

Chronic Aichi virus infection in a patient with X-linked agammaglobulinemia.

Giorgia Buccioli, MD^{a,b}, Leen Moens, PhD^a, Kathryn Payne, BSci^c, Djalila Mekahli, MD, PhD^{b,d}, Elena Levtschenko, MD, PhD^{b,d}, François Vermeulen, MD, PhD^b, Thomas Tousseyn, MD, PhD^e, Paul Gray, MD, PhD^{f,g}, Cindy S. Ma, PhD^{c,h,i}, Stuart G. Tangye, PhD^{c,h,i}, Marc Van Ranst, PhD^j, Julianne R. Brown, PhD^k, Judy Breuer, MD^{kl}, Isabelle Meyts, MD, PhD^{a,b}

^aLaboratory of inborn errors of immunity, Department of Immunology and Microbiology, KU Leuven, Belgium

^bDepartment of Pediatrics, University Hospitals Leuven, Belgium

^cImmunology Division, Garvan Institute of Medical Research, Darlinghurst, NSW, Australia

^dDepartment of Development and Regeneration, KU Leuven, Leuven, Belgium

^eDepartment of Imaging & Pathology, KU Leuven and University Hospitals Leuven, Leuven, Belgium

^fSchool of Women's and Children's Health, Faculty of Medicine, University of New South Wales, Sydney, NSW, Australia

^gDepartment of Immunology and Infectious Disease, Sydney Children's Hospital, Sydney, NSW, Australia

^hSt Vincent's Clinical School, UNSW Sydney Faculty of Medicine, Sydney, NSW, Australia

ⁱCIRCA (Clinical Immunogenomics Research Consortia Australia), Sydney, NSW, Australia

^jDepartment of Microbiology and Immunology, KU Leuven, Belgium

^kGreat Ormond Street Hospital for Children NHS Foundation Trust, London, UK

^lDivision of Infection and Immunity, University College London, London, UK

Corresponding author: Isabelle Meyts, Department of Pediatrics, University Hospitals Leuven

Herestraat 49, 3000 Leuven, Belgium; Tel +32 16 343841; Fax +32 16 343842

Isabelle.Meyts@uzleuven.be

Funding: GB is supported by the Research Foundation - Flanders (project G0C8517N). LM is supported by the CSL Behring Chair in Primary Immunodeficiencies, by the KID-FONDS charity of KU Leuven and by the Jeffrey Modell Foundation. EL is supported by the Research Foundation - Flanders (Clinical Investigator grant 1801110N). IM is supported by the Jeffrey Modell Foundation and by the Research Foundation - Flanders (project G0C8517N).

Key words: X-linked agammaglobulinemia, XLA, Bruton, Aichi virus, AiV1

Compliance with Ethical Standards

The authors declare that they have no conflict of interest.

All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

Informed consent was obtained from all individual participants included in the study.

To the Editor,

X-linked agammaglobulinemia (XLA) is caused by mutations in *BTK*, the gene encoding Bruton's tyrosine kinase. *BTK* is critical for human B cell development and maturation, and hemizygous loss-of-function (LOF) mutations result in peripheral B cell lymphopenia, rudimentary tonsils and lymph nodes, and severely reduced to absent levels of serum immunoglobulins (Ig). Affected boys present in infancy with recurrent and often life-threatening respiratory tract and skin infections with encapsulated bacteria. Gastrointestinal infections with pathogens such as *Giardia lamblia* are also common [1]. XLA patients also have extreme susceptibility to viral infections, especially with Enteroviruses, which can cause severe central nervous system (CNS) disease [2]. The risk of Enteroviral infection is particularly high in patients who experience a diagnostic delay. Chronic Norovirus infection of the gut is another problematic viral infection [3–5], while Astrovirus and Cache Valley virus have been described to cause progressive encephalitis in XLA patients [6,7].

Here we describe a boy with XLA due to a deleterious mutation in *BTK* (c82C>T, p. R28C), diagnosed at the age of three years. Between the age of six and thirteen years, following a holiday in northern Italy, he developed a fever of unknown origin, intermittent bloody diarrhea, generalized lymphadenopathy, progressive hepatosplenomegaly, progressive nephromegaly and refractory temporal lobe epilepsy (detailed clinical history can be found in this article's supplemental data). Blood analysis showed progressive pancytopenia, progressive elevation of liver enzymes, and chronic kidney failure with a glomerular filtration rate as low as 50 ml/min/1.73m², as measured by ⁵¹Cr-EDTA clearance. On ultrasound multiple hypodense focal lesions were evident in the liver, spleen and kidneys, which were enlarged. Histological examination of a liver biopsy specimen demonstrated severe chronic hepatitis with initial fibrosis. Kidney histology showed pronounced diffuse interstitial oligoclonal cytotoxic T cell infiltrates with changing TCR V β repertoires on sequential biopsies. Colon biopsy showed chronic colitis. The patient

developed severe growth delay. At the age of thirteen years he underwent splenectomy due to a mechanical gastrointestinal obstruction; this resulted in resolution of his pancytopenia.

No microbe was isolated from blood, urine, cerebrospinal fluid (CSF) or biopsied tissues, and multiple treatments, such as antimicrobial therapies *ex juvantibus*, elevation of the Ig supplementation dose, and corticosteroids, were ineffective in controlling fever, diarrhea, vomiting, progressive kidney and liver anomalies and convulsions. Only during empirical treatment with liposomal amphotericin B (on the suspicion of *Leishmania* infection) did the patient become afebrile, while antibiotics and nitazoxanide gave no benefit. Immunophenotyping of his peripheral blood showed an increase in proportions of CD8⁺ T cells, mostly displaying a memory phenotype (i.e. high numbers of CD45RA⁻CCR7⁺ central and CD45RA⁻CCR7⁻ effector memory cells and reduced numbers of CD45RA⁺CCR7⁺ naïve and terminally differentiated CD45RA⁺CCR7⁻ effector memory cells), compared to healthy donors and a control XLA patient. Furthermore, his memory CD8⁺ T cells displayed increased expression of markers of activation (HLA-DR, CD38 and HLA-ABC) and exhaustion/senescence (CX3CR1, CD95, CD57; Fig. 1). In contrast, the control XLA patient had decreased CD8⁺ and increased CD4⁺ T cells, with an increased proportion of naïve and terminally differentiated effector memory cells and corresponding decreased proportions of central and effector memory cells compared to healthy controls. Moreover, expression of activation and exhaustion/senescence markers on CD8⁺ T cells from the control XLA patient was comparable to or lower than that of healthy controls (Fig. 1, D-I). This is in agreement with the subtle defects previously described in T cell maturation in XLA patients [8,9].

RNA-sequencing detects the presence and quantity of RNA in a biological sample at a given moment; it is mainly applied to the study of changes in the cellular transcriptome over time or in different groups following treatment or disease. RNA-seq is also a powerful tool in the field of infectious diseases, as it allows for the detection of viral RNA in pathological tissue samples [10,11], and we therefore applied it in a snap-frozen sample from a kidney biopsy performed at twelve years of age in our patient. The whole-

tissue RNA-seq was found to be positive for Aichi virus (AiV1), and subsequently samples of liver, spleen, urine and sputum were also found positive for AiV1 on PCR. AiV1 is a Kobuvirus of the Picornaviridae family and causes self-limiting gastroenteritis in immunocompetent humans. It was first isolated in 1989 during a gastroenteritis outbreak associated with the consumption of raw oysters in Aichi prefecture, Japan, and was later genetically characterized [12]. AiV1 is a common contaminant of water ponds, sewages and shellfish all around the world and its seroprevalence reaches almost 100% by the age of 30 years in all published studies [13–18]. Despite the high prevalence of anti-AiV1 antibodies in children and adults, AiV1 is found rarely in stool samples from patients with acute sporadic or epidemic gastroenteritis and, on average, is isolated in fewer than 1% of cases [13,19]. However, the prevalence of AiV1 in stool specimens of HIV-positive patients is higher, probably indicating an opportunistic behavior in patients with underlying T cell defects [19,20].

Enteroviruses, like AiV1, belong to the Picornavirus family, and are a cause of chronic infection in XLA patients. Chronically infected patients commonly develop meningoencephalitis, sometimes showing the acute flaccid paralysis that is characteristic of poliovirus infections, but arthritis, hepatitis, dermatomyositis, polyradiculitis and myocarditis have also been described [2]. Of note, CSF is often normal, and both culture and PCR for Enteroviruses yield a high false negative rate. Consequently, the gold standard for diagnosis is to perform PCR or NGS techniques on a brain biopsy [2]. Our patient presented with severe multi-organ involvement. A classical microbe-isolation approach, including the use of pan-bacterial PCR, was insufficient to identify the etiologic agent, while RNA-seq obviated the limitations of standard methods in identifying AiV1. Chronic Norovirus gastrointestinal infection is another difficult-to-treat infectious complication of XLA, sometimes severe enough to prompt hematopoietic stem cell transplantation (HSCT) [3,5].

Although well described, the susceptibility of XLA patients to chronic viral infections is unexpected and ill-explained. Various studies have demonstrated abnormal Toll-like receptor (TLR) function and dendritic cell

(DC) maturation and function in XLA patients. Impaired DC activation and cytokine production has been correlated with defective TLR8 and TLR9 signaling in BTK-deficient cells [21,22]. A defect in CD4⁺ T cell maturation has also been described in XLA, with significant reductions in CD4⁺CD45RO⁺ and CD4⁺CD45RO⁺CXCR5⁺ memory T cells and impaired delayed cutaneous hypersensitivity reaction [8,9,23]. Together, these subtle defects could contribute to susceptibility to viral and opportunistic infections in these patients. Various therapeutic strategies have attempted to control and/or eradicate Enterovirus and Norovirus in XLA patients, mostly with little success. High dose Ig substitution is recommended, and local administration (intrathecal or enteral) has been described to yield better outcomes than parenteral infusion alone [2,5]. Antiviral drugs such as cidofovir, ribavirin and interferon (IFN)- α or - β have mostly been ineffective [2,5]. Nitazoxanide, a broad-spectrum antimicrobial agent with antiviral and Norovirus replication-inhibiting properties in vitro, has been used in immunodeficient patients with chronic Norovirus infection with mixed results [4,5]. Pleconaril (an anti-Picornavirus medication) was the most frequently used antiviral in chronic enteroviral meningitis, but is no longer available, while other direct-acting antiviral molecules are currently under development (Vapendavir, Pocopavir and ViroD7000) [2]. Also, itraconazole and fluoxetine showed anti-Enterovirus properties in vitro, and successful treatment of chronic enteroviral meningitis with fluoxetine has been reported in a XLA patient [2,24]. Antiviral effects of immunosuppressive regimens in transplanted and other immunodeficient patients have been reported, specifically the anti-Cytomegalovirus and anti-Norovirus efficacy of the mTOR (mechanistic target of rapamycin) inhibitors sirolimus and everolimus [5,25–27]. Recently, amphotericin B was found to be a potent inhibitor of two different Enteroviruses, in addition to its known anti-fungal indication and the described anti-parasitic and anti-viral effect on *Leishmania*, vesicular stomatitis virus, herpes simplex virus, Sindbi virus, human immunodeficiency virus and others [28]. However, no specific data is available for AiV1.

In the present patient, various courses of treatment were attempted. High doses of Ig were poorly tolerated from a neurological point of view; nitazoxanide treatment gave no results, and itraconazole therapy was discontinued after a few days due to hepatic toxicity. Only liposomal amphotericin B resulted in a temporary afebrile status, and a new trial with Ambisome is envisaged. We tried to introduce fluoxetine in increasing doses with some difficulty due to adverse effects. Immunosuppression with low dose tacrolimus was also attempted to reduce the hyperactivation of his CD8⁺ T cells (observed both in blood and as infiltrates in the kidney biopsies). Many studies of patients suffering chronic viral infections showed that CD8⁺ T cells prematurely undergo exhaustion and senescence, thereby becoming dysfunctional due to constant activation by persistent viral stimuli [29–32]. This was also observed in our patient. Given the severity of his presentation and progressive kidney failure due to diffuse T cell infiltrate in the renal interstitium, HSCT using the patient's HLA-matched sister as a donor is being considered. Indeed, given the possibility of an additional subtle DC/T cell immune defect in XLA, HSCT may be the only curative treatment for XLA patients with chronic viral infections. In conclusion, we describe AiV1 as the cause of severe chronic multi-organ disease and chronic kidney failure in a patient with XLA, and we invite to consider HSCT as a treatment option.

REFERENCES

1. Conley ME, Rohrer J, Minegishi Y. X-linked agammaglobulinemia. *Clin Rev Allergy Immunol*. 2000 Oct 1;19(2):183–204.
2. Bearden D, Collett M, Quan PL, Costa-Carvalho BT, Sullivan KE. Enteroviruses in X-Linked Agammaglobulinemia: Update on Epidemiology and Therapy. *J Allergy Clin Immunol Pract*. 2016 Dec;4(6):1059–65.
3. Frange P, Touzot F, Debré M, Héritier S, Leruez-Ville M, Cros G, et al. Prevalence and Clinical Impact of Norovirus Fecal Shedding in Children with Inherited Immune Deficiencies. *J Infect Dis*. 2012 Oct 15;206(8):1269–74.
4. Kempf B, Edgar JD, Mc Caughey C, Devlin LA. Nitazoxanide Is an Ineffective Treatment of Chronic Norovirus in Patients With X-Linked Agammaglobulinemia and May Yield False-Negative Polymerase Chain Reaction Findings in Stool Specimens. *J Infect Dis*. 2017 Feb 1;215(3):486–7.
5. Brown L-AK, Clark I, Brown JR, Breuer J, Lowe DM. Norovirus infection in primary immune deficiency. *Rev Med Virol*. 2017 Mar 8;27(3):e1926.
6. Frémond M-L, Pérot P, Muth E, Cros G, Dumarest M, Mahlaoui N, et al. Next-Generation Sequencing for Diagnosis and Tailored Therapy: A Case Report of Astrovirus-Associated Progressive Encephalitis. *J Pediatr Infect Dis Soc*. 2015 Sep 1;4(3):e53–7.
7. Wilson MR, Suan D, Duggins A, Schubert RD, Khan LM, Sample HA, et al. A novel cause of chronic viral meningoencephalitis: Cache Valley virus. *Ann Neurol*. 2017 Jun 19;82(1):105–14.
8. Martini H, Enright V, Perro M, Workman S, Birmelin J, Giorda E, et al. Importance of B cell co-stimulation in CD4+ T cell differentiation: X-linked agammaglobulinaemia, a human model. *Clin Exp Immunol*. 2011 Apr 13;164(3):381–7.
9. Ma CS, Wong N, Rao G, Avery DT, Torpy J, Hambridge T, et al. Monogenic mutations differentially affect the quantity and quality of T follicular helper cells in patients with human primary immunodeficiencies. *J Allergy Clin Immunol*. 2015 Oct 1;136(4):993-1006.e1.
10. Kawada J, Okuno Y, Torii Y, Okada R, Hayano S, Ando S, et al. Identification of Viruses in Cases of Pediatric Acute Encephalitis and Encephalopathy Using Next-Generation Sequencing. *Sci Rep*. 2016 Sep 14;6:33452.
11. Wilson MR, Naccache SN, Samayoa E, Biagtan M, Bashir H, Yu G, et al. Actionable diagnosis of neuroleptospirosis by next-generation sequencing. *N Engl J Med*. 2014 Jun 19;370(25):2408–17.
12. Yamashita T, Sakae K, Tsuzuki H, Suzuki Y, Ishikawa N, Takeda N, et al. Complete Nucleotide Sequence and Genetic Organization of Aichi Virus, a Distinct Member of the Picornaviridae Associated with Acute Gastroenteritis in Humans. *J Virol*. 1998 Oct 1;72(10):8408–12.
13. Kitajima M, Gerba CP. Aichi Virus 1: Environmental Occurrence and Behavior. *Pathogens*. 2015 May 19;4(2):256–68.

14. Yamashita T, Sakae K, Ishihara Y, Isomura S, Utagawa E. Prevalence of newly isolated, cytopathic small round virus (Aichi strain) in Japan. *J Clin Microbiol.* 1993 Nov 1;31(11):2938–43.
15. Goyer M, Aho L-S, Bour J-B, Ambert-Balay K, Pothier P. Seroprevalence distribution of Aichi virus among a French population in 2006–2007. *Arch Virol.* 2008 Jun 1;153(6):1171–4.
16. Ribes JM, Montava R, Téllez-Castillo CJ, Fernández-Jiménez M, Buesa J. Seroprevalence of Aichi Virus in a Spanish Population from 2007 to 2008. *Clin Vaccine Immunol.* 2010 Apr 1;17(4):545–9.
17. Sdiri-Loulizi K, Hassine M, Bour J-B, Ambert-Balay K, Mastouri M, Aho L-S, et al. Aichi Virus IgG Seroprevalence in Tunisia Parallels Genomic Detection and Clinical Presentation in Children with Gastroenteritis. *Clin Vaccine Immunol.* 2010 Jul 1;17(7):1111–6.
18. Oh D-Y, Silva PA, Hauroeder B, Diedrich S, Cardoso DDP, Schreier E. Molecular characterization of the first Aichi viruses isolated in Europe and in South America. *Arch Virol.* 2006 Jun 1;151(6):1199–206.
19. Portes SAR, de Mello Volotao E, Rose TL, Rocha MS, Trindade Pinheiro Xavier M da P, de Assis RM, et al. Aichi Virus Positivity in HIV-1 Seropositive Children Hospitalized with Diarrheal Disease. *Curr HIV Res.* 2015;13(4):325–31.
20. Oude Munnink BB, Canuti M, Deijs M, de Vries M, Jebbink MF, Rebers S, et al. Unexplained diarrhoea in HIV-1 infected individuals. *BMC Infect Dis.* 2014 Jan 13;14:22.
21. Lougaris V, Baronio M, Vitali M, Tampella G, Cattalini M, Tassone L, et al. Bruton tyrosine kinase mediates TLR9-dependent human dendritic cell activation. *J Allergy Clin Immunol.* 2014 Jun 1;133(6):1644-1650.e4.
22. Sochorová K, Horváth R, Rožková D, Litzman J, Bartůňková J, Šedivá A, et al. Impaired Toll-like receptor 8-mediated IL-6 and TNF- α production in antigen-presenting cells from patients with X-linked agammaglobulinemia. *Blood.* 2007 Mar 15;109(6):2553–6.
23. Jongco AM, Gough JD, Sarnataro K, Rosenthal DW, Moreau J, Ponda P, et al. X-linked agammaglobulinemia presenting as polymicrobial pneumonia, including *Pneumocystis jirovecii*. *Ann Allergy Asthma Immunol.* 2014 Jan 1;112(1):74-75.e2.
24. Gofshteyn J, Cárdenas AM, Bearden D. Treatment of Chronic Enterovirus Encephalitis With Fluoxetine in a Patient With X-Linked Agammaglobulinemia. *Pediatr Neurol.* 2016 Nov 1;64:94–8.
25. Engelen MA, Gunia S, Stypmann J. Elimination of norovirus in a chronic carrier under immunosuppression after heart transplantation – effect of everolimus. *Transpl Int.* 2011;24(11):e102–3.
26. Andrassy J, Hoffmann VS, Rentsch M, Stangl M, Habicht A, Meiser B, et al. Is Cytomegalovirus Prophylaxis Dispensable in Patients Receiving an mTOR Inhibitor-Based Immunosuppression? A Systematic Review and Meta-Analysis. *Transplantation.* 2012 Dec 27;94(12):1208.
27. Blanco NB, Kuonen R, Bellini C, Manuel O, Estrade C, Mazza-Stalder J, et al. Chronic norovirus gastroenteritis in a double hematopoietic stem cell and lung transplant recipient. *Transpl Infect Dis.* 2011;13(2):213–5.

28. Xu F, Zhao X, Hu S, Li J, Yin L, Mei S, et al. Amphotericin B Inhibits Enterovirus 71 Replication by Impeding Viral Entry. *Sci Rep.* 2016 Sep 9;6:33150.
29. Edwards ESJ, Bier J, Cole TS, Wong M, Hsu P, Berglund LJ, et al. Activating PIK3CD mutations impair human cytotoxic lymphocyte differentiation, function and EBV immunity. *J Allergy Clin Immunol.* 2018 May 22; doi:10.1016/j.jaci.2018.04.030.
30. Wherry EJ, Kurachi M. Molecular and cellular insights into T cell exhaustion. *Nat Rev Immunol.* 2015 Aug;15(8):486–99.
31. Yamamoto T, Price DA, Casazza JP, Ferrari G, Nason M, Chattopadhyay PK, et al. Surface expression patterns of negative regulatory molecules identify determinants of virus-specific CD8+ T-cell exhaustion in HIV infection. *Blood.* 2011 May 5;117(18):4805–15.
32. Randall KL, Chan SS-Y, Ma CS, Fung I, Mei Y, Yabas M, et al. DOCK8 deficiency impairs CD8 T cell survival and function in humans and mice. *J Exp Med.* 2011 Oct 24;208(11):2305–20.

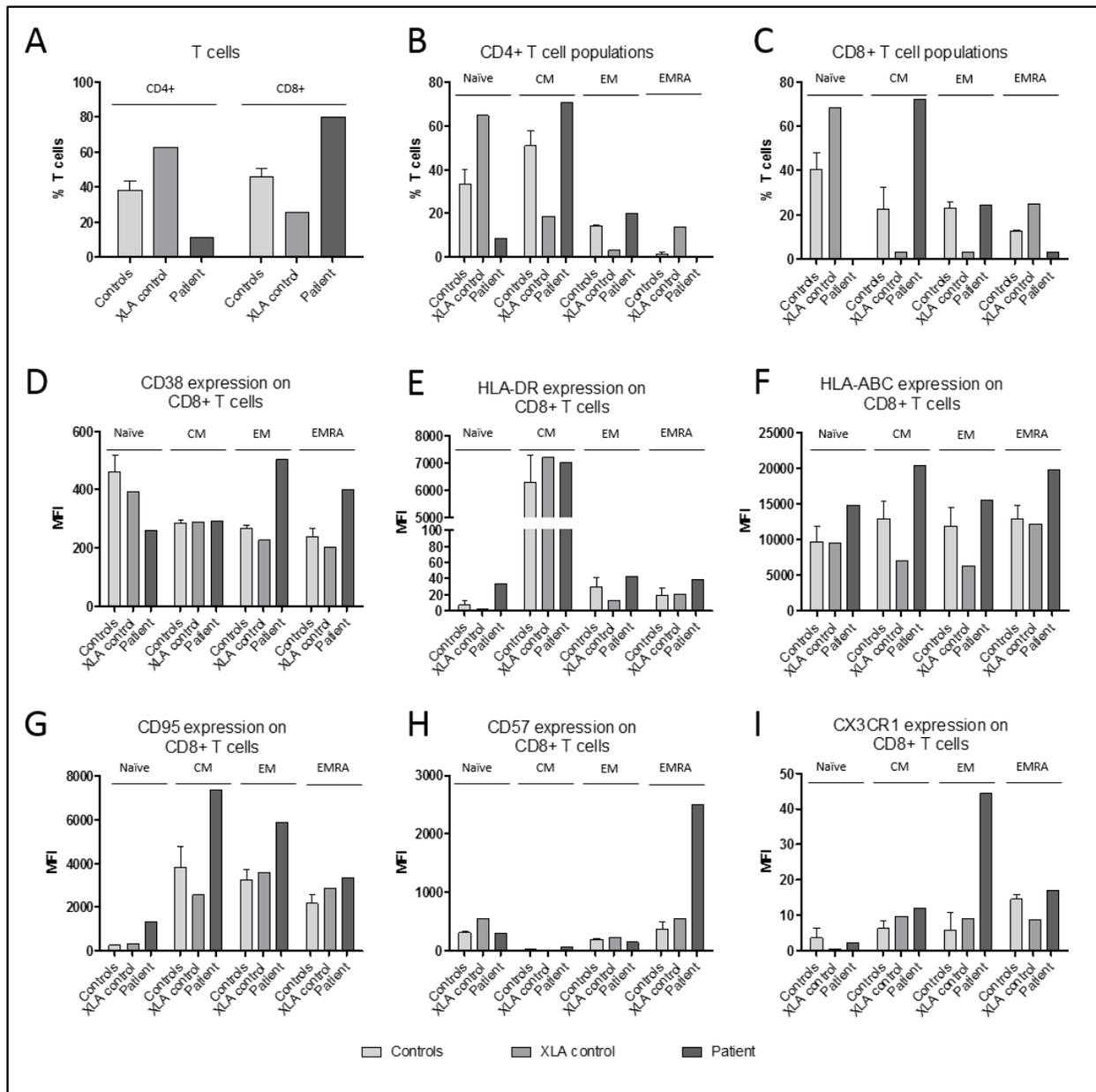


Fig. 1. CD8+ T cells in the proband XLA patient are skewed towards a memory phenotype with evidence of exhaustion and/or senescence. PBMCs from healthy controls, an XLA control and our XLA patient were labeled with mAbs against CD4, CD8, CD45RA, CCR7, CD38, HLA-DR, HLA-ABC, CD95, CD57, and CX3CR1. Proportions of (A) CD4⁺ and CD8⁺ T cell populations, as well as subsets of naïve (CD45RA⁺CCR7⁺), central memory (CM, CD45RA⁺CCR7⁺), effector memory (EM, CD45RA⁺CCR7⁺), and terminally differentiated effector memory cells expressing CD45RA (EMRA, CD45RA⁺CCR7⁺), (B) CD4⁺, and (C) CD8⁺ T cells were delineated. Differential expression of the activation markers CD38 (D), HLA-DR (E), and HLA-ABC (F), as well as the exhaustion/senescence markers CD95 (G), CD57 (H) and CX3CR1 (I) on naïve, T_{CM}, T_{EM} and T_{EMRA} CD8⁺ T cells were determined. Values represent the geometric MFI.