Dear Editor,

Pituitary gigantism is caused by chronic growth hormone (GH) hypersecretion by a pituitary lesion before epiphyseal fusion. Genetic causes have been identified in nearly 50% of patients with pituitary gigantism, with germline mutations in the AIP gene being the most frequent cause (Rostomyan et al. 2015). Recently, a new form of pituitary gigantism, X-linked acrogigantism (X-LAG), was described (Trivellin et al. 2014). X-LAG is due to chromosome Xq26.3 duplication, and GPR101 is the disease-associated gene (Trivellin et al. 2014, Iacovazzo et al. 2016). X-LAG is characterized by mixed GH/prolactin-secreting pituitary macroadenomas and/or hyperplasia in early childhood (Beckers et al. 2015). X-LAG typically occurs sporadically in females, but somatic mosaicism also occurs in males; familial mother-to-son transmission of the Xq26.3 duplication has been reported in three familial isolated pituitary adenoma families (Trivellin et al. 2014, Daly et al. 2016, Gordon et al. 2016, Iacovazzo et al. 2016). The clinical presentation of X-LAG syndrome differs from other genetic forms of pituitary gigantism (Rostomyan et al. 2015), and many well-known historical cases of gigantism share the clinical characteristics of X-LAG syndrome (Beckers et al. 2015, Rostomyan et al. 2015). If untreated during childhood, X-LAG leads to established extreme gigantism (>1.9 m) before puberty (Daly et al. 2016).

We studied a historical case of severe acrogigantism. The subject, J.K., was born in 1872 in Reutlingen, in what is now Baden-Württemberg, Germany. His parents and brother were of normal size. It was reported by his doctor that J.K. had always been 'very large' and he was reputed to have a huge appetite; he measured 1.94 m at the age of 14 and never stopped growing thereafter (Launois & Roy 1904). The long bones and extremities were elongated, and the proximal humeral epiphyses remained unfused (Launois & Roy 1904). Concomitant hypogonadism (testicular atrophy) was present on examination and post-mortem. Current forensic analysis of the skeleton demonstrates bleaching of the bones consistent with reported preservation by prolonged boiling.

Given the clinical history of early-onset acrogigantism, an underlying genetic cause was thought to be likely. DNA extraction from teeth was unreliable as the skull was edentulous when originally photographed in 1904 (Launois & Roy 1904); subsequently, teeth were added to the skull but they could not be confirmed as having been obtained from the subject himself. The skeleton was fragile after preservation by prolonged boiling, and DNA extractions from a metatarsal and the femur were unsuccessful. Based on results obtained from ancient skeletal remains, tissue from the cochlea was obtained via the petrous temporal bone (Pinhasi et al. 2015) as reported in Supplemental materials (see section on supplementary data given at the end of this article). His history of early-onset, severe pituitary gigantism led us to suspect X-LAG (Trivellin et al. 2014, Daly et al. 2016).

The DNA sample was assayed using a ddPCR technique as previously described (Daly et al. 2016). Briefly, the ddPCR compared copy number variations (CNV) at the GPR101 gene as compared with ZIC3, a gene that is not duplicated in X-LAG syndrome. Daly and coworkers recently showed this method was
reliable for both confirming the results of known Xq26.3 duplication carriers and non-carriers, while also identifying the duplication of GPR101 during screening of previously undiagnosed cases of X-LAG syndrome; the false-positive rate in the acrogigantism population was low (1/64 cases was borderline above the CNV minimal and maximal thresholds for duplication) (Daly et al. 2016). The study was approved by the Natural History Museum of Mons, and genetic studies regarding causes of gigantism and endocrine tumors were conducted under approval of the Ethics Committee of the Centre Hospitalier Universitaire de Liège, Belgium.

We extracted DNA from cochlear core powder, and then sequenced and post-processed it on a custom bioinformatics platform for ancient DNA. The DNA had deamination patterns consistent with that of an approximately 100-year-old sample. No pathological variants in AIP and other gigantism-associated genes were noted on DNA read

Figure 1
Panel A shows a photograph of the subject J.K. printed for publicity during his visit to Belgium immediately before his hospitalization in November 1901 at which time he was approximately 259 cm in height. Gigantism and acral enlargement are evident. Panel B shows the subject’s skull in a lateral photograph from 1904 and illustrates marked prognathism and enlargement of the malar bones. Panel C shows an interior view of the subject’s cranium and the markedly enlarged pituitary fossa/sella turcica (open arrow). DNA was obtained by drilling a small core in the petrous part of the temporal bone to reach the cochlea (inset); the process is minor and non-destructive (filled arrow).
sequences. DNA was insufficient for array comparative genomic hybridization studies. The subject's DNA exceeded the statistical thresholds for GPR101 duplication (copy number variation (CNV) value: 3.49 vs >2.0; Poisson CNV minimum value: 3.28 vs >2.0), indicating X-LAG as a likely cause of his severe pituitary acrogigantism.

The increased copy number for GPR101 strongly suggests that J.K. suffered from X-LAG syndrome. The clinical and tumoral characteristics of X-LAG are supportive of this proposed diagnosis. To achieve a height of 194 cm at the age of 14 years in the 1870s required a significant period of uninterrupted overgrowth. Apart from X-LAG, few other conditions are associated with early childhood-onset pituitary gigantism. The skeleton had no evidence of McCune–Albright syndrome, and AIP mutation-associated gigantism typically begins in mid-adolescence (Rostomyan et al. 2015). X-LAG syndrome is associated typically with mixed GH and prolactin-secreting pituitary adenomas (with variable hyperplasia); tumors are usually macroadenomas and can be large and invasive as in the subject's case (Trivellin et al. 2014, Beckers et al. 2015). Testicular atrophy was also present in the current case that indicates significant effects of the pituitary tumor on gonadal function. This would have kept the epiphyses open, thereby contributing to the final height of 259 cm. His proximal humeral epiphyses never fused, and his femoral epiphyses only fused late in his life, according to the autopsy accounts (Launois & Roy 1904). Hypogonadism could also have been compounded by hyperprolactinemia, which is common in X-LAG syndrome (Trivellin et al. 2014, Beckers et al. 2015). Few adult cases of X-LAG have been characterized, so the prevalence of hypogonadism due to tumor impingement is not known, but it is a logical effect of a large tumor mass.

Advances in sequencing technologies, DNA extraction methods and bioinformatic analysis have allowed researchers to overcome challenges such as DNA authentication and contamination from modern and ancient sources and have provided access to genetic material from samples thousands of years old and originating in tropical and desert environments (Pinhasi et al. 2015, Skoglund et al. 2016). These extreme conditions are not optimal to the preservation of DNA. DNA has been retrieved from skeletal remains of other giants using techniques that have relied on DNA recovery from molar teeth. In a recent historic gigantism case, no AIP mutation was found; X-LAG was not studied but ddPCR specific to GPR101 and ZIC3 similar to that reported here could be informative (Radian et al. 2016). As no DNA could be derived from teeth or other bony sites, we utilized the novel approach of retrieving cochlear DNA from within the petrous temporal bone as developed and validated recently in population studies (Pinhasi et al. 2015, Skoglund et al. 2016). Even in skeletal samples that are preserved by suboptimal means during preservation, retrieval of DNA and strongly supportive information that is consistent with the patient's clinical history can be obtained. This approach could be applied more generally to other possible genetic disorders that are evident on ancient or preserved skeletal remains. Also, studies similar to this case could address the utility of other approaches to CNV analysis such as quantitative PCR on non-amplified genomic material. The increasing number of ancient genomes that have been retrieved from phenotypically normal skeletons could provide a useful database of normal ancient DNA features and highlight the interference by environmental factors and age on the analysis and assessment of potential disease-related variants.

Valuable medical information can be gained from the genetic study of skeletal remains. Although J.K. died before the era of pituitary tumor treatment, pituitary gigantism remains a difficult-to-treat condition even today (Beckers et al. 2015, Rostomyan et al. 2015). Better understanding of the natural disease history and potential genetic causes of historical giants should reinforce the need for early effective disease control.

Albert Beckers1
Daniel Fernandes2,3
Frederic Fina4
Mario Novak2
Angelo Abati5
Liliya Rostomyan1
Albert Thiry6
L'Housine Ouafik4
Bertrand Pasture7
Ron Pinhasi2
Adrian F Daly1

1Department of Endocrinology, Centre Hospitalier Universitaire de Liège, University of Liège, Domaine Universitaire du Sart-Tilman, Liège, Belgium
2School of Archaeology and Earth Institute, University College Dublin, Belfield, Dublin, Ireland
3Department of Life Sciences, Centro de Investigação em Antropologia e Saúde, University of Coimbra, Coimbra, Portugal
References


Received in final form 22 December 2016
Accepted 3 January 2017
Accepted Preprint published online 3 January 2017