

#### Hypomorphic mutation in SFTPB leads to adult pulmonary fibrosis

Tifenn Desroziers, Grégoire Prévot, Aurore Coulomb, Valérie Nau, Florence Dastot-Le Moal, Philippe Duquesnoy, Mélanie Héry, Aurélie Le Borgne, Serge Amselem, Marie Legendre, et al.

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# Hypomorphic mutation in SFTPB leads to adult pulmonary fibrosis

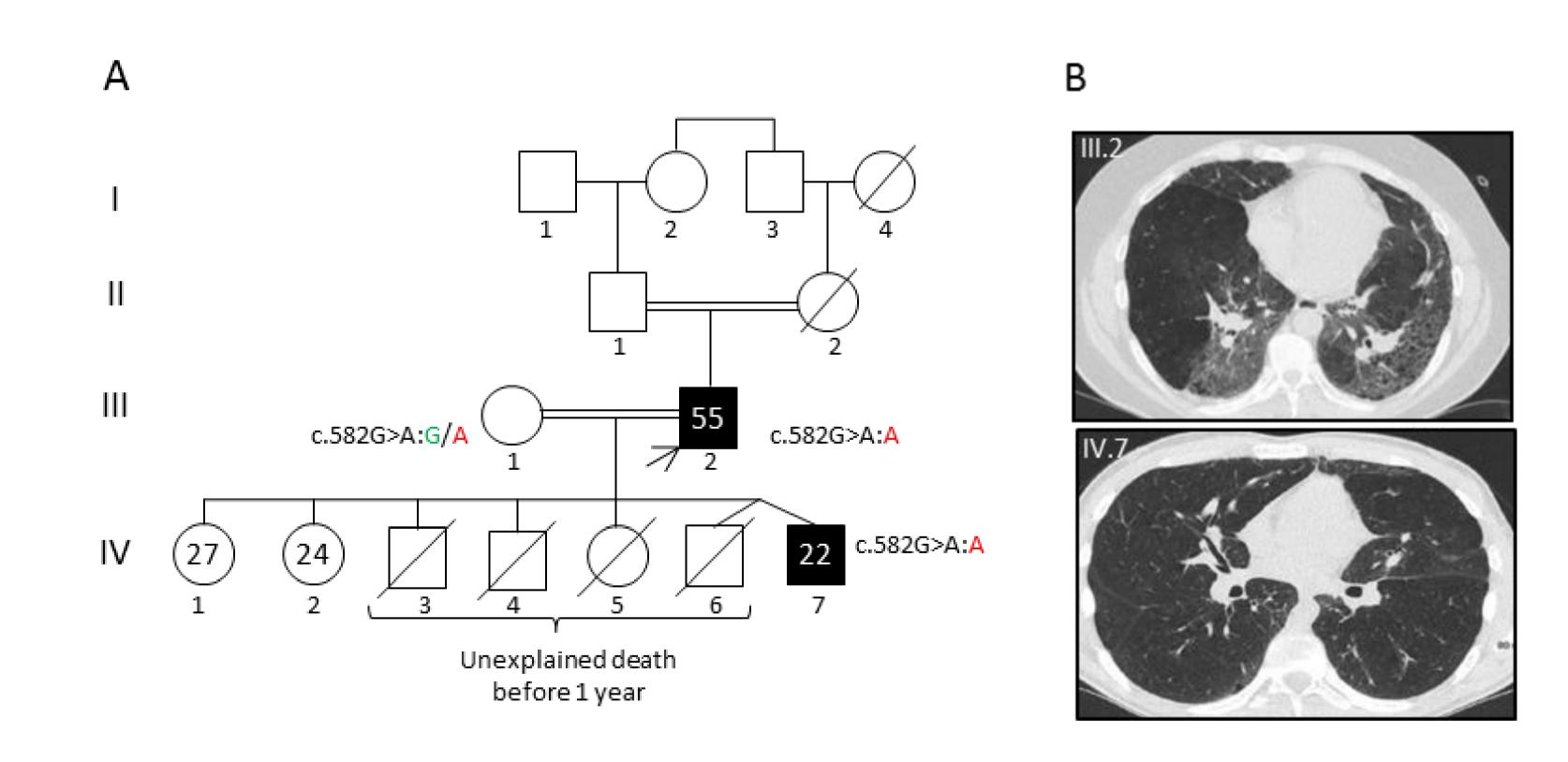
## Introduction -

- interstitial lung diseases (ILD) in **newborns** and exceptional survival in young children.
- The aim of the study was to investigate **the pathogenicity effect** of the variation **c.582G>A p.(Gln194=)**

## Results

**Case presentation** 

- The proband, homozygous for the variation (Fig.1A), presented a fibrosing ILD at 34 years. His thoracic CT-scan showed ground glass opacities and fibrosis. He received a bi-pulmonary transplantation at 51 years (Fig.1B).
- **Proband's son, also homozygous for the variation (Fig.1A) presented** alveolo-interstitial opacities and mild alveolar hemorrhage at 10 months. His thoracic CT-scan at 16 years showed mild ground glass opacities and mild ILD at 22 years (Fig.1B).



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Figure 1: (A) Genealogical tree of the described family. The genotypes are provided with the reference nucleotide in green and the pathogenic variant in red. (B) The CT-scan of the proband at 51 years (left) and his son at 16 years (right)

## **Conclusion and perspectives**

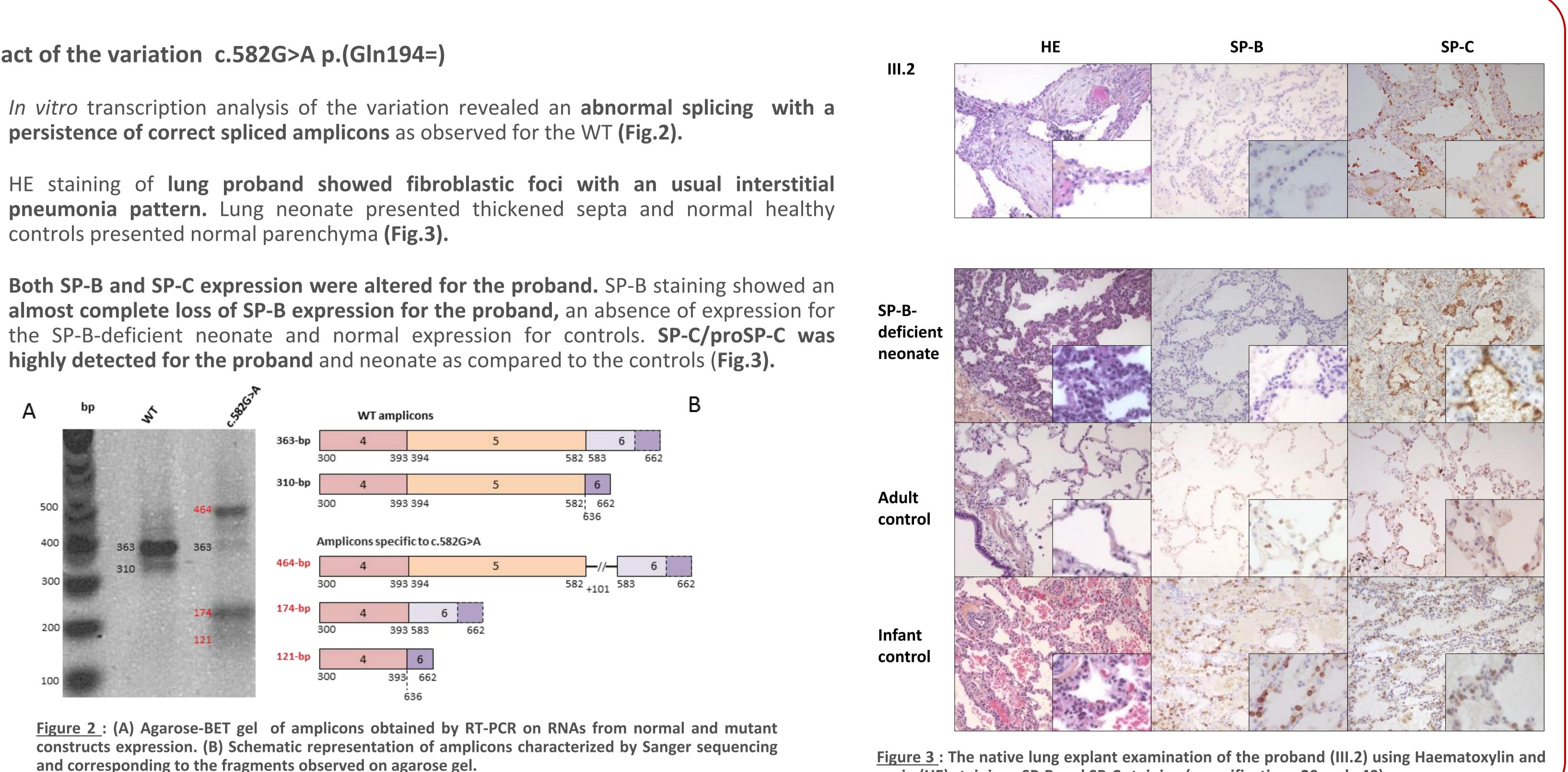
- explain the long-term survival with fibrosing ILD in the two related adults.

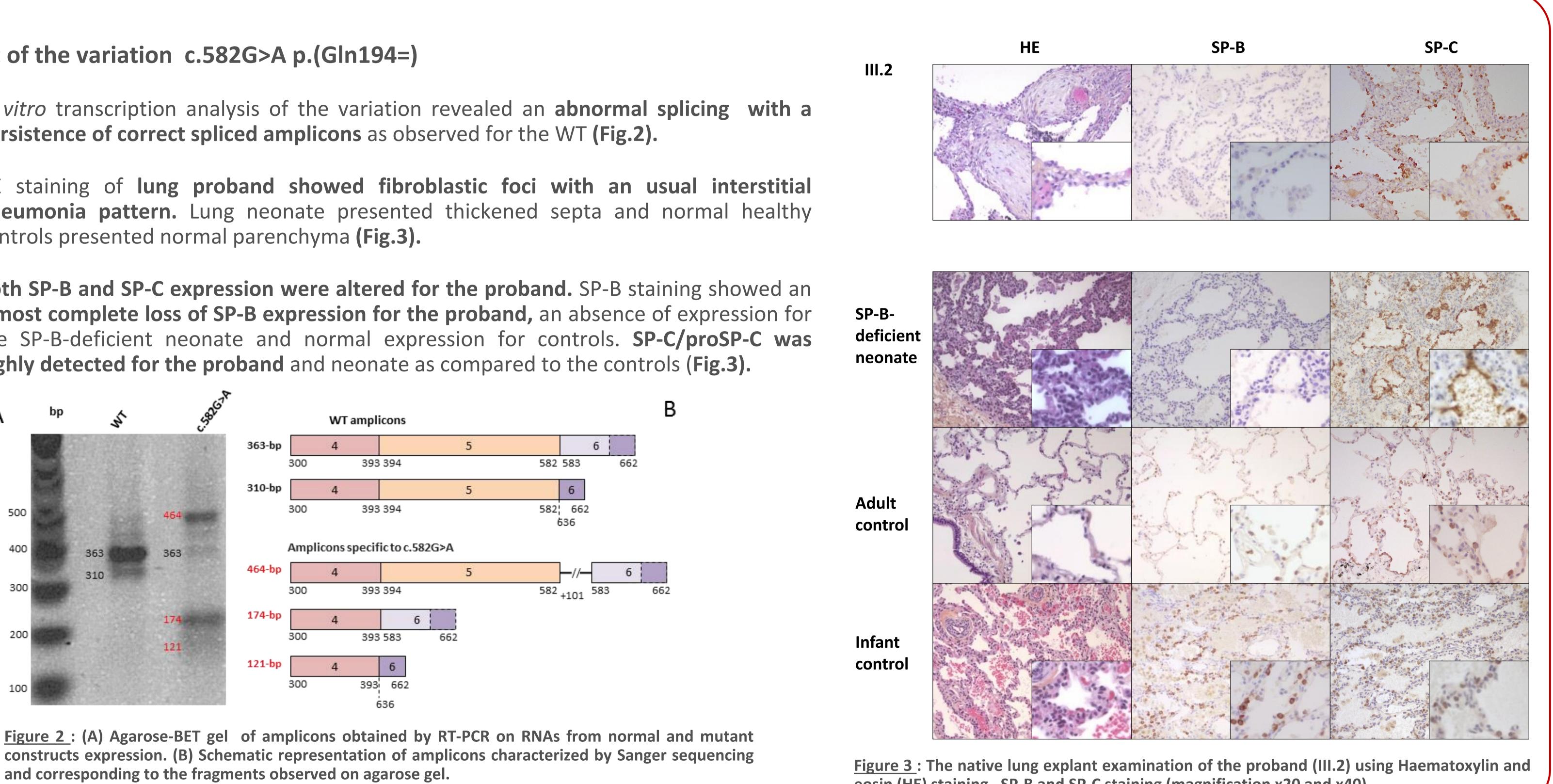


Surfactant protein (SP)-B deficiency due to bi-allelic pathogenic variation has been associated with fatal forms of

We herein report for the first time two related adults with pulmonary fibrosis due to a homozygous SFTPB variation.

### Impact of the variation c.582G>A p.(Gln194=)





The pathogenic variation c.582G>A p.(Gln194=) affected drastically splicing and leading to a residual SP-B expression, could

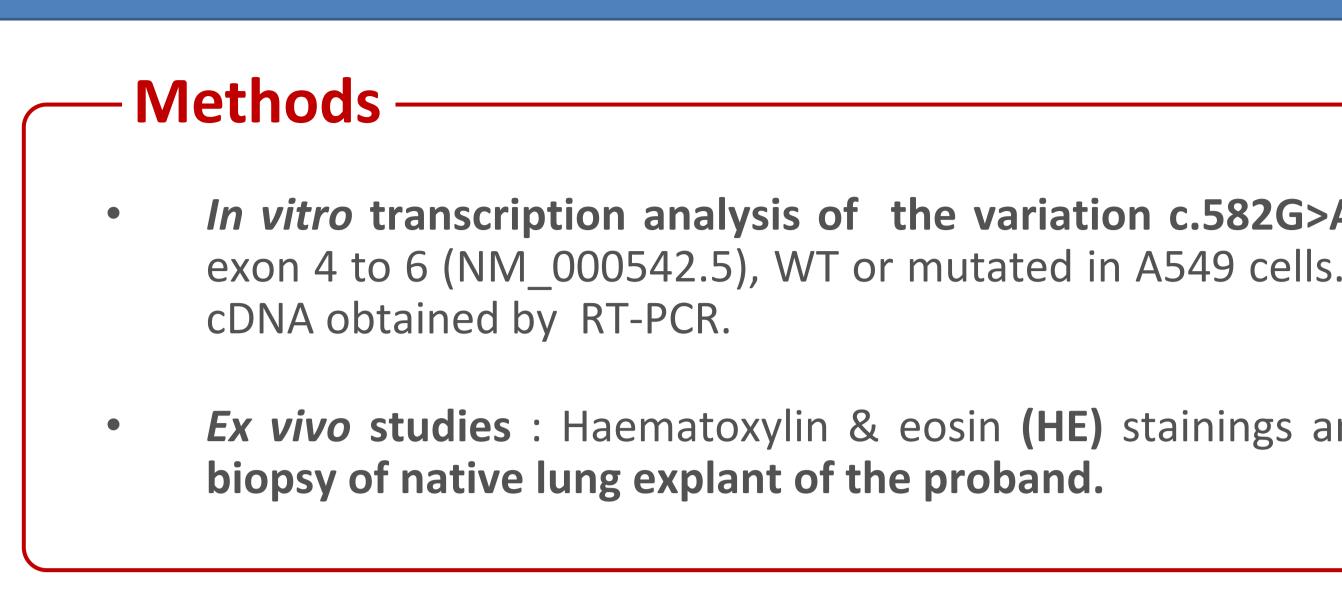
These findings lead to considered SFTPB variants in atypical presentations of ILD particularly in a familial context.





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In vitro transcription analysis of the variation c.582G>A : overexpression of SFTPB genomic constructs spanning exon 4 to 6 (NM\_000542.5), WT or mutated in A549 cells. Analysis by electrophoresis and Sanger sequencing of the

*Ex vivo* studies : Haematoxylin & eosin (HE) stainings and immunostaining targeting SP-B and SP-C assessed on

eosin (HE) staining, SP-B and SP-C staining (magnification x20 and x40).



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