CASE REPORT

Williams–Beuren syndrome: pitfalls for diagnosis in limited resources setting

Aimé Lumaka1,2,3,4, Rita Lukoo3, Gerrye Mubungu1,2,3,4, Paul Lumbala2,3, Gloire Mbayabo2,3, Aimée Mupuala3, Prosper Lukusa Tshilobo1,2,3,4 & Koenraad Devriendt1

1Center for Human Genetics, University Hospitals Leuven, Katholieke Universiteit Leuven, Leuven, Belgium
2Center for Human Genetics, Faculty of Medicine, University of Kinshasa, Kinshasa, Democratic Republic of the Congo
3Department of Pediatrics, Faculty of Medicine, University of Kinshasa, Kinshasa, Democratic Republic of the Congo
4Institut National de Recherche Biomedical, Kinshasa, Democratic Republic of the Congo

Correspondence
Koenraad Devriendt, Center for Human Genetics, University Hospitals Leuven, Herestraat 49 BUS 602, 3000 Leuven, Belgium. Tel: + 32 16 34 59 03; Fax: + 32 16 34 60 60; E-mail: koenraad.devriendt@uzleuven.be

Funding Information
No sources of funding were declared for this study.

Received: 24 June 2015; Revised: 20 September 2015; Accepted: 19 November 2015

Clinical Case Reports 2016; 4(3): 294–297
doi: 10.1002/ccr3.476

Key Clinical Message
Patients with Williams–Beuren Syndrome can be recognized clinically, given the characteristic dysmorphism, intellectual disability, and behavior. We report on a Congolese boy with typical WBS facial characteristics. He suffered meningitis and coma at the age of 2 years then subsequently presented with profound intellectual disability and atypical behavior. The WBS was only made at age 8.2 years and confirmed with FISH testing and microarray-CGH. The present report aims to warn clinicians that infections may associate and/or modify a genetic disease as this may be observed in developing countries given the prevalence of infectious diseases.

Keywords
Central Africa, Democratic Republic of Congo, phenotype, Williams syndrome

Introduction
Williams–Beuren Syndrome (WBS) is one of the most common recurrent microdeletion syndromes, with an estimated incidence varying from 1 in 7500 to 1 in 20,000 births [1, 2]. Patients can be recognized clinically, given their characteristic features, encompassing congenital heart defect, typically supravalvular aortic stenosis (SVAS) and/or peripheral pulmonic stenosis, short stature/growth retardation, recognizable facial dysmorphism, developmental delay and intellectual disability, distinct behavioral phenotype; and, in some instances, hypercalcemia. Facial dysmorphism includes a wide mouth, short nose with bulbous or upturned nasal tip, long philtrum, thick everted vermillion of the lips, bitemporal narrowing, peri-orbital fullness, full cheeks, small jaw, and the characteristic stellate pattern of the iris [3, 4]. The cognitive profile in WBS ranges from mild to severe ID [3, 5, 6]. Although acquisition of language is often delayed in WBS patients, they later develop good language skills and become talkative [7, 8]. Behavioral issues in WBS comprise anxiety, hyperacousis, attention problems, hypersociability or friendly behavior, and strong affinity and interest to music [7, 9, 10]. Recognition of the typical dysmorphism and characteristic behavior lead to a clinical diagnosis.

Some cases are more difficult to diagnose, since they present with atypical features [11], or because they have a different ethnicity where the features may be different from those observed in Caucasians. For instance, African patients with WBS from Cameroon have been reported with incomplete WBS facial dysmorphism [8]. Finally, cases with a deletion smaller or larger than the recurrent 1.5 Mb 7q11.23 microdeletion may also present with a milder or more severe phenotype [12].

We here report a WBS patient from Central Africa carrying the recurrent WBS microdeletion who presented with a typical facial dysmorphism but more severe intellectual disability and atypical behavioral phenotype due to a confounding factor, that is, meningitis with coma supposedly resulting in brain damage.
Case History and Examination

An 8.2 years old Congolese boy was referred by his pediatrician to the genetic clinic at the University Hospitals of the University of Kinshasa, DR Congo, for evaluation of intellectual disability, allegedly resulting from an episode of meningitis and coma. He was born from unrelated, healthy parents after an uneventful pregnancy and delivery. At the age of 2 years, he wasn’t able to stand-up without support or build simple sentences although he could call names of his relatives. Four months later, he suffered from meningitis and remained in coma for 2 days. After recovery, he presented severe intellectual disability and complete loss of intelligible speech. His behavior changed, including hyperactivity, irritability, hypersensitivity to noise and episodes of aggressiveness against strangers. In contrast, he was friendly toward familiar persons and demonstrated a strong attraction to music and musical sounds. He was diagnosed with sequels from meningitis and was followed by a neurologist until the age of 8 years, when he was referred for a genetic evaluation.

At that time, he had severe global growth delay with microcephaly (OFC 47 cm; −3.82 SD), short stature (length 109 cm; −3.43 SD) and low weight (17 kg; <-3 SD). Family history revealed that two siblings of the patient (Fig. 1, upper panel) died from infectious diseases. There were two other males with ID in the pedigree, related through their mothers. Unfortunately, these were not available for evaluation. Patient’s phenotype consisted of a triangular face with low set ears, long philtrum, flat nasal bridge, bulbous nose tip, anteverted nostrils, pronounced nasolabial folds, large mouth, thick and everted vermilion of the upper and lower lip and a small jaw (Fig. 1, lower panel). He also had widely spaced nipples, a small penis, bilateral cryptorchidism, and deep palmar creases. Cardiac ultrasound revealed bicuspid aorta with aortic insufficiency.

Differential Diagnosis and Investigations

Although the pedigree was consistent with X-linked ID, a clinical diagnosis of WBS was retained.
Fluorescent in situ hybridization (FISH) was performed on nuclei from peripheral white blood cells using the commercially available Vysis LSI ELN Kit (Vysis; Abbott Laboratories, Abbott Park, IL). A total of 100 nuclei were examined and a single ELN was observed in all of them, whereas the control probe always showed two signals. This confirmed the diagnosis of WBS (Fig. 2A). Microarray-CGH was performed using a stripped CytoSure™ ISCA v2 array 8×60 k format (OGT, Oxford UK) slide, the CytoSure™ Genomic DNA Labelling Kits (Oxford, UK), and the Oligo aCGH & ChIP-on-Chip Hybridization Kits (Agilent, Santa Clara, CA) following manufacturers’ procedures. This revealed the presence of a recurrent 1.57 Mb deletion arr[hg19]7q11.23 (72,634,874–74,203,685)×1 (Fig. 2B).

**Discussion**

We report on a central African patient with typical gestalt for WBS. Retrospectively, we uploaded facial photographs of the patient to Face2Gene (http://www.fdna.com/face2gene/). Basically, this tool captures gestalt signatures from patient’s facial photograph and compares these to patients in the database, then reports the 10 most likely matches. Interestingly, WBS was reported as the fifth match. This indicates that our patient’s gestalt did not vary from common WBS gestalt.

It is surprising that the patient was only diagnosed at age 8.2 years, despite the typical facial appearance. Mean age of diagnosis for WBS in Western countries is 3.5 years [13, 14]. Several factors may have contributed.
to a delayed diagnosis. First, the facial dysmorphism in WBS evolves over age and may be discrete in infancy and early childhood [4, 13, 14]. Second, our patient had no SVAS, one of the most characteristic features often leading to the diagnosis [13, 15]. Finally, severe phenotypes have been associated with larger deletions. As the deletion in our patient corresponds to the classical WBS microdeletion, we assume that the severe speech deficit and profound intellectual disability in the patient are more likely the complications of the meningitis and coma. It is thus undeniable that an acquired condition modified the clinical course of the patient toward the more severe and atypical phenotype for WBS causing the delay in diagnosis.

Such overlap of genetic disease with an infectious disease is expected to be frequent in the developing world, where infections represent the major medical problem. As previously reported, infections are major cause for misdiagnosis of genetic diseases in developing countries [16]. The present report supports the assumption that infections constitute a major pitfall for genetic diagnosis and warns for clinicians working in developing world that a genetic disease may be masked by an intercurrent infectious disease.

To date, only one report exists concerning WBS in Central Africa [8]. Given the incidence of 1/7500 for WBS [2], this suggests that most cases remain undiagnosed. This is most likely due to a lack of training in the field of dysmorphology and lack of access to genetic facilities in that part of the world.

Acknowledgments

AL acknowledge the contribution of the FWO travel grants for research abroad (Ref: V405213N and K210115) to his research. The authors are grateful to Reinhilde Thoelen for the FISH analysis; Bijou Myndyo, Ervie-Winnter, Eben, and Etsa-Edi Lumaka for their support and contribution to this work.

Conflict of Interest

None declared.

References

1. Morris, C. A., and C. B. Mervis. 2000. Williams syndrome and related disorders. Annu. Rev. Genomics Hum. Genet. 1:461–484.
2. Stromme, P., P. G. Bjornstad, and K. Ramstad. 2002. Prevalence estimation of Williams syndrome. J. Child Neurol. 17:269–271.
3. Merla, G., N. Brunetti-Pierri, L. Micale, and C. Fusco. 2010. Copy number variants at Williams-Beuren syndrome 7q11.23 region. Hum. Genet. 128:3–26.
4. Patil, S. J., B. G. Madhusudhan, S. Shah, and P. V. Suresh. 2012. Facial phenotype at different ages and cardiovascular malformations in children with Williams-Beuren syndrome: a study from India. Am. J. Med. Genet. Part A 158A:1729–1734.
5. Martens, M. A., S. J. Wilson, and D. C. Reutens. 2008. Research Review: Williams syndrome: a critical review of the cognitive, behavioral, and neuroanatomical phenotype. J. Child Psychol. Psychiatry 49:576–608.
6. Mervis, C. B., and A. E. John. 2010. Cognitive and behavioral characteristics of children with Williams syndrome: implications for intervention approaches. Am. J. Med. Genet. Part C Semin. Med. Genet. 154C:229–248.
7. Pegoraro, L. F., C. E. Steiner, E. H. Celeri, C. E. Banzato, and P. Dalgalarrondo. 2014. Cognitive and behavioral heterogeneity in genetic syndromes. J. Pediatr. 90:155–160.
8. Tekendo-Ngongang, C., S. Dahoun, S. Nguefack, S. Saharan Africans: case reports from Cameroon. Mol. Syndromol. 5:287–292.
9. Doyle, T. F., U. Bellugi, J. R. Korenberg, and J. Graham. 2004. “Everybody in the world is my friend” hypersociability in young children with Williams syndrome. Am. J. Med. Genet. Part A 124A:263–273.
10. Ng, R., P. Lai, D. J. Levitin, and U. Bellugi. 2013. Musicality correlates with sociability and emotionality in Williams syndrome. J. Ment. Health Res. Intellect. Disabil. 6:268–279.
11. Sakhuja, P., H. Whyte, B. Kamath, N. Martin, and D. Chitayat. 2015. Williams syndrome presenting with findings consistent with Alagille syndrome. Clin. Case Rep. 3:24–28.
12. Fusco, C., L. Micale, B. Augello, M. Teresa Pellico, D. Menghini, P. Alfieri, et al. 2014. Smaller and larger deletions of the Williams Beuren syndrome region implicate genes involved in mild facial phenotype, epilepsy and autistic traits. Eur. J. Hum. Genet. 22:64–70.
13. Ferrero, G. B., E. Biamino, L. Sorasio, E. Banaudi, L. Peruzzi, S. Forzano, et al. 2007. Presenting phenotype and clinical evaluation in a cohort of 22 Williams-Beuren syndrome patients. Eur. J. Med. Genet. 50:327–337.
14. Huang, L., L. Sadler, M. A. O’Riordan, and N. H. Robin. 2002. Delay in diagnosis of Williams syndrome. Clin. Pediatr. 41:257–261.
15. Eronen, M., M. Peippo, A. Hiippala, M. Raatikka, M. Arvio, R. Johansson, et al. 2002. Cardiovascular manifestations in 75 patients with Williams syndrome. J. Med. Genet. 39:554–558.
16. Lumaka, A., G. Mubungu, C. Nsibu, B. P. Tady, T. Lukusa, and K. Devriendt. 2012. X-linked adrenal hypoplasia congenita: a novel DAX1 missense mutation and challenges for clinical diagnosis in Africa. Eur. J. Pediatr. 171:267–270.