Short Communication

Adiponectin differentially affects gene expression in human mammary epithelial and breast cancer cells

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Adipose tissue acts as an endocrine organ secreting a variety of proteins into the blood known as adipocytokines (Matsuzawa et al., 1999). In contrast to other adipocytokines, serum adiponectin concentration is inversely correlated with obesity (Abbasi et al., 2004). Low serum adiponectin levels were shown to serve as a significant risk factor for breast cancer, particularly in postmenopausal women (Miyoshi et al., 2003).

In this study, we examined the molecular mechanisms underlying the antitumoural action of adiponectin. For this purpose, we studied adiponectin-triggered changes in the oestrogen receptor expression and attempted to identify other genes regulated by this adipocytokine in MCF-7 breast cancer cells and in MCF-10A-immortalised human mammary epithelial cells.

MATERIALS AND METHODS

Materials
Full-length low-weight adiponectin expressed in insect H5 cells as recently described (Neumeier et al., 2006) was kindly provided by Christa Büchler, Department of Internal Medicine I, University of Regensburg. Serum replacement 2 (SR2) was obtained from Sigma (Deisenhofen, Germany). RNeasy Mini Kit, RNase Free DNase Set and Quantitect SYBR Green PCR Kit were obtained from Qiagen (Hilden, Germany).

Cell culture and proliferation assay
MCF-7 (HTB-22) breast cancer cells and MCF-10A-immortalised breast epithelial cells (ATCC, Manassas, VA, USA) were maintained in DMEM/F12 medium containing 10% fetal calf serum (FCS), supplemented with 0.01 mg ml−1 insulin and 1 mm sodium pyruvate (MCF-7) or with 1 mm epidermal growth factor (EGF) (MCF-10A) and cultured with 5% CO2 at 37°C in a humidified incubator. For analysis of adiponectin effects on cell growth, both cell lines were serum starved and cultured in DMEM/F12 containing 1 × SR2. Cells were seeded in 96-well plates in triplicate (2500 cells per well) and were treated with 10 μg ml−1 full-length adiponectin. After 0, 3, 4, 5 and 6 days, relative numbers of viable cells were measured in comparison with the untreated control using the fluorimetrical, resazurin-based Cell Titer Blue assay (Promega, Mannheim, Germany) according to the manufacturer’s instructions at 560nm/590nm in a Victor3 multilabel counter (PerkinElmer, Massachusetts, USA). Cell growth was expressed as a percentage of the untreated medium control. Statistical analysis of the data was performed by one-way ANOVA using Prism 2.0 Software (GraphPad, San Diego, CA, USA), with statistical significance accepted at P<0.05.

RNA preparation and hybridisation of DNA microarrays
Total RNA was isolated by means of the RNeasy kit (Qiagen) according to the manufacturer’s instructions. After DNase treatment and subsequent precipitation, 20 μg of total RNA was used for cDNA synthesis using a cDNA kit (Roche, Mannheim, Germany). After purification of the cDNA, Cy3- and Cy5-labelled cRNA were generated using the MEGAscript™ T7 Kit (Ambion, Woodward, USA). After purification of the labelled cRNA using the High Pure RNA Tissue Kit (Roche) and its fragmentation, 10 μg of Cy3- and Cy5-labelled cRNA were used for the hybridisation of the human 10k-A array (MWG Biotech, Ebersberg, Germany) containing oligonucleotides that represent approximately 10 000 human genes with known functions. The measurement of the slides was performed by the scanning service of MWG Biotech. Data are presented as a relationship between gene expression in untreated MCF-7 or MCF-10A cells (defined as 100%) and the respective expression 48 h after treatment with 10 μg ml−1 adiponectin.

Reverse transcription and real-time PCR
From 1 μg of total RNA, cDNA was synthesised using 100 U M-MLV-P reverse transcriptase (Promega), 2.5 mm dNTP mixture...
and 50 pm random primers (Invitrogen, Karlsruhe, Germany). All PCR primers were designed intron-spanning to avoid amplification of genomic DNA; primer sequences are indicated in Table 1. For real-time PCR analysis, 4 μl of 1:5 diluted cDNA was amplified using the Quantitect SYBR Green PCR Kit (Qiagen) and the LightCycler PCR device (Roche Diagnostics, Mannheim, Germany). The PCR programme was 95°C for 10 min, followed by 40 PCR cycles (95°C for 10 s, 58°C for 10 s, 72°C for 15 s) and a final extension for 5 min at 72°C, followed by a standard melting curve analysis. In all RT–PCR experiments, a 200 bp β-actin fragment was amplified as a reference gene using intron-spanning primers. After testing that the efficiencies of all primer pairs were approximately equal (Stahlberg et al., 2003), data were analysed using the comparative ΔΔCt method (Livak and Schmittgen, 2001) by calculating the difference between the threshold cycle (Ct) values of the target and reference gene of each sample and then comparing the resulting ΔCt values among different samples.

RESULTS
As a pre-requirement for this study, we first tested the expression of adiponectin receptors (AdipoR) 1 and 2 in the cell lines used. By means of real-time RT–PCR, we detected notable amounts of both transcripts in MCF-7 and MCF-10A cells (Figure 1). Receptor mRNA levels were in a similar range in both cell lines (data not shown).

The effect of a physiological concentration of adiponectin on growth of MCF-7 and MCF-10A cells was tested in serum-free, defined culture medium and assessed over a cultivation period of 6 days. Adiponectin significantly inhibited the growth of MCF-10A mammary epithelial cells, but not of MCF-7 breast cancer cells. In relation to the untreated control, this effect was significant after 4 or more days, with a maximum growth reduction by 30% evident after 5 days (Figure 2).

A total of six 10k DNA microarrays (MWG, Martinsried, Germany) were used to analyse adiponectin effects on gene expression. Total RNA was isolated from MCF-7 and MCF-10A cells cultured in serum-free, defined culture medium treated with adiponectin (10 μg ml⁻¹) for 48 h. Microarray data confirmed by 2-fold replication of the experiment including dye swap revealed a more than 5-fold increase of transcript levels of caspase 1 (CASP1) (5.94-fold), TR2 orphan nuclear hormone receptor (6.31-fold) and ubiquitin-specific peptidase 2 (USP2) (5.37-fold) in MCF-10A cells. These effects were not observed in microarray experiments comparing the transcriptome of MCF-7 cells before and after adiponectin treatment (Table 2).

More than 5-fold expression changes after adiponectin treatment observed in the microarray experiments in case of the genes CASP1, TR2 and USP2 were now confirmed by means of RT–PCR.

Figure 1 Expression of adiponectin receptors (AdipoR) 1 and 2 in MCF-7 and MCF-10A cells detected by means of RT–PCR. After real-time RT–PCR analysis (35 cycles, LightCycler, Roche), PCR products were subjected to 1.5% agarose gel electrophoresis.

Figure 2 The effect of adiponectin (10 μg ml⁻¹) on growth of MCF-7 and MCF-10A cells over a cultivation time of 6 days. Data are expressed in percentage of the untreated control (n = 3). *P < 0.05 vs untreated control.

Table 1 Sequences of the PCR primers used in this study

| Primer | Sequence                        |
|--------|---------------------------------|
| AdipoR1| 5'-GGGGAATTCTCCTCCACAAGGATCGTG-3' |
|        | 5'-GGGGTGAGTGAAGTGTTCATGTGAG-3'   |
| AdipoR2| 5'-GGGGATCTTACAGGTCAGAAAATTGCT-3' |
|        | 5'-GGGGTTGACGTCATCTTGCTTGTGG-3'   |
| ERα    | 5'-TCATGAAAGGTGGGATTGGATC-3'      |
|        | 5'-AAGGGTGGCAGTCCTGTCAT-3'        |
| ERβ1   | 5'-TTTGGTTAGTGGCGAGAGG-3'         |
|        | 5'-AGGACTGGTGCCATTACG-3'          |
| ERβ2   | 5'-CCCAAGGGGAGAACACTGAAAG-3'      |
|        | 5'-AACTTCTCCTAGTCTCCTGCTG-3'      |
| ERβ5   | 5'-CACTTTTCCAAAATACCTCACTCACC-3' |
|        | 5'-TTTGGATGTTGCGAGAAGG-3'         |
| USP2   | 5'-CATCCAGGAGTTCCTCCAACAGA-3'     |
|        | 5'-CCAGGCTCTTACTAGGGAAGAAG-3'     |
| CASP1  | 5'-TCTTTTCTCTCCACGAGCCTGTT-3'     |
|        | 5'-CTGCAGGATTTCTTGTCCATAC-3'      |
| TR2    | 5'-CTACACACCGCTGAGCCTCTCTA-3'     |
|        | 5'-TGCAGCTCTGATGTTGGTTCG-3'       |
| β-actin| 5'-CTGGGGATCACAGGAAAACA-3'        |
|        | 5'-CGCTCAAGGAGGCAATG-3'           |

Table 2 Gene expression changes detected by means of microarray analysis (10k-A array, MWG Biotech, Martinsried, Germany) in MCF-7 breast cancer cells and MCF-10A mammary epithelial cells after treatment with adiponectin (NS = not significant) (n = 3 including dye swap)

| GENE | MCF-10A Factor | MCF-7 Factor |
|------|----------------|--------------|
| TR2  | 6.31           | NS           |
| CASP1| 5.94           | NS           |
| USP2 | 5.37           | NS           |
| ZNF235| 4.49         | NS           |
| SERPINB13| 3.67       | NS           |
| MGST2| 3.40           | NS           |
| RNF13| 3.28           | NS           |
| SLC25A20| 0.22         | NS           |
| DDX59| NS             | 4.17         |
| HIP5P| NS             | 3.64         |
| NRG1| NS             | 3.22         |
| PSCD3| NS             | 0.27         |

Microarray analysis based on ANOVA was performed and the cut off value was a 3-fold difference.
Increased levels of these genes in adiponectin-treated MCF-10A cells were also detected by means of real-time RT–PCR, although this increase was only 2.5-fold in case of TR2 and USP2 (Figure 3). Given that our recent results suggested interactions between oestrogen receptor (ER) and adiponectin signalling (Pfeiler et al., 2008), adiponectin effects on expression of ERα and ERβ-isoforms were also examined in these RT–PCR experiments. In MCF-10A cells, we observed a more than 5-fold increase of ER/β and an approximately 2-fold increase of ER/β mRNA levels after adiponectin treatment. As expected from our microarray data, adiponectin did not trigger elevated transcript levels of TR2, CASP1 and USP2 in MCF-7 breast cancer cells, but a significantly reduced USP2 expression by approximately 60%. Similar to the mammary epithelial cell line, in MCF-7 cells ER/β mRNA levels were also upregulated by more than 5-fold after adiponectin treatment.

**DISCUSSION**

Adiponectin levels are inversely related to the prognostic factors of breast cancer (Miyoshi et al., 2003; Mantzoros et al., 2004). Furthermore, direct antitumoural effects of adiponectin such as inhibition of proliferation have been reported from in vitro studies (Kang et al., 2005, 2007; Nakayama et al., 2007). These observations suggested that adiponectin might be able to protect from breast cancer development. In this study, we examined the effects of adiponectin on proliferation and gene expression of immortalised mammary epithelial cells in comparison with breast cancer cells.

Recently, expression of both AdipoR was reported to be present in various benign mammary epithelial and breast cancer cell lines (Takahata et al., 2007). In this study, mRNA levels of AdipoR1 and AdipoR2 were shown to be expressed in a similar range in MCF-7 and MCF-10A cells. However, treatment with adiponectin exerted different effects on these cell lines. Physiological concentrations of this adipocytokine significantly slowed down the proliferation of MCF-10A mammary epithelial cells, but not of MCF-7 breast cancer cells. The results of earlier studies examining the effect of adiponectin on MCF-7 breast cancer cells are contradictory. Our group and others had earlier reported the absence of adiponectin effects on the growth of MCF-7 breast cancer cells (Kang et al., 2005; Pfeiler et al., 2008). Although some studies reported small apoptotic effects of adiponectin on this cell line not significantly affecting the total cell numbers (Arditi et al., 2007; Grossmann et al., 2008), in another study, growth-inhibitory effects on MCF-7 cells were observed (Dieudonne et al., 2006). However, this inconsistency might result from different experimental conditions used in the studies mentioned. In an earlier study, an antiproliferative effect of adiponectin on MCF-7 cells was not present under serum-free culture conditions as used in this study and was suggested to be dependent on the presence of steroid hormones such as 17-β estradiol (Pfeiler et al., 2008).

To our knowledge, this is the first study demonstrating a significant growth inhibitory effect of adiponectin on mammary epithelial cells. This observation might reveal at least one mechanism underlying the association between high adiponectin serum levels and reduced breast cancer risk. It is tempting to speculate that adiponectin might be able to exert protective effects on mammary epithelial cells. The lack of any antiproliferative effect on MCF-7 breast cancer cells suggests that adiponectin-triggered growth control might be lost during carcinogenesis. Given that MCF-7 cells expressed AdipoR to a similar extent, the molecular mechanisms of this different response remained to be elucidated.

Given that earlier data of our group suggested the presence of a cross talk between adiponectin and oestrogen signalling systems (Pfeiler et al., 2008), we first analysed the effect of an adiponectin treatment on ERα and ERβ expression. In MCF-10A cells, adiponectin triggered a significant increase of ER/β and ERβ mRNA levels. ERβ is the dominant oestrogen receptor in normal breast tissue (Speirs et al., 2000, 2002; Shaw et al., 2002), but its expression declines during tumorigenesis (Leygue et al., 1998; Roger et al., 2001). This study and a variety of in vitro studies of our group and others suggested a role for ERβ as a tumour suppressor in breast tissue (Skliris et al., 2003; Treeck et al., 2008). ERβ2 and ERβ5 proteins both are known to form heterodimers with ERz, thereby inhibiting the transcriptional activity of ERz (Peng et al., 2003). As ERz is known to mediate tumour-promoting effects of 17-β estradiol, our data suggest that adiponectin-triggered increase of ERβ isoform expression might be one important molecular mechanism underlying the protective effects of this adipocytokine. This seems to be particularly true for ERβ5 splice variant, as the expression of this isoform was exclusively increased in MCF-10A cells.

Then, we studied the gene expression changes triggered by adiponectin in MCF-7 and MCF-10A cells by means of DNA microarrays. After treatment with adiponectin, in MCF-10A mammary epithelial cells, we observed elevated transcript levels of several genes known to exert antitumoural effects in breast tissue.

Adiponectin treatment led to increased transcript levels of TR2 orphan nuclear hormone receptor in MCF-10A, but not in MCF-7 cells. Recent results of in vitro studies of cultured cells ectopically expressing TR2 suggested potential roles for TR2 in a variety of...
cellular processes (Park et al., 2007). Interestingly, the transcriptional activity of ERα, leading to enhanced proliferation of breast epithelial cells and to tumour progression, can be inhibited not only by ERβ, but also by TR2. Furthermore, TR2-siRNA was reported to enhance ERα-dependent proliferation of MCF-7 cells (Hu et al., 2002). Our data suggest that adiponectin-triggered increase of TR2 expression could also lead to the suppression of ERα-mediated proliferation in mammary epithelial cells, thereby exerting protective, antitumoural effects.

TR2 has also been reported to be upregulated during apoptotic processes and its overexpression resulted in an increased apoptosis rate (Lee and Wei, 2000). It is tempting to speculate that adiponectin-triggered increase of TR2 might be able to confer an increased apoptotic potential to mammary epithelial cells, thereby also exerting protective effects.

In MCF-10A mammary epithelial cells treated with adiponectin, we also found a significant increase of USP2 mRNA levels, but a decrease in MCF-7 breast cancer cells. In breast cancer, an enhanced activity of the ubiquitin–proteasome system has been reported (Deng et al., 2007). However, the exact role of USP in breast tissue remains unknown. Ubiquitination may be a candidate for either a tumour-suppressive or an oncogenic activity (Fujisawa et al., 1998).

Adiponectin treatment of MCF-10A cells also resulted in elevated transcript levels of CASP1, a key enzyme of apoptosis. This effect again was not observed in MCF-7 breast cancer cells. Our data suggest that adiponectin-triggered increase of CASP1 expression might be another molecular mechanism underlying the antitumoural effect of this adipocytokine, which is present in mammary epithelial cells, but has been lost in MCF-7 breast cancer cells. Adiponectin-triggered activation of caspase expression has also been reported from endothelial cells, resulting in decreased angiogenesis (Brakenhielm et al., 2004). Given that decreased neovascularisation exerts antitumoural effects, the antiangiogenic effect of this adipocytokine mediated by CASP1 upregulation might be another mechanism underlying the inverse association between adiponectin serum levels and breast cancer risk.

In this study, we observed an adiponectin-triggered growth inhibition of MCF-10A cells accompanied by specific upregulation of genes with known antitumoural properties. Our data suggest that adiponectin might be able to exert protective effects on mammary epithelial cells only, whereas MCF-7 breast cancer cells might have lost the ability to respond to this adipocytokine in a similar manner.

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