# Bruton's agammaglobulinemia in an adult male due to a novel mutation: a case report

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**Abstract:** X-linked agammaglobulinemia (XLA) is caused by mutation in the gene coding for Bruton's tyrosine kinase (BTK), which impairs peripheral B cell maturation and hypogammaglobulinemia. In this report, we present a case of XLA in a 22-year-old adult male. Genetic testing revealed a novel mutation located at the conserved region (c.383T>C). The patient had a history of recurrent respiratory tract infection which eventually progressed to chronic type II respiratory failure. Several pathogenic bacteria were isolated on culture of respiratory secretions obtained on bronchoscopy. The patient improved on treatment with antibiotics.

Keywords: X-linked agammaglobulinemia (XLA); novel mutation

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# Introduction

X-linked agammaglobulinemia (XLA), also known as Bruton's disease, is a primary immunodeficiency disorder caused by the deficiency of Bruton's tyrosine kinase (BTK) (1). BTK is critical in the maturation of pre-B cells to mature B cells (2,3). Patients with XLA have significantly reduced levels of mature B lymphocytes (<1% of normal) in their peripheral blood (4). According to the BTK gene mutation database (http://bioinf.uta.fi/BTKbase/), missense mutations underlie most alterations. Once the passively transferred protective maternal IgG immunoglobulins fade, these patients become susceptible to recurrent infections, and more than 50% patients have had serious infections by 2 years of life (5-7). The incidence of XLA is estimated to be 1 in 379,000 live births in USA (8). Robust data regarding its prevalence in China is lacking. In this report, we present a case of XLA caused by a novel missense mutation (c.383T>C).

# **Case presentation**

A 22-year-old man presented at our hospital with a history of recurrent respiratory tract infections since the age of five years. At the age of eight years, he developed pyogenic conjunctivitis which led to loss of vision in the left eye. He had normal intelligence and was able to independently carry out his routine daily activities.

About two years ago, he developed respiratory tract infection during his stay in Kunming, a city located at an altitude of 2,000 meters, for which he required frequent hospitalization. His symptoms improved on antibiotic treatment, but he had to receive long-term domiciliary oxygen therapy. In Kunming 2 years back, the PaCO<sub>2</sub> on admission to the hospital averaged 88.6 mmHg, and on discharge averaged 74 mmHg.

One week prior to presenting to our hospital, the patient developed exacerbation of symptoms, and upon admission, he developed obnubilation, for which non-invasive ventilator

Table 1 Results of the tests conducted

Etiology	Results	Reference range
CMVmal (sputum)	1.8×10 <sup>6</sup> copies/mL	<1×10 <sup>3</sup> copies/mL
CMVpies (blood)	<1×10 <sup>3</sup> copies/mL	<1×10 <sup>3</sup> copies/mL
CMVpies (urine)	<1×10 <sup>3</sup> copies/mL	<1×10 <sup>3</sup> copies/mL
EBV-DNA (blood)	<1×10 <sup>3</sup> copies/mL	<1×10 <sup>3</sup> copies/mL
CMVlgG	61.9 AU/mL	0–6 AU/mL
CMVIgM	0 AU/mL	0 AU/mL
CMVpp65 (granulocytic)	21.2% (positive)	0
CMVpp65 (lymphoid)	7.7% (weak positive)	0
CMVpp65 (monocytic)	40.4% (positive)	0
Mycoplasma antibody (blood)	Negative	Negative
Chlamydia antibody (blood)	Negative	Negative
TB-DNA (sputum)	Negative	Negative
Deep sputum	Staphylococcus epidermidis: 1×10 <sup>5</sup> cfu	Negative
Deep sputum	Candida albicans: 1×10 <sup>3</sup> cfu	Negative
Pneumocystis carinii	Negative	Negative

Table 2 The patient's immune function

Immune function	Results	Reference range
CD3/CD4 (Th)	48.6%	28.5-60.5%
CD3/CD8 (Ts)	42.6%	11.1–38.3%
Th/Ts ratio	1.14	0.9–3.6
NK (natural killer cell)	4.3%	5.6-30.9%
lgG	5.55 g/L	7.23–16.85 g/L
IgM	0.226 g/L	0.63–2.77 g/L
IgA	0.255 g/L	0.69–3.82 g/L
C3	0.713	0.9–1.5 g/L
C4	0.185	0.2–0.4 g/L
CD3 (total T-lymphocyte)	93.5%	59.4-84.6%
CD19 CD20 (total B-lymphocyte)	0.0%	6.4–22.6%

was initially applied. Subsequently, oral tracheal intubation and invasive mechanically ventilation were administered.

The patient underwent an extensive diagnostic workup, and the results of microbiology tests are summarized in *Table 1*, and those of cellular and humoral immunity in *Table 2.* There was no evidence of autoimmune disease. Tumor-related parameters were negative. Patients with both FACS analysis, Western blotting analysis, analysis of memory B cells were negative, T cells and NK cell activity in compensatory increase.

Chest computed tomography (CT) revealed obvious emphysema with bilateral bronchiectasis (*Figure 1*). Echocardiography showed an enlarged right ventricle (21 mm) with mild-to-moderate tricuspid regurgitation. Other cardiac chambers and valves were normal. His mean pulmonary arterial pressure was 57 mmHg and ejection fraction (EF) was 68%. The clinical picture was consistent with *cor pulmonale*.

Sputum specimens were obtained at the time of endotracheal intubation and various pathogenic bacteria, especially intracellular Gram-positive cocci were isolated. The patient was administered intravenous teicoplanin, voriconazole and ganciclovir. The serum levels of IgG were below normal, and intravenous gamma globulin (10 g/day) was administered. The mechanical ventilation was successfully retracted after 14 days, and he was discharged after a total hospital stay of 28 days. The partial pressure of arterial carbon dioxide at discharge was 55 mmHg, which was significantly lower than average value of 74 recorded over several time points over the past two years.

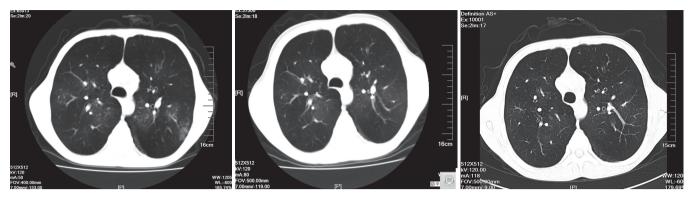
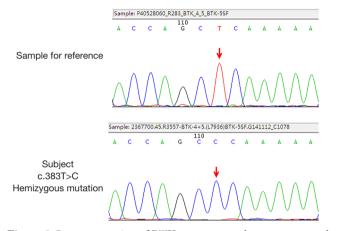


Figure 1 Chest CT radiographs at the same level at different time-points.



**Figure 2** Gene sequencing of BTK gene exon codes area compared with the reference sequence (NM\_000061.2 with NG\_009616.1). BTK, Bruton's tyrosine kinase.

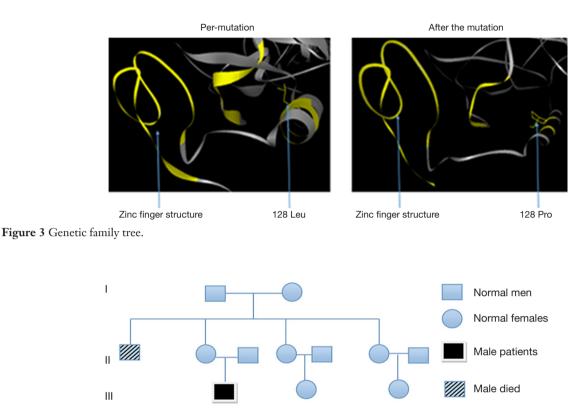
### Discussion

The patient had a history of recurrent pulmonary infections since childhood, and developed full blown respiratory failure, likely as a result of repeated pulmonary infections in the past two years. He had a B (CD19+, CD20+) lymphocyte count of nearly 0% and low levels of IgG, IgM and IgA. Since his maternal uncle had died at a young age, a genetic basis was suspected, and the patient was tested for *BTK* gene mutation after obtaining consent of his family, and appropriate counselling. The specimens obtained were sent to the Guangzhou KingMed Diagnostics for direct sequencing of exonic coding region of the *BTK* gene. A hemizygous *BTK* mutation at (c.383T>C) was detected by comparing with the reference sequence (NM\_000061.2 and NG\_009616.1). This point mutation was a missense mutation [p.(Leu128Pro)] at amino acid residue 128 in the translating protein, that caused leucine to be substituted by proline (*Figure 2*). This point mutation was located at the conserved region of the *BTK* gene, and the bioinformatics analysis software LTR and PolyPhen-2 suggested that this alteration was likely to engender high pathogenicity, while the analysis of the protein structure suggested that the substitution of leucine at amino acid 128 by proline would affect the structure of zinc fingers and functional tetramers (*Figure 3*), thus leading to serious dysfunction.

The patient was diagnosed with Bruton's disease in accordance with the diagnostic criteria of the Pan-American Group for Immuno-Deficiency (PAGID) and the European Society of Immunodeficiency Disease (ESID) (9). The same hemizygous BTK mutation was also detected in his mother; but not in his second and third aunts (*Figure 4*).

In the BTK gene mutation database (http://bioinf. uta.fi/BTKbase/), missense mutation, deletion and nonsense mutations were the top three types of mutations observed. TK, pH and SH2 domains were involved in a vast proportion of cases. Lee *et al.* (10) found that patients with missense mutations on average live longer than those with other types of mutations. A study of 110 cases of XLA by Broides *et al.* (11) revealed that 56% of patients with missense mutation were diagnosed at the age of  $\geq 6$  years. Further, missense mutations accounts for a higher proportion of patients who are diagnosed at a later age. The age of this patient at diagnosis was 22 years, which indicates a significantly delayed diagnosis in this case. This case highlights the need for a high degree of suspicion for XLA on the part of clinicians to avoid missed diagnosis.

Patients with XLA are particularly susceptible to respiratory infections (12,13). In our patient, Lower respiratory tract secretion specimens were obtained on bronchoscopy at the time of endotracheal intubation for respiratory failure. On culture, several pathogenic specimens



**Figure 4** Peripheral blood FACS analysis of the patient compared to that of a normal adult. The upper panel represents the BTK-mutant patient. The B cell group is <1% (<10% of that in the normal adult). FACS, fluorescent-activated cell sorting; BTK, Bruton's tyrosine kinase.

were isolated such as Staphylococcus epidermidis, Candida albicans and CMV virus. In his previous hospitalizations, Lower respiratory tract secretion specimens were not obtained, and no pathogenic bacteria were isolated on routine sputum culture. BTK mutations lead to impaired humoral immunity, thus rendering the patient susceptible to Streptococci, Neisseria, Enterovirus and other pathogens (14-16). In XLA patients with long-standing impairment of immunity, the possibility of occult opportunistic infections should be considered despite negative sputum culture.

Peripheral blood FACS analysis, Western blotting and memory B cell ELISPOT analysis displayed peripheral B cells were all negative. A complementary increase in T cells, NK cells (mainly composed of CD14 monocytes) and CD56 NK cells was noted (*Figures 5-7*).

The human body relies mainly on cellular immunity to protect against CMV (17). CMV is known to evade the immune system through a variety of mechanisms. There was still a risk of acute exacerbation of chronic bronchial CMV infection although the cellular immune function was normal or even increased (18-21). The fact that the patient has grown into manhood is closely related to his long-term use of IgG replacement therapy on a regular basis. In this case, the level of serum IgG was nearly normal, while the level of IgM and IgA was around 10% of normal. Thus, whether IgM-rich immunoglobulin should be used in the acute stage of infection is a matter of conjecture, and no evidence-based guidelines exist (22-24). This also highlights that the human body is an organized whole, and weakness of one defense mechanism (such as humoral immunity) triggers compensatory increase in other mechanisms (such as cellular immunity, antigen presentation, and NK cells).

From the clinical point of view, physicians should not only consider suppurative bacterial infection, but also opportunistic infection, such as fungi and CMV, and also institute a plan for dynamic monitoring of chronic persistent infection, and make appropriate adjustments to the therapeutic regimen on an ongoing basis.

In conclusion, genetic analysis confirmed the diagnosis of our patient, and identified a previously unreported mutation. In adult patients with repeated episodes of

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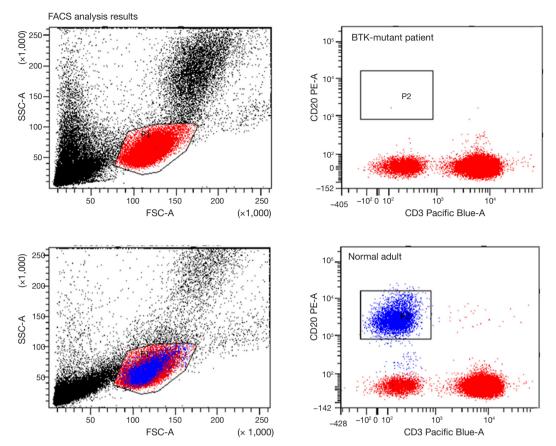
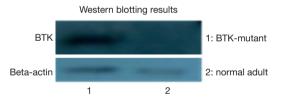


Figure 5 Results of Western blotting showing expression of BTK antibody in the patient. BTK, Bruton's tyrosine kinase.



**Figure 6** Memory B cell ELISPOT analysis of peripheral blood of BTK patient and normal adult. Expression of BTK-mutant memory B cells approximates zero as against 3,000 cells/million in a normal adult. ELISPOT, enzyme-linked immunospot assay; BTK, Bruton's tyrosine kinase.

Wemovidg cells Wemovidg cells 1000-BTK-mutant Normal adult

Memory B cell ELISPOT analysis

3000

Figure 7 Simulation of the spatial structure of point mutation.

infection, clinicians should consider the possibility of XLA.

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#### Footnote

*Conflicts of Interest*: The authors have no conflicts of interest to declare.

*Informed Consent*: Written informed consent was obtained from the patient for publication of this manuscript and any accompanying images.

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