EXPERIMENTAL STUDY

PPARγ inhibits GH synthesis and secretion and increases apoptosis of pituitary GH-secreting adenomas

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Abstract

Objective: The objective of the study was to evaluate the expression and functional activity of Peroxisome proliferator-activated receptor (PPAR) γ in pituitary adenomas from 14 consecutive acromegalic patients and to establish its role in apoptosis.

Subjects and methods: Fourteen consecutive acromegalic patients were enrolled in the study. Wistar-Furth rats were used for in vivo studies. Expression of PPARγ was evaluated by RT-PCR and Western blot. Apoptosis and cell cycle were assessed by FACS analysis. The effects of PPARγ ligands on transcriptional regulation of GH gene were evaluated by RT-PCR and electromobility shift assay.

Results: PPARγ was expressed in all human GH-secreting adenoma (GH-oma), in normal pituitary tissue samples (39±24% and 78±5% of immunostained nuclei respectively; P, 0.0002; ANOVA), and in rat GH-secreting (GH3) cells. A PPRE-containing reporter plasmid transfected into GH3 cells was activated by ciglitazone or rosiglitazone (TZDs), indicating that PPARγ was functionally active. Treatment of GH3 cells with TZDs increased apoptosis in a dose-dependent manner (P = 0.0003) and arrested cell proliferation, reducing the number of cells in the S-phase (P < 0.0001 vs untreated cells). TZDs increased the expression of TRAIL, leaving unaffected that of p53 and Bax. TZDs reduced GH concentrations in the culture media from 43.7 ± 5.4 ng/ml to 2.1 ± 0.3 ng/ml (P < 0.0001) and in cell extracts (P < 0.004). PPARγ–RXRa heterodimers bound to GH promoter, inhibiting its activity and reducing GH mRNA levels (1.8 × 10^6 vs 5.7 × 10^6 transcripts respectively vs untreated cells; P < 0.002). Subcutaneous GH-oma developed in rats injected with GH3 cells; tumor growth increased in placebo-treated rats and to a lesser extent in TZDs-treated animals (24.1 ± 2.0 g, and 14.8 ± 4.2 g respectively, P < 0.03). Serum GH concentrations were lower in TZDs-treated rats than in controls (871 ± 67 ng/ml vs 1.309 ± 238 ng/ml; P < 0.05).

Conclusions: The results of this study indicate that PPARγ controls GH transcription and secretion as well as apoptosis and growth of GH-oma; thus, TZDs have the potential of a useful tool in the complex therapeutic management of acromegalic patients.

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Introduction

Acromegaly is a syndrome characterized by excessive growth hormone (GH) secretion from pituitary adenomas, which in turns leads to overexpression of insulin-like growth factor (IGF)-1, the main effector of GH action (1, 2). The treatment of acromegaly is a therapeutic challenge (3). In spite of combination of surgery, external X-ray therapy and medical therapy, many patients continue to have uncontrolled GH secretion and active disease (4–15).

Peroxisome proliferator-activated receptor (PPAR) γ is a ligand-dependent transcription factor belonging to the nuclear receptor superfamily (16); it is highly expressed in adipose tissue, where it plays a key role in the regulation of adipocyte differentiation and fat metabolism (17, 18), and in colonic mucosa, where it exerts important functions in the differentiation process (19, 20). It is a functional receptor for the thiazolidinedione (TZDs) class of antidiabetic drugs and may function as a tumor-suppressor gene (21, 22); its activation has been reported to induce differentiation of liposarcoma (23), prostate cancer (24) or several transformed cells (19, 25).

PPARγ activation induced apoptosis in human and rat glioma with a transient upregulation of Bax and...
Bad proteins levels (26); in human papillary thyroid carcinoma cells, in which it enhanced expression of c-myc (27); and in the colonic cancer HT-29 cell line, where it downregulated c-myc and upregulated c-jun and gadd153 (28).

Recently, Heaney et al. (29, 30) reported that expression of PPARγ in normal pituitary gland seemed to be restricted to ACTH-secreting cells; nevertheless, this receptor was expressed in almost all pituitary adenomas.

However, PPARγ expression was studied in only three pure pituitary GH-secreting tumors (30). Activation of PPARγ induced apoptosis in various pituitary tumor cell lines (29, 30) and reduced the levels of ACTH, prolactin (PRL) and GH in the culture media. Nude mice injected with the pituitary GH-secreting GH3 cells harbored subcutaneous tumors, the development of which was partially prevented or retarded by the simultaneous treatment with TZDs (30). The increased TZDs-dependent apoptosis of rat pituitary cell lines was associated with the reduced expression of the anti-apoptotic protein Bcl2 and increased levels of proapoptotic Bax or p53 expression (29). However, spontaneous GH-secreting adenomas developed before a chemopreventive treatment could be initiated. In other words, therapies can be started only after acromegaly has become clinically manifest. Thus, it is more appropriate to evaluate the effects of TZDs on a pituitary tumor after its development.

TZDs treatment reduced cell viability and reduced the expression of pro-opiomelanocortin mRNA of ACTH-secreting cells. However, it is not known whether reduction of GH levels in the culture media of GH3 cells or in whole animals carrying subcutaneous GHomas was due to the antiapoptotic effects of TZDs or to decreased transcription of the GH gene.

Taking into account the reported expression of PPARγ in the pituitary and the anti proliferative effects of PPARγ ligands, we performed a series of experiments to evaluate the effect of this nuclear receptor on cell viability and GH secretion of pituitary GH-secreting adenomas. The aim of the study was to examine the expression of PPARγ in the pituitary adenoma from 14 consecutive patients with acromegaly and in normal pituitary tissue samples, to determine the effects of PPARγ activation in promoting apoptosis, and reducing tumor growth and GH secretion of adenoma either in vitro or in vivo, and to study the PPARγ-dependent transcriptional regulation of human GH gene.

Materials and methods

**Pituitary GH-secreting adenoma and normal pituitary tissue**

Pituitary GH-secreting adenomas samples were obtained from 14 consecutive acromegalic patients submitted to trans-sphenoidal adenomectomy. All acromegalic patients included in the present study had a pure GH-secreting adenoma and received a 6-month trial with somatostatin analogs before surgery. Normal pituitary tissue samples comprised the tissue outside pituitary adenomas obtained during neurosurgical excision of nine pituitary adenomas (five GH-secreting adenomas of the present series and four nonfunctioning adenomas). Pituitary tissue surrounding pituitary adenomas was considered normal at gross analysis by the neurosurgeon and the pathologist: in addition, all pituitary tissue samples surrounding pituitary adenomas were positively immunostained with anti-ACTH, anti-TSH (thyroid-stimulating hormone), anti-PRL, anti-LH and anti-FSH (follicle-stimulating hormone) antibody (data not shown). Samples of normal adult human skin (n = 5) were obtained during plastic surgery. All patients gave their informed consent, and the study was approved by the institutional review committee. Tissue samples were either immediately stored in liquid nitrogen for subsequent analysis or embedded in paraffin. Sections of pituitary tissue samples were immunostained with an anti-human GH antibody or an anti-PPARγ antibody, as previously described (31). Negative controls were obtained by omitting the anti-PPARγ or anti-GH-specific primary antibodies, which were replaced by rabbit non-immune serum.

**Cell culture and cell preparation**

GH3 and NIH3T3 cell lines were obtained from Data Bank ISDT (Genua, Italy) and cultured in F10 (GH3) or DMEM (NIH3T3), 10% FCS, 2 mM glutamine and penicillin-streptomycin solution, in a humidified 5% CO2 atmosphere at 37°C. Under starvation conditions, cells were extensively washed in PBS, and then incubated for 12 h in serum-free medium.

The effect of PPARγ activation on apoptosis and cell cycle was evaluated either in starved cells or in cells cultured in complete medium in the absence or presence of 1–50 μM ciglitazone (cig) (Alexis Biochemicals, San Diego, CA, USA) or 1–50 μM rosiglitazone (ros) (kindly provided by Glaxo-Kline-Beecham, Milan, Italy). Assessment of apoptosis by Annexin V or DNA fragmentation assay was done after up to 72 h of culture (see below).

**Annexin V binding assay**

Cells were washed in PBS, trypsinized, centrifuged and washed with PBS before incubating with FITC-labeled monoclonal annexin V antibody and propidium iodide for 15 min at RT according to the supplier’s instructions (Sigma-Aldrich, Milan, Italy) with few modifications. Cells were analyzed on a FACS (FACS Calibur) Becton Dickinson (Franklin Lakes, NJ, USA) apparatus. For each sample, 500 000 cells were analyzed. Data were analyzed using Cell Quest software (Becton Dickinson, Franklin Lakes, NJ, USA) and
represent the mean ± S.D. of five different experiments, each performed in triplicate.

**DNA fragmentation assay**
Quantitative determination of fragmented DNA in cytoplasm was done by Cell Death Detection ELISA Plus (Roche Diagnostics, Indianapolis, IN, USA), which detects the amount of histone-associated DNA fragments. The assay was performed according to the supplier’s manual; briefly, 10^6 cells were grown in a 96-well culture plate for 24 h, either starved for 12 h or maintained in complete medium for an additional 12 h, and then incubated with either ros or cig for up to 72 h, as indicated in the cell culture and cell preparation section. Supernatant culture media were removed, lysis buffer was added to each vial and histone-associated DNA fragments were colorimetrically quantified. Data represent the mean ± S.D. of five independent experiments, each performed in triplicate.

**Cell-cycle analysis**
GH3 or NIH3T3 cells either starved or cultured in complete grown medium were treated with either 1 – 50 μM ros or cig, or were untreated for 24 h; cells were trypsinized, collected by centrifugation, washed with PBS and treated with RNase A (Promega, Madison, WI, USA). DNA was stained with propidium iodide for 15 min at 4 °C and analyzed with a FACS Calibur (Becton Dickinson, Franklin Lakes, NJ, USA) apparatus. Analysis was performed by ModFit LT software (Becton Dickinson, Franklin Lakes, NJ, USA). Experiments were carried out in triplicate and repeated three times.

**Antibodies**
Human polyclonal anti-p53, anti-TRAIL, anti-Bax, anti-PPARγ and anti-β-actin antibody were obtained from Santa Cruz Biotechnology (Santa Cruz, CA, USA).

**Immunoblotting**
Cells (9 × 10^6) were washed in PBS and lysed in lysis buffer (50 mM TRIS–HCl, pH 6.8, 10% glycerol, 2.5% SDS, 10 mM DTT and 1 mM PMSF). Nuclear extracts were prepared as previously reported (32). Protein concentration was measured by Bradford assay using the Bio-Rad reagent (Bio-Rad Laboratories, Hercules, CA, USA). Proteins (25 μg) were resolved by 12% SDS–PAGE, transferred onto nitrocellulose membrane and stained with red ponceau to verify the amount of protein per lane. Nuclear extracts or whole-cell extracts were incubated overnight at 4 °C in 50% TBS (200 mM TRIS–HCl, pH 7.6, and 1.4 M NaCl) and 50% TTBS (TBS, 0.05% Tween-20), containing 5% nonfat dry milk, and subsequently incubated with the appropriate primary antibody for 1 h at RT. After extensive washing in TTBS, a horseradish peroxidase-conjugated antirabbit IgG was added for 1 h. After four washings of the membranes with TTBS and TBS, proteins were detected using an enhanced chemiluminescence detection system (Amersham Pharmacia Biotech, Piscataway NJ, USA). Experiments were performed three times with different nuclear or cell extracts.

**Transfections and plasmids**
The pBLCAT2PPRE vector containing a single copy of the PPAR response element of the acylCoA-oxidase gene has been described (33). Some 2 × 10^5 GH3 cells were transiently transfected with 10 μg of the pBLCAT2PPRE plasmid or the empty parental vector and 1 μg of the PCH110 plasmid containing the β-galactosidase gene, using the calcium phosphate method; the latter plasmid was used to account for the variation of transfection efficiency. NIH3T3 cells were transiently transfected under the same conditions or cotransfected with pBLCAT2PPRE and pSG5-PPARγ plasmid. Cell extracts were prepared 48 h later, and chloramphenicol acetyl transferase (CAT) assay and β-gal assay were performed as previously described (33). Results were expressed as arbitrary units considering 100% of the value obtained in the absence of cig or ros. Cig or ros were used at 1 – 10 μM. GH3 cells were transfected with the pBLCAT3-hGH plasmid containing the −397/+1 DNA sequence of the 5’ flanking region of the hGH gene, which was obtained by amplifying the corresponding sequence from the phGH plasmid (a gift of Dr N. Eberhardt, Mayo Clinic, USA). The integrity of the DNA fragment was confirmed by sequencing analysis. Data represent the mean of four independent experiments performed in triplicate.

**Gel mobility shift assay**
Expression vectors pSG5-PPARγ and pSG5-RXRα containing the cDNA of human PPARγ (kindly provided by Dr B. Desvergne, Lausanne, Switzerland) or mouse RXRα (kindly provided by Dr R.M. Evans, San Diego, CA, USA) were used for in vitro transcription-translation by the reticulocyte lysate system. The unprogrammed reticulocyte lysate was used as a negative control. The −397/−144, −397/+1 and −144/+1 fragments of the hGH promoter were obtained by PCR amplification of the original phGH plasmid, purified and 32P-labeled with T4 polynucleotide kinase. The binding reaction was carried out as previously reported (33); the DNA-protein complexes were resolved on a 5% nondenaturing polyacrylamide gel at 4 °C; gels were dried and exposed to autoradiography at −70 °C.

**RT-PCR for GH**
An amount of 1 μg of total RNA obtained from 30 × 10^6 GH3 or NIH3T3 cells was reverse-transcribed...
Figure 1 Expression of PPARγ in human pituitary GH-secreting tumors, in normal pituitary tissue and in GH3 cells. Panel 1) Tissue samples were obtained from 14 consecutive acromegalic patients with pure GH-secreting adenoma undergoing surgical excision of the tumor. Normal pituitary tissue samples were obtained during tumor excision of five GH-secreting adenomas (from the present series) and four nonfunctioning adenomas (panel 1H). Immunohistochemistry was performed using a specific antibody against human GH (panel 1A and C (GH-omas), and E (normal tissue)) or the N-terminus of PPARγ (panel 1B and D (GH-omas), and F (normal tissue)). Negative control was a GH-secreting tumor incubated with nonimmune serum (not shown). Original magnification × 250. Mean PPARγ expression, evaluated as percentage of stained nuclei, was 39 ± 24 in GH-secreting adenomas and 78 ± 5 in normal tissue (P < 0.0002); representative colocalization of PPARγ and GH in normal pituitary tissue (panel 1G). Panel 2) Representative Western blot of three human pituitary GH-secreting adenomas (ACRO 5, 4 and 9) and three normal pituitary tissue samples (N Pit 5, 4 and 9). Nuclear extracts from human tissue samples, rat GH3 and mouse NIH3T3 cells were immunoblotted with a specific antibody against PPARγ. In vitro synthesized PPARγ was used as a positive control; nuclear extracts from human skin and from NIH3T3 cells were used as a negative control. Panel 3) GH3 cells were transiently transfected with the pBLCAT2PPRE vector containing the PPAR response element of the acylCoA-oxidase gene and exposed to 1 μM ciglitazone (cig) or rosiglitazone (ros). NIH3T3 cells were transfected with the same reporter gene alone or cotransfected with pBLCAT2PPRE plasmid and pSG5-PPARγ. Data were expressed as relative CAT activity, assuming that the CAT activity obtained with the vehicle was considered 100%. Results represent the mean of four experiments carried out in triplicate.
in 50 mM Tris HCl (pH 8.3), 75 mM KCl, 3 mM MgCl₂, 10 mM DTT, 200 μM each dNTP, 10 mM Oligo (dT) primer, 20 U Rnasin (Promega) and 200 U M-MLV RT, in a final volume of 20 μl. The reaction was terminated by heating at 95°C for 5 min. A volume of 5 μl of the reaction was then amplified in a final volume of 25 μl in 1 x PCR buffer (Taq Man Universal PCR Master Mix, supplied by Applied Biosystems (Foster City, CA, USA), 300 mM of each primer and 200 nM of fluoresceinate probe for 34 cycles, as
Rats were killed by chloralum hydrate, and the treatment was ineffective. Ros was administered daily by gavage. On preliminary results with 40 mg/kg per day, which was the dose of the drug based on the animal care committee of our institution; after subcutaneous tumor development, the animals were randomized to receive ros (120 mg/kg per day) or vehicle. The dose of the drug was based on previous reports (34). Primers and the probe were as follows: primer sense, $5'$-gctgctgctgcatccacctac-3'; primer antisense, $5'$-ctggcattctgaattccga-3'; and probe, $5'$-6FAM-aggctcctgctgcatccaggg-3'. The primers set amplify a 80 bp fragment, the identity of which was confirmed by DNA sequencing. A control in which reverse transcription was omitted before PCR amplification was always included to exclude the possibility that any amplification was due to contaminating genomic DNA. Each experiment was performed in triplicate; data are expressed as number of copies of GH and represent the mean of three independent experiments.

**GH assays**

Cell-culture media were collected after exposure of cells to 1, 10, 50 μM cig or ros; cell lysates were obtained by treating the cells with lysis buffer (50 mM TRIS–HCl, pH 6.8, 10% glycerol, 2.5% SDS and 10 mM DTT); after centrifugation, supernatants were sonicated; blood samples were obtained from animals at baseline (before GH3 cells injection) and when rats were killed. GH was determined by radioimmunoassay (Amersham Biosciences). Measurement of GH in the cell lysates was normalized by protein concentrations. Sensitivity of GH assay was 1.0 ng/ml.

**Animals**

Five-week-old female Wistar-Furth rats were obtained from Harlan (Udine, Italy) and inoculated with $2 \times 10^6$ rat pituitary GH-secreting tumor cells, in accordance with the animal care committee of our institution; after subcutaneous tumor development, animals were randomized to receive ros (120 mg/kg per day) or vehicle. The dose of the drug was based on preliminary results with 40 mg/kg per day, which was ineffective. Ros was administered daily by gavage. Rats were killed by chloralum hydrate, and the tumors were weighed, and stored in liquid nitrogen.

**Statistics**

Data were expressed as mean±s.d. Comparison of parameters was performed by the analysis of variance (ANOVA).

**Results**

**PPARγ expression in pituitary GH-secreting tumors and normal pituitary tissues**

The expression of PPARγ was assessed in 14 consecutive surgically resected pituitary GH-secreting adenomas (GH-omas) and in nine normal pituitary tissue samples. Expression of PPARγ was analyzed by immunohistochemistry. PPARγ protein was intensively expressed, albeit showing heterogeneous distribution, in all pituitary GH-omas (Fig. 1, panel 1). Immunostained nuclei ranged 5–80% (mean 39±24%) in GH-omas, and 70–85% (mean 78±5%) in normal pituitary tissue samples ($P < 0.0002$). PPARγ and GH colocalized in the same pituitary cells as shown by double immunostaining (Fig. 1, panel 1). The presence of PPARγ was confirmed in nuclear extracts from pituitary normal tissue, GH-oma and GH3 cell extracts by Western blot; in vitro synthesized PPARγ was used as a positive control, while extracts of normal adult human skin and extracts of NIH3T3 cells were used as a negative control (35) (Fig. 1, panel 2). To evaluate the functional integrity of the receptor in GH3 cells, a reporter plasmid containing a PPRE fused to a heterologous promoter and a reporter gene (pBLCAT2PPRE), was transiently transfected into either GH3 or NIH3T3 cells. Addition of 1 μM cig or ros increased the basal level of expression of the reporter plasmid by 1.8-fold and 2.2-fold respectively only in GH3 cells, indicating that this cell line expressed a functionally integer PPARγ (Fig. 1, panel 3). Cotransfection of PPARγ was necessary to activate the pBLCAT2PPRE plasmid in NIH3T3 cells, indicating the absence of a significant amount of functionally active PPARγ in this cell line.

**PPARγ ligands increase apoptosis in GH3 cells**

GH3 cells exposed to 1–50 μM cig or ros for up to 72 h exhibited an increased apoptosis rate (Fig. 2). Apoptosis increased from 17.0±2.3% to 36.0±6.5% ($P < 0.004$) and to 44.3±6.0% ($P = 0.0007$) in cells cultured in complete medium after exposure to 50 μM cig or 50 μM ros respectively, as assessed by the annexin V iodide propidium method (Fig. 2, panel 1); treatment of starved cells with either 1–50 μM cig or ros gave similar results (data not shown). Treatment of NIH3T3 cells with either cig or ros did not affect the apoptosis rate, thus indicating the specificity of action of PPARγ agonists (Fig. 2, panel 2). Apoptosis increased in a dose-dependent manner in GH3 cells treated with either cig or ros up to 2.5-fold, as assessed by measuring the amount of histone-associated DNA fragments ($P = 0.0003$) (Fig. 2, panel 3).

**Effects of PPARγ ligands on GH3 proliferation**

Treatment with 1–50 μM cig or ros for up 72 h increased the number of GH3 cells in the G1 phase (92±4% vs 73±3% in controls, $P < 0.001$) and reduced the cells in the S-phase (8.5±0.9 vs 18.5±0.5% in controls, $P < 0.0001$) of cells grown in complete medium, thus indicating a G0–G1 cell-cycle arrest (Fig. 3). Similar results were obtained in starved GH3 cells (data not shown). Treatment with either cig or ros did not affect the cell cycle of NIH3T3 cells (Fig. 3).
Effects of PPARγ ligands on apoptosis-associated proteins

GH3 cells were treated with either 50 μM cig or ros for up to 72 h. Cellular extracts were immunoblotted with specific antibody against p53, TRAIL and Bax. Expression of p53 and Bax was not affected by either cig or ros treatment, while activation of PPARγ by these drugs was associated with increased expression of TRAIL proteins (Fig. 4). NIH3T3 cell extracts were not used because the apoptosis rate of this cell line was not affected by cig or ros treatment as reported in the above sections.

PPARγ ligands reduce the secretion of GH from GH3 cells

GH3 cells were treated with 1–50 μM ros for up 72 h, and culture media and cellular extracts were collected.
for GH determinations. An amount of 10 μM ros reduced GH concentrations in culture media from $8.3 \pm 1.4$ to $1.9 \pm 0.7$ ng/ml after 24 h ($P < 0.04$), from $16.5 \pm 3.7$ to $3.6 \pm 0.9$ ng/ml after 48 h ($P < 0.03$) and from $43.7 \pm 5.4$ to $2.1 \pm 0.3$ ng/ml after 72 h ($P < 0.0001$). Ros at 1–50 μM reduced GH concentrations in culture media in a dose-dependent manner ($P < 0.04$) (Fig. 5, panel 2). GH concentrations decreased in cell extracts from ros-treated cells from $233 \pm 13$ to $109 \pm 11$ ng/ml ($P < 0.004$) (Fig. 5, panel 2).

**PPARγ ligands reduce transcriptional activity of human GH promoter**

The levels of expression of GH mRNA were measured in either untreated or 1–10 μM cig- or ros-treated GH3 cells by RT-PCR. GH transcripts reduced from $5.7 \times 10^6$ in untreated cells to $3.0 \times 10^6$ and $1.8 \times 10^6$ copies in 1 and 10 μM cig-treated cells respectively ($P < 0.002$). Cells treated with 1–10 μM ros had a similar response (Fig. 6, panel 1). RNA obtained from NIH3T3 cells was used as a negative control. Since cig and ros had very similar effects on the transcription of hGH gene, in the next experiments we used only one TDZ.

The −397/+1, −397/−144, and −144/+1 DNA fragment of the 5'-flanking region of the GH gene were used as 32P-labeled probes in gel mobility shift assays: PPARγ–RXRα heterodimers bound to the −397/+1 and the −397/−144 DNA fragment (data not shown); the latter and shorter probe was then used for further experiments. The PPARγ–RXRα heterodimer formed a specific complex with the GH promoter, indicating...
the presence of a putative PPRE in the hGH gene (Fig. 6, panel 2). Addition of 10 μM cig reduced the binding of the PPARγ–RXRα heterodimer to the hGH promoter (Fig. 6, panel 2). The pBLCAT3-hGH−397/+1 plasmid was transfected into GH3 cells alone or with pSG5-PPARγ and/or pSG5–RXRα. Treatment of the transfected cells with cig decreased the CAT activity by 50% (P < 0.0005 vs untreated cells). Cotransfection with PPARγ and RXRα further reduced the CAT activity by up to 65% (P < 0.0001 vs untreated cells; P = NS vs cig-treated cells not transfected with PPARγ and RXRα) (Fig. 6, panel 3).

**Ros treatment slows progression of pituitary GH-secreting tumors and reduces GH secretion in rats**

Five-week-old female Wistar-Furth rats were injected subcutaneously with 2 × 10⁶ GH3 cells. After 3 weeks, all animals developed 2 cm subcutaneous tumors. Rats were randomly assigned to receive either ros (120 mg/kg per day) (n = 5) or vehicle (n = 5). The dose of ros was based on that reported to be effective in other studies (29, 30) and on experiments with lower doses of ros (40 mg/kg per day), which were ineffective (Fig. 7). After 3–4 weeks, tumor size increased in vehicle-treated rats; in contrast, ros-treated animals had a slower increase of tumor weight (14.8 ± 4.2 g vs 24.1 ± 2.0 g in placebo-treated rats, P < 0.03) (Fig. 7, panels 1–2). According to the smaller weight of TZDs-treated rats, serum GH concentrations were lower in rats treated with 120 mg/kg per day TZDs (871 ± 67 ng/ml) than in placebo-treated rats (1.309 ± 238) or rats treated with 40 mg/kg per day TZDs (1.289 ± 327) dose (*P < 0.05) (Fig. 7, panel 3).

**Discussion**

PPARγ is considered a tumor-suppressor gene, the activation of which leads to cellular differentiation, apoptosis and arrest of tumor progression (23–28, 36). Recently, we showed a possible link between GH-IGF-I hypersecretion and PPARγ in the colonic mucosa of acromegalic patients (31, 34). The involvement of PPARγ in pituitary diseases has been indicated by recent reports by Heaney et al. (29, 30): its expression...
seems to be restricted to ACTH-secreting cells only in the normal pituitary, while pituitary adenomas have an abundant expression of this nuclear receptor. PPARγ may induce cell-cycle arrest and apoptosis in pituitary hormone-secreting tumors and nonfunctioning adenomas (29, 30). Treatment with TZDs given simultaneously with the injection of GH3 cells retarded the development of subcutaneous GH-secreting tumors in nude mice (30). We believe that, albeit important, this observation has some limitations. The expression of PPARγ was studied in only three pituitary pure GH-secreting adenomas, while our series included 14 consecutive acromegalic patients showing a variable expression of this nuclear receptor; in conflict with the report of Heaney et al. (29, 30), normal pituitary tissue showed a high expression of PPARγ which colocalized with GH. One explanation might be the different origin of normal pituitary tissue: autoptic in the study of Heaney et al. but fresh tissue in the present paper. The variation in the expression of PPARγ in GH-secreting adenomas was not related to serum GH or IGF-1 levels nor to the weight of adenomas (data not shown); however, it could not be ruled out that the long-standing exposure to high GH-IGF1 concentrations may play a role in reducing the expression of PPARγ in the pituitary adenomas, as reported for the

**Figure 6** PPARγ ligands reduce the transcriptional activity of the hGH promoter. Panels 1 and 2) Expression of GH mRNA in GH3 cells treated with either 1–10 μM cig (filled columns) or ros (empty columns); analysis of GH expression was made by RT-PCR. The representative standard curve for quantitative RT-PCR for GH is shown in panel 1; the threshold cycle (Ct; that is, the PCR cycle number at which the fluorescence signal reached above baseline) was −3.269 (starting copy number) + 37.265 (r = 0.995). Panel 2) GH levels in ros-treated GH3 cells decreased from to 5.7 × 10⁶ to 1.8 × 10⁶ transcripts (P < 0.002), similarly to those observed in cig-treated cells. Panel 3) Gel mobility shift assay used the −397/–144 DNA sequence of the human GH promoter as a 32P-labeled probe: the specific complex of PPARγ-RXRα heterodimer with hGH promoter is shown by arrows. Panel 4) The pBLCAT3-hGH plasmid containing the −397/+1 flanking region of the hGH gene was transfected into the GH3 cells. Exposure of the transfected cells to 1–10 μM ros decreased the CAT activity by up to 70%. CAT activity was expressed as a percentage of the value obtained in absence of drug, which was arbitrarily considered 100%.

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colonic mucosa (31, 34) of some acromegalic patients. More important, our data clearly showed that TZDs treatment can slow the growth of well-established GH-secreting tumors. Heaney et al. showed the preventive effects of TZDs on tumor development. In fact, nude mice injected with GH3 cells and simultaneously treated with TZDs had delayed development of subcutaneous tumors, which grew more slowly than those of controls. We started the treatment with TZDs only after complete development of subcutaneous tumors, the growth of which was only partially reduced. This is the situation occurring in vivo in humans, because the treatment is initiated only after tumors have developed and grown. GH concentrations, measured either in the culture media of GH3 cells treated with ros or in the blood from rats treated with this drug, clearly decreased. The fact that we did not observe a significant tumor shrinkage after ros treatment might also be due to the greater volume of the tumors with respect to those reported by Heaney et al. (29, 30).

Our data clearly demonstrated that TZDs also affected the synthesis of GH; in fact, GH3 cells treated with TZDs had a more reduced intracellular content of GH than controls. Moreover, PPARγ ligands might effectively reduce GH hypersecretion, either reducing GH transcription and secretion or increasing the apoptosis and arresting the growth of pituitary GH-secreting cells. Whether the maximum dose of ros allowed in humans will cause GH deficiency remains to be elucidated; however, this outcome seems to be unlikely either because this dose might not be sufficient in reducing GH secretion or because of the compensatory mechanisms of the complex signaling network regulating GH secretion.
Our data also showed that increased TRAIL protein expression might be a common effector of PPARγ–induced apoptosis in GH3 cells, as reported for other transcription factors (37). Unlike other pituitary cell lines, in which TZDs treatment was associated with downregulation of Bcl-2 and increased expression of Bax (30), GH3 cells did not exhibit variations of these apoptotic proteins under TZDs exposure. These effects of PPARγ ligands were specific since they were not observed in NIH3T3 cells, which did not express detectable amount of PPARγ.

Ros is currently employed in the treatment of type 2 diabetes as a monotherapy or in combined therapy. Its metabolism in the liver differs from that of troglitazone, and this might explain its low incidence of hepatic abnormalities (38). However, the dose effective in lowering GH secretion and to induce apoptosis in GH3-induced tumors was higher than those used in the treatment of type 2 diabetes. Therefore, a note of caution is necessary. It also should be pointed out that similar doses of TZDs have been used in preventing atherosclerosis or to treat thyroid tumors in mice (24, 38, 39). TZDs have also been used for treating human liposarcoma or prostate cancer at doses higher than those employed for glycemic control (23, 40). In conclusion, although our results need to be validated in clinical trials, they provide the molecular basis for the use of TZDs as adjunct therapeutic agents in the challenging management of acromegalic patients.

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