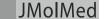
ORIGINAL ARTICLE





Glucocorticoids dexamethasone and prednisolone suppress fibroblast growth factor 23 (FGF23)

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Abstract

Fibroblast growth factor 23 (FGF23) is a hormone mainly secreted by bone cells. Its most prominent effects are the regulation of renal phosphate reabsorption and calcitriol (active vitamin D, 1,25(OH)₂D₃) formation, effects dependent on its co-receptor α Klotho. Besides these actions, further paracrine and endocrine effects exist. The production of FGF23 is regulated by 1,25(OH)₂D₃, parathyroid hormone, dietary phosphate intake, iron status, as well as inflammation. Glucocorticoids are hormones with anti-inflammatory properties and are, therefore, widely used for acute and chronic inflammatory diseases, autoimmune disorders, and malignancies. The present study explored whether glucocorticoids influence the production of FGF23 in vitro as well as in mice. Fgf23 transcription was analyzed by semi-quantitative real-time PCR. Serum concentrations of FGF23 and 1,25(OH)₂D₃ were measured by ELISA. Urinary phosphate and Ca²⁺ excretion were determined in metabolic cages. As a result, in UMR106 rat osteoblast-like cells and in MC3T3-E1 cells, both, dexamethasone and prednisolone, downregulated Fgf23 transcription and FGF23 protein synthesis. Dexamethasone increased Dmp1 and Phex (encoding FGF23-regulating genes) as well as Nfkbia (encoding NF κ B inhibitor I κ B α) transcription in UMR106 cells. In mice, a single injection of dexamethasone or prednisolone was followed by a significant decrease of serum C-terminal and intact FGF23 concentration and bone Fgf23 mRNA expression within 12 h. These effects were paralleled by increased renal phosphate excretion and enhanced 1,25(OH)₂D₃ formation. We conclude that a single glucocorticoid treatment strongly downregulates the FGF23 plasma concentration.

Key messages

- Glucocorticoids dexamethasone and prednisolone suppress the formation of bone-derived hormone fibroblast growth factor 23 (FGF23) in vitro.
- The effect is accompanied by an upregulation of Dmp1, Phex, and IκBα, negative regulators of FGF23, in UMR106 osteoblast-like cells.
- Glucocorticoid receptor antagonist RU-486 attenuates the effect of dexamethasone on FGF23, Dmp1, and Phex.
- In mice, a single glucocorticoid dose suppresses FGF23 and enhances 1,25(OH)₂D₃ (active vitamin D).

Keywords Phosphate $\cdot 1,25(OH)_2D_3 \cdot \alpha Klotho \cdot Inflammation$

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Introduction

The phosphaturic hormone fibroblast growth factor 23 (FGF23) is mainly synthesized in bone cells (osteoblasts/osteocytes) and affects renal phosphate handling and the production of calcitriol (1,25(OH)₂D₃), the active form of vitamin D [1, 2]. These effects are mediated by the concerted action of the receptor for FGF23 (FGFR) and transmembrane protein α Klotho which serves as a co-receptor [2]. Binding of FGF23 to this receptor results in the internalization of secondary-active phosphate transporter NaP_iIIa residing in the apical cell



membrane of the proximal tubule and in downregulation of CYP27B1, the key enzyme for the generation of 1,25(OH)₂D₃ [1]. Hence, FGF23 lowers the plasma concentration of both, phosphate and 1,25(OH)₂D₃, thereby depending on αKlotho [1], an anti-aging protein [3]. Another target organ of FGF23 is the parathyroid gland where FGF23 regulates the production of parathyroid hormone, PTH, in a aKlotho-dependent manner [4, 5]. Apart from this, FGF23 also exerts paracrine and further endocrine effects without the contribution of αKlotho: It induces hypertrophy of the left ventricle [6], regulates neutrophil recruitment [7], or stimulates the secretion of pro-inflammatory cytokines from the liver [8]. The induction of left ventricular hypertrophy may, at least in part, be due to indirect effects of FGF23, too, e.g., elevation of blood pressure [9–11]. Similarly, the FGF23 effect on inflammation in hepatocytes may also involve indirect actions, e.g., the induction of interleukin-6 production in Kupffer cells that affect hepatocytes [12].

The dramatic phenotype of FGF23- or αKlotho-deficient mice is mostly characterized by sequelae of massive hyperphosphatemia which are far-reaching and affect most organs and tissues: The animals suffer from rapid aging with a markedly shortened life span and exhibit a wide range of aging associated diseases [13, 14]. A low phosphate or low vitamin D diet prevents such consequences [15]. Similar to FGF23- or αKlotho-deficient mice, patients with chronic kidney disease (CKD), which are typically characterized by high FGF23 levels and low αKlotho expression, suffer from the sequelae of hyperphosphatemia which drastically increase their cardiovascular mortality [16]. The FGF23 plasma concentration rises early in CKD and correlates well with disease activity, progression and outcome making FGF23 a valuable disease biomarker in CKD [17, 18]. In addition, high FGF23 plasma levels are found in cardiovascular, metabolic, and inflammatory disorders, suggesting that FGF23 may be a general disease biomarker [19].

The regulation of FGF23 production is only partly understood: Important regulators include 1,25(OH)₂D₃ [20] and PTH [21]. Both strongly affect the plasma phosphate concentration as does FGF23. Therefore, and since FGF23 also regulates 1,25(OH)₂D₃ and PTH, all three hormones are part of a complex feedback loop regulating plasma phosphate. Further regulators are alimentary phosphorus intake [22], inflammation [23], and inflammatory cytokines including tumor necrosis factor- α [24], interleukin-6 [25], the iron status [26], or erythropoietin [27]. Mechanistically, the transcription of the FGF23 gene is dependent on transcription factor Nurr1 [21], NFkB upregulating Ca²⁺ release-activated Ca²⁺ channel (CRAC) Orail [28, 29], as well as insulin-dependent phosphoinositide 3-kinase (PI3K)/protein kinase B (PKB)/Akt/forkhead box O1 (FOXO1) signaling [30]. Orai1-mediated store-operated Ca²⁺ entry (SOCE) is a fundamental driver of FGF23 gene expression [28, 31]. The aforementioned signaling has, thus far, only been verified for FGF23 production in bone, not for other tissues.

Cortisol is the main glucocorticoid in humans produced by the adrenal gland [32]. Glucocorticoids are stress hormones with a long list of metabolic (particularly affecting glucose and lipid metabolism) and immunological effects [32]. Synthetic glucocorticoids such as prednisolone or dexamethasone are widely used in the therapy of inflammatory diseases owing to their strong and multiple anti-inflammatory properties [33]: Glucocorticoids inhibit the production of proinflammatory prostaglandins through annexins [34], induce apoptosis of T cells [35], or suppress the transcriptional activity of pro-inflammatory transcription factor complex NFκB [36]. As steroid hormones, glucocorticoids bind to an intracellular receptor, the glucocorticoid receptor (GR), which translocates to the nucleus, binds DNA, and induces or represses the transcription of target genes [37].

Glucocorticoid therapies are initiated for a plethora of frequent inflammatory and autoimmune diseases including glomerulonephritis, rheumatoid arthritis, asthma, or inflammatory bowel disease to name a few. They are limited due to a variety of adverse effects. These effects are dependent on the dose and duration of therapy, but are often observed and include steroid diabetes, hypertension, thin skin, muscle atrophy, obesity, and loss of bone mass due to direct and indirect effects [38]: Glucocorticoids affect osteoblast and osteoclast activity [39] but also impact on phosphate metabolism by regulating renal phosphate transporter NaP_iIIa [40].

Recently, a study suggested an inhibitory effect of glucocorticoids on FGF23 [41], whereas another study suggested a stimulatory effect [42]. Both studies focused on long-term effects of glucocorticoids which may include secondary cellular responses owing to the broad spectrum of glucocorticoid effects. We therefore sought to study the cellular effects of an acute glucocorticoid treatment, both in vitro and in vivo. To this end, we analyzed the production of FGF23 in UMR106 osteoblast-like cells and in MC3T3-E1 cells as well as in mice treated with glucocorticoids.

Materials and methods

Cell culture

Experiments were performed in UMR106 rat osteoblastic osteosarcoma cells (CRL-1661; ATCC, Manassas, VA, USA). UMR106 cells were cultured in Dulbecco's Modified Eagle Medium (DMEM) high glucose (Gibco, Life Technologies, Darmstadt, Germany) supplemented with 10% fetal bovine serum (FBS) (Gibco, Life Technologies), 100 U/ml penicillin, and 100 μ g/ml streptomycin (Gibco, Life Technologies). Twenty-four hours after seeding, the cells were treated with the indicated concentrations of dexamethasone, prednisolone



(both from Sigma-Aldrich, Schnelldorf, Germany; stock dissolved in dimethyl sulfoxide (DMSO)), or glucocorticoid receptor antagonist RU-486 (Tocris Bioscience, Bristol, UK; stock dissolved in ethanol) for 24 or 48 h, as indicated. Control cells were incubated with the appropriate volume of DMSO or ethanol (both from Sigma-Aldrich), respectively. Untreated UMR106 cells do not express appreciable amounts of Fgf23 mRNA as reflected by the cycle threshold (Ct) of Fgf23 in semi-quantitative RT-PCR analysis. Calcitriol potently upregulates Fgf23 expression in UMR106 cells. In line with this, the arithmetic means \pm SEM (n = 3) of the Ct values of UMR106 cells were 34.0 ± 0.75 (untreated) and 28.0 ± 0.06 (treated with 10 nM calcitriol for 24 h). Therefore, all cell culture treatments were carried out in the presence of 10 nM calcitriol (Tocris Bioscience) [43].

MC3T3-E1 Subclone 4 mouse pre-osteoblast cells (CRL-2593; ATCC) were cultured in alpha-Minimum Essential Medium (α -MEM) containing nucleosides and 2 mM L-glutamine (Gibco, Life Technologies) and supplemented with 10% FBS, 100 U/ml penicillin, and 100 µg/ml streptomycin. The cells were used from passages 20 to 23. The cells were seeded on rat tail type I collagen-coated 12-well plates (80,000 cells per well) for 24 h. Differentiation was induced by culturing the cells in osteogenic medium containing 50 µg/ml ascorbic acid (Sigma-Aldrich), and 4 mM β -glycerophosphate (AppliChem, Darmstadt, Germany). Cells were differentiated for 6 or 13 days and then treated for 24 h with dexamethasone (30 nM or 100 nM) or vehicle in the presence of calcitriol (10 nM) throughout the experiment or 24 h before harvesting the cells, as indicated.

Analysis of FGF23 protein in the cell culture medium

The cell culture supernatants of UMR106 cells treated with 30 nM or 100 nM dexamethasone or vehicle in the presence of 10 nM calcitriol for 48 h were collected and stored at -70 °C. The medium was concentrated utilizing Vivaspin 6 concentrators (Sartoritus, Göttingen, Germany). C-terminal FGF23 and intact FGF23 were determined by commercial ELISA Kits (Mouse/Rat FGF-23 (C-Term), Mouse/Rat FGF-23 (Intact); both from Immutopics, San Clemente, CA, USA) according to the manufacturer's instructions.

Animal experiments and tissue collection

All animal experiments were conducted in accordance with the federal law for the welfare of animals and were approved by the respective state government. The mice had free access to tap water and standard rodent chow mainly based on wheat, soybeans, barley, oat hulls, and sugar beet pulp with a rough composition owing to the nature of the plant products (V1534: app. 0.7% phosphorus, 1.0% calcium; Ssniff, Soest, Germany;

more details can be found at https://www.ssniff.de/) ad libitum.

Adult wild-type mice (C57BL/6; Charles River, Sulzfeld, Germany) of both sexes were injected intraperitoneally with a single dose of dexamethasone (20 mg/kg body weight; Fortecortin®, Merck, Darmstadt, Germany; formulation: 4.37 mg/ml dexamethasone dihydrogen phosphate disodium, sodium edetate, creatinine, sodium citrate, 1 mol/l sodium hydroxide and water for injections) or prednisolone (20 mg/kg body weight; PredniGalen®, GALENpharma, Kiel, Germany; formulation: 10 mg/ml prednisolone acetate, macrogol 4000, polysorbate 80, disodium phosphate dodecahydrate, sodium dihydrogen phosphate dihydrate, water for injections and 15 mg/ml benzyl alcohol). As control, 0.9% saline was chosen since an exact formulation of the drugs was not available. After 12 h, blood was taken by retro-orbital puncture under isoflurane anesthesia. The animals were sacrificed and kidneys were immediately isolated and snap-frozen in liquid nitrogen. Bones were also removed and prepared by thoroughly scrubbing the femur and tibia to remove other tissues. Next, both epiphyses were cut, and the bone marrow was flushed out with 0.9% saline.

Serum biochemistry

Photometric methods were used for the analysis of the serum phosphate (Analyticon Biotechnologies, Lichtenfels, Germany) and Ca²⁺ (Fuji DRI-CHEM NX500i, Fujifilm, Düsseldorf, Germany) concentration. For the serum concentration of 1,25(OH)₂D₃ (IDS, Boldon, UK), C-terminal FGF23, intact FGF23, and intact PTH (all three from Immutopics) ELISA kits were employed.

Renal parameters

To assess renal phosphate and Ca²⁺ excretion, the mice were placed individually in siliconized metabolic cages (Tecniplast, Hohenpeissenberg, Germany). Standard chow pellets tend to crumble when eaten by mice. Hence, they cannot be used in metabolic cages since they would contaminate the collected urine. This is why we used a control diet with pellets with different consistency and precisely defined ingredients (C1000: 0.8% phosphorus, 0.9% calcium; Altromin, Lage, Germany; more detailed information can be found at https:// altromin.de/). Following a 3-day habituation period, mice were treated with 20 mg/kg body weight dexamethasone or vehicle. After 12 h, a small amount of blood was withdrawn. Additionally, 24-h urine was collected under water-saturated oil starting from the time of injection. The urinary phosphate and Ca²⁺ concentrations were measured by means of commercial assays (both from Analyticon Biotechnologies, Lichtenfels, Germany). Prior to the determination of Ca²⁺, 24-h urine was acidified with hydrochloric acid in order to



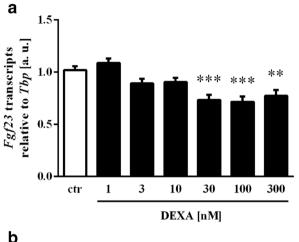
dissolve calcium salts. A creatinine detection kit (Jaffé method, Labor + Technik, Berlin, Germany) was employed for the measurement of the urinary creatinine concentration. Serum creatinine was determined by an enzymatic method (Chrystal Chem, Zaandam, Netherlands).

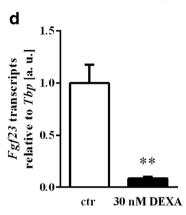
Semi-quantitative real-time PCR

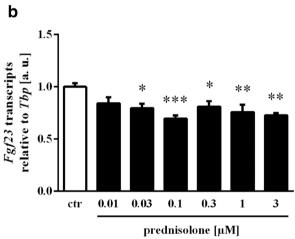
Total RNA was extracted from UMR106 cells, MC3T3-E1 cells, or mouse kidneys using peqGold Trifast™ reagent

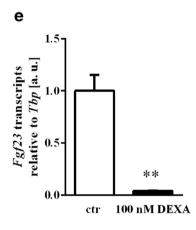
(Peqlab, Erlangen, Germany). Total RNA (1.2 µg) was reverse-transcribed with the GoScript™ Reverse Transcription System (Promega, Mannheim, Germany) and random primers (Promega). One µg of total RNA was used for first-strand cDNA from MC3T3-E1 cells with persistent calcitriol stimulation.

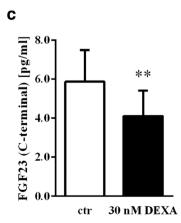
Total bone RNA was initially extracted with peqGold TrifastTM reagent followed by a DNase treatment and a purification step using the RNase-free DNase Set and the RNeasy Mini Kit (both from QIAGEN, Hilden, Germany). First-strand

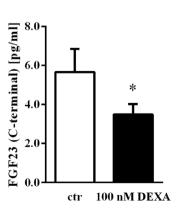


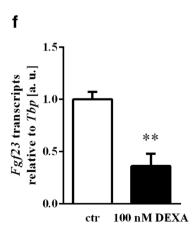














▼ Fig. 1 Effect of glucocorticoids on Fgf23 expression and protein synthesis in UMR106 and MC3T3-E1 cells. Arithmetic means \pm SEM (n = 6, arbitrary units (a. u.)) of relative Fgf23 mRNA expression normalized to Tbp transcript levels in UMR106 cells incubated without (white bars) or with (black bars) dexamethasone (DEXA, a) or prednisolone (b) at the indicated concentrations in the presence of calcitriol (10 nM) for 24 h. *p < 0.05, **p < 0.01, ***p < 0.001 indicate significant differences compared to UMR106 cells treated with 10 nM calcitriol only. (c) Arithmetic means ± SEM of C-terminal FGF23 protein concentration in the cell culture supernatant of UMR106 cells treated with vehicle (white bars) or with 30 nM (n = 8) or 100 nM (n = 6) dexamethasone (DEXA, black bars) in the presence of calcitriol (10 nM) for 48 h. *p < 0.05, **p < 0.01, indicate significant difference compared to UMR106 cells treated with 10 nM calcitriol only. Arithmetic means \pm SEM (n = 5, arbitrary units (a. u.)) of relative Fgf23 mRNA expression normalized to Tbp transcript levels in MC3T3-E1 cells at day 7 of osteoblast differentiation incubated with vehicle (white bars) or 30 nM (d) or 100 nM (e) dexamethasone (DEXA, black bars) in the presence of calcitriol (10 nM) for 24 h. **p < 0.01 indicates significant difference compared to control cells. (f) Arithmetic means \pm SEM (n = 4, arbitrary units (a. u.)) of relative Fgf23 mRNA expression normalized to Tbp transcript levels in MC3T3-E1 cells at day 14 in osteogenic differentiation medium containing calcitriol (10 nM) and incubated with 100 nM dexamethasone (DEXA, black bar) or vehicle (white bar) for 24 h. **p<0.01 indicates significant difference compared to vehicle-treated MC3T3-E1 cells. (a and b: one-way ANOVA followed by Dunnett's multiple comparisons test; **c**–**f**: paired *t* test)

cDNA was synthesized from 300 ng of total bone RNA using the GoScript™ Reverse Transcription System.

For the analysis of relative transcript levels, semi-quantitative real-time PCR (qRT-PCR) using GoTaq® qPCR Master Mix (Promega) was carried out. The reaction mix for the amplification of 2 μ l cDNA templates (1.5 μ l cDNA in the case of bone Fgf23 expression analysis) contained 10 μ l 2x GoTaq® qPCR Master Mix (Promega), primers, and sterile water up to 20 μ l. The qRT-PCR program was 95 °C for 2 min, 40 cycles of denaturation at 95 °C for 10 s, annealing at primer-specific temperature for 30 s, and extension at 72 °C for 25–30 s.

The following primers $(5' \rightarrow 3')$ orientation and annealing temperatures were used:

rat Dmp1 (62 °C):
CGCCCATGGCAAATAGTGAC;
CGTGCTGTCTTCACTGGACT;
rat Fgf23 (57 °C):
TAGAGCCTATTCAGACACTTC;
CATCAGGGCACTGTAGATAG;
rat Nfkbia (58 °C):
AGACTCGTTCCTGCACTTGG;
TCTCGGAGCTCAGGATCACA;
rat Phex (58 °C):
ATGGCTGGATAAGCAATAAC;
GCTTTTTCAATCGCTTTCTC;
rat Tbp (57 °C):
ACTCCTGCCACACCAGCC;
GGTCAAGTTTACAGCCAAGATTCA;

mouse Cyp24a1 (58 °C): CCCAAAGGAACAGTCTTAAC; GGTCTAAACTTGTCAGCATC; mouse Cyp27b1 (58 °C): AGTGTTGAGATTGTACCCTG: CGTATCTTGGGGAATTACATAG; mouse Dmp1 (58 °C): GAACAGTGAGTCATCAGAAG: AAAGGTATCATCTCCACTGTC; mouse Fgf23 (58 °C): TCGAAGGTTCCTTTGTATGGA: AGTGATGCTTCTGCGACAAGT; mouse Gapdh (58 °C): GGTGAAGGTCGGTGTGAACG: CTCGCTCCTGGAAGATGGTG; mouse K1 (56 °C): CCTTAAAAGCAATCAGACTGG; GAAAGCCATTGTCCTCTATC; mouse Nfkbia (57 °C): GGAGACTCGTTCCTGCACTTGG; CTCAGCAATTCCTGGCTGGT; mouse Phex (57 °C): TCATTGATACCAGACTCTACC; CAATGGTTTTCTTCCTCTCG; mouse Slc34a1 (58 °C): AATGCAACCATATCTTCGTG: GGAAAGTCTGTGTTGATGAC; mouse Tbp (60 °C): CCAGACCCCACAACTCTTCC; CAGTTGTCCGTGGCTCTCTT.

Calculated mRNA expression levels were normalized to the expression levels of *Tbp* (rat UMR106 cells and mouse MC3T3-E1 cells) and *Gapdh* (mouse tissue). Relative quantification of gene expression was performed using the $2^{-\Delta\Delta Ct}$ method.

Statistics

Data are shown as arithmetic means \pm SEM, and n represents the number of mice per group or the number of independent cell culture experiments, respectively. Normality was tested with Shapiro-Wilk test. Two groups were compared by paired t test or unpaired two-tailed t test (Welch's correction was applied for data with different standard deviations) or Mann-Whitney U test (for data not passing normality test). Statistical testing of more than two groups was performed by one-way ANOVA followed by Dunnett's or Tukey's multiple comparisons test (homoscedastic data) or Games-Howell test (heteroscedastic data) as indicated in the figure legends. Only differences with P < 0.05 were considered statistically significant.



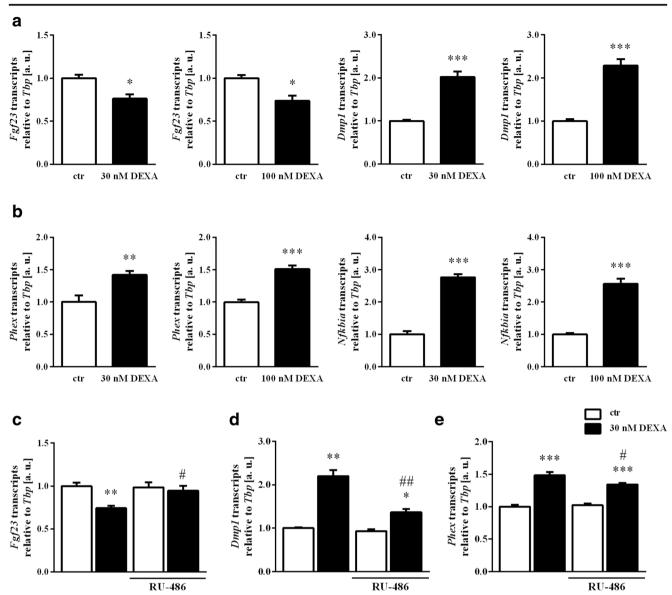


Fig. 2 Impact of dexamethasone and glucocorticoid antagonist RU-486 on regulators of Fg/23 expression. Arithmetic means \pm SEM (arbitrary units (a. u.)) of relative mRNA transcript levels of Fg/23, Dmp1 (a), Phex, and Nfkbia (b), all normalized to Tbp, in UMR106 cells treated with 30 nM (n=9) or 100 nM dexamethasone (n=7) (DEXA, black bars) or vehicle only (white bars) in the presence of 10 nM calcitriol for 24 h. *p < 0.05, **p < 0.01, ***p < 0.001 indicate statistically significant difference from vehicle-treated cells. Arithmetic means \pm SEM (n=6, arbitrary units (a. u.)) of relative mRNA transcript levels of Fg/23 (c), Dmp1

(d), and *Phex* (e), all normalized to *Tbp* in calcitriol-stimulated (10 nM for 24 h) UMR106 cells treated without (white bars) or with 30 nM dexamethasone (DEXA, black bars) in the absence or presence of the glucocorticoid receptor antagonist RU-486 (30 nM) for 24 h. *p < 0.05, **p < 0.01, ***p < 0.001 indicate statistically significant difference from vehicle-treated cells (first bar). #p < 0.05, ##p < 0.01 indicate statistically significant difference from the absence of RU-486 (2nd bar vs. 4th bar). (a and b: paired t test; c-e: one-way ANOVA followed by Tukey's multiple comparisons test (c and e) or Games-Howell test (d))

Results

We used UMR 106 osteoblast-like cells to study whether dexamethasone, a synthetic glucocorticoid with high potency that is widely used in clinical practice, influences FGF23 production. The cells were incubated with 1–300 nM dexamethasone for 24 h, and *Fgf23* transcripts were determined by qRT-PCR. The dose dependence of the dexamethasone effect on *Fgf23* gene expression is depicted in Fig. 1a. Similarly, the

expression of Fg/23 mRNA was reduced by the synthetic glucocorticoid prednisolone that is also widely used in the treatment of autoimmune diseases (Fig. 1b). In order to test whether dexamethasone also affects FGF23 secretion, we analyzed the FGF23 protein concentration into the cell culture supernatant. As a result, the concentration of C-terminal FGF23 in the cell culture medium of cells incubated with 30 nM (Fig. 1c, left panel) or 100 nM (Fig. 1c, right panel) dexamethasone for 48 h was significantly lower than in the



medium of vehicle-treated cells. We also tried to determine intact FGF23, however, for n=7 out of n=14 experiments, the concentration of intact FGF23 was below the detection limit. For the remaining n=7 measurements, the concentration of intact FGF23 was 3.49 ± 0.51 pg/ml in the supernatant of UMR106 cells incubated for 48 h without and 2.52 ± 0.21 pg/ml in the supernatant of cells incubated with 30 nM dexamethasone (p=0.12).

MC3T3-E1 cells were used in further experiments to verify the glucocorticoid effect on FGF23. Also in these cells dexamethasone potently lowered *Fgf23* gene expression (Figs. 1d–f). The effect could be observed in cells differentiated for 6 days (Figs. 1d, e) or 13 days (Fig. 1f) and then treated with 30 nM (Fig. 1d) or 100 nM (Figs. 1e, f) dexamethasone for 24 h.

A new series of experiments addressed the impact of dexamethasone on regulators of Fgf23 expression. Again, 30 nM or 100 nM dexamethasone suppressed Fgf23 gene expression in UMR106 cells (Fig. 2a). Importantly, Dmp1 (Fig. 2a, right panels) and Phex (Fig. 2b, left panels) mRNA transcript levels were significantly higher in dexamethasone-treated (30 nM or 100 nM) UMR106 cells than in control cells. Glucocorticoids have strong anti-inflammatory properties owing to their inhibitory effect on NF κ B. Hence, we studied the expression of I κ B α (encoded by Nfkbia), a cytoplasmic protein that inhibits nuclear translocation of NF κ B. As shown in Fig. 2b, right panels, Nfkbia expression was significantly enhanced following treatment with 30 or 100 nM dexamethasone for 24 h.

We also investigated the contribution of the glucocorticoid receptor to the dexamethasone effect on FGF23 by performing experiments with glucocorticoid receptor antagonist RU-486. As a result, 30 nM RU-486 significantly attenuated the dexamethasone (30 nM) effect on *Fgf23* transcripts (Fig. 2c), *Dmp1* transcripts (Fig. 2d) and *Phex* transcripts (Fig. 2e). Thus, the dexamethasone effect is—at least in part—mediated by the glucocorticoid receptor.

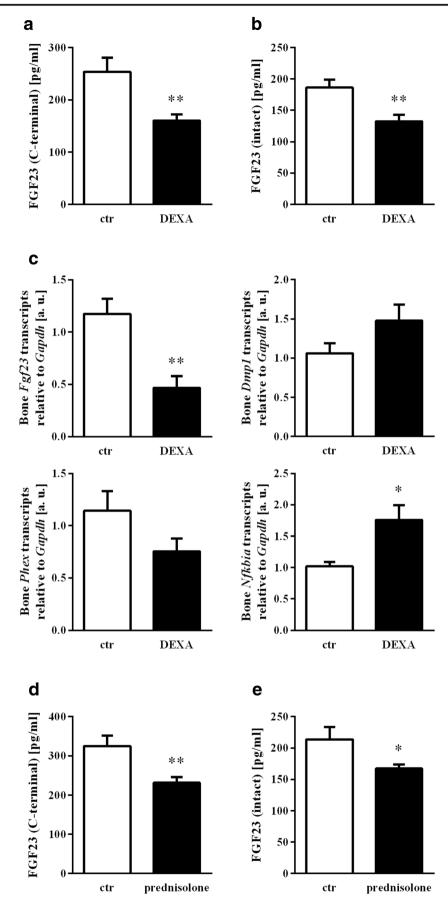
Since the cells had to be treated with $1,25(OH)_2D_3$ to upregulate Fgf23 gene expression, the dexamethasone effect could, at least in theory, have been due to an interaction of dexamethasone with $1,25(OH)_2D_3$ signaling. Therefore, we tested whether the dexamethasone effect on upstream regulators of Fgf23 gene expression was different between cells treated with $1,25(OH)_2D_3$ or left untreated. As demonstrated in suppl. Fig. 1, the effect of 30 nM dexamethasone on Dmp1 expression (suppl. Fig. 1a), Phex expression (suppl. Fig. 1b), or Nfkbia expression (suppl. Fig. 1c) was comparable between cells treated with $1,25(OH)_2D_3$ or left untreated.

Our in vitro results suggest an inhibitory effect of glucocorticoids on FGF23 synthesis. We therefore sought to investigate whether the regulation of FGF23 by glucocorticoids has in vivo relevance given their frequent use for medical purposes. Mice were sham-treated or treated with a single dose of dexamethasone. Twelve hours later, blood was collected, and the serum concentration of C-terminal and intact FGF23 was determined. As demonstrated in Fig. 3, dexamethasone is given only once significantly and markedly reduced the serum level of both, C-terminal (Fig. 3a) and intact (Fig. 3b) FGF23. To check whether the lower FGF23 serum concentration is paralleled by reduced gene expression, we determined Fgf23 transcript levels in the bone of dexamethasone-treated animals, the main site of FGF23 production. As a result, dexamethasone downregulated Fgf23 gene expression in the bone, pointing to a highly relevant inhibitory effect of glucocorticoids on the production of this hormone even after a single dose (Fig. 3c, first panel). The effect was paralleled by upregulation of Nfkbia expression (Fig. 3c, last panel), whereas Dmp1 and Phex expressions were not significantly modified (Figs. 3c, 2nd and 3rd panel). Moreover, a single injection of prednisolone similarly lowered the C-terminal (Fig. 3d) and intact FGF23 serum level (Fig. 3e).

One of the main actions of FGF23 is the suppression of the formation of 1,25(OH)₂D₃ by downregulation of *CYP27B1*, the key renal enzyme for the hydroxylation of 25(OH)D₃. Dexamethasone-dependent inhibition of FGF23 formation is therefore likely to also affect the serum concentration of 1,25(OH)₂D₃. As expected, the single dose of dexamethasone resulted in a surge of the serum concentration of 1,25(OH)₂D₃ (Fig. 4a). PTH was not affected by dexamethasone treatment (Fig. 4b). Enhanced 1,25(OH)₂D₃ formation upon short-term glucocorticoid administration was paralleled by increased renal expression of *Cyp27b1* (Fig. 4c). The expression of *Cyp24a1*, the renal enzyme catalyzing the inactivation of 1,25(OH)₂D₃ was, however, not significantly affected by dexamethasone (Fig. 4d).

FGF23 directly induces renal phosphate excretion by targeting NaP_iIIa (encoded by Slc34a1), the main renal Na⁺dependent phosphate transporter. To study the impact of a single dose of dexamethasone on renal phosphate handling, metabolic cage studies were conducted. Dexamethasone significantly increased creatinine clearance (Fig. 5a), a finding in line with the literature [44]. As demonstrated in Fig. 5b and d, dexamethasone treatment resulted in an increase in renal phosphate excretion, which was paralleled by a lower serum phosphate level, within 12 h. This finding is also in line with an earlier report demonstrating a direct effect of dexamethasone on NaP_iIIa [40]. The serum Ca²⁺ concentration (Fig. 5c) was not affected by the glucocorticoid within 12 h, whereas dexamethasone treatment tended to increase urinary Ca²⁺ excretion; however, this effect did not reach statistical significance (Fig. 5e). In line with phosphaturia upon dexamethasone treatment, the renal expression of Slc34a1 was reduced in animals receiving the glucocorticoid (Fig. 5f), as revealed by qRT-PCR. FGF23-mediated regulation of urinary phosphate excretion is dependent on its co-receptor αKlotho. As shown in Fig. 5g, treatment with dexamethasone did not significantly affect Kl mRNA expression in the kidney.







◄ Fig. 3 The glucocorticoid effect on serum FGF23 and bone Fg/23 expression. Arithmetic means \pm SEM of the serum concentration of C-terminal FGF23 (**a**, n = 11) and intact FGF23 (**b**, n = 13) as well as relative osseous mRNA expression (**c**) of Fg/23 (n = 7, arbitrary units (a. u.), Dmp1 (n = 9), Phex (n = 9), and Nfkbia (n = 9) normalized to housekeeping gene Gapdh in control mice (ctr, white bars) and dexamethasonetreated mice (20 mg/kg body weight, DEXA, black bars) 12 h after injection. Arithmetic means \pm SEM of the serum concentration of C-terminal (**d**, n = 10) and intact FGF23 (**e**, n = 10) in mice 12 h after treatment with vehicle (ctr, white bars) or prednisolone (20 mg/kg body weight; black bars). *p < 0.05, **p < 0.01 indicate significant difference compared to vehicle-treated mice. (**a**, **c** (right lower panel) and **e**: unpaired t test with Welch's correction; **b** and **c** (right upper panel): unpaired t test; **c** (left upper panel and left lower panel) and **d**: Mann-Whitney U test)

Discussion

Our study discloses glucocorticoids as powerful suppressors of the production of FGF23. We demonstrated in vitro and in vivo that the widely used glucocorticoids dexamethasone and prednisolone downregulate *Fgf23* gene expression and lower the serum concentration of FGF23.

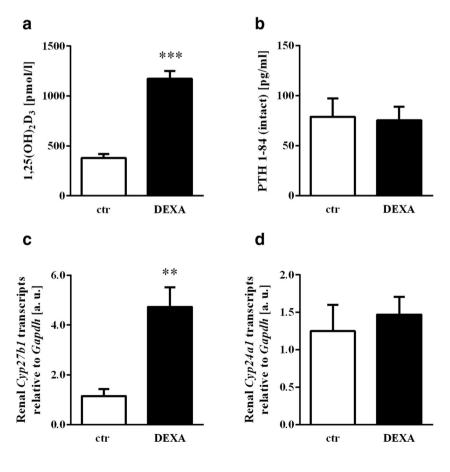
Dexamethasone is a synthetic glucocorticoid with high potency that is widely used in the treatment of inflammatory conditions, but also in cerebral edema and to prevent chemotherapy-induced nausea and vomiting [38]. In contrast

to some other glucocorticoids, it has no considerable mineralocorticoid activity [45]. Prednisolone is widely used in autoimmune diseases including rheumatoid arthritis, polymyalgia rheumatica, or inflammatory bowel disease. Mineralocorticoid receptor agonists including aldosterone induce the synthesis of FGF23, as seen in primary and secondary hyperaldosteronism [46]. In contrast, our study reveals the GR as a negative regulator of FGF23.

The regulation of renal Na⁺-dependent phosphate transport by dexamethasone has been described before [40]. In their study, an inhibitory effect on the membrane expression of NaP_i paralleled by phosphaturia was found after 4 days of dexamethasone treatment [40]. We also observed phosphaturia in response to acute dexamethasone treatment. It is therefore possible that, apart from direct suppression of FGF23 as suggested by our cell culture experiments, dexamethasone-induced phosphaturia also contributes to the suppression of FGF23 production.

We observed a strong and significant reduction of the FGF23 serum concentration as early as 12 h after a single dose of dexamethasone or prednisolone. Notably, both, C-terminal and intact FGF23 were reduced, suggesting that glucocorticoids indeed suppress Fgf23 gene expression rather than merely affecting post-translational processing. The latter would have been expected if C-terminal FGF23 had only been

Fig. 4 Impact of dexamethasone on 1,25(OH)₂D₃ formation, metabolism and PTH serum level. Arithmetic means \pm SEM (n = 6) of the 1,25(OH)₂D₃ (a) and intact PTH (b) serum concentration in control mice (ctr, white bars) and mice treated with 20 mg/kg body weight dexamethasone (DEXA, black bars) 12 h after the injection. Arithmetic means \pm SEM (n = 6, arbitrary units (a. u.)) of relative mRNA transcript levels of Cyp27b1 (c) and Cyp24a1 (d) normalized to Gapdh in kidneys from mice 12 h after the injection of vehicle only (ctr, white bars) or 20 mg/kg body weight dexamethasone (DEXA, black bars). **p < 0.01, ***p < 0.001 indicate statistically significant differences from control-treated mice. (a, b and d: unpaired t test; c: unpaired t test with Welch's correction)





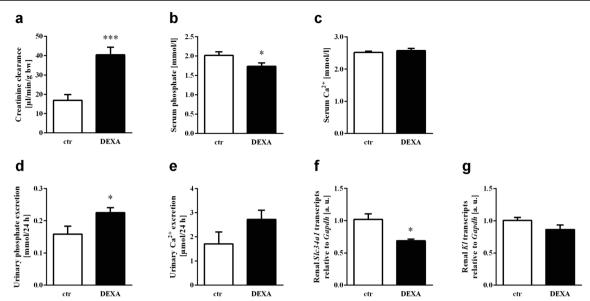


Fig. 5 Effect of dexamethasone on phosphate and Ca^{2+} metabolism. Arithmetic means \pm SEM of creatinine clearance (**a**, n = 6), the serum concentration of phosphate (**b**, n = 17; 12 h after injection) and Ca^{2+} (**c**, n = 12; 12 h after injection), as well as phosphate (**d**, n = 6) and Ca^{2+} (**e**, n = 6) excretion in mice in 24 h urine after injection of vehicle (ctr, white bars) or 20 mg/kg body weight dexamethasone (DEXA, black bars). Arithmetic means \pm SEM (n = 6, arbitrary units (a. u.)) of relative

mRNA transcript levels of Slc34a1 (f) and Kl (g) normalized to Gapdh in kidneys from mice 12 h after the injection of vehicle (ctr, white bars) or 20 mg/kg body weight dexamethasone (DEXA, black bars). *p<0.05, ****p<0.001 indicate statistically significant differences from control mice. (a, b, d, e and g: unpaired t test; c: Mann-Whitney U test; f: unpaired t test with Welch's correction)

affected by the glucocorticoid. Short-term treatments even with very high doses of glucocorticoids are frequent in daily clinical practice. Therefore, our results may be of high relevance and glucocorticoid-dependent suppression of FGF23 a frequent phenomenon.

Due to the inhibitory effect of FGF23 on the formation of 1,25(OH)₂D₃, dexamethasone-dependent suppression of FGF23 could be expected to elevate the 1,25(OH)₂D₃ serum concentration. In line with this, we saw a surge in the 1,25(OH)₂D₃ serum level 12 h after the single dose of dexamethasone. In contrast, it is well-established textbook knowledge that particularly a long-term glucocorticoid therapy clinically indicated in various chronic autoimmune diseases requires the patient to take additional vitamin D to prevent bone loss [47], and steroid therapy is indeed associated with low 25(OH)D₃ levels [48]. A glucocorticoid-induced increase in active vitamin D may therefore come as a surprise. It is, however, the logical consequence of the suppression of FGF23 formation by glucocorticoids. In addition, it must be kept in mind that our study only reflects the acute effects of a single dose of glucocorticoids within 12 h. It is well conceivable that the immediate increase in the serum 1,25(OH)₂D₃ concentration due to a glucocorticoid-dependent downregulation of FGF23 is overridden by further direct and indirect effects of glucocorticoids on the bone and vitamin D metabolism in the long run resulting in the well-documented bone loss and vitamin D deficiency.

According to our results, neither Ca²⁺ metabolism nor PTH secretion was acutely affected by glucocorticoid administration.

In contrast, a long lasting steroid therapy favors renal Ca²⁺ loss and secondary hyperparathyroidism [49]. It appears to be likely that the acute effect of a glucocorticoid is therefore the downregulation of FGF23, whereas other effects on bone and Ca²⁺ and phosphate metabolism come later. Similar to the cell culture experiments, Nfkbia was upregulated in vivo, whereas Dmp1 and Phex were not significantly affected by dexamethasone treatment. In our in vivo study, glucocorticoid effects on blood parameters were determined after 12 h whereas urine was collected for 24 h in order to cover an entire day-night period. Hence, this may be a limitation of our study since glucocorticoids may affect circadian rhythm as well as urinary Ca²⁺ and phosphate excretion may be influenced by circadian rhythm. For technical reasons, pellets with different consistency had to be used in the metabolic cages. This diet had a slightly different composition compared to standard chow. However, since all ingredients were present in adequate amounts in both diets, this difference is unlikely to significantly affect the results.

A short-term glucocorticoid treatment is particularly relevant when strong anti-inflammatory (e.g., anaphylactic shock) or anti-emetic effects (e.g., chemotherapy) are needed. According to the results of this study, such a therapy could be paralleled by a decrease of FGF23 and a surge in 1,25(OH)₂D₃. The exact relevance of these anticipated changes in blood parameters needs to be defined in future clinical studies. However, it is intriguing to speculate that a decrease in the FGF23 serum level may indeed be beneficial as FGF23 has been shown to go up in various cardiovascular and renal



diseases and in particular in those associated with inflammation.

Our cell culture studies disclosed that dexamethasone and prednisolone suppress Fgf23 gene expression in two bone cell lines. Moreover, this effect was paralleled by a lower C-terminal FGF23 protein concentration in the cell culture supernatant, i.e., secretion of FGF23 was also reduced by dexamethasone. Also the concentration of intact FGF23 tended to be lower in the supernatant from dexamethasone-treated cells although the values were below the detection limit for 50% of the samples. In both cell lines, Fgf23 gene expression was induced by 1,25(OH)₂D₃ as described by others before [43]. The glucocorticoid effect on FGF23 could have been, at least in part, due to the crosstalk of 1,25(OH)₂D₃ and glucocorticoid signaling. However, the dexamethasone effect on upstream regulators of Fgf23 gene expression Dmp1, Phex, or Nfkbia was not altered by 1,25(OH)₂D₃. Therefore, it appears to be likely that glucocorticoids are not merely effective through suppression of 1,25(OH)₂D₃ signaling. However, these findings do not completely rule out that dexamethasone also downregulates 1,25(OH)₂D₃ signaling, at least to some extent. It must be kept in mind that the in vivo experiments clearly indicate that glucocorticoids indeed suppress FGF23, a finding suggesting that also the cell culture experiments reveal a relevant effect of glucocorticoids on FGF23.

Many of the multiple cellular effects of glucocorticoids can directly or indirectly be referred to their anti-inflammatory properties that are also the reason for their therapeutic potential in various diseases. Notably, inflammation in the form of transcription factor complex NFkB is a major driver of FGF23 production [28]. In an attempt to further characterize the molecular mechanism of the dexamethasone effect on FGF23 we found that dexamethasone induces gene expression of $I\kappa B\alpha$, an inhibitor of NFkB that mediates anti-inflammatory effects of glucocorticoids [36]. Moreover, our experiments revealed that gene expression of FGF23 inhibitors dentin matrix acidic phosphoprotein 1 (Dmp1) and phosphate regulating endopeptidase homolog X-linked (Phex) [50] are also upregulated by dexamethasone [51] as has been shown by others before [52, 53]. Hence, the glucocorticoiddependent inhibition of FGF23 is likely to be, at least in part, mediated by the glucocorticoid effects on $I\kappa B\alpha$, Dmp1, and Phex. Using glucocorticoid receptor antagonist RU-486, we could further demonstrate that the dexamethasone effects were, at least in part, mediated by the glucocorticoid receptor.

Taken together, our study demonstrates that glucocorticoids acutely downregulate the production of FGF23.

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Authors' contribution MFe, GIS, and MFö designed research; MFe, JS, HS, DK-T, and MFö analyzed data; MFe, FE, JS, and PG performed

research; and MFö and MFe wrote the paper. All authors revised and approved the final version of the manuscript.

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Data availability The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

Compliance with ethical standards

Conflict of interests All authors state they have no conflicts of interest.

Ethics approval The animal experiments were approved by the state governments of Saxony-Anhalt (Landesverwaltungsamt Halle/Saale) and Baden-Württemberg (Regierungspräsidium Stuttgart).

Consent for publication Not applicable.

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