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# Polyacetylenes from *Notopterygium incisum*—New Selective Partial Agonists of Peroxisome Proliferator-Activated Receptor-Gamma

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## Abstract

Peroxisome proliferator-activated receptor gamma (PPAR $\gamma$ ) is a key regulator of glucose and lipid metabolism and therefore an important pharmacological target to combat metabolic diseases. Since the currently used full PPAR $\gamma$  agonists display serious side effects, identification of novel ligands, particularly partial agonists, is highly relevant. Searching for new active compounds, we investigated extracts of the underground parts of *Notopterygium incisum*, a medicinal plant used in traditional Chinese medicine, and observed significant PPAR $\gamma$  activation using a PPAR $\gamma$ -driven luciferase reporter model. Activity-guided fractionation of the dichloromethane extract led to the isolation of six polyacetylenes, which displayed properties of selective partial PPAR $\gamma$  agonists in the luciferase reporter model. Since PPAR $\gamma$  activation by this class of compounds has so far not been reported, we have chosen the prototypical polyacetylene faltarindiol for further investigation. The effect of faltarindiol (10  $\mu$ M) in the luciferase reporter model was blocked upon co-treatment with the PPAR $\gamma$  antagonist T0070907 (1  $\mu$ M). Faltarindiol bound to the purified human PPAR $\gamma$  receptor with a  $K_i$  of 3.07  $\mu$ M. *In silico* docking studies suggested a binding mode within the ligand binding site, where hydrogen bonds to Cys285 and Glu295 are predicted to be formed in addition to extensive hydrophobic interactions. Furthermore, faltarindiol further induced 3T3-L1 preadipocyte differentiation and enhanced the insulin-induced glucose uptake in differentiated 3T3-L1 adipocytes confirming effectiveness in cell models with endogenous PPAR $\gamma$  expression. In conclusion, we identified faltarindiol-type polyacetylenes as a novel class of natural partial PPAR $\gamma$  agonists, having potential to be further explored as pharmaceutical leads or dietary supplements.

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¶ These authors contributed equally to this work.

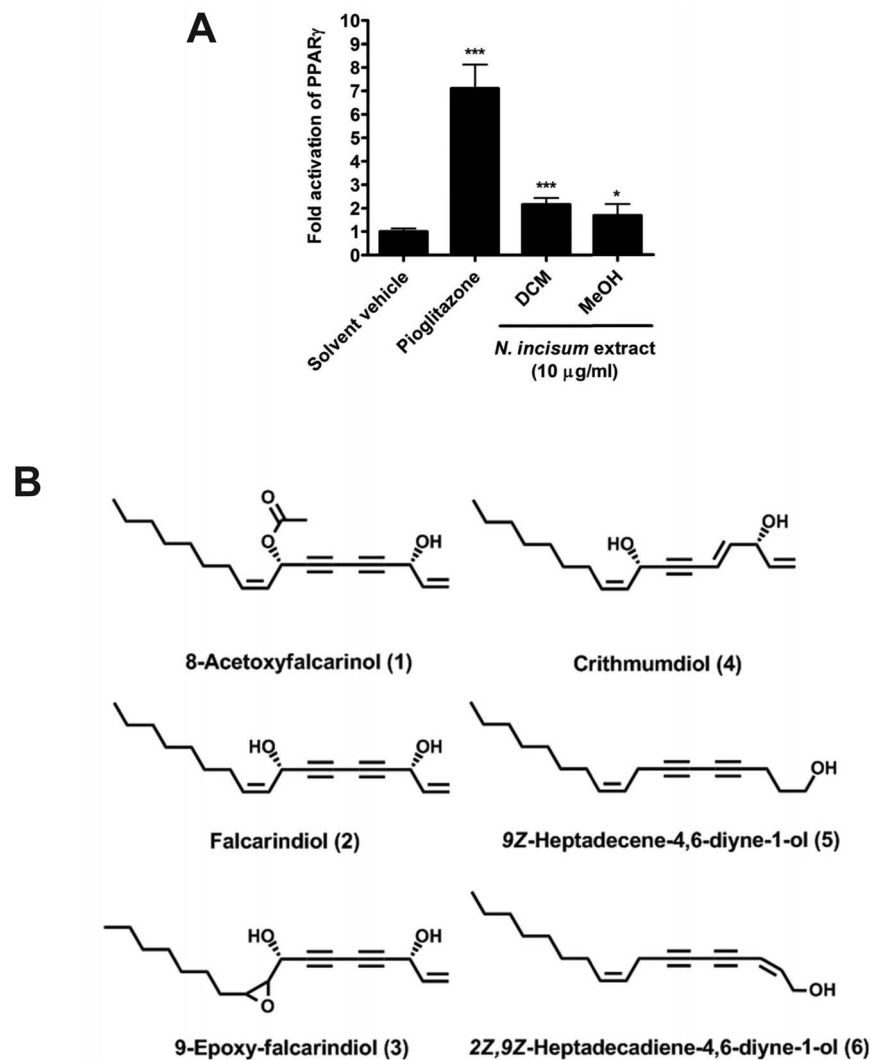
¶ These authors also contributed equally to this work.

## Introduction

Peroxisome proliferator activated receptors (PPARs) are ligand-dependent nuclear receptors which play a major role in lipid and glucose metabolism [1–5]. Three subtypes of PPAR, namely PPAR $\alpha$ , PPAR $\beta/\delta$ , and PPAR $\gamma$ , have been identified in human and other species [6,7]. Upon ligand binding, PPARs form heterodimers with another nuclear receptor, the retinoid X receptor (RXR) and subsequently bind to response elements located in the promoter region of their target genes [8]. The binding of a PPAR/RXR heterodimer to its PPAR response element (PPRE) triggers the recruitment of nuclear receptor

coactivators [9,10], and subsequent chromatin rearrangement allowing initiation of the transcription of target genes [11].

The three PPAR subtypes are expressed in different tissues and regulate distinct physiological functions. PPAR $\alpha$  is mainly expressed in muscle, liver, heart, and kidney, and is involved in the regulation of genes that play a role in lipid and lipoprotein metabolism [12,13]. The PPAR $\beta/\delta$  subtype is ubiquitously expressed in various tissues and predominantly associated with lipid metabolism and energy expenditure [14]. Among the three PPAR subtypes, PPAR $\gamma$  is the best studied one. PPAR $\gamma$  is expressed in adipose tissue, lung, large intestine, kidney, liver, heart, and macrophages [15], and is engaged in the regulation of adipogenesis, insulin sensitivity, and inflammation [16]. The



**Figure 1. PPAR $\gamma$  activation by *N. incisum* extracts and chemical structures of the isolated polyacetylenes.** (A) HEK-293 cells were transiently co-transfected with a plasmid encoding full-length human PPAR $\gamma$ , a reporter plasmid containing PPRE coupled to a luciferase reporter and an EGFP plasmid as internal control. After re-seeding, cells were treated as indicated for 18 h. Since the positive control pioglitazone (5  $\mu$ M), as well as the dried DCM- and MeOH-extract were reconstituted in DMSO, cells were treated with equal amount of the solvent vehicle DMSO (0.1%) as negative control. The luciferase activity was normalized to the EGFP-derived fluorescence, and the result is expressed as fold induction compared to the solvent vehicle control. The data shown are means  $\pm$  SD of three independent experiments each performed in quadruplet. \*\*\* $p$ <0.001, \* $p$ <0.05 (compared to the solvent vehicle group; ANOVA/Bonferroni). (B) Chemical structures of the PPAR $\gamma$ -activating polyacetylenes isolated from *N. incisum*. doi:10.1371/journal.pone.0061755.g001

important functions of PPAR $\gamma$  in the regulation of glucose and lipid metabolism make it an attractive pharmacological target for combating metabolic diseases [17–19]. It is established that the activation of this receptor mediates the action of the glitazone-type drugs clinically used for the treatment of type 2 diabetes [20]. Furthermore, recent findings supporting a role of PPAR $\gamma$  in inflammation and cell growth have promoted the investigation of PPAR $\gamma$  agonists as experimental drugs for some chronic diseases such as atherosclerosis and cancer, among others [21–23]. The glitazones currently used in clinics (e.g., pioglitazone) potently activate PPAR $\gamma$  as full agonists. In spite of being effective drugs, their use is limited due to undesirable side effects [24], urging the identification and characterization of new PPAR $\gamma$  agonists. A number of recent studies revealed that partial PPAR $\gamma$  agonists inducing submaximal receptor activation are demonstrating great promise by exerting good hypoglycaemic activity with reduced side effects [25–27].

Natural products encompass a broad structural diversity of secondary metabolites, which often represent privileged structures serving a variety of biological functions [28]. Therefore, we focused on the identification of novel PPAR $\gamma$  activators derived from natural sources. As part of an ethnopharmacology-based screening approach, extracts from medicinal plants used in traditional Chinese medicine to treat inflammation-related conditions were tested for PPAR $\gamma$  activation. Extracts of the rhizomes and roots of *Notopterygium incisum* displayed significant PPAR $\gamma$  activation. The plant has been traditionally used in China for the treatment of rheumatism, cold, and headache [29], and its lipophilic extracts have shown inhibitory effects on 5-lipoxygenase and cyclooxygenase [30]. Pure compounds isolated from the dichloromethane extracts were now examined for PPAR $\gamma$  activation, leading to the identification of falcarindiol-type polyacetylenes as a novel group of natural selective partial PPAR $\gamma$  agonists.

## Materials and Methods

### Chemicals, Cell Culture Reagents, and Plasmids

Calf serum, L-glutamine, and Dulbecco's modified Eagle's medium (DMEM) containing 4.5 g/l glucose was purchased from Lonza (Basel, Switzerland). Fetal bovine serum (FBS) was supplied from Gibco (Lofer, Austria). PPAR $\alpha$  and PPAR $\beta/\delta$  agonists GW7647 and GW0742, respectively, as well as the PPAR $\gamma$  antagonist T0070907, were purchased from Cayman (Missouri, USA); pioglitazone was purchased from Molekula Ltd (Shaftesbury, UK). All other chemicals were obtained from Sigma–Aldrich (Vienna, Austria). The tested compounds, or dried plant extracts, were dissolved in dimethyl sulfoxide (DMSO), aliquoted, and stored at  $-20^{\circ}\text{C}$  until use. The final concentration of DMSO in all experiments was 0.1% or lower. An equal amount of DMSO was always tested in each experiment to assure that the solvent vehicle does not influence the results. The three human PPAR subtype expression plasmids (pSG5-PL-hPPAR-alpha, pSG5-hPPAR-beta, pSG5-PL-hPPAR-gamma1) were a kind gift from Prof. Beatrice Desvergne and Prof. Walter Wahli (Center for Integrative Genomics, University of Lausanne, Switzerland), and the luciferase reporter plasmid (tk-PPREx3-luc) was kindly provided by Prof. Ronald M. Evans (Salk Institute for Biological Studies, San Diego, California).

### PPAR Luciferase Reporter Gene Transactivation

HEK-293 cells (ATCC, USA) were cultured in DMEM with phenol red, supplemented with 100 U/ml benzylpenicillin, 100  $\mu\text{g}/\text{ml}$  streptomycin, 2 mM L-glutamine, and 10% FBS. Cells were maintained in 75  $\text{cm}^2$  flasks containing 10 ml medium at  $37^{\circ}\text{C}$  and 5%  $\text{CO}_2$ . The cells were seeded in 10 cm dishes at a density of  $6 \times 10^6$  cells/dish, incubated for 18 h, and transfected by the calcium phosphate precipitation method [31] with 4  $\mu\text{g}$  of the reporter plasmid (tk-PPREx3-luc), 4  $\mu\text{g}$  from the respective PPAR subtype expression plasmid, and 2  $\mu\text{g}$  green fluorescent protein plasmid (pEGFP-N1, Clontech, Mountain View, CA) as internal control. After 6 h, the transfected cells were harvested and re-seeded ( $5 \times 10^4$  cells/well) in 96-well plates containing DMEM

without phenol red, supplemented with 100 U/ml benzylpenicillin, 100  $\mu\text{g}/\text{ml}$  streptomycin, 2 mM L-glutamine, and 5% charcoal-stripped FBS. The cells were further treated with the indicated extracts or compounds or the solvent vehicle and incubated for 18 h. The medium was then discarded and the cells were lysed with a reporter lysis buffer (E3971, Promega, Madison, USA). Luciferase activity of the cell lysates was evaluated using a GeniosPro plate reader (Tecan, Grödig, Austria). The luminescence signals obtained from the luciferase activity measurements were normalized to the EGFP-derived fluorescence, to account for differences in the transfection efficiency or cell number. Luciferase reporter gene assays were performed with polyacetylene concentrations up to 30  $\mu\text{M}$  as higher dosages led to a downregulation of the internal normalization control (EGFP, not shown).

### Extraction and Isolation

Plant material of *Radix et Rhizoma Notopterygii* was purchased from Plantasia Austria in 2008. A voucher specimen (No. 650107) is kept at the Institute of Pharmaceutical Sciences, Department of Pharmacognosy at the University of Graz. The plant material was grounded and extensively percolated with dichloromethane (DCM) to obtain a crude DCM extract. Bioassay-guided fractionation of the DCM extract resulted in the isolation and identification of the following compounds (Fig. 1B): 8-acetoxycarindiol (1, 3R,8S,9Z-3-hydroxyheptadeca-1,9-diene-4,6-diyne-8-yl acetate), faltarindiol (2, 3R,8S,9Z-heptadeca-1,9-diene-4,6-diyne-3,8-diol), 9-epoxy-faltarindiol (3, 1R,6R-1-(3-heptyloxiran-2-yl)octa-7-ene-2,4-diyne-1,6-diol), crithmundiol (4, 3R,4E,8S,9Z-heptadeca-1,4,9-triene-6-yne-3,8-diol), 9Z-heptadecene-4,6-diyne-1-ol (5) and 2Z,9Z-heptadecadiene-4,6-diyne-1-ol (6). The purity of the isolated compounds was over 95% for 1-4 and above 90% for 5 and 6. All analytical HPLC experiments were performed using an Agilent 1100 series system (Agilent 1100, Series Degaser G1311 A, Quat Pump G1311, Autosampler G1313 A, Colcom G1316 A, DAD 1315 B) equipped with a diode-array detector. Analyses were performed using a SB-C18 Zorbax column (3.5  $\mu\text{m}$ ; 150 $\times$ 2.1 mm; Agilent Technologies) at a flow rate of 300  $\mu\text{l}/\text{min}$  and a gradient elution program. The extract

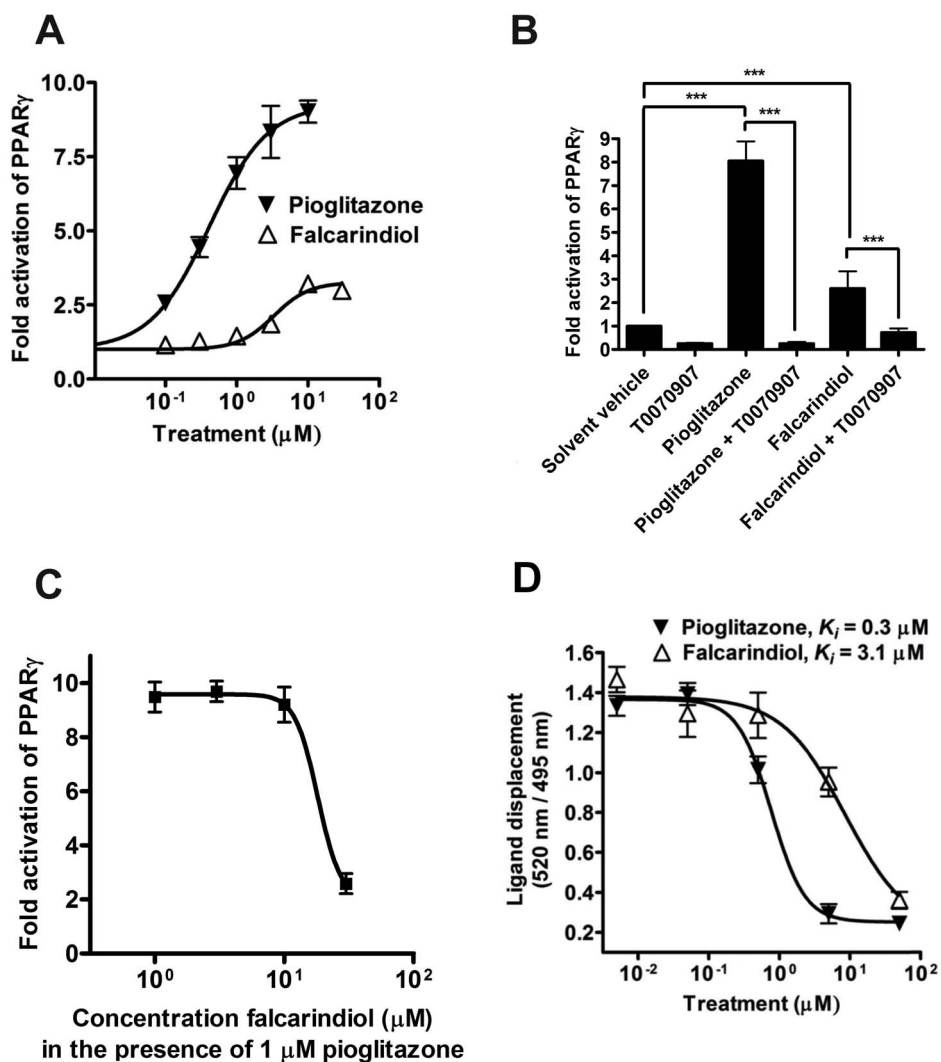
**Table 1.** Activity of the isolated polyacetylenes towards the three subtypes of human PPAR ( $\alpha$ ,  $\beta/\delta$ ,  $\gamma$ ) in a luciferase reporter transactivation assay.

	PPAR $\gamma$ activation		PPAR $\alpha$ activation		PPAR $\beta/\delta$ activation	
	EC $_{50}$ ( $\mu\text{M}$ )	maximal fold activation	EC $_{50}$ ( $\mu\text{M}$ )	maximal fold activation	EC $_{50}$ ( $\mu\text{M}$ )	maximal fold activation
Pioglitazone	0.41	9.28	–	–	–	–
GW7647	–	–	0.0021	3.08	–	–
GW0742	–	–	–	–	0.0017	20.2
(1) 8-Acetoxyfaltarindiol	3.59	2.36	n.d.	n.d.	n.d.	n.d.
(2) Faltarindiol	3.29	3.26	n.d.	n.d.	n.d.	n.d.
(3) 9-Epoxy-faltarindiol	2.03	1.88	n.d.	n.d.	n.d.	n.d.
(4) Crithmundiol	4.58	2.29	n.d.	n.d.	n.d.	n.d.
(5) 9-Heptadecene-4,6-diyne-1-ol	11.31	1.92	n.d.	n.d.	n.d.	n.d.
(6) 2Z,9Z-Heptadecadiene-4,6-diyne-1-ol	4.18	1.73	n.d.	n.d.	n.d.	n.d.

n.d. not detected up to 30  $\mu\text{M}$ .

HEK-293 cells transiently co-transfected with the respective PPAR subtype expression plasmids, a luciferase reporter plasmid (tk-PPREx3-luc), and EGFP as internal control were treated with six different concentrations of the polyacetylenes (0.1–30  $\mu\text{M}$ ) for 18 h. The luciferase activity was expressed as fold induction above the vehicle control (DMSO, 0.1%) after normalization to the EGFP-derived fluorescence. To verify the specificity of the performed assays, GW7647, GW0742, and pioglitazone were used as a selective agonist for PPAR $\alpha$ , PPAR $\beta/\delta$ , and PPAR $\gamma$ , respectively. EC $_{50}$  and maximal fold activation were determined with a non-linear regression in the GraphPad Prism software version 4.03 (GraphPad Software Inc, USA). The data shown are means of three independent experiments performed in triplicate.

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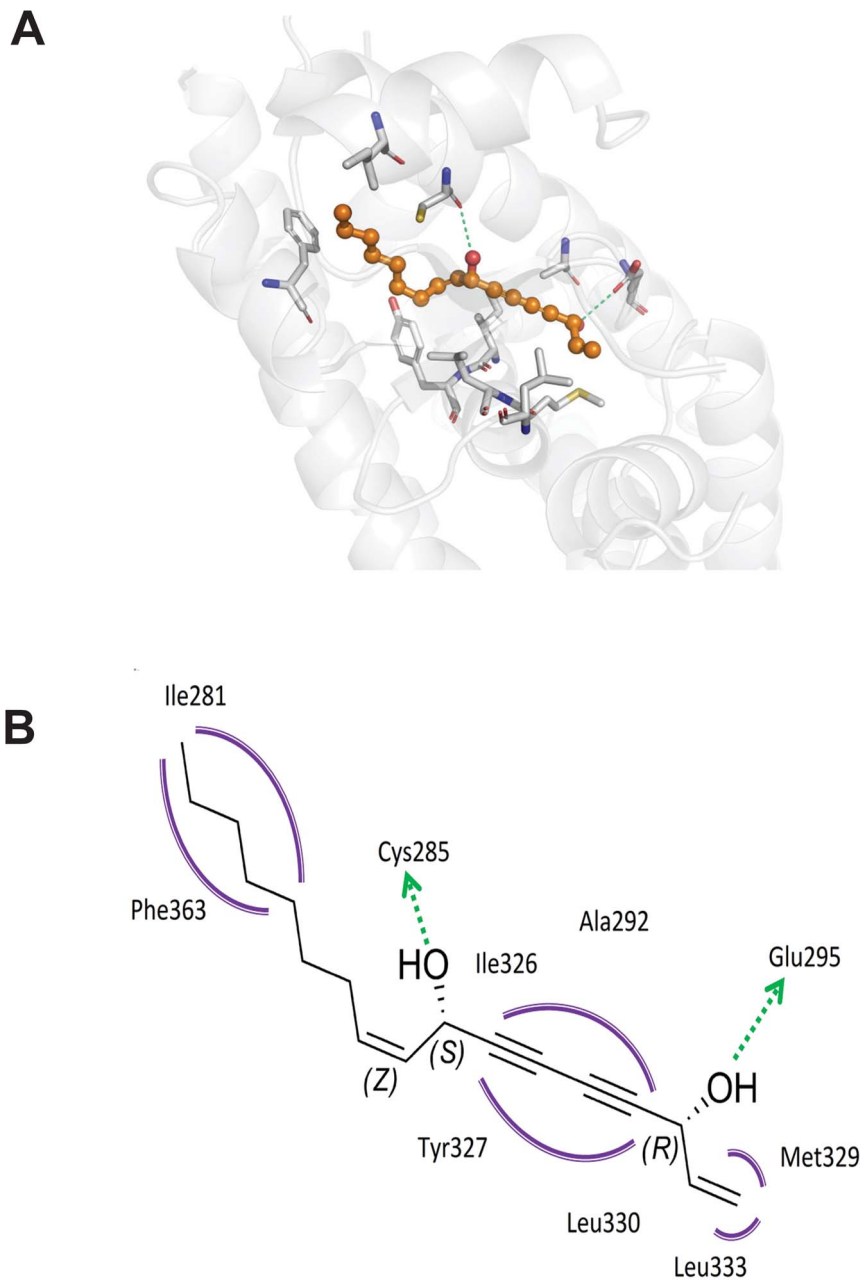
**Figure 2. PPAR $\gamma$ -mediated transactivation activity as well as receptor binding activity of falcarindiol.** (A) HEK-293 cells, transiently transfected with a human PPAR $\gamma$  expression plasmid, a luciferase reporter plasmid (tk-PPRE3-luc) and EGFP as internal control, were treated with different concentrations of pioglitazone or falcarindiol (0.1–30  $\mu$ M) for 18 h. Luciferase activity was normalized by the EGFP-derived fluorescence, and the result is expressed as fold induction compared to the solvent vehicle control (DMSO, 0.1%). The data points shown are means  $\pm$  SEM of three independent experiments each performed in quadruplet. (B) Cells were transfected and treated as indicated above. Pioglitazone was applied at 5  $\mu$ M, falcarindiol at 10  $\mu$ M, and the PPAR $\gamma$  antagonist T0070907 at 1  $\mu$ M. The data shown are means  $\pm$  SD of six independent experiments each performed in quadruplet. \*\*\* $p$ <0.001 (ANOVA/Bonferroni). (C) The cells were prepared as indicated above and treated for 18 h with different concentrations of falcarindiol, always in the presence of 1  $\mu$ M pioglitazone. The data shown are means  $\pm$  SD of three independent experiments each performed in quadruplet. (D) Dilutions of the two investigated compounds were prepared in DMSO and mixed with a buffer containing the hPPAR $\gamma$  LBD tagged with GST, terbium-labelled anti-GST antibody, and fluorescently-labelled PPAR $\gamma$  agonist. After 6 h of incubation, the ability of the test compounds to bind to the PPAR $\gamma$  LBD and thus to displace the fluorescently labelled ligand was estimated from the decrease of the emission ratio 520 nm/495 nm upon excitation at 340 nm. Each data point represents the mean  $\pm$  SEM from four independent experiments performed in duplicate.  
doi:10.1371/journal.pone.0061755.g002

was phytochemically investigated using a combination of different chromatographic techniques such as LC, SPE (RP-C18), and preparative HPLC (RP-C18). Isolated constituents were characterized and identified by multidimensional NMR spectroscopy ( $^1\text{H}$ ,  $^{13}\text{C}$ , COSY, HMBC, HSQC) and mass spectrometry.

### Receptor Binding Study

The LanthaScreen<sup>®</sup> time-resolved fluorescence resonance energy transfer (TR-FRET) PPAR $\gamma$  competitive binding assay (Invitrogen, Lofer, Austria) was performed according to the manufacturer's protocol. The test compounds dissolved in DMSO or the blank solvent vehicle (DMSO) were incubated together with

the human PPAR $\gamma$  ligand-binding domain (LBD) tagged with GST, a terbium-labeled anti-GST antibody, and a fluorescently labeled PPAR ligand (Fluormone Pan-PPAR Green, Invitrogen). In this assay, the fluorescently labeled ligand is binding to the human PPAR $\gamma$  LBD, which brings it in close spatial proximity to the terbium-labeled anti-GST antibody. Excitation of the terbium at 340 nm results in energy transfer (FRET) and partial excitation of the fluorescently labeled PPAR ligand, followed by emission at 520 nm. Test-compounds binding to the human PPAR $\gamma$  LBD are competing with the fluorescently labeled ligand and displacing it, resulting in a decrease of the FRET signal. The signal obtained at 520 nm is normalized to the signal obtained from the terbium



**Figure 3. Proposed molecular binding mode of falcariindiol in the PPAR $\gamma$  LBD.** Predicted binding mode of falcariindiol shown as 3D depiction in which predicted hydrogen-bonds are shown as dashed lines (A), and 2D depiction including chemical features of the interaction pattern derived from the docking pose (B). Chemical features in the 2D depiction are color-coded: green arrow – hydrogen-bond donor; magenta – hydrophobic contacts.

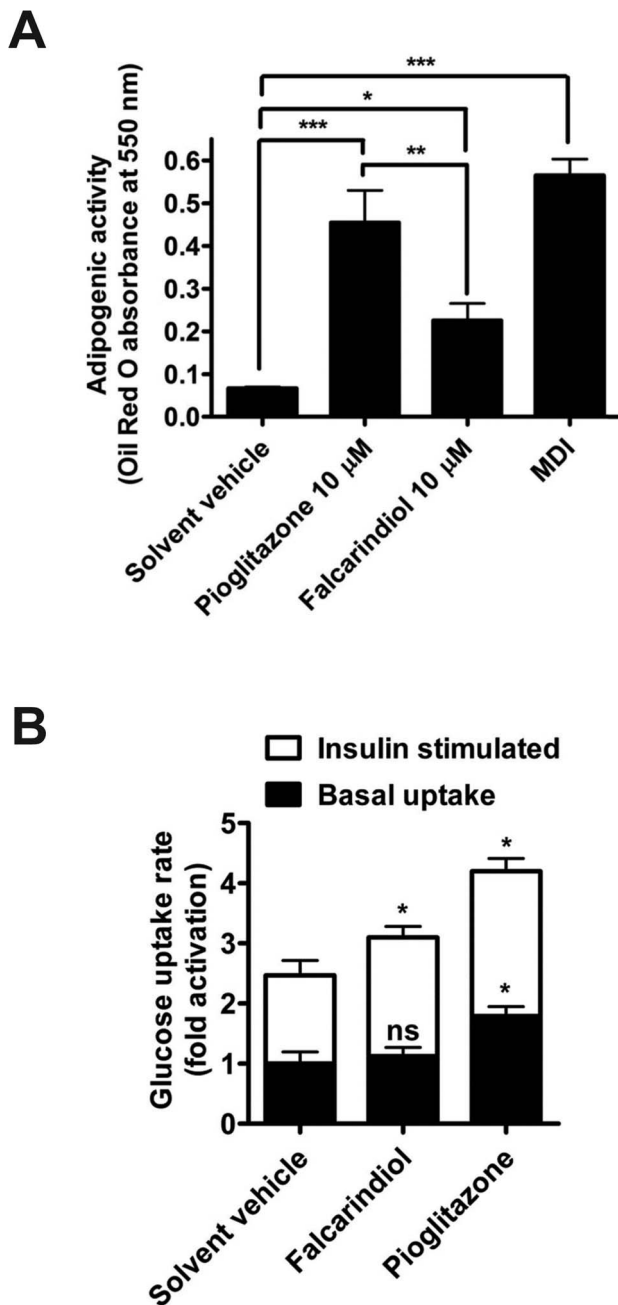
doi:10.1371/journal.pone.0061755.g003

emission at 495 nm; therefore, the decrease in the 520 nm/495 nm ratio is used as a measure for the ability of the tested compounds to bind to the human PPAR $\gamma$  LBD. All measurements were performed with a GeniosPro plate reader (Tecan, Grödig, Austria).

### Molecular Docking

To predict binding modes of the falcariindiol-type polyacetylene PPAR $\gamma$  partial agonists, a docking study was performed, using the quantum mechanics-polarized ligand docking (QPLD) workflow [32] in Maestro version 9.2.112 (Schrodinger, LLC, New York,

NY, 2011; <http://www.schrodinger.com>). In this docking study, the X-ray crystal structure of PPAR $\gamma$  with two co-crystallized molecules of the natural product magnolol was used (Protein Data Bank entry: 3r5n; <http://www.rcsb.org/pdb/>) [33]. The ligand and protein preparation were conducted using the Ligprep module with default settings for the ligands and the Protein Preparation Wizard for the protein. The protein was prepared by adding hydrogens, removing water molecules, assigning atom and bond types, and a refinement by completing missing side chains. Then, an exhaustive exploration of the hydrogen bond assignment was conducted. The protein was minimized applying the OPLS-AA 2005 force field with a threshold of 0.3 Å RMSD. Afterward,



**Figure 4. Assessment of adipogenicity and glucose uptake-enhancing properties of falcариндиол.** (A) 3T3-L1 preadipocytes were differentiated to adipocytes as described in the *Materials and Methods* section for 10 days in the presence of pioglitazone, falcариндиол, solvent vehicle (0.1% DMSO), or a standard differentiation mixture (MDI; containing 10% FBS, 500 µM IBMX, 500 nM dexamethasone, and 1 µg/ml insulin) as a positive control. For an estimate of accumulated lipids, and thus for the adipogenic potential of the test compounds, Oil Red O staining and photometric quantification at 550 nm was performed. The data shown are means  $\pm$  SD of three independent experiments. \*\*\* $p$ <0.001, \*\* $p$ <0.01, \* $p$ <0.05 (ANOVA/Bonferroni). (B) 3T3-L1 mature adipocytes were incubated with falcариндиол (10 µM) or solvent vehicle (0.1% DMSO) for 48 h and 2-deoxy-D-(1H<sup>3</sup>)-glucose cellular uptake was determined for 10 min in the presence or absence (basal uptake) of insulin (500 pM). The data shown are means  $\pm$  SEM of five independent experiments. \* $p$ <0.05 (compared to the solvent vehicle group; two-tailed paired  $t$ -test).

doi:10.1371/journal.pone.0061755.g004

atomic partial charges of the ligands were calculated within the QPLD workflow, applying the semi-empirical method AM1. Furthermore, the initial docking was conducted using Glide in standard precision (SP) mode, basically default settings, and returning up to five diverse poses for each ligand. These intermediate poses were submitted to the second docking procedure, after calculating the atomic partial charges of the ligands based upon the QM/MM approach performed by the module QSite. Mulliken population analysis was performed for the calculation of ligand atomic partial charges, which is based on the proposed orientation of the ligands relative to the receptor, and, therefore, influenced by the electrostatic field of the receptor. Afterwards, the final docking poses were retrieved using Glide in SP mode. For the final ranking of the docking poses, several scoring functions were evaluated, from which the scoring from Emodel calculations achieved the best results in several runs. To theoretically validate the docking workflow, the docking poses of magnolol were compared to its native pose in the X-ray crystal structure by calculating the geometric RMSD. The highest-ranked pose had an RMSD of 0.555 Å and the mean RMSD of all 10 calculated poses was 0.554 Å, respectively. Finally, the docking poses of falcариндиол were visually inspected and analyzed within the program LigandScout 3.03 [34] (Inte:Ligand GmbH, Maria Enzersdorf, Austria, 2012; <http://www.inteligand.com>).

#### Adipogenicity of Falcариндиол

3T3-L1 preadipocytes (ATCC, USA) were propagated in DMEM supplemented with 10% calf serum. For differentiation, preadipocytes were grown to confluence (day -2) and kept for two additional days before the medium was changed to DMEM supplemented with 10% FBS, 1 µg/ml insulin, and falcариндиол (10 µM) or the full PPAR $\gamma$  agonist pioglitazone (10 µM) (day 0). The medium was renewed every two days until day 8. For an estimate of lipid accumulation, Oil Red O staining was performed. For this, cells were fixed in 10% formaldehyde for 1 h and stained with Oil Red O for 10 min. After washing off excessive dye, bound dye was solubilized with 100% isopropanol and photometrically quantified at 550 nm. For better appreciation of the adipogenic potential of the investigated PPAR $\gamma$  agonists, a subset of preadipocytes was treated according to a common standard differentiation protocol (2 days in DMEM/10% FBS/500 µM IBMX/500 nM dexamethasone and 1 µg/mL insulin (MDI), followed by 2 days in DMEM/10% FBS/1 µg/ml insulin and 6 days in DMEM/10% FBS).

#### 2-Deoxy-D-(1H<sup>3</sup>)-glucose Uptake

Confluent 3T3-L1 preadipocytes (ATCC, USA) were differentiated (day 0) into mature adipocytes for 10 days (2 days in DMEM supplemented with 10% FBS, 50 µM IBMX, 1 µg/ml insulin and 500 nM dexamethasone followed by 2 days in DMEM/10% FBS/1 µg/ml insulin and 6 days in DMEM/10% FBS), and incubated with the test compounds (10 µM) or 0.1% DMSO as vehicle control for 48 h at 37°C. Before the experiment, cells were incubated for 4 h in DMEM/0.1% BSA followed by 1 h in KRH/BSA buffer (50 mM HEPES, 136 mM NaCl, 23.5 mM KCl, 1.25 mM MgSO<sub>4</sub>, 1.25 mM CaCl<sub>2</sub> and 0.1% BSA). The cells were stimulated with insulin (500 pM) for 15 min, or not stimulated (for the “basal uptake” evaluation), and then glucose uptake was initiated by addition of 2-deoxy-D-glucose spiked with 2-deoxy-D-(1H<sup>3</sup>)-glucose (final concentrations 0.1 mM and 0.45 µCi/ml, respectively). After 10 min the reaction was stopped by three rapid washes with ice-cold PBS. The glucose uptake rate was determined by liquid scintillation counting (Perkin Elmer, Waltham, MA, USA) of cell lysates (20 mM HEPES, 1% Igepal

CA-630), and normalized to protein content assessed by the Bradford protein assay [35].

### Novelty Evaluation

The novelty evaluation of the findings was performed with PubMed (<http://www.ncbi.nlm.nih.gov/pubmed/>) and SciFinder (<https://scifinder.cas.org/>) using “falcarindiol” as keyword, resulting in 99 and 364 literature references, respectively (as of 13<sup>th</sup> of December, 2012). None of the retrieved references was related to PPAR modulation. To further evaluate the novelty of the chemical scaffold of falcarindiol as PPAR ligand, chemical similarity search was performed with SciFinder, using the query structure of falcarindiol with a Tanimoto similarity score >80%. As outcome 714 similar chemical structures, associated with 1367 literature references were retrieved. None of the obtained literature references was related to PPAR.

### Statistical Methods and Data Analysis

All statistical analyses were done with the GraphPad Prism software version 4.03. One-way analysis of variance (ANOVA) with Bonferroni post hoc test, or two-tailed paired t-test was used to determine statistical significance. Nonlinear regression (sigmoidal dose response) was used to calculate the EC<sub>50</sub> values and maximal fold activation.  $K_i$  values of the investigated compounds in the receptor binding assay were calculated with the Cheng-Prusoff equation:  $(K_i) = IC_{50} / (1 + L / K_D)$ , where IC<sub>50</sub> is the concentration of competitor that displaces 50% of the ligand, L is the concentration of Fluormone™ Pan-PPAR Green used in the assay (5 nM), and K<sub>D</sub> is the binding constant (2.8 nM) of Fluormone™ Pan-PPAR Green to PPAR $\gamma$  LBD [36].

## Results

### Identification of Polyacetylenes from *Notopterygium Incisum* as Selective Partial PPAR $\gamma$ Agonists

Aiming to identify novel PPAR $\gamma$  ligands from natural sources with an ethnopharmacological background, we used a PPAR $\gamma$ -driven luciferase reporter gene assay (Fig. 1A) and observed a significant PPAR $\gamma$  activation exerted by the DCM (2.16±0.28-fold activation, p<0.001) and MeOH (1.68±0.49-fold activation, p<0.05) extracts from *N. incisum* roots and rhizomes applied at 10  $\mu$ g/ml (Fig. 1A). A bioactivity-guided approach using the DCM extract led to the isolation of six active polyacetylenes: 8-acetoxyfalcarinol (**1**), falcarindiol (**2**), 9-epoxy-falcarindiol (**3**), crithmumdiol (**4**), 9Z-heptadecene-4,6-diyne-1-ol (**5**), and 2Z,9Z-heptadecadiene-4,6-diyne-1-ol (**6**) (Fig. 1B). All polyacetylenes activated PPAR $\gamma$  as partial agonists with maximal activation ranging from 1.73- to 3.26-fold compared to the blank treatment control, while the full agonist pioglitazone activating PPAR $\gamma$  up to 9.28-fold. The observed activating effect was specific for PPAR $\gamma$  with no activation of PPAR $\alpha$  or PPAR $\beta/\delta$  (Table 1). Falcarindiol (**2**) is one of the most abundant polyacetylenes that we identified in *N. incisum*. Furthermore, it is one of the best investigated polyacetylenes in general, and is broadly found in many other plant species. Since all active compounds that we identified from *N. incisum* were polyacetylenes, and since PPAR $\gamma$  activating properties of this class of compounds have not been reported so far (see also “Novelty evaluation” in the “Materials and Methods” section), we have chosen the prototypical polyacetylene falcarindiol for further pharmacological characterization.

### Transactivation Activity of Falcarindiol in the Luciferase Reporter Model

As a partial PPAR $\gamma$  agonist, falcarindiol transactivated the luciferase reporter with an EC<sub>50</sub> of 3.29  $\mu$ M (95% CI: 2.33–4.64  $\mu$ M) inducing a maximum of 3.26 (95% CI: 2.90–3.61)-fold activation, whereas the full PPAR $\gamma$  agonist pioglitazone displayed an EC<sub>50</sub> of 0.41  $\mu$ M (95% CI: 0.25–0.67  $\mu$ M) and a maximum of 9.28 (95% CI: 8.08–10.48)-fold activation (Fig. 2A). The stimulating effect of 10  $\mu$ M falcarindiol was blocked upon co-treatment with the PPAR $\gamma$  antagonist T0070907 confirming the PPAR $\gamma$  dependence of the observed activation (Fig. 2B). Noteworthy, the antagonistic effect of T007090907 was more pronounced when the receptor was activated with pioglitazone than with falcarindiol, an observation which is in agreement with previous studies demonstrating that the antagonistic potency of T007090907 is dependent on the type of ligand used to activate PPAR $\gamma$  [37]. Furthermore, co-treatment with falcarindiol blocked up to 70–80% of the stimulating effect of pioglitazone (1  $\mu$ M) with an IC<sub>50</sub> of 18.14  $\mu$ M (95% CI: 14.32 to 22.97  $\mu$ M), suggesting competition of the two compounds for the same binding site of the receptor (Fig. 2C).

### Receptor Binding and Molecular Modeling

To further demonstrate the direct receptor binding of falcarindiol we performed a TR-FRET competitive binding assay with purified human PPAR $\gamma$ -LBD (Fig. 2D). In this model falcarindiol and pioglitazone displayed  $K_i$  values of 3.07  $\mu$ M and 0.27  $\mu$ M, respectively. Noteworthy, there was a very good correlation between the EC<sub>50</sub> values obtained with falcarindiol and pioglitazone in the cell based luciferase reporter model (3.29  $\mu$ M and 0.41  $\mu$ M, respectively) and the  $K_i$  values obtained in the receptor binding assay with purified PPAR $\gamma$ -LBD (3.07  $\mu$ M and 0.27  $\mu$ M, respectively).

In order to gain insight into the molecular interaction of the tested polyacetylene-type PPAR $\gamma$  agonists with the receptor binding pocket of the protein, docking studies were performed (Fig. 3). The composition and overall structure of the PPAR $\gamma$  LBD have been analyzed previously, describing the PPAR $\gamma$  LBD with a Y-shaped topology [38]. The entrance includes several polar residues (e.g., Arg288, Glu291, Glu295, and Glu343), while the remaining two branches of the binding pocket, arm I and arm II, are composed mostly of hydrophobic residues, except a few moderately polar residues in arm I (e.g., Cys285, Ser289, His323, His449, and Tyr473) [38]. Consistent with the relatively similar structure (Fig. 1B) and bioactivity (Table 1) of the investigated polyacetylenes, all compounds (**1–6**) docked to the PPAR $\gamma$  LBD in a quite similar manner as depicted for falcarindiol (Fig. 3A and 3B). The molecular binding mode of the falcarindiol-type polyacetylene PPAR $\gamma$  partial agonists is proposed to consist of hydrophobic contacts of the ligand alkyl chains with residues of arm II (e.g., Ile281) and arm I (e.g., Ile326, Tyr327, Phe363) as well as residues of the entrance region (e.g., Ala292, Met329, Leu330, and Leu333). The hydroxyl groups form hydrogen bonds with the backbone amide of Cys285 in arm I, and the carboxylate group of Glu295 in the entrance (Fig. 3A and 3B).

### Effectiveness of Falcarindiol in Cell Models with Endogenous Expression of PPAR $\gamma$

A main target site involved in the insulin-sensitizing action of the clinically used PPAR $\gamma$  agonists is the fat tissue [39,40]. To further confirm the effectiveness of falcarindiol in a functionally relevant cell model with endogenous expression of PPAR $\gamma$ , we examined the effect of this compound on 3T3-L1 adipocyte



differentiation and glucose uptake. As evident by the accumulation of lipid droplets and subsequent Oil Red O staining, the treatment with 10  $\mu$ M falcariindiol induced the differentiation to adipocytes as expected, although to a smaller extent than pioglitazone at the same concentration (Fig. 4A). Furthermore, falcariindiol significantly enhanced the insulin-mediated glucose uptake in differentiated 3T3-L1 adipocytes (Fig. 4B).

## Discussion

In this study we identify polyacetylenes from *N. incisum* as representatives of a novel group of natural products activating PPAR $\gamma$  as partial agonists, and characterize in more detail the mode of action of the prototypical polyacetylene falcariindiol.

C(17)-polyacetylenes of the falcariindiol-type are found in many plants of the Apiaceae family, including commonly used vegetables and seasonings such as carrots, celery, fennel, dill, parsley, and parsnip [41,42]. A variety of bioactivities have been reported so far for this compound class which may contribute to the health promoting effects of certain edible plants as well as medicinal plants used in traditional medicine [43]. The possibility that this compound class may modulate the activity of PPAR $\gamma$ , however, was so far not investigated. Since PPAR $\gamma$  is a key regulator of a number of pathways related to lipid and carbohydrate metabolism, this nuclear receptor represents an important pharmacological target for the treatment of diabetes type 2 and the metabolic syndrome. Furthermore, PPAR $\gamma$  appears to be a suitable target for modulation by diet or food supplements, since PPAR $\gamma$  activity was reported to be enhanced by a number of common food constituents or natural products present in certain seasonings or medicinal plants [44–46]. All polyacetylenes from *N. incisum* that were investigated in this study displayed a partial PPAR $\gamma$  agonism with a maximal activation several folds lower than the activation induced by the full PPAR $\gamma$  agonist pioglitazone (Table 1, Fig. 1B, and Fig. 2A). Docking studies (Fig. 3A and 3B) suggest receptor binding in a manner similar to other partial PPAR $\gamma$  agonists, which have been co-crystallized with the receptor, indicating that a hydrogen-bonding network formed by full agonists to residues of arm I is not formed in a comparable manner or even lacking in

case of partial agonists [47,48]. The partial agonism makes the polyacetylenes an interesting class of novel PPAR $\gamma$  ligands, since it is currently considered that the development of new partial agonists that do not activate PPAR $\gamma$  as strongly as the glitazone-type of full agonists might be a way to retain the needed pharmaceutical effectiveness with reduced side effects [49,50]. Furthermore, the investigated polyacetylenes selectively activated PPAR $\gamma$  without activating PPAR $\alpha$  or PPAR $\beta/\delta$  (Table 1). This is another rather favorable profile of action since the PPAR $\gamma$  agonists currently approved on the market are isoform-specific PPAR $\gamma$  activators, although there is some research evidence suggesting possible advantages for PPAR dual agonists or pan-agonists [50]. Very interestingly, previous studies have reported an antidiabetic activity of polyacetylenes isolated from *Bidens pilosa* upon oral administration in mice [51], possibly indicating effective modulation of PPAR $\gamma$  activity *in vivo*.

In summary, we report the identification of polyacetylenes as novel class of natural products activating PPAR $\gamma$ . The observed partial activation mode makes these compounds highly interesting PPAR $\gamma$  ligands, having the potential to be further explored as pharmaceutical leads or dietary supplements.

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## Author Contributions

Conceived and designed the experiments: AGA MB NF XL SMN CM MPK AC OK AS EHH DS VMD RB. Performed the experiments: AGA MB NF XL SMN CM MPK AC OK AS EHH. Analyzed the data: AGA MB NF XL SMN CM MPK OK AS EHH DS VMD RB. Wrote the paper: AGA MB NF XL SMN.

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