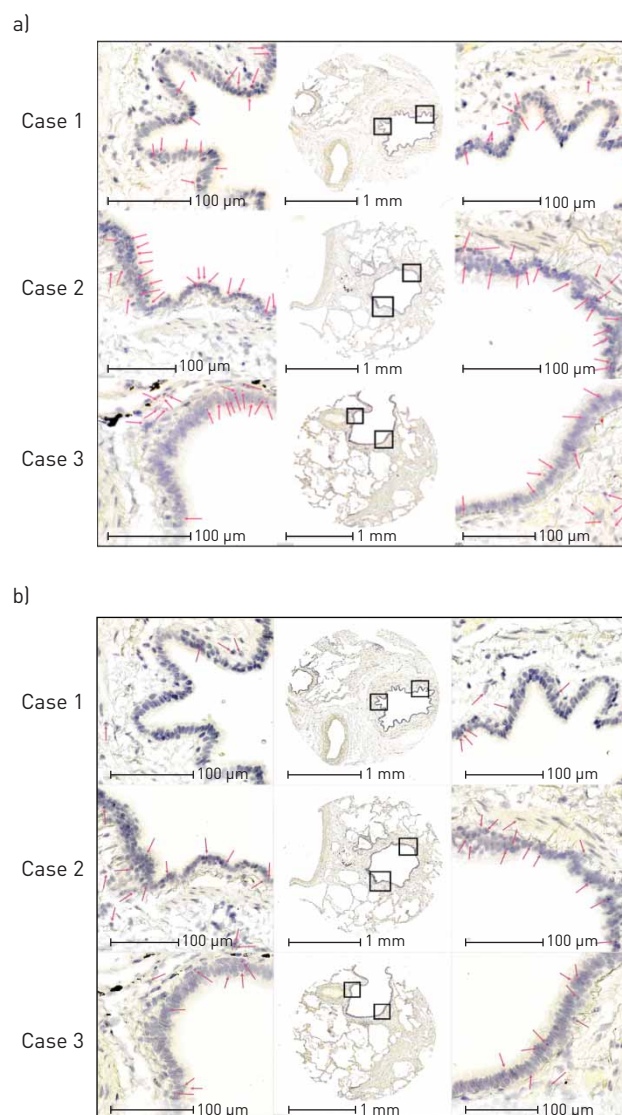


FIGURE 3 *In situ* hybridisation detection of *CNR1* and *CNR2* gene transcripts in human airway epithelial cells. *In situ* hybridisation of a) *CNR1* and b) *CNR2* in three patient donors representative of n=10. Serial sections of each of the three cases were stained for *CNR1* and *CNR2*. Low magnification images are in the centre of a and b, with high-powered magnification regions of interest on either side highlighted by black boxes. Pink arrows correspond to positive puncta representative of mRNA transcript.

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