

Induction stage-dependent expression of vascular endothelial growth factor and aquaporin-1 in diethylstilbestrol-treated rat pituitary

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The anterior pituitary gland undergoes tumorigenic changes in response to oestrogen treatment in several breeds of rats. We administered diethylstilbestrol (DES) to female Wistar rats and assessed whether the expression of vascular endothelial growth factor (VEGF) and aquaporin-1 (AQP-1) was altered at different time points following DES administration. *In vivo* magnetic resonance imaging (MRI) scans showed that the mass index corresponding to the mid-sagittal area of DES-treated pituitary was significantly higher than the vehicle-controlled pituitary ($p < 0.01$) at three specific time points, accompanied by a significant reduction in body weight. Haematoxylin and eosin (HE) staining and immunohistochemical analysis demonstrated that during early stages of induction, DES increased cell proliferation and sprouting of endothelial cells, and VEGF expression transitioned from a vessel-surrounding pattern to a diffuse pattern. During later stages, angiogenesis was predominant, and VEGF expression decreased. In contrast to the early abundant expression of VEGF, endothelial expression of AQP-1 increased during later stages. Our data indicated a dynamic scenario of biological alterations in DES-treated pituitary tissue, in which VEGF and AQP-1 exert their functions at different stages of induction, and we provide novel insights into understanding oestrogen-related tumorigenesis in the anterior pituitary gland.

Key words: diethylstilbestrol, angiogenesis, vascular endothelial growth factor, aquaporin-1.

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Morphological changes of the pituitary tumour induced by oestrogen *in vitro* predominantly include increased cell proliferation and angiogenesis. In contrast to the large number of studies assessing the effects of oestrogen on mature pituitary tumour tissue, the morphological changes and differential expression of relevant molecules at specific stages of induction have not been thoroughly investigated.

Angiogenesis, a condition of neovascularisation that arises from existing vasculature, plays pivotal roles in foetal development, injury repair, and reproductive cycle regulation. Additionally, it is a prerequisite for the induction of tumour growth and invasion (Gimbrone *et al.* 1972, Gimbrone *et al.* 1973, Folkman *et al.* 1990). Angiogenesis critically contributes to tumorigenesis and invasion of pituitary tumours, and it affects the response of these tumours to therapeutic intervention (Goth *et al.* 2003). High-density vascularisation is associated with increased invasion potential in human mammary tumours, as well as bladder and gastric neoplasias; thus, it contributes to a poorer prognosis and low survival rates (Weidner *et al.* 1991, Weidner *et al.* 1992, Maeda *et al.* 1995, Bochner *et al.* 1995). Angiogenesis results from signalling mediated by multiple factors (Guinebretiere, 2005); particularly vascular endothelial growth factor (VEGF), which in turn mediates the biological effects of other growth factors. Aquaporins (AQPs) are a family of proteins that shuttle water across the cellular membrane, and a subset of these proteins can counteract glioma-associated lactic acidosis by clearing glycerol and lactate from the extracellular space (Warth, 2007). AQP-1 mRNA and protein are expressed at very low levels in rat brain primary microvascular endothelial cells, and its expression increases with passaging (Dolman, 2005). AQP-1 has also been detected in the pituitary gland and is expressed in vascular endothelial

cells of the adenohypophysis and neurohypophysis (Kuwahara, 2007). However, expression of AQP-1 in pituitary in response to oestrogen is still not clear.

In the current study, DES was intraperitoneally administered to rats to induce chronic tumourigenesis in the anterior pituitary gland.

Magnetic resonance imaging (MRI) and haematoxylin and eosin (HE) staining were applied to monitor *in vivo* tumour growth and morphological changes. Electron microscopy was utilised to assess the final ultrastructural attributes of the pituitary gland. Immunohistochemistry was used to evaluate the expression of VEGF and AQP-1 and their localisation at specific stages following DES administration.

Materials and Methods

A total of 60 female Wistar rats, 3 weeks old and weighing 70–80 g, were used. The animals were housed with free access to tap water and standard pellet food. They were kept at a controlled temperature ($24 \pm 1^\circ\text{C}$) and humidity ($55 \pm 5\%$), and a 12 hour day-night cycle (10 p.m.–10 a.m.) was maintained. The induction was based on a previously published method (Zhao et al. 2007). Rats were randomly allocated into two groups for each time point. In the vehicle-controlled group, rats were injected intraperitoneally with sunflower seed oil (1 mL/kg, twice a week) for 12 weeks, whereas animals in the DES group were administered DES intraperitoneally (5 mg/kg, twice a week) for 12 weeks. The procedures undertaken were in strict compliance with the guidelines on the care and use of laboratory animals at our institution. DES was purchased from Sigma Chemical Co. (St Louis, MO, USA). Anti-VEGF antibody was purchased from Boster Biotech (Wuhan, China), and anti-AQP-1 antibody was purchased from Chemicon (Temecula, CA, USA). The DES injection was prepared by dissolving DES powder in sunflower seed oil to obtain a final concentration of 5 mg/mL.

MRI scanning

At week 4, week 8, and week 12, three rats were randomly selected from each group for MRI scanning. The T_2 WI mid-sagittal scanning technique was used to monitor the changes in the pituitary gland at each time point. Intraperitoneal administration of hydration chloraldehyde at 30 mg/100 g of body

weight was applied to minimise animal movement during scanning. The rats were placed in a specifically designed saddle-coil probe, and a GEMSL \times 3T superconductive MR (General Electric, Fremont, CA) medical imaging instrument was used to obtain a mid-sagittal T_2 WI image with a TR of 3100 ms and a TE of 115 ms. Data collection matrix sizes were 320 \times 288 for T_2 WI. The mid-sagittal section was defined as the in-line position of the olfactory bulb, corpus callosum, cerebellum, and brain stem. The software Image Tool II was used to obtain the mid-sagittal area indexed by the number of pixels within each rat pituitary gland.

Morphological methods

At each time point, rodents were weighed and anaesthetised by hydration chloraldehyde at 30 mg/100 g body weight and transcardially perfused with a 40 g/L solution of paraformaldehyde diluted in PBS (0.01 M, pH 7.4). The tissues were embedded in paraffin and sectioned at 4 μm for HE staining and immunohistochemistry.

Immunohistochemical staining was applied to demonstrate the expression and localisation of VEGF and AQP-1. In brief, deparaffinised sections were rehydrated through a graded series of ethanol to phosphate buffered saline (PBS). Antigen retrieval was performed, and endogenous peroxidase clearance was performed by incubation in 3% H_2O_2 . Then sections were blocked with 10% normal goat serum for 30 minutes. Afterwards, these sections were individually incubated with anti-VEGF and anti-AQP-1 antibodies (diluted at 1:50) at 4°C overnight. The antigen-antibody complexes were visualised using the avidin-biotin-peroxidase complex (ABC) method. Counterstaining was performed with Mayer's haematoxylin.

Transmission electron microscopy

After induction, one rat from the vehicle control group and two rats from the DES group were randomly selected for transmission electron microscopic analysis. For electron microscopy, pituitary tissues were fixed in 2% glutaraldehyde in 0.1 M PBS (pH 7.4) and then subjected to postfixation in 2% osmium tetroxide. The tissues were embedded in Araldite, and the ultra thin sections were examined using electron microscopy (Philips EM208).

Statistical analysis

Results were expressed as means \pm standard devi-

ation (SD), and one-way ANOVA analysis was utilised to compare the differences between the two groups in terms of body weight, mid-sagittal area, with post hoc testing using the Bonferroni procedure. P values less than 0.05 were considered significant.

Results

Body weight

Fur loss in the DES group was initially observed as early as 2 weeks after the initiation of DES treatment and was sustained throughout the remainder of the study (data not shown). At weeks 4, 8, and 12, the body weights in the vehicle control group were 189.6 ± 15.6 g, 256.4 ± 27.0 g, and 244.5 ± 23.8 g, respectively, whereas in the DES group they were 164 ± 11.4 g, 209.2 ± 13.5 g and 208.4 ± 21.0 g, respectively. Therefore, body weight was significantly reduced in the DES group at each time point ($n=10$ in each group, and $p < 0.05$) compared with the vehicle-controlled group.

MRI changes

At three time points, T_2 WI scanning of the pituitary gland in all tested rats in the vehicle-controlled group indicated a typical triangular pattern with sharp margins and uniform image signal intensity lighter than that of the surrounding brain tissue. The peripheral meninges and cerebral spinal fluid (CSF) were

also continuous. No abnormal changes in the shape were observed in the vehicle-controlled group throughout the course of the study (Figure 1 A1-3). At week 4, the pituitary gland in the DES-treated rats started to enlarge, losing its typical triangular shape, and the dura mater showed heterogeneous signals under T_2 WI, with a blurred subarachnoid space (Figure 1 B1). At week 8, a discontinuous dura mater, further enlarged pituitary gland, and heterogeneous intrahypophyseal signal intensity were observed (Figure 1 B2). At week 12, the pituitary gland crept over the sella turcica in a hillock-like shape. The continuity of the dura mater was severely destroyed, and the enlarged pituitary gland expanded into the subarachnoid space, resulting in a malformed pituitary gland in the DES-treated rats (Figure 1 B3). MRI scanning has been proved to be a practical method for monitoring the growth and morphological alterations of the pituitary gland induced by oestrogen treatment, and the mid-sagittal area closely correlates with the mass of the pituitary gland. During the progression of these studies, the pituitary gland in the control group exhibited slow growth; the indexed number of pixels was 67.0 ± 4.6 , 75.7 ± 7.6 , and 88.3 ± 3.2 at weeks 4, 8, and 12, respectively. In contrast, the mid-sagittal areas of the pituitary tissue in the DES-treated group were 131.0 ± 11.5 , 160.7 ± 11.7 and 315.67 ± 11.6 , respectively, which was significantly higher than in the control group ($n=3$ in each group and $p < 0.01$ in comparison to the controls) (Figure 1 C).

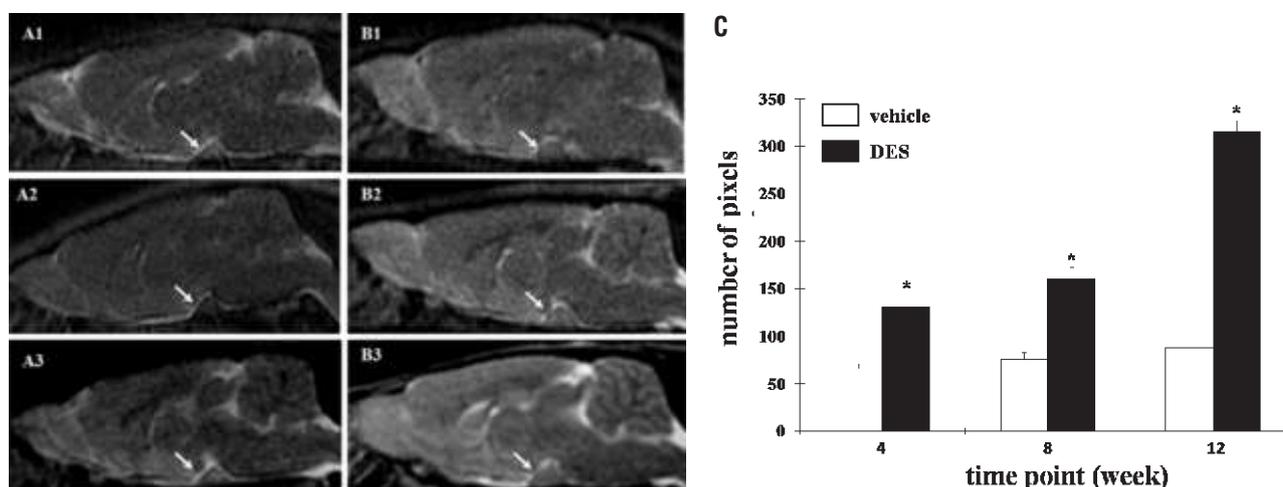


Figure 1. Representative pictures showing the rat pituitary gland under mid-sagittal T_2 WI. *In vivo* MRI scanning showed that DES induced gradual enlargement of the pituitary gland (indicated by the arrow) at 4, 8, and 12 weeks (B 1-3), in contrast to the lack of change in the vehicle-treated pituitary (A1-3). As shown in C, long-term treatment of DES resulted in significantly larger mid-sagittal areas of the pituitary gland compared with the vehicle controls ($*p < 0.01$, compared with controls, respectively).

Histology and immunohistochemistry

Initially after the administration of DES, localised cell proliferation was observed (Figure 2A), and at 8 weeks, there was further spreading of proliferative cells. A typical characteristic of this stage is the sprouting of spindle-like endothelial cells lining the proliferative cells and the formation of large cord-like structures (Figure 2B). At this stage, no vascularised lumens were apparent in the area of proliferation. At week 12, vascular lakes were predominant in the anterior pituitary gland (Figure 2C); however, there was still a discernable borderline between

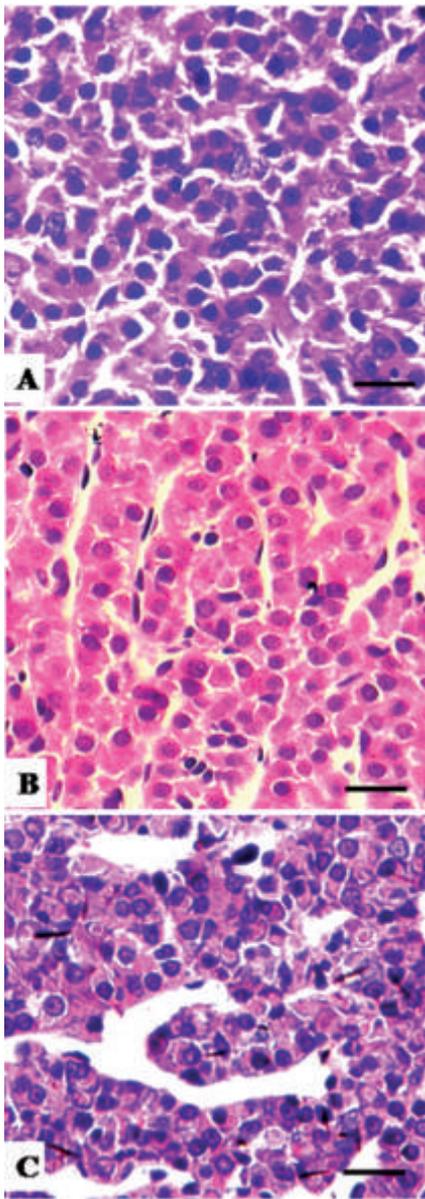


Figure 2. HE staining showing the morphological changes of the pituitary tissues subjected to DES treatment at week 4 (A), week 8 (B), and week 12 (C), respectively. (Scale bars, 20 μ m).

areas with and without angiogenesis. Notably, in the area without angiogenesis, cells with condensed nuclei were visible, which was rarely observed in vascular lake-enriched areas (*data not shown*). Based on the dynamic alterations observed from HE staining of DES-treated rat pituitary tissue, we found that angiogenesis characterised by vascular lake formation was a subsequent event to increased cell proliferation during pituitary tumourigenesis. We hypothesised that different factors may function during specific stages, so we investigated the expression of VEGF, a widely accepted angiogenic growth factor, and aquaporin-1 (AQP-1), a potential angiogenic stimulator, in response to DES administration over a 12-week time course. Throughout tumour induction and progression, expression of VEGF was weak and sparse in the vehicle-controlled group (*data not shown*). We could not detect the expression of VEGF in the endothelial cells in either group at any time point. Initially, in the DES-treated group, cytoplasmic VEGF-positive cells were mainly localised to the areas surrounding vessels (Figure 3A). At week 8, a diffuse expression pattern of VEGF was observed, whereas a subset of cells exhibited a perinuclear pattern (Figure 3B). When angiogenesis was complete at week 12, the intensity of VEGF expression was reduced (Figure 3C); however, VEGF-positive cells were still prevalent in angiogenic areas in comparison to less angiogenic areas, resulting in sharp separation of the two niches (Figure 3D). Prior to the completion of angiogenesis, we could not recognise differential expression of AQP-1 between the two groups (*data not shown*), but by week 12, expression of AQP-1 was enhanced in the endothelial cells that lined the vascular walls (Figure 4). In contrast, there was no apparent expression of AQP-1 in proliferating cells.

Transmission electron microscopy

The intracellular membrane localisation of a secretory particle was detected in DES-treated pituitary tissue at week 12 (Figure 5A). The rough endoplasmic reticulum (RER) was well developed into a concentric-circled structure, a typical characteristic for the development of prolactinoma (Figure 5B). Concurrently, secretory granules with diameters ranging from 100 nm to 300 nm accumulated in areas adjacent to the vascular wall (Figure 5C). The vascular wall surrounding neovascularised vessels was thin and poorly developed and had endothelial cells attached (Figure 5D).

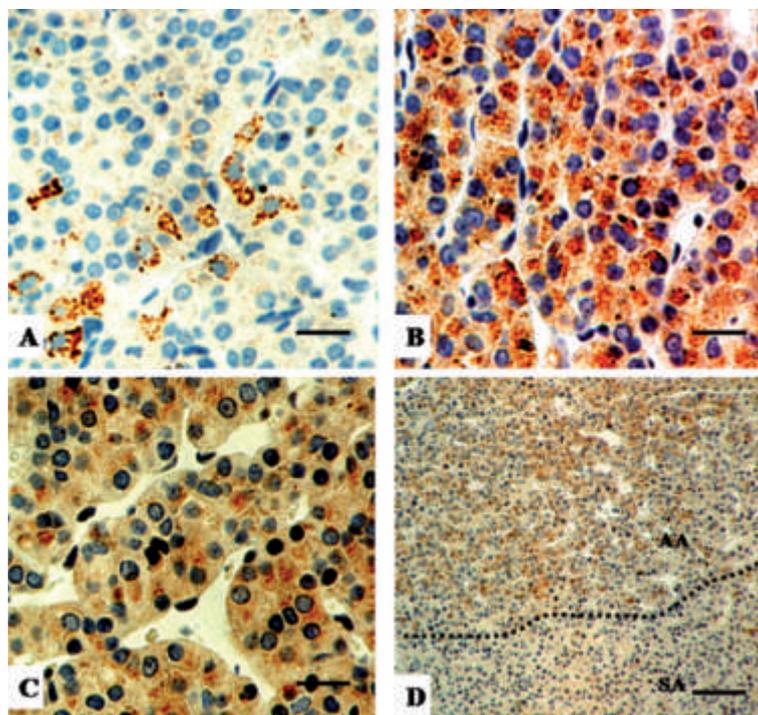


Figure 3. Immunostaining showing the expression of VEGF at week 4 (A), week 8 (B) and week 12 (C and D). Initially, expression of VEGF was localised to the perivascular zone (A), and then its expression transitioned to a diffuse pattern (B). VEGF expression was reduced dramatically during angiogenesis (C), although its expression in the angiogenic area still exceeded that in the area with less angiogenesis (D) (scale bars: A-C , 20 µm; D, 80 µm).

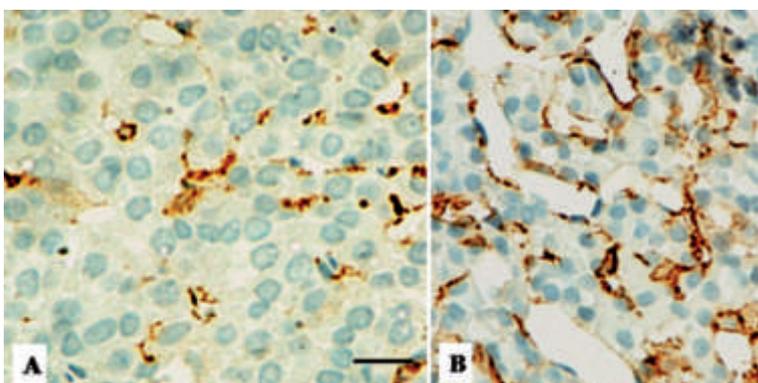


Figure 4. Immunostaining for the expression of AQP-1 in the vehicle-controlled group (A) and DES group (B) at week 12. A weak signal of AQP-1 expression was detected in the normal hypophyseal vascular walls (A), and DES-induced expression of AQP-1 was exclusively localised to the endothelial cells of neo-vascularised vessels (B) (scale bars: A and B, 20 µm).

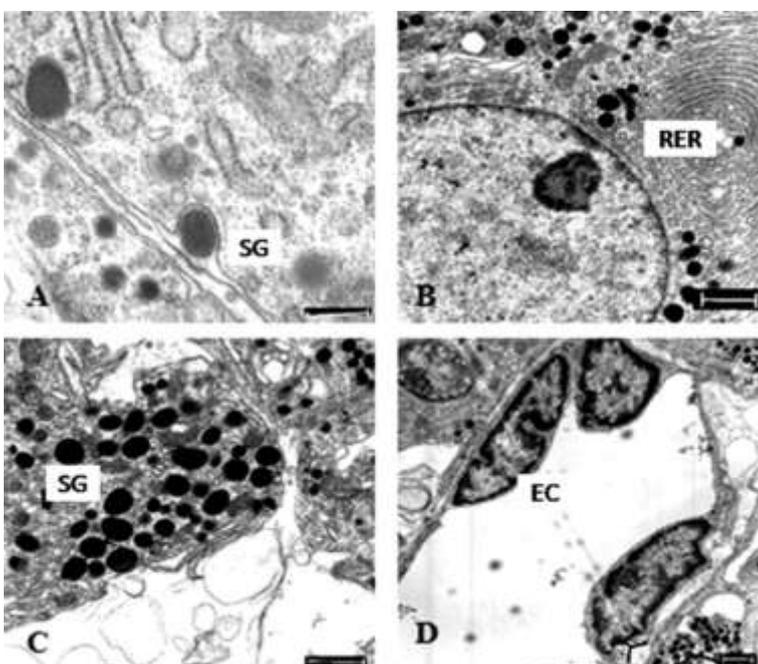


Figure 5. Representative image showing the ultrastructural results of the anterior pituitary gland treated with DES (SG: secretion granule; RER: rough endoplasmic reticulum; EC: endothelial cell) (scale bars: A, 300 nm; B, C, and D, 1 µm).

Discussion

Oestrogen has proven to be an efficient tumourigenic inducer for the pituitary gland, and several events contribute to this process. Intraperitoneal injection of oestrogen provided chronic tumourigenesis induction with real time dose modulation according to body weight. This method facilitates monitoring of the dynamic changes in neoplastic growth *in vivo* through MRI and immunohistochemistry, enabling the identification of morphological and expression alterations.

The mid-sagittal accumulated pixel number under T2WI highly correlated with pituitary body mass, suggesting that it is a high fidelity tool to evaluate rat pituitary mass *in vivo* (Rudin, 1988). Van Nesselrooij (van Nesselrooij, 1992) found that MRI could detect enlarged pituitary glands in rats as early as 16 days after treatment with estradiol-17. Our results showed that DES accumulation correlated with increased mid-sagittal area, which reached its peak at week 12, suggesting increased pituitary volume and mass.

MRI investigation also detected a blurred and discontinuous CSF signal representing the narrowing of the interstitial space between the pituitary tissue and dura mater at week 12. This suggests that enlargement of pituitary tissue may oppress adjacent structures such as the dura mater and pons.

In previous studies, non-canonical proliferation of pituitary cells that occurred with the emergence of blood-enriched space was defined as a pituitary tumour (Sato, 1997). In our study, *in vivo* MRI test and HE staining provided detailed information on the dynamic morphological changes that occurred during tumourigenesis of the pituitary gland. The pituitary cell proliferation transitioned from a local pattern to a diffuse pattern with the development and completion of neo-vascularisation. Angiogenesis was apparent at week 12, which may have contributed to the rapid increase in volume of the pituitary gland, as confirmed by MRI at week 12.

The dynamic morphological changes led us to hypothesise that specific growth factors may be expressed in a distinct pattern during pituitary tumourigenesis. To assess this, we performed immunostaining for VEGF, a widely accepted key regulator of angiogenesis, and AQP-1, a candidate stimulator of angiogenesis, proliferation and

metastasis. VEGF, a homodimeric heparin-binding affinity protein with a molecular weight of 45 kDa, was initially extracted from bovine pituitary follicular cells by Gospodarowicz *et al.* (Gospodarowicz, 1989). VEGF interacts with its receptor to signal in a paracrine fashion, specifically exerting mitogenic effects on nearby endothelial cells. Additionally, VEGF can increase the osmotic ability of the vascular endothelia, leading to the extravasation of serum proteins, fibre sedimentation, damage to the blood-brain-barrier due to enhanced endothelial proliferation. VEGF expression was detected under conditions of ischaemia, anoxia, high blood sugar levels, and hormone exposure. VEGF expression has been observed in rat and human growth hormone (GH) pituitary tumours, prolactinoma, and adrenocorticotrophic hormone (ACTH) adenoma, which can be stimulated by transformation factors including oestrogen and inhibited by dexamethasone (Loyhrer, 2001). In the normal pituitary gland, expression of VEGF was mainly restricted to endothelial cells, but here, we detected expression of VEGF in proliferative pituitary cells, but not in endothelial cells. This is consistent with Pan's report that VEGF was strongly expressed in pituitary adenomas and correlated with the tumour grade (Pan, 2003). Banerjee *et al.* (Banerjee, 2000) determined that methoxyestradiol could downregulate VEGF expression and potentially inhibit tumour growth by reducing expression of the pituitary tumour transforming gene (PTTG) (McCabe, 2002; Heaney, 2002).

Expression of VEGF was initially limited to areas surrounding the vessels. We hypothesise that these cells initially sensitise and react to the stimulation of DES by upregulating the expression of VEGF because they are in direct contact with the vascular wall. At week 8, VEGF was expressed in a diffuse pattern due to continuous stimulation of DES. Notably, perinuclear localisation of VEGF was noted in a subset of proliferative cells. The nuclear localisation of VEGF was additionally confirmed by immune electron microscopy, which indicated that VEGF associates with the nucleolus and euchromatin (Jorge, 2005). The nuclear translocation of VEGF endows the tumour cells with stronger proliferative ability and increased transcription of pro-tumourigenic and angiogenic factors. During the final stages of induction, the expression of VEGF

was sharply reduced, but the differential expression of VEGF prevailed between areas where angiogenesis was present and absent. Although the pituitary VEGF level correlates well with mass (Cracchiolo, 2002), high serum levels of VEGF are not a prerequisite, suggesting that additional factors are involved in this process.

We present the first study examining the expression of aquaporin-1 in response to DES stimulation, and we evaluate its potential role as a stimulator of angiogenesis and executor of metastasis. Although it has been reported that transfection of AQP-1 into the pituitary endocrine cell line AtT-20 affects dense core secretory granule (DCSG) biogenesis and normal levels of hormonal secretion (Arnaoutova, 2008), we failed to detect the expression of AQP-1 in the pituitary cells in both groups at each time point. In contrast, at week 12, AQP-1 was strongly expressed in the vascular endothelial cells, suggesting that AQP-1 may function as an executor of angiogenesis and possibly indirectly affect cell proliferation, tumour cell migration, and invasion. AQP1-knockout mice exhibited reduced tumour angiogenesis, suggesting that AQP-1 facilitates cell migration by affecting the transport of water to cell protrusions formed during migration and that it is, therefore, a potential therapeutic target to treat tumours (Verkman, 2005; Hu, 2006; Papadopoulos, 2008). It was also reported that AQP-1 was strongly expressed in endothelial cells of the capillary and postcapillary venules in Lewis-lung-carcinoma tumour tissues in mice, which could be inhibited by the mild diuretic Acetazolamide (Xiang, 2004).

In summary, our results showed progressive changes in the pituitary gland upon DES treatment. Anatomically, the vault-shape enlargement of the pituitary gland and the discontinuous CSF signal in the mid-sagittal section evident upon MRI strongly suggested that hypophyseal proliferation and angiogenesis resulted from DES treatment. Non-endothelial hypophyseal VEGF expression appears to function during angiogenic initiation, whereas subsequent sustained angiogenesis may partially rely on the endothelial expression of AQP-1. Disrupting the action of VEGF and AQP-1 depending on the stage of angiogenesis may provide an important therapeutic strategy to control tumour cell growth, both *in vitro* and clinically.

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