





Generalised mosaicism for *TSC2* mutation in isolated lymphangioleiomyomatosis

Barbara Ogórek¹, Lana Hamieh¹, Kathryn Lasseter¹, Shefali Bagwe¹, Tania Machado², Carmen Herranz-Ors³, Aaron R. Thorner⁴, Anwesha Nag⁴, Peter Gulleman⁵, Krinio Giannikou¹, Lisa R. Young⁵, Miquel Àngel Pujana³, Thomas N. Darling⁶, Souheil El-Chemaly¹, Joel Moss², Elizabeth P. Henske¹ and David J. Kwiatkowski¹

Affiliations: ¹Dept of Medicine, Brigham and Women's Hospital, Boston, MA, USA. ²Pulmonary Branch, National Heart, Lung, and Blood Institute (NHLBI), Bethesda, MD, USA. ³ProCURE, Catalan Institute of Oncology (ICO), Bellvitge Institute of Biomedical Research (IDIBELL), Barcelona, Spain. ⁴Center for Cancer Genome Discovery, Dana-Farber Cancer Institute, Boston, MA, USA. ⁵Division of Pediatric Pulmonary Medicine, Dept of Pediatrics, Vanderbilt University School of Medicine, Nashville, TN, USA. ⁶Dept of Dermatology, Uniformed Services University of the Health Sciences, Bethesda, MD, USA.

Correspondence: David J. Kwiatkowski, Division of Pulmonary and Critical Care Medicine and Genetics, Dept of Medicine, Brigham and Women's Hospital, 20 Shattuck Street Thorn 826, Boston, MA 02115, USA. E-mail: dk@rics.bwh.harvard.edu

@ERSpublications

Analysis of plasma cell-free DNA from 61 sporadic lymphangioleiomyomatosis (LAM) patients identified generalised mosaicism for a TSC2 mutation in one, suggesting that some sporadic LAM patients are occult generalised mosaics for TSC2 mutations http://bit.ly/2yLr0Ls

Cite this article as: Ogórek B, Hamieh L, Lasseter K, *et al.* Generalised mosaicism for *TSC2* mutation in isolated lymphangioleiomyomatosis. *Eur Respir J* 2019; 54: 1900938 [https://doi.org/10.1183/13993003.00938-2019].

This single-page version can be shared freely online.

To the Editor:

Lymphangioleiomyomatosis (LAM) is a rare, slowly progressive pulmonary disease causing cystic lung destruction and respiratory failure. It affects predominantly premenopausal women, and rarely men. It can occur as a sporadic condition (sporadic LAM) or in association with tuberous sclerosis complex (TSC) [1]. LAM is caused by biallelic inactivation of the tumour suppressor gene *TSC2* in LAM cells, which leads to hyperactivation of mammalian target of rapamycin complex (mTORC)1, resulting in anabolism and LAM cell proliferation [2]. Sirolimus and everolimus, mTORC1 allosteric inhibitors, have been shown to retard progression of LAM [3].

Copyright ©ERS 2019