



# NKX2.1 mutation revealed by a lymphoid interstitial pneumonia in an adult with rheumatoid arthritis

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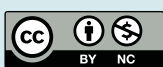
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To the Editor:

*NKX2.1* encodes the thyroid transcription factor 1 (TTF1), which is implicated in lung development and control of the expression of surfactant proteins [1]. This transcription factor is also expressed in the brain and the thyroid, and *NKX2.1* heterozygous mutations are responsible for the “brain–lung–thyroid syndrome”. This rare syndrome is characterised by central nervous system abnormalities, interstitial lung disease (ILD) and hypothyroidism, though the triad is inconstant with heterogeneity of organs involved and clinical presentation. A severe ILD can be the only manifestation in up to 25% of *NKX2.1* mutation carriers [2]. Heterogeneous phenotypes may be present in the same family without a clear explanation for the variable penetrance [3]. Most of the cases have been reported in newborns with fatal disease due to neonatal respiratory distress syndrome or ILD in childhood [2–5], and only a few cases are described in adulthood [2, 3, 5]. Recently it has been shown that familial pulmonary fibrosis and rheumatoid arthritis (RA)-associated ILD (RA-ILD) share the same genetic background, but no association with an *NKX2.1* mutation has been reported to date [6, 7]. We describe the case of an adult patient with hypothyroidism and RA-ILD with *NKX2.1* heterozygous pathogenic variation.

A 37-year-old nonsmoker female was referred to our department for the evaluation of ILD. A diagnosis of RA was given 6 years earlier. She also reported hypothyroidism of unknown aetiology (treated with synthetic thyroxin), obesity, hypertension and diabetes. ILD was detected at RA diagnosis (figure 1a). RA treatment required low-dose steroids with different disease-modifying antirheumatic drugs including methotrexate, tocilizumab, rituximab and abatacept. At time of referral, she complained of severe shortness of breath (modified Medical Research Council (mMRC) dyspnoea scale 3) and dry cough. Clinical examination revealed bilateral crackles. Neurological examination was normal. Chest high-resolution computed tomography (HRCT) showed diffuse ground-glass opacities and thickening of the bronchovascular bundles and small nodules in a centrilobular and subpleural location, associated with traction bronchiectasis and honeycombing predominating in the left upper lobe (figure 1b). These features had developed in areas with previous ground-glass opacities. Diffuse enlargement of mediastinal and hilar lymph nodes was also present. Pulmonary function tests showed a severe restrictive defect with reduced forced vital capacity (forced vital capacity (FVC) 2020 mL, 54% predicted) and reduced diffusion capacity for carbon monoxide ( $D_{LCO}$ , 25% of the predicted value) with normal forced expiratory volume in 1 s ( $FEV_1$ )/FVC ratio (0.87, 107% predicted). During the 6-min walk test she walked 380 m (76% predicted) with desaturation at 80%.

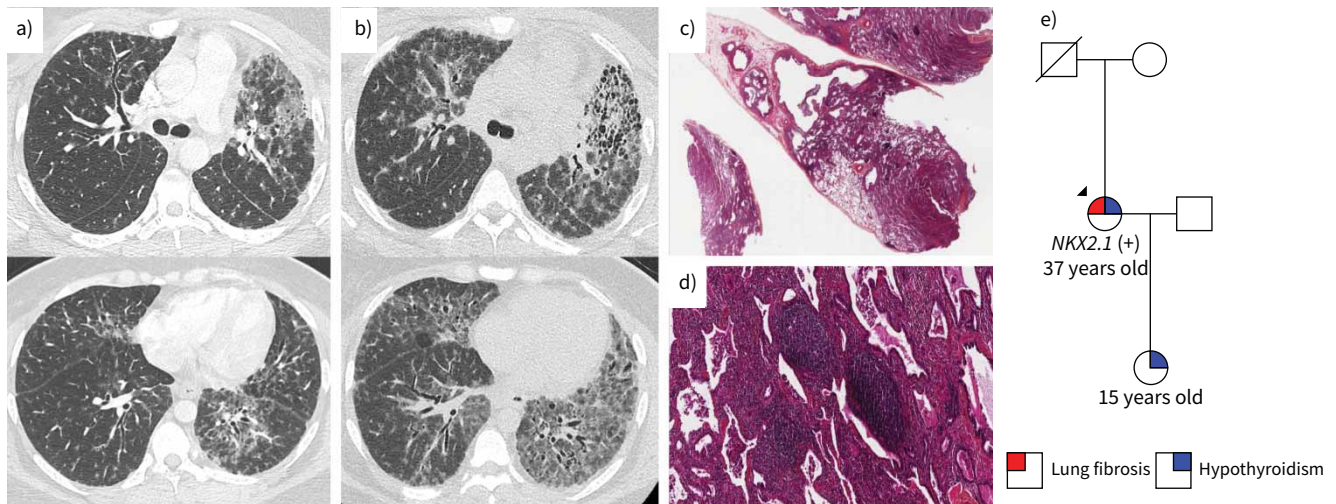
HIV testing was negative. C-reactive protein was  $13 \text{ mg}\cdot\text{L}^{-1}$ . Immunological investigations showed positive anti-citrullinated peptide antibodies ( $35 \text{ U}\cdot\text{L}^{-1}$ ) and rheumatoid factor ( $>200 \text{ IU}\cdot\text{L}^{-1}$ ). A search for anti-extractable nuclear antigen (ENA), myositis-specific antibodies and thyroiditis antibodies was negative. Serum protein electrophoresis showed restrictions of gamma heterogeneity but serum protein immunoelectrophoresis was normal. Cytological examination of the bronchoalveolar lavage fluid revealed  $410\,000 \text{ cells}\cdot\text{mL}^{-1}$  with 68% alveolar macrophages, 10% lymphocytes and 22% neutrophils. Microbiological evaluation was negative. Endobronchial ultrasound-guided mediastinal lymph node sampling showed normal lymph node cytology. After multidisciplinary discussion and due to the very atypical HRCT pattern with suspicion of pulmonary lymphoma, surgical lung biopsy was performed. Three biopsy specimens were taken from the left lung (apex, lingula and lower lobe). Histology showed a diffuse infiltration of the pulmonary interstitium by small lymphocytes (mainly T-cells) and polytypic



Shareable abstract (@ERSpublications)

This is the first case of a 37-year-old female patient carrier of a heterozygous *NKX2.1* mutation associated with RA-ILD with a histological pattern of LIP. This case illustrates the wide panel of ILD subtypes associated with *NKX2.1* mutations. <https://bit.ly/3F49OTS>

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**FIGURE 1** a) First thoracic computed tomography (CT) scan for diagnosis of rheumatoid arthritis, showing ground-glass opacities and traction bronchiectasis. b) Evolution of thoracic CT scan 6 years later showing extension of ground-glass opacities and thickening of the bronchovascular bundles and small nodules, associated with traction bronchiectasis and honeycombing in the left upper lobe. c) Histology of surgical lung biopsy: diffuse lymphoid infiltrate expanding the pulmonary interstitium and distorting architecture with cysts; haematoxylin, eosin, saffron (HES) stain; scanning magnification; d)  $\times 20$  objective. e) Index case (black arrowhead) was diagnosed with the heterozygous p.(Gln317\*) mutation. Her daughter has hypothyroidism.

plasmacytes. Some reactive B-cell follicles were present. A molecular analysis failed to detect any monoclonal population (figure 1c, d). The histological pattern was consistent with lymphocytic interstitial pneumonia (LIP) with some fibrotic remodelling. Histology of the minor salivary glands was normal. We proposed treatment with methylprednisolone pulses ( $250 \text{ mg}\cdot\text{day}^{-1}$  for 3 days) then prednisone  $60 \text{ mg}\cdot\text{day}^{-1}$  with progressive tapering, and rituximab (1000 mg on day 1 and day 15). After 6 months of follow-up, dyspnoea was stable (mMRC 3) as was her 6-min walk test with a distance of 380 m (76% predicted). Lung function test results remained stable (FVC 1900 mL, 52% predicted;  $D_{\text{LCO}}$  24% predicted).

Because of the early onset of the lung disease and the association with hypothyroidism, a germline genetic analysis was prescribed. A heterozygous nonsense mutation was identified in *NKX2.1* (NM\_001079668) (c.949C>T p.(Gln317\*)). No mutation was detected in other genes associated with genetic ILD, including telomere-related genes (*TERT*, *TERC*, *PARN*, *RTEL1*, *DKC1*, *TINF2*, *NOP10*, *NHP2*) and other surfactant-related genes (*SFTPC*, *STFPA1*, *STFPA2*, *ABCA3*).

The *NKX2.1* p.(Gln317\*) nonsense mutation was considered as likely pathogenic. This mutation creates a premature stop codon in the last exon of the gene, and the corresponding transcripts probably escape nonsense-mediated mRNA decay (transcripts degradation before protein production), but it probably leads to the production of a non-functional truncated protein lacking its last 85 amino acids (out of 401). It is not described in the gnomAD database that encompasses about 15 500 control subjects with sequencing data at this genomic position. The patient did not have any siblings and there was no parental history of miscarriage. Neither her mother nor her mother's sibling had a history of ILD or neurological disease. Her father died with myocardial infarction without a history of lung disorder, whereas her 15-year-old daughter had an abnormal thyroid test (figure 1e). In conformity with French legislation, her minor daughter was not tested because she had no respiratory or neurological involvement with a normal clinical evaluation. A genetic diagnosis will be offered to her once she gets her majority. Unfortunately, we did not have access to the DNA of our patient's parents to determine if the mutation was *de novo*.

This is the first report of a pathogenic *NKX2.1* mutation associated with RA-ILD revealed in adulthood with a confirmed diagnosis of LIP. This case illustrates the variable phenotypic expression associated with *NKX2.1* mutations and expands the lung phenotype associated with *NKX2.1* mutations to secondary ILD.

In paediatric patients, genetic analysis of surfactant-related genes is recommended in the diagnosis of cryptogenic ILD [8], but guidelines do not recommend any genetic analysis in ILD adult patients. In our

centre, we propose genetic analysis for patients with fibrotic ILD before 50 years of age or presenting with extrapulmonary manifestation suggestive of specific genetic syndrome or before lung transplantation [9, 10], in accordance with the recent European Respiratory Society statement on familial pulmonary fibrosis [11]. According to the laboratory and the phenotype, the genetic analysis may include telomere-related genes, surfactant-related genes analysed through next generation sequencing panel or clinical whole exome sequencing. In this case, the genetic analysis was indicated because of: 1) the unusually young age for severe RA-ILD; 2) the fibrotic ILD pattern; and 3) the association with hypothyroidism of unknown aetiology.

The other interesting point concerns the histological pattern. LIP is a very rare entity in adults. LIP can be idiopathic or associated with HIV infection or with autoimmune diseases, especially Sjögren's syndrome [12]. Cases of LIP with RA without Sjögren's syndrome are very rare [13]. One case of LIP associated with *NKX2.1* mutation has been suspected in a 5-year-old child but without histological confirmation [2]. There is no lung histology reported in adult cases associated with *NKX2.1* mutation [2, 3, 5]. In children, nonspecific interstitial pneumonia and desquamative interstitial pneumonia have been observed [2].

The link between RA, LIP and *NKX2.1* mutation is still unclear. Our main hypothesis is that RA just happened to overlap with the *NKX2.1* gene mutation; RA triggered the development of the ILD and the genetic predisposition facilitated the development of a particularly severe and fibrotic ILD. However, some authors found surfactant proteins in the synovial membrane of RA patients and have discussed the role of surfactant protein in RA inflammation [14, 15]. This case report raises the question that dysfunction of TTF1 protein may affect synovial homeostasis and facilitate the development of RA. Alternatively, dysfunction of TTF1 may promote a pathological immune response targeting the lung, triggering a diffuse lymphocytic infiltration with secondary protein citrullination in the lung, a condition that could promote the development of RA [16, 17].

In conclusion, we report the first case of a 37-year-old female carrier of a heterozygous *NKX2.1* mutation associated with RA-ILD with a histological pattern of LIP. This case illustrates the wide panel of ILD subtypes associated with *NKX2.1* mutations.

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