CLINICAL STUDY

**Soluble α-Klotho: a novel serum biomarker for the activity of GH-producing pituitary adenomas**

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**Abstract**

*Objective:* Klotho is a lifespan-influencing gene expressed mainly in the kidneys. Soluble α-Klotho (αKL) is released into the circulation. In this study, we present baseline αKL serum levels of patients with acromegaly compared with controls with other pituitary adenomas and assess changes following transsphenoidal surgery.

*Design:* Prospective controlled study.

*Methods:* We measured soluble αKL (sandwich ELISA) and IGF1 (RIA) in sera of 14 patients (eight females and six males) with active acromegaly and in 22 control patients (13 females and nine males) operated for non-GH-producing pituitary adenomas. Immunohistochemical staining for Klotho was performed in resected adenomas and in normal pituitary tissue samples.

*Results:* Soluble αKL was high in the acromegaly group preoperatively (median 4217 pg/ml, interquartile range (IQR) 1812–6623 pg/ml) and declined after surgery during early follow-up (2–6 days; median 645 pg/ml, IQR 550–1303 pg/ml) (P<0.001) and during late follow-up (2–3 months post-operatively, median 902 pg/ml, IQR 497–1340 pg/ml; P<0.001). In controls, preoperative soluble αKL was significantly lower than in acromegalics, 532 pg/ml (400–677 pg/ml; P<0.001). Following surgery, soluble αKL remained low during early and late follow-up — changes over time within the control group were not statistically significant. These results were independent of age, sex and kidney function. Klotho staining was equal or slightly decreased in GH-positive adenomas compared with controls.

*Conclusion:* High soluble αKL serum levels were specific to GH-producing adenomas and decreased rapidly following adenoma removal. Thus, soluble αKL appears to be a new specific and sensitive biomarker reflecting disease activity in acromegaly. Similar Klotho staining patterns in controls and acromegalics suggest that the rise in serum αKL is caused by systemic actions of pituitary GH rather than due to increased expression of Klotho by the pituitary (adenoma).

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Introduction

The α-klotho gene (αKL) was serendipitously discovered as a lifespan-influencing gene in mice after recognition that its disruption caused accelerated ageing (1). Lifespan extension was achieved by overexpression (2). In humans, variants of αKL were associated with ageing (3) and the phenotype of a rare homozygous nonsense mutation was described (4). The gene and protein were named after the Greek goddess Klotho who spins the thread of life. αKL encodes αKL protein, a 130 kDa type 1 membrane protein (1014 amino acids long) considered the founder of the Klotho family that is predominantly expressed in the kidneys, choroid plexus and several endocrine organs including the pituitary, parathyroid, testis, ovary, placenta and pancreas (1). The αKL protein exists in two forms with distinct functions. Membrane-bound αKL is a co-receptor for fibroblast growth factor 23, a bone-derived phosphaturic hormone that inhibits renal phosphate reabsorption and calcitriol production (5, 6). Soluble αKL attenuates insulin/insulin-like growth factor 1 (IGF1) signalling and regulates calcium homoeostasis (7, 8). The extracellular domain of the membrane-bound form can be enzymatically cleaved (ectodomain shedding) and released as soluble αKL into blood, urine and cerebrospinal fluid (9, 10). Recently, a sandwich ELISA for the measurement of soluble αKL was established and is now commercially available (11). Soluble αKL was inversely related to age in healthy subjects (11) and
with mortality in the elderly (12). High levels of soluble zKL were found in human umbilical cord blood (13), and based on a positive correlation between plasma levels of soluble zKL and growth and metabolic parameters in premature and term neonates, it has been speculated that Klotho may play a role in the stimulation of growth (14).

GH excess due to benign adenomas of the pituitary gland is the major cause for acromegaly with an incidence of approximately four cases per 1 million persons per year (15). Acromegaly is usually diagnosed with a considerable delay and therefore possibly associated with increased mortality even after curative transsphenoidal surgery (16). Clinical features develop slowly over many years and include metabolic derangements and pathognomonic changes in the patient’s appearance, mainly soft tissue swelling and skeletal bone growth, resulting in typical acral enlargement and coarse facial features. Metabolic changes include increased plasma glucose in the wake of insulin resistance despite reduced visceral fat and high levels of serum phosphate accompanied by higher than normal renal glomerular filtration (17, 18, 19). Both insulin resistance and elevated serum phosphate are associated with increased mortality in the general population (20, 21). In the setting of reduced visceral fat and increased glomerular filtration, insulin resistance and serum phosphate elevation are unusual, and the underlying mechanisms are unclear. Currently, GH (the hormone directly produced by the adenoma) and IGF1 (a GH-dependent, predominantly liver-derived hormone) are the classical biochemical markers of disease activity in acromegaly.

We recently found that serum-soluble zKL is markedly elevated in patients with acromegaly and that this zKL excess reversed following adenoma removal (22). In order to assess whether high serum levels of soluble zKL are specific to patients with GH-producing pituitary adenomas, we now present a prospective controlled study that documents the baseline serum levels of zKL in patients with acromegaly compared with a control group of patients with other pituitary adenomas. Additionally, we monitor the temporal changes in zKL levels following transsphenoidal surgery, and we report detailed immunohistochemical analysis of the adenomas removed.

Materials and methods

Patient characteristics

We included 14 consecutive patients with active acromegaly (eight females and six males) with a mean age of 48 years (range 29–81 years) who underwent transsphenoidal surgery at the University Hospital Zurich. The preoperative diagnosis of acromegaly was based on pathognomonic clinical findings and biochemical markers (excess IGF1 and GH, nonsuppressible during a 75 g oral glucose tolerance test (oGTT)). Patients were excluded when histopathological examination failed to identify a GH-producing adenoma. As a control group, we included 22 patients (13 females and nine males) with a mean age of 48 years (range 14–81 years) operated for pituitary adenomas not producing GH (13 non-functioning adenomas (NFAs) and nine prolactinomas). All patients provided written informed consent and the study was approved by the Local Ethics Committee. Detailed patient characteristics are summarized in Tables 1 and 2. Tumour volume calculation was based on preoperative magnetic resonance imaging (MRI) and the diameter method (tumour volume = 4/3πx½y²z², where x, y and z are the maximum diameters within the three axes. Surgical strategy was transnasal transsphenoidal using micro-surgical technique and intraoperative MRI (PoleStar N20, 0.15T, Medtronic Navigation, Minneapolis, MN, USA). All procedures were performed by the senior author (R-L Bernays) – details on the surgical strategy have been described previously (23).

Histopathology

All pathology materials, consisting of H&E and reticulin-stained sections, a full panel of anterior pituitary hormone immunohistochemistry, and the MIB-1 (Ki-67) proliferation marker were reviewed to confirm the diagnosis. Pituitary adenomas were then categorized as either GH-positive adenomas, adenomas without hormone expression (NFAs) or prolactinomas. GH antibodies were obtained from Thermo scientific (Waltham, MA, USA; MA5-11926, 1:3000) and Klotho antibodies were purchased from Abcam (Cambridge, UK; ab68208, 1:40). Immunohistoenotypic analysis was performed using a Leica Bond-Max automated immuno-nostainer employing 2 µm-thick, formalin-fixed, parafin-embedded sections. GH signal detection was performed using 3,3′-diaminobenzidin (brown) and 3-amino-9-ethylcarbazole (red) for Klotho detection respectively. Sections were counterstained with haematoxylin. For co-localization analysis, the sections were treated with citric acid for 15 min at 95 °C and immunofluorescent co-stainings were performed according to the Bond staining protocol. GH-positive areas were detected using the secondary antibody Alexa 488 anti-mouse (1:1000) and Alexa 594 anti-rabbit antibody (1:1000) for Klotho signals respectively.

Assays

All blood samples were drawn around the same time in the morning after overnight fasting. Soluble zKL was determined using a sandwich ELISA described by Yamazaki et al. (11) (Kyowa Hakko Kirin Co. Ltd., Tokyo, Japan) according to the manufacturer’s instructions. IGF1 was measured by RIA after the removal of carrier proteins as described elsewhere (22) and...
GH by IRMA (hGH-RIATC; CIS Bio International, Oris Industries, Gif-Sur-Yvette, France). Creatinine was measured using the kinetic Jaffe method on a Roche COBAS 8000 analyzer (Roche Diagnostics) and glomerular filtration rate (GFR) was estimated using the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) formula. These values were recorded as soluble αKL is associated with renal function (11). Both soluble αKL and IGF1 were measured preoperatively, at least once shortly after surgery (2–6 days post-operatively, before discharge) and again at the first outpatient follow-up (2–3 months after surgery). In all control patients and in a subgroup of seven acromegaly patients, multiple short-term post-operative measurements were performed. GH was measured preoperatively in all patients.

**Table 1** (A) Patient characteristics of the acromegaly group (n=14); immunohistological reactivity to other hormones than GH is shown in parentheses. (B) Patient characteristics of the control group (n=22); minor immunohistological hormonal reactivity is shown in parentheses. Preoperative GH, α-Klotho and IGF1 serum levels in controls with minor GH positivity were within the normal range (see Supplementary Table 1, see section on supplementary data given at the end of this article) and the latter two markers did not show the post-operative decrease characteristic for the acromegaly group.

### Statistical analyses

Statistical analyses were performed using commercially available software IBM SPSS Statistics 20 (SPSS, Inc.) and Matlab (www.mathworks.com). Continuous variables are presented as median with interquartile range (IQR). Preoperative variables were compared between groups using the Mann–Whitney U test or the c² test as appropriate. The Wilcoxon signed-rank test was used for comparing pre- and post-operative values within one group. To corroborate the results of bivariate testing, we constructed a general linear model (GLM) for repeated measures. The dependent variable was soluble αKL after log-transformation. The within-subjects factor was time with the levels preoperative, early post-operative and late post-operative. Between-subjects
factors were group and sex, and covariates were age and preoperative GFR. Two-tailed P values < 0.05 were considered statistically significant.

**Results**

**Patient characteristics**

Both study groups, acromegalics (eight females and six males; mean age 48 years) and controls (13 females and nine males; mean age 49 years), had successful transsphenoidal removal (at least debulking) of their adenomas – detailed histopathological results are shown in Fig. 1. Within the group of patients with active acromegaly, median estimated tumour volume was 1484 mm$^3$ (IQR 503–3786 mm$^3$). Of all 14 acromegalics patients, two (14%) patients presented with microadenomas, whereas 12 (86%) presented with macroadenomas. In the control group ($n$ = 22; 13 females and nine males; 13 NEAs and nine prolactinomas; mean age of 49 years), median preoperative tumour volume was 2301 mm$^3$ (580–6444 mm$^3$). Microadenomas were seen in two cases (9%). When preoperative tumour volume was compared between both groups, no statistical difference was identified ($P = 0.45$, Mann–Whitney $U$ test).

**Soluble serum zKL**

Soluble zKL was high in the acromegaly group before surgery (Figs 1 and 2) with a soluble zKL median of 4217 pg/ml (1813–6624 pg/ml), then levels declined after removal of the GH-producing adenoma to a median of 646 pg/ml (550–1303 pg/ml) ($P < 0.001$, Wilcoxon signed-rank test) during early follow-up (2–6 days post-operatively), then to a median of 902 pg/ml (498–1341 pg/ml; $P < 0.001$) during late follow-up (2–3 months post-operatively) – soluble zKL kinetics are shown in Fig. 1. Compared with acromegals, the preoperative median of soluble zKL in controls was significantly lower, 532 pg/ml (400–678 pg/ml; $P < 0.001$). Following surgery, soluble zKL stayed low with 404 pg/ml (320–635 pg/ml) during early follow-up and 524 pg/ml (359–621 pg/ml) during late follow-up – changes over time within the control group were not statistically significant. The relative drop of soluble zKL (early follow-up/preoperative) was more pronounced in the acromegaly group, 0.25 (range 0.1–0.5) compared with controls, 0.95 (range 0.5–1.6) ($P < 0.001$). Short-term kinetics of zKL levels of individual acromegalic patients compared with controls is plotted in Fig. 2A. To corroborate the results of bivariate testing, we constructed a GLM for repeated measures after log-transformation of our data. The only significant interaction was found between TIME and GROUP ($P = 33$, hypothesis degrees of freedom (df) 2, error df 17, $P < 0.001$) – the temporal changes of the two groups on a log-scale are not parallel. Group differences and significant post-operative decrease in the acromegaly group concerning soluble zKL levels were independent of age, sex and kidney function (GFR).

**IGF1**

As expected, preoperative median IGF1 levels were higher in the acromegaly group, 483 ng/ml (367–640 ng/ml) compared with the control group, 86 ng/ml (53–136 ng/ml) ($P < 0.001$). Within the acromegaly group, median preoperative IGF1 levels of 483 ng/ml returned to median early post-operative (2–6 days post-operatively) IGF1 levels of 182 ng/ml (144–229 ng/ml; $P < 0.001$), whereas no significant difference was found between preoperative and post-operative IGF1 levels within the control group. The long-term IGF1 time course within the acromegaly group (Fig. 1A) and controls (Fig. 2A) is illustrated in Fig. 1. The short-term time course of IGF1 levels of individual acromegalic patients compared with controls are plotted in Fig. 2B.
disease features improved as judged by all patients and treating physicians. Median preoperative GH levels in the control group were 0.3 ng/ml (0.1–0.7 ng/ml) – post-operative measurements of GH were only performed in the acromegaly group. The covariation of soluble α-Klotho, IGF1 and GH is shown in Supplementary Figure 1, see section on supplementary data given at the end of this article.

BMI and kidney function

BMI was significantly higher in the acromegaly group (P=0.010) with a median BMI of 28.9 kg/m² (24.6–34.7 kg/m²) compared with a median BMI of 24.1 kg/m² (20.2–27.8 kg/m²) in the control group. In terms of preoperative kidney function, there was no significant difference regarding preoperative GFR (P=0.35) with a median preoperative GFR of 104 ml/min per 1.73 m² (97–118 ml/min per 1.73 m²) and 98 ml/min per 1.73 m² (81.25–120 ml/min per 1.73 m²) in acromegals and in controls respectively. However, distinct changes were observed in response to surgery: post-operative GFR decreased in patients undergoing transphenoidal surgery for GH-producing adenomas, median ΔGFR (post-operative GFR–preoperative GFR) was −4.50 ml/min per 1.73 m² (−9.00 to −0.75 ml/min per 1.73 m²), whereas GFR increased in controls with a median ΔGFR of 6.50 ml/min per 1.73 m² (3.50–11.50 ml/min per 1.73 m²) (P<0.001). The difference between preoperative and post-operative GFR was statistically significant in both acromegals (P=0.006) and controls (P=0.001).

Immunohistochemistry

GH-producing adenomas (Fig. 3A, B and C) showed variable GH expression within the tumour (brown) from samples with strong, diffuse immunoreactivity to those with weaker and/or focal paranuclear staining. These staining patterns corresponded to either densely or sparsely granulated subtypes. In cases with focal or weak GH staining, effacement of the normal lobular architecture confirmed the presence of tumour. The Klotho expression pattern (red) in corresponding areas was more diffuse and independent of GH-positive cells. The cells in the GH-negative control group, consisting of hormone-inactive pituitary adenomas (Fig. 3D and E), and a prolactinoma (Fig. 3F) showed diffuse and strong positivity for Klotho. There is no stringent

![Figure 1](https://example.com/figure1.png)  
**Figure 1** Time course of serum-soluble α-Klotho in acromegals (A, red boxes (n=12)) and controls (B, blue boxes (n=12)) and IGF1 (white boxes) in both groups. Preoperative, early (2–6 days post-operatively, at discharge from hospital) and late (2–3 months post-operatively) follow-up values are plotted. Note the different scales in (A) and (B).

![Figure 2](https://example.com/figure2.png)  
**Figure 2** Individual short-term time course of soluble α-Klotho (A) and IGF1 (B) in acromegaly patients (n=7) (coloured lines) compared with box-plots of all control patients.
co-localization of these two markers. Immunofluorescence analysis (Fig. 4) emphasizes that Klotho (Alexa 594, red) is independently expressed from GH-positive cells (Alexa 488, green). Original magnification 400×.

Klotho and GH staining in normal pituitary is shown in Fig. 5.

Discussion

This is the first prospective controlled study that documents serum levels of zKL in patients with active acromegaly compared with a control group of patients with other pituitary adenomas. Moreover, we monitored zKL levels over time (short- and long-term follow-up) following transsphenoidal surgery. Our results show a highly significant difference in preoperative zKL levels between acromegals and controls, indicating that zKL excess is specific for GH-positive adenomas. Reversal of zKL occurs rapidly after surgery and there are no significant differences between short- and long-term follow-up, suggesting that zKL is a very sensitive marker for disease activity of acromegaly. Both the preoperative group difference and the rapid post-operative decrease in zKL levels in acromegals remained significant after adjusting our results for age, sex and kidney function. The mechanisms leading to soluble zKL excess in the serum in active acromegaly remain unclear. Soluble zKL could arise either from a distinct transcript (24) or from ectodomain shedding of membrane Klotho (9, 10, 25). It remains unclear whether in acromegals membrane-associated Klotho (mainly found in the kidneys) and soluble zKL (as detected in the serum) rise concurrently or whether elevated soluble zKL is unrelated or possibly inversely related to the abundance of plasma membrane Klotho in the kidneys – possibly resulting from enhanced enzymatic activity. Admittedly, we cannot provide our own experimental data to support this favoured hypothesis. zKL (130 kDa) appears to result from proteolytic ectodomain clipping (10). Two members of the ‘A Disintegrin and Metalloproteinase’ (ADAM) family, ADAM10 and ADAM17, have been suggested as the responsible enzymes (9), and the activity of secretases (25, 26) shedding the ectodomain from the integral membrane Klotho may be increased in acromegaly, either directly by GH or indirectly by factors or a proteolytic activity induced by GH.

For obvious reasons, renal biopsies to check for changes in membrane-bound Klotho abundance were not feasible in our study population. However, we analysed Klotho staining in GH-producing adenomas, in controls (NFAs and prolactinomas) and in normal pituitary tissue samples. The immunohistochemical staining pattern presented suggests that the rise in serum zKL is not explained by increased pituitary (adenoma) Klotho expression but rather due to an increase in pituitary GH secretion.
Until now, GH (a hormone produced by the adenoma itself) and especially IGF1 (a GH-stimulated, predominantly liver-derived peptide) have been the ‘classical’ biomarkers for diagnosing and monitoring disease activity during the treatment of patients with acromegaly. In fact, their normalization has been linked to decreased mortality (27, 28, 29). However, it has been recognized that both parameters entail various shortcomings.

Serum levels of IGF1 are influenced not only by GH status but also by age, moreover, by gender (estrogens), race, genetic makeup, liver function, nutritional status, portal insulin, thyroid hormones and by concomitant inflammatory disease. Some of these influences, particularly the former, may also have an impact on zKL (to an as yet unknown extent). Serum IGF1 is mainly derived from the liver and tightly bound to IGF binding proteins (IGFBPs) (30, 31). Changes in IGFBP concentrations contribute to the limitations known for a variety of IGF1 assays (32). To circumvent these problems, we used a classical and time-consuming assay in which carrier proteins are removed before the samples are incubated with the antibodies (33, 34).

Dynamic testing using oGTT to suppress GH is widely used; however, patients with acromegaly can demonstrate normal oGTT GH suppression despite elevated IGF1 levels (27, 35). In patients receiving non-surgical treatment of acromegaly, such as long-acting somatostatin analogues (LA-SRIFs) (36), pegvisomant (PEG-V) (37) or radiotherapy for GH-producing adenomas, GH values may be misleading due to highly irregular GH secretion pattern and flattened GH pulses (38).

The limitations (both biological and technical) of the assays used to measure GH (39) and IGF1 are well known (40), and it was stated that additional, possibly more specific and sensitive, biomarkers are desperately needed (41).

Our study has some notable limitations. The assay for zKL has been introduced only recently; therefore, knowledge of its shortcomings is limited. Additionally, the normal range for serum levels of soluble zKL has not been established. In our laboratory, we measured soluble zKL in the sera of 26 healthy volunteers (11 females and 15 males; mean age 39 years) as previously reported (22): zKL (median and IQR) was 596 (506–734) pg/ml. Similar to IGF1, serum-soluble zKL also decreases with increasing age; moreover, it may be low in patients with renal failure (11), which prompted us to routinely check creatinine. In the
context of NFA, some of our patients may have been GH deficient, but the design of our study did not allow us to determine whether these patients had lower than normal IGF1 and sKlotho serum levels. Furthermore, the molecular mechanisms of sKlotho excess in active acromegaly and the functional impact of sKlotho in acromegaly disease biology remains unknown.

**Conclusions**

Acromegaly is (thus far, to our knowledge, up to this writing) the only acquired disease known to man with excessively elevated levels of soluble sKlotho. Highly elevated soluble sKlotho is specific to GH-producing adenomas of the pituitary gland and rapidly decreases following adenoma removal. Thus, soluble sKlotho appears to represent a new, quite specific and fairly sensitive biomarker reflecting disease activity in patients with acromegaly.

**Supplementary data**

This is linked to the online version of the paper at http://dx.doi.org/10.1530/EJE-12-1045.

**Declaration of interest**

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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**References**


