RESEARCH

TAAR1 induces a disturbed GSK3 β phosphorylation in recurrent miscarriages through the ODC

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Abstract

Objectives: Thyroid hormones play an important role in the maintenance of pregnancy. Their derivatives, endogenous amines, act via binding to the trace amine-associated receptor (TAAR1). The aim of our study was to analyse the regulation of TAAR1, serine/threonine kinase (pGSK3 β) and ornithine decarboxylase (ODC) in placentas of healthy pregnancies, spontaneous (SM) and recurrent miscarriages (RM) and to investigate the influence of thyroid hormone derivatives on TAAR1 expression in trophoblast model cells *in vitro*.

Methods: Patients with SM ($n=15$) and RM ($n=15$) were compared with patients with healthy pregnancies ($n=15$) (pregnancy weeks 7–13 each). Immunohistochemistry was applied to analyse placental TAAR1, pGSK3 β and ODC expression. Protein expression of the receptors after stimulation with T₃, T₁AM and RO5203548 in BeWo trophoblast model cells was determined via Western blot. Double-immunofluorescence was used to determine placental expression of TAAR1 and ODC.

Results: Levels of TAAR1, pGSK3 β and ODC were higher in placentas of RM in comparison to healthy controls. Stimulation of BeWo cells with T₃, T₁AM and RO5203548 significantly increased TAAR1 expression. ODC expression in BeWo cells was upregulated through T₃. Via double-immunofluorescence, TAAR1 and ODC-positive EVT could be detected.

Conclusions: Upregulation of placental TAAR1 may indicate an increased decarboxylation of thyroid hormones in miscarriages. Patients with RM may have a lack of T₃ through an enhanced transformation of T₃ into T₁AM induced by the ODC. Future investigations could be carried out to analyse what role a prophylactic T₃ substitution plays for patients.

Key Words

- ▶ TAAR1
- ▶ pGSK3 β
- ▶ ODC
- ▶ 3-iodothyronamine
- ▶ RO5203548
- ▶ spontaneous miscarriages
- ▶ recurrent miscarriages

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Introduction

Thyroid disorders such as hypothyroidism and hyperthyroidism are associated with increased miscarriage and stillbirth rates (1). Untreated hypothyroidism can result in a twofold increase of miscarriages; therefore, the maintenance of an euthyroid state is crucial for a healthy

pregnancy (2). Thyroid hormones, which bind to the thyroid hormone receptor (THR), play an important role during trophoblast differentiation and the development of the foetal nervous system (3). THR belongs to the nuclear receptor superfamily, and it is involved in various

physiological processes (4). In a former study, we were able to show that mRNA and protein expression of the THR isoforms THR α 1, THR α 2, THR β 1 and THR β 2 are reduced in spontaneous and recurrent miscarriages (RM) (5). The majority of THR seems to be expressed by decidual stromal cells (5).

Degradation of thyroid hormones results in the formation of trace amines (TAs), also referred to as thyronamines or endogenous amines (6). Transformation of thyroid hormones into TAs is conducted by the enzyme ornithine decarboxylase (ODC) through a deiodisation and a decarboxylation process (7). Thyronamines (3-T₁AM, T₀AM) are endogenous compounds, which can possibly be derived from L-thyroxine (T₄) (7). Decarboxylated thyroid hormones act through rapid actions like rapid lowering of body temperature and heart rate, while classical thyroid hormones act epigenetically (8). Ornithine decarboxylase (ODC) is a rate-limiting enzyme for the biosynthesis of polyamines, and it is responsible for the decarboxylation of thyroid hormones (9, 10). Changes in the brain ODC/polyamine system can have an influence on the development of catecholaminergic nerve pathways (11). Bulgaroni and colleagues found that ODC upregulation may be linked to cytotrophoblast proliferation and placental repair in compensation to injuries induced by organophosphate pesticides (12).

Thyroid hormone derivatives function by binding to the G-protein-coupled receptor trace amine-associated receptor 1 (TAAR1) (8). TAAR1 was first described in 2001 by Borowsky and coworkers (13). Upon activation of adenylyl cyclase, TAAR1 induces a rise of the intracellular cAMP level (13, 14). TAAR1 is expressed by the placenta, brain, spinal cord, stomach, pancreatic β -cells and immune cells like macrophages, leukocytes and dendritic cells (15, 16, 17). Different compounds, including dopaminergic, adrenergic and serotonergic classes and a wide spectrum of agonists, such as trace amines, common biogenic amines, amphetamine-like stimulants and 3-iodothyronamine can activate TAAR1 (18, 19, 20). The only known TAAR1 antagonist to date is RO5212773 (EPPTB) (21).

TAAR1 has so far been investigated in the field of neuroscience whereby it could be shown that TAAR1 is able to modulate the serotonergic and dopaminergic system in the brain (15). As TAAR1 is a negative regulator of dopamine transmission, TAAR1 and its agonists as well as antagonists represent a target for psychiatric disorders like Parkinson's disease, depression and schizophrenia (15). Stimulation of TAAR1 can furthermore have a beneficial effect on glucose and obesity control in mice with type 2 diabetes mellitus (22).

In a recent study, we analysed the regulation of TAAR1 in breast cancer tissue of primary breast cancer patients and identified that TAAR1 seems to be an independent predictor for breast cancer survival (23). TAAR1 overexpression (IRS $\geq$ 6) was associated with a significantly longer overall survival (OS) of breast cancer patients ($P=0.02$) than reduced TAAR1 expression (IRS < 6) during a maximum follow-up of 14 years, demonstrating that TAAR1 has a favourable effect on OS of early breast cancer patients (23).

It is speculated that TAAR1 directly or indirectly interferes with the β -arrestin2/Akt/GSK3 pathway (24, 25). GSK3 is a serine/threonine protein kinase, and it is encoded by the two genes GSK3A and GSK3B (26). Besides cellular proliferation, migration, glucose regulation and apoptosis, the isoform GSK3 β is a key enzyme in the wnt-signalling pathways (27). Those are a group of signal transduction pathways that play an important role in carcinogenesis and embryo development. The latter is controlled through cell proliferation and cell migration, and wnt pathways regulate pre-implantation, blastocyst implantation and uterine decidualisation (27). Wnt-signalling proteins participate in numerous developmental events during embryogenesis and differentiation of diverse reproductive tissues (28, 29).

A crosstalk between TAAR1 and dopamine D2L receptor (D2R) has formerly been described (6, 30, 31). Harmeier and colleagues found that the activation of the TAAR1-D2R complex negatively modulates GSK3 β signalling *in vitro* and *in vivo*. D2R is a target receptor for various antipsychotic drugs, and TAAR1 can interact with D2R forming a functional heteromer (6, 32).

Aim of our study was to investigate the placental expression of TAAR1, GSK3 β and ODC in healthy pregnancies and spontaneous as well as RM and to further analyse the crosstalk between these receptors. Additionally, the influence of TAAR1 ligands on the expression of TAAR1 in trophoblast model cells *in vitro* was assessed. As we assume that the processing of thyroid hormones through the ODC plays an important role during miscarriages, regulation of the ODC was a further component of the study.

Materials and methods

Tissue samples

Samples collected from spontaneous (SM, 15 cases) and RM, 15 cases) in the time period between 2011 and 2012 were obtained from the archives of the Gynaecology and Obstetrics Department of the Ludwig-Maximilians-

University Munich, Germany. Samples from elective terminations of pregnancy (ETP, 15 cases) collected in the same time period were provided by outpatients clinic in Munich. Tissue samples were attained by curettage without any prior pharmacological induction. In cases of miscarriages, curettages were performed within 24h after diagnosis. The average gestational age for ETP was 10.6 ± 2.1 , for SM 10.1 ± 1.5 weeks and for RM 10 ± 1.5 weeks. Hypothyroidism as well as hyperthyroidism and TPO antibody positivity was excluded in all women. History taking was comprehensive, aiming to exclude—apart from common disorders—potential involvement of clotting disorders and autoimmune diseases, which are already known as aggravating factors for increased risk for miscarriages. In all samples, analysis of chromosomal abnormalities was performed. In this study, PowerPlex 16HS multiple PCR system was used, which detects 16 STR loci, with the limitation of this method in picking up chromosomal abnormalities at base pair level. Analyses were carried out for all samples including controls. DNA extraction and profiling were performed as described by Ziegelmüller and coworkers (33). The PCR products were also analysed as described (33). Additionally, in all samples, microbiology analysis excluded possible intra-uterine infection. All women had a normal first trimester vaginal swab.

Ethical approval and informed consent

All procedures that involved human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the Helsinki declaration of 1964 and its later amendments or comparable ethical standards. The study was approved by the Local Ethics Committee of the Ludwig-Maximilians University of Munich, Germany, and was assigned the reference number 337-06. Consent has been obtained from each patient or subject after full explanation of the purpose and nature of all procedures used.

Immunohistochemistry

After tissue preparation, tissue samples were fixed in 3.7% formalin and then embedded in paraffin. Formalin-fixed paraffin-embedded sections ($3 \mu\text{m}$) were deparaffinised in Roticlear (Carl Roth, Karlsruhe, Germany), blocked and rehydrated with 3% H_2O_2 /methanol for 20 min. Rehydration of the slides was carried out by using an alcohol gradient up to distilled water. The slides were placed in a pressure cooker containing sodium citrate ($\text{pH}=6.0$). A blocking solution was used (Polymer Kit, Zytomed Systems, Berlin, Germany) to prevent any non-specific binding of the primary antibodies. Tissue sections were then incubated with the primary antibodies for 16h at 4°C (Table 1). Reactivity was detected by using the ZytoChem Plus HRP Polymer System (Mouse/Rabbit) (Zytomed Systems; catalogue-ID: POLHRP-100), in accordance with the manufacturer's protocol. Staining was performed with DAB and counterstaining was performed with hemalaun. Appropriate positive controls (brain tissue for GSK3- β and breast cancer tissue (23) and intestinal tissue for TAAR1 (34)) and negative controls (negative control serum added on third trimester placenta) (Negative Control for Super Sensitive Rabbit Antibodies, Rabbit IgG, BioGenex, Fremont, USA) were included in each experiment (Supplementary Fig. 1, see section on supplementary data given at the end of this article). Positive and negative control experiments of TAAR1 expression are shown in the Supplementary figure. Per slide, ten fields were examined by two independent observers with a Leitz Diaplan microscope (Leitz, Wetzlar, Germany) by applying the semi-quantitative immunoreactive score (IRS) (35). The IRS is calculated by multiplying the intensity of cell staining (0: none; 1: weak; 2: moderate; 3: strong) with the percentage of positively stained cells (0: no staining; 1: $<10\%$ of the cells; 2: $11\text{--}50\%$; 3: $51\text{--}80\%$; 4: $>80\%$).

The pattern of the immunochemical staining reaction was evaluated by two independent blinded observers including a gynaecological pathologist (D M). In one

Table 1 Features of the antibodies used for staining.

Antibody	Incubation	Blocking solution	Blocking condition (min)
TAAR1, polyclonal (Rabbit IgG, Abcam)	1:800 in PBS 16 h at 4°C	Reagent I (Polymer Kit, Zytomed Systems, Berlin, Germany)	15
Phosphorylated GSK3 β , polyclonal (phosphoY216) (Rabbit IgG, Abcam)	1:100 in TBS 16 h at 4°C	Reagent I (Polymer Kit, Zytomed Systems, Berlin, Germany)	5
ODC, monoclonal (Mouse IgG, Novus Biologicals)	1:4000 in PBS 16 h at 4°C	Reagent I (Polymer Kit, Zytomed Systems, Berlin, Germany)	5

case, the evaluation of the two observers of the GSK β expression differed. This case was re-evaluated by both observers together. After the re-evaluation, both observers achieved the same result.

Cell lines and cell stimulation

The choriocarcinoma cell lines BeWo (ECACC, Salisbury, UK) and JEG-3 (ATCC), which are useful models of human trophoblasts, were used for the study. The cells were cultured in DMEM (3.7 g/L NaHCO₃, 4.5 g/L D-glucose, 1.028 g/L stable glutamine, and Na-Pyruvate; Biochrom, Berlin, Germany). 10% heat-inactivated FCS was added to the medium, and the solution was incubated at an atmospheric CO₂ concentration of 5% and at 37°C.

BeWo and JEG-3 cells were separately grown on sterile 12 multiwell slides at a density of 500,000 cells/mL DMEM with 10% foetal cow serum. After 4 h, medium was changed to pure DMEM. The cells were stimulated with 0.01 nM or 0.1 nM triiodothyronine (T₃) (Sigma T2877; Lot:106K1157V, Sigma-Aldrich), thyronamine (T₀AM) (Fluka 80345, Lot: BCBL2185V, Buchs, Switzerland), 3-Iodothyronamine (T₁AM) (Cayman Chemical) and RO5203548 (Glaxo Laboratories, Hopkinton, MA, USA) for 2 h (PCR samples) and 48 h (Immunocytochemistry and Western blot samples), respectively. Control cells were incubated without stimulants. As a control solvent, the following buffers were used: The stimulant T₃ has been diluted in 1 M NaOH in DMEM (Dulbecco's Modified Eagle Medium) and T₁AM and RO5203548 were diluted in DMSO (Dimethylsulfoxide).

Immunocytochemistry

After incubation, the Quadriperm slides were fixated in an ethanol/methanol solution for 15 min. Between each step, the slides were washed in PBS (Biochrom). Next, the primary antibody (anti-TAAR1) was applied for 16 h at 4°C. Incubation with the secondary antibody (ZytoChem Plus HRP Polymer System (mouse/rabbit) Tytomed: Nr. POLHRP-100) for 30 min followed. Staining of the slides was carried out with 3-amino-9-ethylcarbazole plus (AEC plus; Dako) for 5–8 min, followed by counterstaining with hemalaun for 30 s and finally washing in tap water and cover-slipping of the slides.

Assessment of the slides was carried out with a Leitz Diaplan light microscope (Leitz) and a digital camera system (JVC, Yokohoma, Japan) by two independent observers. For each slide, six visual fields were examined

at a magnification of $\times 10$ and 25. Per field, the cells were counted and assigned to three groups (negative, low, medium/strong staining).

TaqMan-real-time PCR

RNA of stimulated and control JEG-3 and BeWo cells was isolated using the NucleoSpinRNAII (Macherey-Nagel, Düren, Germany). Quantification and evaluation of RNA samples was carried out with the NanoPhotometer (Implen, Munich, Germany).

Reverse transcription was performed using the High-Capacity cDNA reverse transcription kit (Applied Biosystems) with a temperature protocol of 25°C for 10 min, 37°C for 2 h, 85°C for 5 s and hold at 4°C.

The PCR was performed in 96-well reaction plates holding a volume of 20 μ L. Each well contained 1 μ L TaqMan Gene Expression Assay 20 $\times$ (Applied Biosystems), 10 μ L TaqMan Universal PCR Master Mix 2 $\times$ (Applied Biosystems), 8 μ L H₂O (DEPC treated DI water, Sigma) and 1 μ L sample. For the PCR assay, ABI PRISM 7500 Fast (Applied Biosystems) was used with thermal cycling conditions of 20 s at 95°C, 40 cycles of amplification with 3 s at 95°C and 30 s at 60°C. For quantification of the expression, the 2^{− $\Delta\Delta$ CT} method was used by applying β -actin as the housekeeping gene. Each experiment was validated three times.

Western blot

Stimulated and control cell samples were lysed for 30 min with 200 μ L buffer solution consisting of a 1:100 dilution of protease inhibitor (Sigma-Aldrich) in RIPA buffer (Radioimmunoprecipitation assay buffer, Sigma-Aldrich). The lysate was centrifuged and a Bradford protein assay of the supernatant followed. The proteins were separated according to their molecular weight by a SDS-PAGE and transferred onto a PVDF membrane (Merck Millipore). Non-specific binding was blocked via a 1-h incubation of the membrane with 1 $\times$ Casein Solution (Vector Laboratories, Burlingame, CA, USA). The primary antibodies anti-TAAR1, anti-ODC and anti- β -actin (Clone AC-15, Mouse IgG, Sigma), diluted in a 1:500 and 1:1000 Casein solution, respectively, were applied for 16 h at 4°C. After rinsing with 1 $\times$ Casein solution, the membrane was incubated with biotinylated anti-mouse IgG antibody, respectively anti-rabbit IgG antibody and ABC-AmP reagent (both VECTASTAIN ABC-AmP Kit for mouse IgG, Vector Laboratories) according to the manufacturer's protocol. Staining was performed

with BCIP/NBT chromogenic substrate (Vectastain ABC-AmP Kit, Vector Laboratories). Detection was carried out with Bio-Rad Universal Hood II (Bio-Rad Laboratories) and the values quantified using Bio-Rad Quantity One software (Bio-Rad Laboratories).

Double-immunofluorescence

To determine the cells that express TAAR1 and ODC in the decidua, double-immunofluorescence analysis with markers for extravillous trophoblast (EVT) and decidual stroma cells was performed. HLAG, which is a marker for EVT, was applied. Placental tissue of both ETP and RM groups were stained.

After tissue retrieval, tissue samples were fixed in 3.7% formalin and embedded in paraffin. Formalin-fixed paraffin-embedded sections (3 µm) were then deparaffinised in Rotoclear (Carl Roth), rehydrated and blocked with 3% H₂O₂/methanol for 20 min for endogenous peroxidase activity. Rehydration of the slides was performed with an alcohol gradient up to distilled water. The slides were then placed in a pressure cooker containing sodium citrate (pH=6.0). To prevent any non-specific binding of the primary antibodies, a blocking solution was applied for 15 min (Ultra V-Block, Thermo Scientific, Lab Vision, Lot: PBQ140925). Afterwards, incubation was performed with the primary antibodies overnight at 4°C (Table 2). The secondary antibodies Cy-2-labelled goat-anti-rabbit IgG, diluted 1:100 (Dianova, Hamburg, Germany), Cy-3-labelled goat-anti-mouse IgG, diluted 1:500 (Dianova) and Cy-3-labelled goat-anti-rabbit IgG, diluted 1:500 (Dianova) were applied on the slides. Finally, the slides were embedded in DAPI containing mounting buffer (Vector Laboratories, Vectashield H-1200). For the identification of TAAR1/HLAG expressing cells, the selected sections were additionally incubated with monoclonal HLAG-FITC mouse IgG1 (Serotec, Bio-Rad) for 30 min, which was diluted 1:50. Afterwards, the slides were analysed with the fluorescent Axioskop photomicroscope (Zeiss). Images were taken with a digital Axiocam camera system (Zeiss).

Statistics

Data collection, processing and analysis of statistical data were performed with IBM SPSS Statistics for Windows, version 22.0. IBM Corp. All statistical tests were two-sided with a 5% significance level. Non-parametric Mann-Whitney *U* tests were performed for comparison of central tendency. Wilcoxon signed-rank tests were performed for the statistical analysis of cell culture experiments. As causes for either spontaneous or RM differ, those two groups cannot be compared with each other. For ODC immunohistochemistry, a score developed from the percentage of low, middle and highly stained decidual cells in each sample was used. The formula is as follows: score=log10 (1×percentage of low stained cells+10×percentage of middle intensity cells+100×percentage of highly stained cells).

Results

TAAR1 expression in the placenta

Via immunohistochemical analysis, a significant upregulation of TAAR1 protein expression could be detected in syncytiotrophoblasts, cytotrophoblasts, decidua and glands of RM in comparison to healthy controls (all *P*<0.001) (Fig. 1). In spontaneous miscarriages, TAAR1 protein expression was also significantly increased in syncytiotrophoblasts, cytotrophoblasts, decidua and glands (decidua and glands: *P*<0.001; syncytiotrophoblasts: *P*=0.002) (Fig. 1).

Phosphorylated GSK3β expression in the placenta

Phosphorylated GSK3β protein (Y216) expression was significantly increased in RM in comparison to GSK3β protein expression in healthy controls in syncytiotrophoblasts (*P*=0.031), cytotrophoblasts (*P*=0.033) and in the decidua (*P*<0.001). The difference between GSK3β expression in glands of RM compared to healthy controls was not significant (*P*=0.286) (Fig. 2). There was no significant difference between GSK3β protein expression of spontaneous miscarriages and

Table 2 Features of the antibodies used for double-immunofluorescence.

Primary antibodies	Antibody type	Company	Dilution
TAAR1	Mouse IgG monoclonal	Santa Cruz, sc-514311	1:300
TAAR1	Rabbit IgG polyclonal	Abcam, ab65633, Lot: GR171220-3	1:800
ODC	Mouse IgG polyclonal	Novus Biologicals, SPM565, Lot: 4953-1XP160908	1:200
HLAG	Mouse IgG1 monoclonal clone: MEM G/9	Serotec, MCA2044F, Lot: 300709	1:50

Figure 1

TAAR1 expression in control placental tissue, spontaneous miscarriages and recurrent miscarriages. (A) Boxplot and scatterplot (all jittered) of TAAR1 protein expression in syncytiotrophoblasts, (B) milder TAAR1 staining in syncytiotrophoblasts of control tissue, (C) stronger TAAR1 staining in syncytiotrophoblasts of recurrent miscarriages, (D) stronger TAAR1 staining in syncytiotrophoblasts of spontaneous miscarriages, (E) Boxplot and scatterplot of TAAR1 protein expression in decidua, (F) milder TAAR1 staining in decidua of control tissue, (G) stronger TAAR1 staining in decidua of RM, (H) stronger TAAR1 staining in decidua of SM, (I) Boxplot and scatterplot of TAAR1 protein expression in glands, (J) milder TAAR1 staining in glands of control tissue, (K) stronger TAAR1 staining in glands of RM, (L) stronger TAAR1 staining in glands of SM (all magnification $\times 25$).

Figure 2

GSK3β expression in control placental tissue, spontaneous miscarriages and recurrent miscarriages. (A) Boxplot and scatterplot (all jittered) of GSK3β protein expression in glands, (B) GSK3β staining in glands of control tissue, (C) GSK3β staining in glands of SM, (D) Boxplot and scatterplot of GSK3β protein expression in decidua, (E) milder GSK3β staining in decidua of control tissue, ×25 magnification, (F) stronger GSK3β staining in decidua of RM, ×25, (G) Boxplot and scatterplot of GSK3β protein expression in cytotrophoblasts, (H) GSK3β staining in cytotrophoblasts of control tissue, ×10, (I) GSK3β staining in cytotrophoblasts of RM, ×10, (J) Boxplot and scatterplot of GSK3β protein expression in syncytiotrophoblasts, (K) milder GSK3β staining in syncytiotrophoblasts of control tissue, ×25, (L) stronger GSK3β staining in syncytiotrophoblasts of RM, ×25.

Figure 3

ODC expression in different pregnancy weeks. ODC expression has been analysed on the basis of a logarithmic scale with the following formula: $\log_{10}(\text{intensity } 1 + 10 \times \text{intensity } 2 + 100 \times \text{intensity } 3)$. ODC expression in decida of recurrent miscarriages and control tissue increases with gestational age.

healthy controls in glands, decida, syncytiotrophoblasts and cytotrophoblasts (Fig. 2).

ODC expression in the placenta

ODC expression in decida tissue of controls and RM from 8th up to 13th week of gestation has been analysed

and compared with each other (Fig. 3). In Fig. 3 it is apparent that ODC expression in decida tissue of the controls and RM increases with gestational age (Fig. 4). There is a significant upregulation ($P=0.028$) of ODC expression in RM in comparison to healthy controls for the subgroup of gestational weeks 8–10 (Fig. 4).

Figure 4

ODC protein expression in decida tissue of placentas from pregnancy weeks 8 until 10. (A) Boxplot and scatterplot (jittered) of ODC protein expression in decida of control (gestational weeks 8–10) and miscarriages (weeks 8–10), showing a significant upregulation of ODC expression in recurrent miscarriages ($P=0.028$). (B, D) ODC staining in control decida in the 9th gestational week, $\times 10$ (B) and $\times 25$ (D). (C, E) ODC staining in decida of recurrent miscarriages in the 9th gestational week, $\times 10$ (C) and $\times 25$ (E). (F) ODC staining in control decida in the 10th gestational week, $\times 10$. (G) ODC staining in decida of recurrent miscarriages in the 10th gestational week, $\times 10$.

Figure 5

Western blot analysis of TAAR1 expression after stimulation with T₃ and RO5203548. (A) Bar graph diagram of TAAR1 expression after stimulation with 0.01 nM and 0.1 nM T₃. (B) Bar graph diagram of TAAR1 expression after stimulation with 0.01 nM and 0.1 nM RO5203548. (C) Picture of Western blot membrane of BeWo cells where (D). BeWo cell control group. (E) BeWo cells stimulated with 0.01 nM RO5203548, (F) BeWo cells stimulated with 0.1 nM RO5203548. (G) BeWo cell control group, (H) BeWo cells stimulated with 0.01 nM T₃, (I) BeWo cells stimulated with 0.1 nM T₃.

TAAR1 protein expression in BeWo trophoblast model cells after stimulation with T₃, T₁AM and RO5203548

Western blot

The stimulant T₃ induced a significant upregulation of TAAR1 protein expression in BeWo cells at a concentration of 0.01 nM ($P=0.011$) and 0.1 nM ($P=0.008$) (Fig. 5). TAAR1 expression in BeWo cells was also upregulated after stimulation with 0.1 nM RO5203548 ($P=0.011$) for 6 h (Fig. 5). At a lower concentration, 0.01 nM, RO5203548 had no significant influence on TAAR1 expression ($P=0.859$) (Fig. 5). The endogenous ligand T₁AM induced a significant upregulation ($P=0.007$) of TAAR1 expression

at a concentration of 10 nM and after an incubation time of 6 h (Fig. 6).

ODC protein expression in BeWo trophoblast model cells after stimulation with T₃

BeWo cells were stimulated with 0.01 nM and 0.1 nM T₃ for 48 h and ODC expression was analysed afterwards. Figure 7 shows a significantly increased ODC expression in BeWo cells after incubation with 0.01 nM T₃ and 0.1 nM T₃ (both $P=0.007$) (Fig. 7).

Double-immunofluorescence

TAAR1 and HLAG

In placentas of electively terminated pregnancies (ETP) TAAR1-positive cells are stained in red (Fig. 8A) and HLAG-positive cells, which were identified as interstitial EVT, are stained in green (Fig. 8B). Triple filter excitation of ETP shows the non-exclusive simultaneous expression of TAAR1 and HLAG in yellow (Fig. 8C), indicating that EVTs express TAAR1. ODC-positive cells are stained in red in control tissue (Fig. 8D), and an increased expression of ODC can be found in placentas of RM (Fig. 8G). Triple filter excitation of ETP shows a coexpression of ODC and HLAG visible in yellow (Fig. 8F). In placentas of RM, more cells stained in yellow can be detected compared to the controls, indicating an increased level of simultaneous expression of ODC and HLAG in RM (Fig. 8I).

Figure 6

Western blot analysis of TAAR1 expression after stimulation with T₁AM. (A) Bar graph diagram of TAAR1 expression after stimulation with 10 nM T₁AM showing a significant upregulation of TAAR1 expression ($P=0.007$). (B) Picture of Western blot membrane of BeWo cells where (A). BeWo cell control group. (B) BeWo cells stimulated with 10 nM T₁AM.

Figure 7

Western blot of ODC expression. (A) Boxplot analysis of ODC expression in BeWo cells after incubation with T_3 . 0.01 nM and 0.1 nM T_3 induce a significant upregulation of ODC protein expression ($P=0.007$). (B) Picture of Western blot membrane of BeWo cells where (A) and (B) BeWo cell control group. (B) and (E) BeWo cells stimulated with 0.01 nM T_3 for 24 h. (C) and (F) BeWo cells stimulated with 0.1 nM T_3 for 24 h.

ODC and HLAG

In control placental tissue of healthy pregnancies, an increased number of HLAG-positive cells, stained in green, can be detected (Fig. 8E), indicating an increased expression of EVT in controls in comparison to RM (Fig. 8H).

Discussion

In this study, the regulation of TAAR1, Phospho-GSK3 β and ODC have been described in the placenta of healthy pregnancies and spontaneous as well as RM. A significant upregulation of TAAR1 in syncytio- and cytotrophoblasts, decidua and glands of spontaneous and RM may presumably be a sign for an increased decarboxylation of thyroid hormones in miscarriages. A lack of thyroid hormones during pregnancy may possibly induce an upregulation of TAAR1 expression, which denotes a metabolism of thyroid hormones. An excess of thyroid hormones may lead to an increase in decarboxylation

of thyroid hormones, which can consequently have a negative effect on the pregnancy.

An interaction between TAAR1 and D2R has recently been described (6). Harmeier and colleagues suggest that a heterodimerisation of TAAR1 and D2R elevates the recruitment of β Arr2 to TAAR1 and activates Akt, which consequently leads to an inhibition of GSK3 β via an alternative pathway (6). Enhanced GSK3 β phosphorylation indicates a silencing of the downstream signalling cascade (36, 37). In our study, we discovered enhanced expression of phosphorylated GSK3 β (pGSK3 β) expression in syncytiotrophoblasts, cytotrophoblasts and the decidua of RM in comparison to healthy controls and thereby a concomitant pGSK3 β expression with TAAR1. An increase in pGSK3 β expression leads to an inhibition of unphosphorylated GSK3 β and consequently to an activation of the Wnt signalling. In a current study, we also investigated a reduced D2R expression in syncytiotrophoblasts and cytotrophoblasts of RM in comparison to trophoblasts of healthy controls (in revision at JCEM). These findings are in line with the results of Espinoza and coworkers, who detected a negative correlation between TAAR1 and D2R in the striatum of TAAR1-KO-mice, where D2 dopamine receptors are overexpressed and thereby selectively activate the G-protein-independent AKT/GSK3 signalling pathway (24). In contrast, Harmeier and coworkers found a concomitant increase of TAAR1 and D2R expression in the neurone cell line HEK293 (6). This difference might be due to the variability of TAAR1 and D2R expression in different cell lines and tissue.

One further aim of the study included the identification of ligands that have the ability to upregulate TAAR1 expression in trophoblast model cells and could thereby be responsible for the induction of miscarriages. Triiodothyronine, also known as T_3 , at a concentration of 0.01 nM and 0.1 nM induced a significant upregulation of TAAR1 expression in BeWo trophoblast model cells. The ligand RO5203548 significantly increased TAAR1 expression in BeWo and JEG-3 cells. RO5203548 has been described as a partial agonist and highly selective for TAAR1 in dopaminergic neurons (15). Partial agonists can either act as an agonist or an antagonist depending on the intrinsic activity of the receptor (15). Endogenous ligands, such as T_1 AM, have a high affinity for TAAR1 (13, 18). In our study, 10 nM T_1 AM induced a significant upregulation of TAAR1 expression after an incubation time of 6 h. We are aware of the fact that TAAR1 has an intron-less sequence. For that reason, we performed intense genomic DNA digestion to avoid that TAAR1 RNA is contaminated

Figure 8

Double-immunofluorescence of TAAR1 and ODC. (A) Red stained TAAR1 expression in elective terminations of pregnancy. (B) Green stained HLAG expression in decidua of ETP placenta. (C) Coexpression of TAAR1 and HLAG in ETP visible in yellow. (D) Red stained ODC expression in decidua of ETP placenta. (E) Green stained HLAG expression in decidua of ETP placenta. (F) Double staining of both ODC and HLAG. A double expression is visible in yellow. (G) Red stained ODC expression in decidua of RM placenta. (H) Green stained HLAG expression in decidua of RM placenta. (I) Double staining of both ODC and HLAG. A double expression is visible in yellow; (all magnification $\times 40$).

with genomic DNA. We used commercially TAAR1 primers from 'Applied Biosystems by Thermo Fisher'. The policy of the company forbids to publish the primer sequences but the company guarantees the specificity of the sequences. An upregulation of TAAR1 is induced through the transformation of T_3 into T_1AM . We could detect that 0.01 nM and 0.1 nM T_3 induced a significant upregulation of ODC expression in BeWo cells via Western blot analysis. An increased expression of ODC after stimulation with T_3 suggests that thyroid hormone derivatives are present during miscarriages. Transformation of thyroid hormones into endogenous amines may therefore partly be carried out by the ODC. In our study, all patients with RM had normal thyroid hormone levels. Individuals with hypothyroidism and hyperthyroidism and patients with TPO antibody positivity were excluded from the study as those pathologies can influence miscarriage rates. We presume that patients with normal

thyroid hormonal levels possibly have a deficiency of T_3 during pregnancy induced through the ODC. In a former study, we found that patients with spontaneous and RM have reduced expression of THR (5). A reduced level of THR during pregnancy may rather be due to an enhanced transformation of thyroid hormones into their derivate T_1AM than to a decrease in thyroid hormone synthesis. One possible approach could therefore be a prophylactic substitution of L-thyroxin to patients with RM, independent of their TSH and thyroid hormone levels. A further possibility to reduce the transformation of T_3 into T_1AM would be an inhibition of the ODC.

In conclusion, we found enhanced placental TAAR1 and pGSK3 β expression in RM, which consequently leads to an inhibition of GSK3 β and activation of the Wnt-signalling pathway. TAAR1 seems to be an inductor of the wnt-signalling pathway, and could play role in the embryonal development. Inhibition of GSK3 β , as

observed in the neurological system through TAAR1, could also be detected in RM, indicating a TAAR1-dependent regulation of GSK3 β in RM. Using endogenous ligands, we demonstrated a T₃, RO5203548 and T₁AM-dependent upregulation of TAAR1 in trophoblast tumour cells. Aims of future studies include investigation of the functional interaction between TAAR1, ODC, GSK3 β and the Wnt-signalling pathway in placentas of miscarriages. Future experiments should elucidate the role of β Arr2 in the signalling cascade in miscarriages. β Arr2 is known to dephosphorylate Akt and thereby activate GSK3 β (6). *In vivo* experiments with a knockout or knockdown of TAAR1 through antisense or siRNA technologies should be considered. TAAR1 may represent a novel target for the treatment of miscarriages. Further studies are required to analyse the signalling pathways of the endogenous ligands of TAAR1 and to possibly re-examine thyroid hormone signalling (38).

Supplementary data

This is linked to the online version of the paper at <https://doi.org/10.1530/EC-17-0272>.

Declaration of interest

The authors declare that we have no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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Author contribution statement

A V, U J and S S conceived and designed the experiments. S S, M G, E T, C K and S H performed the experiments. A V, U J, S S, M G, E T, C K, S H and K H analysed the data. A V, U J, S S, M G and K H wrote the research article. All authors read and approved the manuscript. S M, K H, H H, M P, S H, B T and D M did the final revision of the manuscript.

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