Telomere length analysis in Cushing’s syndrome

Anna Aulinas1, María-José Ramírez2, María-José Barahona3,4, Elena Valassi1,4, Eugenia Resmini1,4, Eugénia Mato1,5, Alicia Santos1,4, Iris Crespo1,4, Olga Bell1,5, Jordi Surrallés2 and Susan M Webb1,4

1Endocrinology/Medicine Departments, Sant Pau Biomedical Research Institute, Hospital de Sant Pau, Universitat Autònoma de Barcelona, Servei d’Endocrinologia, C/Sant Antoni Maria Claret, 167, 08025 Barcelona, Spain, 2Department of Genetics and Microbiology, Center for Biomedical Network Research on Rare Diseases (CIBERER Unit 745), Universitat Autònoma de Barcelona, Bellaterra, Barcelona, Spain, 3Endocrinology Department, Hospital Universitari Mutua Terrassa, Terrassa, Barcelona, Spain, 4Center for Biomedical Network Research on Rare Diseases (CIBER Unit 747), ISCIII, Hospital de Sant Pau, Barcelona, Spain and 5Endocrinology Department, Center for Biomedical Network Research on Bioengineering, Biomaterials and Nanomedicine (CIBER-BBN), Hospital de Sant Pau, Barcelona, Spain

Abstract

Introduction: Hypercortisolism in Cushing’s syndrome (CS) is associated with increased morbidity and mortality. Hypercortisolism also occurs in chronic depressive disorders and stress, where telomere length (TL) is shorter than in controls. We hypothesized that shortening of telomere might occur in CS and contribute to premature aging and morbidity.

Aim: To investigate TL in CS patients compared with controls.

Methods: Seventy-seven CS patients (14 males, 59 pituitary, 17 adrenal, and one ectopic; 21 with active disease) were compared with 77 gender-, age-, and smoking-matched controls. Fifteen CS were evaluated longitudinally, during active disease and after remission of hypercortisolism. Leukocyte TL was measured by telomere restriction fragment–Southern technique. Clinical markers were included in a multiple linear regression analysis to investigate potential predictors of TL.

Results: Mean TL in CS patients and controls was similar (7667 vs 7483 bp, NS). After adjustment for age, in the longitudinal evaluation, TL was shorter in active disease than after remission (7273 vs 7870, \(P < 0.05\)). Age and dyslipidemia were negative predictors \(\left(\begin{array}{l} P < 0.05 \end{array}\right)\), and total leukocyte count was a positive predictor for TL \(\left(\begin{array}{l} P < 0.05 \end{array}\right)\). As expected, a negative correlation was found between TL and age (CS, \(R = -0.400\) and controls, \(R = -0.292\); \(P < 0.05\)). No correlation was found between circulating cortisol, duration of exposure to hypercortisolism or biochemical cure and TL.

Conclusion: Even though in the cross-sectional comparison of CS and controls no difference in TL was found, in the longitudinal evaluation, patients with active CS had shorter TL than after biochemical cure of hypercortisolism. These preliminary results suggest that hypercortisolism might negatively impact telomere maintenance. Larger studies are needed to confirm these findings.

Introduction

Cushing’s syndrome (CS), a rare disease due to excessive cortisol secretion, is associated with increased mortality and severe morbidity (increased cardiovascular risk and fatigability, osteopenia, neuropsychological alterations, and impaired health-related quality of life), not completely reversible after biochemical control (1). The mechanisms by which these abnormalities do not recover completely appear to be complex and are not currently well understood. Hyperstimulation of the hypothalamic–pituitary–adrenal (HPA) axis also resulting in hypercortisolism may also occur in psychiatric diseases such as acute and chronic stress and post-traumatic stress...
disorder (2, 3). These situations are associated with poor health indexes, and telomere length (TL) has been found to be shorter than that in matched controls (4).

Telomeres are repetitive DNA sequences, located at the end of linear chromosomes, essential to maintain genomic stability. Without telomeres, genetic material could be lost after every cell division; thus, when telomeres are critically short, cell division stops and senescence and apoptosis are induced (5). To avoid telomere attrition and to maintain TL, germ-line cells and a few somatic cells produce an enzymatic complex called telomerase. Telomerase function can be regulated by genetic, epigenetic, environmental, and hormonal factors (5). These include mainly stress hormones such as cortisol, catecholamines, estrogens, and growth factors.

In this line, accelerated telomere shortening, higher levels of urinary catecholamines, and free urinary cortisol have been observed in situations with high perceived psychological stress (in sisters of patients with cancer, in acute mental stress) (6). In vitro studies have shown a 50% reduction in telomerase activity in lymphocytes after exposure to high levels of hydrocortisone (7) and a rapid and dynamic loss of telomeric sequences after exposure of mice thymocytes to dexamethasone (8). Shorter leukocyte TL has been described to be associated with elevated cortisol responses and dysregulated patterns of daily cortisol secretion in women who are patient caregivers (9). Recently, a longitudinal study evaluating the association between coexisting changes in cortisol and telomerase activity in peripheral blood mononuclear cells has been published (10). The authors examined whether participation in mindfulness-based interventions and improvements in psychological distress and metabolic factors were associated with increases in telomerase activity. They observed that serum cortisol levels were negatively correlated with changes in telomerase activity, suggesting that changes in stress-related cortisol might be one of the signals regulating telomerase levels in humans.

This evidence led us to hypothesize that telomere shortening may be behind the increased morbidity and features of premature aging in patients with CS. Hypercortisolemia could contribute to premature aging by inducing accelerated telomere shortening, which in turn could be implied in the persistent morbidity and clinical consequences associated with CS, even years after biochemical remission. As TL is an indicator of chromosome stability, proliferative capacity, and cellular aging, measuring TL could contribute to the understanding of its clinical and biological significance. To the best of our knowledge, telomere dysfunction has not been evaluated in CS patients before.

The aim of this study was to investigate TL in patients diagnosed with CS compared with sex-, age-, and smoking-matched healthy controls and to evaluate whether normalization of the hypothalamic-pituitary-adrenal axis after treatment reverses possible abnormalities.

**Subjects and methods**

**Subjects**

In this case-control study, patients with endogenous CS followed in our institution since 1982 were eligible. Patients with adrenal carcinoma were excluded. Seventy-seven CS patients and 77 controls, matched for gender, age, and smoking participated in the study. Fourteen were men (18.2%) and 63 women (81.8%). Mean age at the time of the study was 48.6 ± 12.8 years. Fifty-nine patients were of pituitary origin (76.6%), 17 of adrenal origin (adrenal adenoma or bilateral macronodular hyperplasia), and in one patient the origin was unknown (ectopic ACTH secretion of unknown source). Twenty-one patients (27.3%) had active disease at the time of the study and 56 (72.7%) were cured; mean time of remission of hypercortisolism was 6.4 ± 7.2 years. Eight active CS patients (38%) were treated with metyrapone, six (28.5%) with ketoconazole, and three (14.2%) with both drugs. Mean duration of endogenous hypercortisolism was 72 months (range 11–264). Duration of hypercortisolism was considered as the period between onset of symptoms (as referred by the patients) and remission of hypercortisolism (in patients in remission) or the time of current analysis (in active patients). The period between onset of symptoms and biochemical diagnosis of CS was 34 months (range 3–120). Twenty-two patients (28.6%) had received pituitary radiotherapy and 71 (92.2%) had undergone surgery. Fifty-three percent (n=41) were cured after initial treatment and had no recurrence and 19.5% (n=15) were cured after further therapies for recurrent disease. Fifteen cured patients (19.5%) were adrenal insufficient at the time of telomere analysis and required substitution therapy with hydrocortisone (mean dose 17.6 ± 3.7 mg and range 10–20). Nine patients (11.7%) were GH deficient (four of which were replaced with recombinant human GH (rhGH)); eight women (10.4%) were gonadotropin deficient (all on estrogen/progesterone hormone replacement therapy); and 15 patients (19.4%) were hypothyroid, ten due to thyroid-stimulating hormone (TSH) deficiency and five due to primary
hypothyroidism (all on L-thyroxine (L-T4) replacement). CS was considered in remission if either adrenal insufficiency was demonstrated (basal morning cortisol <100 nmol/l (<4 μg/dl) and/or undetectable 24-h free urinary cortisol) or morning cortisol suppression (<50 nmol/l, <1.8 μg/dl) after 1 mg dexamethasone overnight was observed. Twenty-five patients (32%) were on antihypertensive medication, 17 (22%) on statin treatment for dyslipidemia, and 12 (16%) were treated with calcium and vitamin D.

In a subgroup of 15 CS (all women) patients studied initially with active disease, a second analysis of TL was carried out once they were in remission. In this longitudinal study, three were of adrenal origin and 12 of pituitary origin. Mean age at the time of active disease was 43.5 ± 12.1 years and at remission was 46.6 ± 11.3 years. The time elapsed between both analyses was 40.1 ± 15.6 months and mean time of remission was 28.5 ± 14.1 months. Three cured patients (20%) were adrenal insufficient at the time of telomere analysis and required substitution therapy with hydrocortisone (mean dose 18.3 ± 2.2 mg and range 10–20); four patients (26.6%) were hypothyroid, two due to TSH deficiency and two due to primary hypothyroidism (all on L-T4 replacement). None of the cured patients were GH-deficient; seven women (46.6%) were postmenopausal at remission, but no gonadotropin deficiency was observed (n=8).

Seventy-seven controls selected from the blood bank donor’s database or from healthy volunteers recruited among hospital employees were matched for gender, age, and smoking status, three features known to affect TL. Namely, age is an important determinant of TL, typically decreasing with advancing age (11). Females usually present longer TL than males, because estrogens stimulate telomerase activity and protect DNA from reactive oxygen species-induced damage (12). Cigarette smoke constituents increase cumulative and systemic oxidative stress and inflammation, which induce increased white blood cell turnover, resulting in accelerated TL shortening (13). Medical history and physical examination excluded any who reported glucocorticoid exposure, severe and/or acute diseases, and severe psychiatric alterations (however, anxiety and mild depression were not exclusion criteria). Four controls (5.7%) were on antihypertensive therapy, another four (5.7%) were receiving statin treatment for dyslipidemia, and three (4.3%) were treated with calcium and vitamin D.

Anthropometry (weight, height, BMI, and waist:hip ratio) was measured in patients and controls. Hypertension was defined as systolic blood pressure >140 mmHg or diastolic blood pressure >90 mmHg or the use of antihypertensive medications. Dyslipidemia was defined as total cholesterol (TC) >220 mg/dl, LDL >130 mg/dl, triglyceride levels ≥150 mg/dl, or treatment with lipid-lowering medication. Diabetes mellitus (DM) was confirmed with fasting glucose levels >126 mg/dl in two consecutive determinations or 2-h glucose after oral glucose tolerance test (OGTT) >200 mg/dl. Adult patients were considered osteopenic when T score was <-1 and >-2.5 or osteoporotic when T score was <-2.5 s.d.

All participants provided a blood sample for DNA extraction and gave their informed consent. The study was approved by the Hospital Ethics Committee.

**Methods**

*Genomic DNA extraction from total leukocytes* 
Genomic DNA extraction from total leukocytes was carried out using an adapted Proteinase K and Phenol protocol (14). Blood samples from the patients were collected in EDTA tubes to reduce DNA degradation. Genomic DNA was isolated from blood buffy coats. The buffy coat and white blood cell pellets were stored frozen at −80 °C before processing. The white blood cell layers were harvested and digested with buffer containing 0.1 M MgCl2, 0.02 M EDTA, 0.5% SDS, 0.01 M Tris, pH 8.0, and 1 mg/ml of proteinase K at 37 °C overnight. The lysates were homogenized by passing through a blunt 20-gauge needle (0.9 mm diameter) at 4 °C temperature and DNA was purified by phenol:chloroform:isoamyl alcohol (25:24:1) extraction, and ethanol precipitation. Finally, genomic DNA was dissolved in Tris–EDTA buffer and quantified by spectrophotometric analysis. The quality of genomic DNA was checked for high molecular weight by 1% agarose gel electrophoresis.

**TL measurements**
TL measurements were carried out by the telomere restriction fragment assay (TRF) using the Telo TAGGG Telomere Length Assay Kit (Roche 12209136001). Briefly, 1 μg of DNA was digested with 20 units of Rsal and HinfI for 2 h at 37 °C. Samples were loaded on a 0.5% Seakem Gold Agarose gel and were run for 21 h at 35 V. The gels were treated with HCl, denaturalized and neutralized, and then transferred to a nylon membrane by capillarity for 12–18 h. After fixation with u.v., hybridization was carried out with a DIG-labeled telomeric probe (3 h at 42 °C). Finally, restriction washes, incubation with anti-DIG-AP antibody, and detection by chemiluminescence were carried out. Images were analysed with the program Quantity One. TRF mean was
calculated using the formula: TRF mean = ΣODi/ Σ(ODi/Li), where ODi is the chemiluminescent signal and Li is the length of the TRF fragment at position i (15). A control sample, 2 μg digested DNA derived from a single batch of HeLa cells, was run on each gel to minimize interassay variation. The mean TL for HeLa cells was 4113 bp, with a s.d. of ±210 bp, which is in the acceptable range of accuracy of the Southern blot technique. The accuracy of southern blot technique is up to ±300 bp (16).

Biochemical, hormone, and bone analyses

Routine serum determinations were carried out by standard automated laboratory methods: fasting glucose, TC, HDL and LDL cholesterol and triglyceride levels. Blood counts were made using automated cell counters. Twenty-four hours urinary free cortisol was measured with a commercial RIA made using automated cell counters. Twenty-four hours serum cortisol, and insulin-like growth factor 1 (IGF1) levels were measured using a commercial chemiluminescent immunometric assay. Lumbar spine and whole-body bone mineral density and bone mineral content were measured by DXA scanning (Delphi QDR 4500; Hologic); the mean precision error (coefficient of variation) was 1%.

Statistical analysis

Statistical analyses were carried out using the SPSS 19.0 Statistical Package for Windows (SPSS, Inc.). Initially a descriptive analysis of all variables was carried out in order to verify correct introduction of data in the database. Quantitative data are expressed as mean and s.d. (Gaussian distribution) or as median and range (non-Gaussian distribution), and categorical data are expressed as percentages. Data distribution was analyzed by the Kolmogorov–Smirnov test. TL variable was normally distributed. Logarithmic transformations were carried out where necessary to normalize the distribution of a particular measure. Comparison between two groups was made using Student’s t-test (Gaussian distribution) or Mann–Whitney’s U (non-Gaussian distribution)-test. A χ² test was performed for categorical variables. Fisher’s exact test was performed when appropriate. Pearson’s correlation coefficient was used to estimate linear association between two quantitative variables. Analysis of covariance was performed to evaluate TL after adjustment for age and for total leukocyte count (as covariates). Multiple linear regression analysis including age, gender, BMI, type 2 DM, dyslipidemia, hypertension, psychiatric history, duration of hypercortisolism, current hypercortisolism, total leukocytes, and 24-h urinary free cortisol as potential predictive factors for TL (as dependent variable) was performed. P values <0.05 were considered significant.

Results

Comparison between CS and matched controls

Main baseline characteristics of CS patients and controls are summarized in Table 1. CS patients had more hypertension, diabetes, dyslipidemia, and osteoporosis than their matched controls (P<0.05). Mean TL values in CS and controls are summarized in Fig. 1. No differences were observed between males and females (7732±1242 vs 7540±1361 bp respectively). TL did not differ between CS and controls (7667±1260 vs 7483±1214 respectively, NS). TL did not differ between active CS, cured CS (with or without secondary adrenal insufficiency) and their matched controls (Fig. 1).

As expected, a negative linear correlation between age and TL in the whole sample was observed (R = −0.341, P<0.001). When both groups were evaluated separately, this negative correlation was maintained in CS patients (R = −0.400, P<0.001) and in controls (R = −0.292, P<0.01) (Fig. 2). A positive correlation was found between IGF1 and TL in CS patients (R =0.331, P<0.05), but was not correlated with the presence or absence of GH deficiency or rhGH replacement therapy. No differences

| Table 1 Baseline characteristics of patients with Cushing’s syndrome (CS) and controls. Data are presented as percentage and mean ± s.d. |
|------------------------|------------------------|------------------------|------------------------|
|                        | CS (n=77)              | Controls (n=77)        | P                      |
| Age (years)            | 48.6 ± 12.8            | 48.4 ± 12.6            | NS                     |
| Smokers (%)            | 24.7                   | 19.4                   | NS                     |
| Alcohol consumption (%)| 26                     | 27.3                   | NS                     |
| Diabetes mellitus (type 2) (%) | 14.3               | 1.4                    | <0.05                  |
| Arterial hypertension (%)| 57.1                  | 12.9                   | <0.001                 |
| Dyslipidemia (%)        | 45.5                   | 20.0                   | <0.05                  |
| Osteoporosis (%)        | 29.9                   | 2.9                    | <0.001                 |
| Psychiatric history (%) | 37.7                   | 11.4                   | <0.001                 |
| BMI (kg/m²)            | 28 ± 5.6               | 26.4 ± 4.9             | <0.05                  |
| Waist:hip ratio        | 0.92 ± 0.07            | 0.85 ± 0.07            | <0.05                  |
| 24-h Urinary free cortisol (nmol/24 h) | 266 ± 180 | 132 ± 59               | <0.001                 |
| Morning serum cortisol (nmol/l) | 450 ± 259 | 375 ± 120              | <0.05                  |
| Leukocytes (×10⁹/l)    | 7.3 ± 2.3              | 5.8 ± 1.7              | <0.05                  |
| Neutrophils (×10⁹/l)   | 4.4 ± 2.0              | 3.5 ± 1.2              | <0.05                  |
| Lymphocytes (×10⁹/l)   | 2.1 ± 0.8              | 1.9 ± 0.4              | NS                     |
Longitudinal analysis in CS patients evaluated both during active disease and in remission

As expected, patients were older once remission was attained. Ten patients (66%) clearly showed an increment of TL upon remission of CS. In five (33%) patients, TL decreased after remission (Fig. 3), but was minimal in two and of doubtful relevance, because it was around the detection limit of 300 bp (around 4%) TL’s variation in our population (17). Moreover, after adjustment for age as covariate, TL was shorter in active disease than after remission (7273 ± 1263 vs 7870 ± 1039, respectively, \( P<0.05 \)) in the same patients (Fig. 3), in sharp contrast with TL shortening usually observed as age increases. No significant differences in the presence of hypertension, dyslipidemia, diabetes, or use of medications were observed between the group of patients who have increased TL during remission and those who did not have increased TL. Patients who incremented TL also decreased their BMI more after remission than those who did not increase TL (−2.3 vs −0.8 kg/m²), although due to the small group size, it did not reach statistical significance \((P=0.19)\). A trend for a positive correlation between TL at remission and duration of remission was also seen \((R=0.494, P=0.061)\).

Discussion

To the best of our knowledge, this is the first study to evaluate TL in this rare disease and with a relatively large series of CS patients. When investigated longitudinally, our preliminary data show that patients with active CS have a shorter TL, which become longer after hypercortisolism disappeared with effective treatment. However, in the cross-sectional case–control study comparing all patients with CS and matched controls, no differences in TL were found. This was also the case when patients with active hypercortisolism, and those considered in...
Table 2  Total leukocyte counts and leukocyte main subsets distribution (neutrophils and lymphocytes) of Cushing’s syndrome (CS) patients during active disease and remission and their matched controls. Data are expressed as mean ± s.d.

<table>
<thead>
<tr>
<th></th>
<th>CS</th>
<th>Controls</th>
<th>P</th>
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</thead>
<tbody>
<tr>
<td>Leukocytes in active disease (×10⁹/l) (n = 21)</td>
<td>8.8 ± 2.3</td>
<td>5.9 ± 1.4</td>
<td>&lt; 0.01</td>
</tr>
<tr>
<td>Neutrophils (%)</td>
<td>64.7 ± 11.0</td>
<td>55.5 ± 6.1</td>
<td>&lt; 0.05</td>
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<tr>
<td>Lymphocytes (%)</td>
<td>24.5 ± 9.1</td>
<td>32.1 ± 7.8</td>
<td>&lt; 0.05</td>
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<tr>
<td>Leukocytes in cured patients without adrenal insufficiency (×10⁹/l) (n = 41)</td>
<td>6.7 ± 2.1</td>
<td>5.8 ± 1.8</td>
<td>&lt; 0.05</td>
</tr>
<tr>
<td>Neutrophils (%)</td>
<td>57.1 ± 8.2</td>
<td>54.9 ± 13.8</td>
<td>NS</td>
</tr>
<tr>
<td>Lymphocytes (%)</td>
<td>31.1 ± 6.6</td>
<td>30.9 ± 7.1</td>
<td>NS</td>
</tr>
<tr>
<td>Leukocytes in cured patients with adrenal insufficiency (×10⁹/l) (n = 15)</td>
<td>6.6 ± 1.5</td>
<td>6.2 ± 2.1</td>
<td>NS</td>
</tr>
<tr>
<td>Neutrophils (%)</td>
<td>58.3 ± 8.7</td>
<td>52.5 ± 7.7</td>
<td>NS</td>
</tr>
<tr>
<td>Lymphocytes (%)</td>
<td>29.6 ± 9.6</td>
<td>34.5 ± 6.6</td>
<td>NS</td>
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CS patients provide a unique opportunity to examine the effects of hypercortisolism on telomere maintenance. CS determines increased morbidity and mortality, especially in the untreated state but also after therapy when compared with background population (1, 18). Severe morbidities are also increased even in the 3 years before diagnosis when compared with the normal population, and are not completely reversible after endocrine cure (18). The mechanisms by which CS patients do not recover completely after biochemical remission are still unknown. It is possible that telomere dysfunctions partially contribute to these abnormalities. In other situations where hypercortisolism is often present, such as chronic stress and some psychiatric conditions, TL has been found to be shorter than that in matched controls (6, 9). These previous evidences took us to hypothesize that TL shortening could contribute to the increased morbidity and features of premature aging observed in endogenous hypercortisolism of CS. Thus, we planned this study in order to investigate the telomere system in these patients.

We have evaluated a significant number of CS patients (n = 77), a rare disease with an incidence ranging from 0.7 to 2.4 cases/million inhabitants per year (19). They were carefully matched for age, gender, and smoking status with controls. These relatively small groups may contribute to explain why no differences in TL were observed between CS and controls. Furthermore, many other factors apart from hypercortisolism may affect TL, both individual and environmental (genetic, epigenetic, socioeconomic status, lifestyle, growth factors, etc.) (5). In addition, TL may be affected by what is known as a ‘pseudolengthening’ mechanism (20); specifically, TL of lymphocytes becomes increasingly shorter than those of granulocytes over the years (17). And as a redistribution of leukocyte cell type is often seen in hypercortisolism (lymphopenia and neutrophilia), this may also affect the measured TL obtained from the total leukocyte count (21). In fact, we did find that in active disease, total leukocyte and neutrophil counts were higher and lymphocytes were lower than that observed in matched controls. We observed that total white blood cell counts in each individual blood sample also affected TL, and CS patients had higher total leukocyte counts compared with healthy controls, similar to other series (21). However, after adjustment for total leukocyte count (as a covariate), no differences in TL between CS and their healthy controls were identified.

In the multiple regression analysis, leukocytes count together with age and the presence of dyslipidemia were the predictive factors for TL, explaining 23% of the TL present in our CS patients. Not surprisingly, age was a negative predictive factor for TL, in the whole sample and in the different subgroups analysed. A positive correlation was also seen between IGF1 levels and TL, as described in healthy population (11, 22). Both findings support the reliability and validity of our results and the methodology used, since similar correlations have been described in much larger populations (but not in CS patients) (14); namely TL was positively correlated with serum IGF1 and negatively associated with age in a cohort of 476 healthy Caucasians aged 16–104 years (22). We also observed a negative correlation between TL and dyslipidemia as described in other paradigms, in which cholesterol has been associated with faster biological aging (23).

As expected, some baseline characteristics differed between CS and controls, such as serum morning cortisol and 24-h urinary free cortisol, certain cardiovascular risk
factors, and psychiatric conditions (anxiety and depression), which were more prevalent in CS patients. Most of these features have recently been related to telomere dysfunctions (9, 24), although not all results published in the literature are concordant (25). Even though they did not seem to have impact on TL in the case–control regression analysis, with the exception of dyslipidemia which negatively affected TL, we cannot rule out that in much larger studies some of these clinical features could determine TL in some way or another. We did not find any influence of medical treatment to reduce cortisol during active disease or glucocorticoid replacement in patients with adrenal insufficiency after CS therapy on TL.

The longitudinal analysis of 15 patients evaluated both during hypercortisolism and in remission, adjusting for age (as a covariate), confirmed our initial hypothesis, because patients with hypercortisolism during active disease did have shorter telomeres than later in remission (average 596 bp). In spite of being 40.1 ± 15.6 months older at remission, TL was longer and positively associated with duration of remission. Although this finding is preliminary based on a small number of patients, and should be confirmed in the future in larger studies, it would support our initial hypothesis of a negative effect of a hyperactive hypothalamic–pituitary–adrenal axis on TL and cell senescence observed in other studies. Accelerated telomere shortening was observed in a group of 647 women (who had a sister with breast cancer) with higher perceived stress and higher levels of urinary free cortisol and cathecolamines (6). Similarly, shorter buccal cell TL was observed in children exposed to laboratory stressors with higher levels of salivary cortisol and higher autonomic reactivity (26). Greater cortisol responses and dysregulated patterns of daily cortisol secretion were associated with shorter leukocyte TL in 14 postmenopausal women caregivers of a partner with dementia compared with matched noncaregiver controls (27). Consistent with this and with our longitudinal results, one in vitro study observed how exposure to high hydrocortisone levels, comparable with those that might be reached in vivo during stress, reduced telomerase activity in lymphocytes (7). As the major pathway for telomere lengthening seems to be through telomerase activation, this could explain why a patient could have shorter TL during hypercortisolism. It is probably that when cortisol normalizes, a recovery of telomerase activity takes place, increasing TL or lowering attrition rates.

Contrary to this evidence and to our results, a recent publication showed that telomere shortening associated with hypocortisolism was observed in patients with high levels of chronic stress exposure or high degrees of inflammation, which could lead to an exhaustion of the HPA axis. It is difficult to identify the mechanism responsible for accelerated telomere shortening in hypocortisolism, often preceded by a hypercortisolemic phase in long-term chronic stress exposure, suggesting that TL could be a measure of cumulative stress (28). We found no differences in TL in our hypocortisolemic patients compared with cured patients without secondary adrenal insufficiency; an explanation could be that all adrenal-insufficient patients were correctly replaced with hydrocortisone.

Lifestyle modifications such as increased physical activity after remission may also increase TL, as reported in some studies, by inducing changes in telomerase activity. The mean fall in BMI in patients who increased TL was greater than in those who decreased TL after
remission (−2.3 vs −0.8 kg/m²), but did not reach statistical significance, probably due to the small sample size in the longitudinal evaluation. This change in BMI may contribute to explain the increase in TL in cured patients, similar to that seen in a recent longitudinal intervention study with Mediterranean diet, where BMI was inversely correlated with changes in TL (29).

A model of dynamic telomere balance under stress has been suggested, in which severe stress first would lead to increased turnover and depletion of circulating cells followed by a compensatory re-population when stress ends (in short stress conditions). This model could also be present in CS patients, but has to be confirmed. It would appear to be important to distinguish between true reversal of telomere shortening and replenishment by younger cells (‘pseudo lengthening’) that probably takes place in CS after remission (20).

The study has several limitations. The sample size, although respectable considering that CS is a rare disease, precludes any analysis in different etiological subgroups of CS. This also did not allow to control for all potential confounders, especially medical treatment during active disease, physical activity, current stress, etc. Especially in hypocortisolemic patients after surgery for CS, a perfect cortisol replacement is an elusive goal. Although the results of the longitudinal evaluation are opposite to what is expected by increasing age, and is an interesting result, it is certainly preliminary based on a small group of patients. We could not include the remaining six active patients, because four of them still had active disease and two were lost to follow-up. A larger group of patients, as well as a longer longitudinal follow-up, would clearly strengthen the conclusion of these preliminary findings. White blood cells, the most characterized tissue source for telomere studies, easily obtainable from peripheral blood, may vary in their cell type’s distribution in blood as seen in CS patients. TL variability even in the same cell and for individuals of similar age complicates any conclusions on telomere biology in CS patients (30). Most studies on telomere biology and aging are much larger and cross-sectional, but large scale, longitudinal, prospective, and well-designed studies are lacking. It would be interesting to evaluate TL in other tissues such as the pituitary or the adrenal in CS, because glucocorticoids induce changes in the immune system; however, this would be even more difficult than obtaining peripheral leukocytes for TL evaluation. In addition, we could not measure telomerase activity (which required fresh processing), which would provide a more direct approach to both the telomere system and its dynamics.

The main conclusion of this study is that in individual CS patients in whom hypercortisolism is controlled after successful treatment, TL increases despite being on average 3 years older. It would appear, therefore, that telomerase activity would be induced once hypercortisolism disappears, and this could be one of the mechanisms by which increased morbidity, mortality, and biological aging improve when disease is controlled. However, in the entire group of CS patients, no difference in TL was observed when compared with healthy controls, pointing to the fact that many other factors determine TL apart from age, including dyslipidemia, healthier lifestyles or differences in leukocyte subsets cell counts. Larger prospective studies are required to confirm these changes in TL in CS patients and to investigate the implications of these abnormalities further.

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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