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# **Upregulation of hypoxia-inducible factors and autophagy in von Hippel-Lindau-associated retinal hemangioblastoma**

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# **Abstract**

**Purpose:** To describe pathological and molecular changes of three patients with clinically severe von Hippel-Lindau (VHL)-associated retinal hemangioblastoma (RH) with rapid progression.

**Methods:** Medical records, ocular histopathology and transmission electron microscopy from three cases of VHL-associated RHs at the National Eye Institute were retrospectively reviewed. One eye of each patient was enucleated. Hypoxia-inducible factor (HIF) 1α and HIF2α expressions were identified by quantitative reverse transcription polymerase chain reaction (qRT-PCR) and immunohistochemistry.

**Results:** All three cases had rapidly growing RHs that were resistant to multiple conventional therapies and two (Patients 1 and 2) were also resistant to multiple intravitreal anti-vascular endothelial growth factor (VEGF) treatments. Macroscopically, all the enucleated eyes had multiple RHs, serous retinal detachment, severe retinal disorganization and focal hemorrhages. Histopathology showed typical RHs composed of vacuolated foamy von Hippel-Lindau cells and capillary networks. Retinal gliosis and hemorrhages were also presented. Additionally, T lymphocytes and macrophages were infiltrated in the tumors of two patients resistant to anti-VEGF therapy. Immunohistochemistry and qRT-PCR found upregulation of HIF1α in the retinal lesions of all eyes. Importantly, upregulation of HIF2α was exclusively detected in the two cases with inflammatory infiltration and resistance to anti-VEGF therapy. Ultrastructural images showed autophagy, lipid droplet, glycogen aggregations and cytoplasmic degeneration in many VHL cells.

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**Conclusions:** Based on the histopathological and molecular pathological findings, autophagy, inflammation and/or upregulation of HIF2α could potentially contribute to the aggressive course of RHs, resulting in the resistance to multiple anti-VEGF and radiation therapies in these patients.

# **Keywords**

von Hippel-Lindau; retina; hemangioblastoma; hypoxia-inducible factors; autophagy; inflammation

# **Introduction**

von Hippel-Lindau (VHL) disease is an autosomal dominant disease with an incidence of one in 36,000 live births [1]. It has multi-systemic tumors and affects the retina and central nervous system (CNS), as well as visceral organs, including the kidney, inner ear, pancreas, adrenal glands and epididymis at varying frequencies [2]. VHL is associated with a germline mutation of the VHL tumor suppressor gene on the chromosome 3p25 [3]. Loss of heterozygosity of VHL gene is well demonstrated in VHL-associated tumor cells including retinal and optic nerve head hemangioblastoma [4, 5]. VHL protein (pVHL) is encoded by VHL gene and acts to degrade the transcription factor called hypoxia-inducible factor (HIF). HIF is composed of  $\alpha$  and  $\beta$  subunits and responds to changes in tissue oxygen concentration. HIF1α and HIF2α dimerize with constitutively expressed HIF1β, also termed aryl hydrocarbon receptor nuclear translocator (ARNT), to activate the transcription of hypoxia-responsive genes. HIF2α, also termed endothelial Per-Arnt-Sim (PAS) domain protein 1, is 48% identical in sequence to HIF1α. Both HIF1α and HIF2α are inducible by hypoxia and associated with angiogenesis, glycolysis and apoptosis under hypoxic conditions [4, 6]. However, HIF1α and HIF2α are believed to have distinct functions, both in normal cellular physiology and in tumorigenesis [6–10].

Retinal hemangioblastoma (RH) is seen in 45-60% of VHL patients [11, 12], with two-thirds of them with multiple lesions [13] and bilateral in 26% of patients with VHL [14]. The ocular tumor appears typically as a round, globular reddish vascular lesion with a dilated feeding artery and a tortuous draining vein. It is found in either the juxtapapillary retina or the peripheral area. Although it usually has a benign nature and may be slow-growing, RH would disrupt retinal architecture or cause total retinal detachment, neovascular glaucoma, resulting in painful vision loss or even blindness in VHL patients. This is especially difficult when the tumor is located on or around the optic nerve as no known therapy is effective for this lesion. Early diagnosis and appropriate treatment reduce the risk of visual loss or blindness, especially for the lesions located peripherally. Several conventional methods have been used to ablate hemangioblastoma, including laser photocoagulation, cryotherapy, radiotherapy and photodynamic therapy. Recently, on the basis of molecular knowledge of tumorigenesis of VHL disease, some highly upregulated molecules, such as vascular endothelial growth factor (VEGF) and platelet-derived growth factor (PDGF), have been targeted in systemic and ocular VHL therapies [15].

Herein, we reported the pathological and molecular features of three enucleated eyes from three VHL patients with rapidly growing RHs that were refractory to therapy. In addition to

multiple VHL-associated RHs in the retina, we found upregulation of  $HIF1\alpha$  in all three patients with VHL-associated RHs. More importantly, the inflammatory infiltration and upregulation of HIF2α are exclusively presented in two cases resistant to anti-VEGF therapy.

# **Materials and Methods**

The study was approved by the National Eye Institute (NEI) Institutional Review Board for human subjects and an informed consent was signed by the patients. This study was conducted at the NEI and followed the tenets stated in the Declaration of Helsinki. Unilaterally enucleated eyes from three patients of VHL-associated RHs were submitted to NEI.

#### **Clinical examinations**

Clinical examinations of the three patients were performed by a NEI ophthalmologist (EYC) and three local ophthalmologists, respectively. Table 1 summarized the brief clinical findings in these three patients.

#### **Histopathology**

The enucleated eyes were fixed in 10% neutral buffered formalin for at least 24 hours before macroscopic and microscopic evaluation. The globes were cut horizontally through the macula along the pupillary-optic nerve head plane, where a VHL lesion was located. In areas of RHs, a segment through the lesion was also obtained for transmission electron microscopy (TEM). The grossed eyes were paraffin embedded and slides were stained with hematoxylin & eosin (H&E) and Periodic acid-Schiff (PAS). Immunohistochemistry staining (avidin-biotin complex method) was also performed with primary antibodies to the following antigens: CD68 (macrophage biomarker, 1:100, BD pharmingen, San Diego, CA, USA), CD45RO (mainly for T lymphocyte, 1:100, Dako, Carpinteria, CA, USA), CD3 (T lymphocyte biomarker, 1:100, Dako), CD20 (B lymphocyte biomarker, 1:40, Dako), glial fibrillary acidic protein (GFAP, 1:100, Dako), HIF1α (1: 100, Sigma-Aldrich, St Louis, MO, USA) and HIF2α (1:200, Novus Biologicals, Littleton, CO, USA).

#### **TEM**

The retina containing the lesion was double-fixed in 2.5% gluteraldehyde and osmium tetroxide (0.5%), dehydrated, and embedded in Spurr's epoxy resin. Ultrathin sections (90 nm) that include retinal VHL tumors were prepared and double-stained with uranyl acetate and lead citrate, and viewed using a JEOL JEM 1010 transmission electron microscope.

#### **Microdissection and qRT-PCR**

Microdissection was performed manually on uncovered, H&E stained glass slides [16–18]. Approximately equal numbers  $(\sim 200)$  of cells from the juxtapapillary and peripheral RH areas were microdissected. The microdissected cells were subjected for qRT-PCR using the technique described previously [4, 5, 19]. Total RNA was isolated from RH cells using Arcturus Paradise RNA isolation kit (Molecular Devices., Sunnyvale, California, USA). Equal amounts of RNA were reverse transcribed with Superscript II RNase H Reverse

Transcriptase (Invitrogen, Grand Island, NY, USA) to cDNA. Quantitative reverse transcription polymerase chain reaction (qRT-PCR) was performed on the resulting cDNA using Brilliant SYBR Green QPCR Master Mix (Stratagene, La Jolla, CA, USA). The comparative cycle threshold (Ct) value method, representing log transformation, was used to establish relative quantification of the fold changes in gene expression using 7500 Real Time PCR System (Life Technologies Co. Carlsbad, CA, USA). β-actin was used as an internal control. Primers of  $\beta$ -actin, HIF1A and HIF2A were purchased from SABiosciences.

#### **Mutational Analysis of HIF2A**

Genomic DNA was extracted from tumor tissues obtained from the patients with the use of a NucleoSpin Tissue Kit (Macherey-Nagel). HIF2A exons were amplified by PCR assays. The primer sets for exon amplification have been described previously [20]. The DNA sequence of each exon was determined by forward and reverse sequencing.

# **Results**

# **Patient 1**

A 40-year-old Caucasian male had a history of systemic VHL disease, including RH, bilateral renal cell carcinoma, and brainstem VHL hemangioblastomas. The patient had VHL-associated RH diagnosed at age 15 (in 1986). His left eye had deteriorated and was eventually enucleated in 2003. His optic nerve head lesion of the right eye was treated with laser photocoagulation in 1993. This optic nerve tumor progressed, causing marked exudation that affected his vision with a decrease to 20/200 in 2004. His vision further deteriorated to 20/500 with increased retinal hard exudates in January 2005 (Fig. 1a). The patient was then enrolled in a trial of intravitreal ranibizumab at the NEI/National Institutes of Health (NIH). He received monthly intravitreal ranibizumab 0.5 mg. His vision had improved to 20/250 in April 2005, but declined to 20/400 in May 2005. Despite additional ranibizumab injections, his vision continued to deteriorate to 20/500 within a month. At the completion of the multiple injections of ranibizumab in August 2005, he developed additional small RHs in the macula in addition to the optic nerve head tumor. His vision was less than 20/500. In February 2006, external beam radiation therapy was given to the large optic nerve head RH and laser therapy to the small RHs. In spite of all the therapeutic regimens, his vision dropped to 20/800. In March 2010, he developed vitreous hemorrhages, serous retinal detachment and persistent lipid accumulation with multiple chorioretinal scars, in addition to RHs in the optic nerve head and retina (Fig. 1b). In May 2011, he developed rubeosis iridis with IOP of 12 mmHg. Multiple intravitreal injections of 1.25 mg of bevacizumab were administered for iris neovascularization. However, there was no response to the treatment and the rubeotic eye had barely light perception and the IOP was controlled with topical anti-glaucoma medication, Cosopt. Because of uncontrollable pain, the patient elected to enucleate the blind right eye in January 2012. The patient also had a history of kidney transplantation for his renal cell carcinoma and has been treated with immunosuppressants, including prednisone, CellCept (mycophenolate mofetil) and parpmune (sirolimus). His father had died with VHL complications.

Macroscopic examination showed a whitish mass with focal hemorrhages around the optic nerve head and several smaller yellow-whitish lesions in the peripapillary retina (Fig. 1c, d). There were thick pre-retinal fibrous tissues with areas of hemorrhages in the vitreous (Fig. 1c, d); the retina outside the mass was detached (Fig. 1c). Small round yellowish chorioretinal lesions and diffuse yellowish crystal lipid deposition were observed in the retina.

Microscopic examination disclosed a large RH located at the optic nerve head, extending to the nasal and temporal peripapillary retina and involving the entire neuroretina (Fig. 1e). The tumor was mainly composed of clear polygonal cells with small nuclei and finely reticulated or foamy cytoplasm. Focal tortuous, thin-walled, capillary-like vessels and small hemorrhages were intermingled in the RH. Within the RH, there were many small and large areas of cystic degeneration, some of which contained a few retinal pigment epithelium (RPE), pigment granules, fibrins, and/or erythrocytes. Some macrophages (CD68 positive, Fig. 1f) and T lymphocytes (CD45RO positive, Fig. 1g) infiltrates were visible in the RH, while B lymphocytes (CD20 positive) were rarely seen. The retina also showed degeneration, disorganization and gliosis (GFAP positive, Fig. 1h). Secondary exudative retinal detachment was noted in the retina adjacent to the hemangioblastoma. A large continuous chorioretinal adhesion with focal inflammatory choroidal infiltrates was also seen in the peripapillary retina. Several other small and large peripheral RHs with similar features involved the neuroretina. Many erythrocytes, scattered lymphocytes and pigmentladen macrophages were mixed with condensed vitreous strands.

#### **Patient 2**

A 40-year-old female had a history of visual changes when she was pregnant with her son in 2000. Her right eye presented with vascular lesions at the optic nerve head surrounding with lipid exudates. Her DNA testing was positive for VHL. In 2004, her vision had continued to decline in the right eye. In 2006, fundus examination showed increasing lipid exudates and focal retinal detachment temporal disc. At her annual VHL screening, she was detected to have a cystic lesion in the right kidney. She continued to have a "questionable" hemangioblastoma in her thoracic spinal cord (T4 and T5). Her son also tested positive for VHL. She received 14 ranibizumab injections from February 2005 to February 2006 as part of the intravitreal trial. Despite the treatment, her ocular lesions progressed to total retinal detachment, developed rubeosis iridis and early phthisis bulbi. The painful blind right eye was enucleated in 2009. A portion of the eye was submitted to NEI.

Macroscopic examination of the specimen showed retinal tumors, total retinal detachment as well as extremely disorganized macula and optic nerve head. Focal hemorrhages were also seen.

Microscopic examination showed rubeosis iridis. Small amount of hemorrhages and fibrins were seen in the vitreous. The retina was totally detached in a funnel shape. The retinal stock completely lost its normal architecture and was replaced by glial tissues. There were a few nodules formed by RH cells, fibroglial tissue, hemosiderin-laden pigmented macrophages, rare giant cells and small vessels. Scattered T lymphocyte (CD3 positive) and moderate macrophage (CD68 positive) infiltrates were visualized in the RHs and

disorganized retina, while B-lymphocytes (CD20 positive) were not seen. The temporal (funnel mouth) peripheral retina was barely recognized, closely attached to an organized blood and vitreous fibrovascular membrane. The nasal peripheral retina (funnel mouth) had slightly better retinal structure of gliosis (GFAP positive) and cystic degeneration. A small focal foamy cellular aggregation (RH) and scattered erythrocytes were observed.

#### **Patient 3**

A 35-year-old female with VHL diagnosed in 1983 had a history of enucleation of the left eye in 1991 and intracranial surgery in 1993. In 2004, she presented with a blind severely painful right eye in spite of conservative therapies. Ocular examination showed no view into the posterior segment of the right eye because of a dense cataract and rubeosis iridis. No anti-VEGF therapy was administered to this patient. The right eye became blind and was enucleated in 2004. The frozen eye was submitted to NEI.

Macroscopy showed that the retina containing several RHs and was totally detached and adherent to the posterior lens. The subretinal space was filled with orange-yellowish fluid. There was a thin layer of bony structure beneath the detached retina and the posterior choroid (inferior > superior). No optic disc was visible due to total retinal detachment.

Microscopy showed the total detached retina with barely visible architecture; degenerative cysts, gliosis (GFAP positive) and several isolated foci of vascular abnormalities were also seen. Some of these vascular abnormalities were typical VHL-associated RHs consisting of thin-walled, capillary-like or small hyalinized vascular walls forming an anastomosing pattern separated by a few plump, vacuolated cells. The other vascular abnormalities were mostly occluded hypocellular vascular structures. Areas of ghost erythrocytes and large cholesterol clefts were also present in the detached retina. Bone formation was noted in the subretinal tissue above and/or in the posterior choroid. Focal lymphocytic infiltration is seen in the choroid.

#### **HIF1**α **and HIF2**α **expressions in RHs**

 $HIF1A$  transcript detected by qRT-PCR was expressed in all retinas of the three patients, especially in Patients 1 and 2 (Fig. 2a). Similarly, HIF1α protein was positively expressed in some VHL-associated RH cells of all retinas in the three patients (Fig. 2b, c, d). In contrast, HIF2A transcript expression was significantly higher in Patients1 and 2, who had retinal inflammatory infiltration and were resistant to anti-VEGF therapy (Fig. 3a). HIF2α protein was also abundant in the VHL-associated RH cells of these two patients compared to that in Patient 3 (Fig. 3b, c, d). HIF2A gene mutation was not detected in the tissues of all three patients.

#### **Ultrastructure of RH**

The viable VHL tumor cells appeared undifferentiated in the three cases. Most VHLassociated RH cells contained large vacuoles, degenerated cytoplasm and abnormal mitochondria (Fig. 4a, b). Some VHL cells contained autophagosomes, glycogen aggregations, lipid droplets and chromatin disintegration; and others appeared apoptotic (Fig. 4c, d). Activated glial cells were intermingled with VHL-associated RH cells.

# **Discussion**

Our study provides molecular evidence that the VHL-associated RH cells express HIF2α in two patients resistant to anti-VEGF therapy. Furthermore, histopathological findings identify the unusual presence of inflammatory infiltrates in the RH cells in these two patients. These suggest that high levels of HIF2α could be another key element that is directly or indirectly associated with development of highly aggressive vascular tumors. Moreover, ultrastructural examination revealed the presence of autophagy, mitochondrial dysfunction, lipid metabolic abnormality and cytoplasmic degeneration in the RH cells. One or all of the pathological and molecular features of the patients may be associated with the refractory VHL clinical course and their resistance to multiple anti-VEGF and radiation therapies.

The pVHL encoded by the *VHL* gene is ubiquitously expressed and regulates the ubiquitination of HIF, thus marking it for degradation [21]. Under hypoxic conditions, intracellular levels of HIF are upregulated and result in increased expression of several proteins reversing hypoxia, such as erythropoietin, VEGF and PDGF [22]. Inactivation of VHL protein causes constitutive stabilization of HIF and ultimately leads to VHL-associated tumor progression. Our previous studies have already illustrated upregulation of VEGF in VHL-associated human RH specimens [4, 19]. Although VEGF is regulated by both HIF1α and 2α subunits, suppression of HIF2α activity is thought to be more critical than HIF1α activity for tumor suppression by pVHL [23, 24]. In Vhl deficient mouse liver, HIF2α is the dominant HIF in the pathogenesis of VHL-associated vascular tumors [9]. Mouse embryos originating from *Hif2a* deficient embryonic stem cells display severe vascular defects [25]. A significant correlation between high VEGF and HIF2α protein levels was found in human neuroblastoma specimens and HIF2α stabilization was detected in well-vascularized regions of the neuroblastoma patient specimens [26]. High HIF2α protein levels have even been correlated with advanced clinical stage and high VEGF expression [26]. Recently, somatic mutations in HIF2A are reported in patients with paragangliomas or somatostationomas together with poycythema [27]. These mutations disrupt the prolyl hydroxylation domain (PHD) of HIF2A and abolish the modification by PHD, resulting in the decrease of VHL recognition and VHL-associated degradation, as well as the stabilization of HIF2α [28].

In our current study, upregulation of HIF2α protein without HIF2A mutation is specifically detected in the two patients who were resistant to anti-VEGF therapy. This result may indicate a feedback loop between HIF2α and anti-VEGF therapy. That is, the VHLassociated RHs cause HIF2α upregulation, which results in high VEGF expression. Consequently, these high VEGF expression tumors are designated to anti-VEGF therapy, which in turn ends with even higher HIF2α expression. Except for the potential contribution of anti-VEGF therapy, long-standing clinical course and combined secondary complications in these VHL patients could lead to upregulation of HIF2α. The fact of upregulated HIF2α also raises another question as whether its elevation would result in resistance of anti-VEGF therapy. However, there is no solid evidence that HIF2α elevation would directly cause resistance of anti-VEGF therapy. The recent IVAN study has not found relationship between HIF2A SNP alteration and treatment responsiveness to VEGF inhibition in patients with neovascular age-related macular degeneration [29].

HIF2α is not only a critical factor regulating erythropoiesis and neovascularization, but also involved in lipid homeostasis and glucose metabolism in the liver [30, 31]. Quantitative analysis of glycogen content in the adult mouse liver revealed that Vhl inactivation leads to abnormal glycogen accumulation in hepatocytes [30]. This is in agreement with our ultrastructural findings of glycogen particles in the RH cells. Additionally, Rankin et al. have found that HIF2α activation in mouse hepatocytes suppresses lipid synthesis and fatty acid β-oxidation, and promotes lipid accumulation [31]. This could explain the presence of these lipid droplets in the VHL tumor cells.

In addition to the molecular upregulation of HIF2α, ultrastructural abnormalities in the RH cells were also prevalent. Most VHL-associated RH cells revealed cytoplasmic degeneration, abnormal mitochondria, autophagy, and nuclear chromatin disintegration, some of these cellular damages could have resulted from radiation therapy. The finding of autophagy or apoptosis in the RH could be due to previous anti-VEGF and/or radiotherapy. Autophagy is a process of degradation and recycling of proteins and intracellular components in response to starvation or stress [32]. With its duality of function, autophagy could either promote cell survival under metabolic situation or cause cell death under stressed conditions [33]. It can even protect certain cancer cells against anticancer treatments by blocking the apoptotic pathway [32]. Based on the clinical findings and ultrastructural features, autophagy in the VHL-associated RHs might indicate a causative role, which promotes growth and progression of these RH cells and contributes to the tumor aggressiveness.

Macrophages and T lymphocytes were observed in the tumor areas in two patients (Patients 1 and 2) who had an aggressive course of the disease. Chronic inflammation, although mild, is an uncommon presentation in VHL-associated RHs. Although some studies show that systemic and local anti-VEGF treatment had anti-inflammatory effects [34–36], preventive VEGF inhibition has been reported to worsen inflammation in a mouse model of acute colitis [37]. Evidence has also demonstrated that tumor resistance to anti-VEGF therapy can accelerate myeloid cell infiltration, which is strongly associated with tumor hypoxia [38]. Myeloid cells can be attracted to regions of tumor hypoxia and play a role in mediating resistance to antiangiogenic therapy [39, 40]. Further investigation is needed to explore the effects of anti-VEGF therapy on the inflammatory cellular infiltration and tumor hypoxia in VHL-associated RHs. The potential role of anti-inflammatory therapy remains to be investigated.

Both ultrastructural and molecular findings demonstrate potential factors that may lead to the refractory course of VHL patients and resistance to both conventional and anti-VEGF therapies. Several antiangiogenic agents, such as SU5416 (inhibitor of VEGF receptor 2), bevacizumab, ranibizumab, pegaptanib (an aptamer inhibits VEGF isoform 165), have been attempted in several clinical trials. However, the general efficacy of antiangiogenic agents in VHL-associated RHs is uncertain and even unfavorable [15]. Our new finding of HIF2α expression in the resistant RH cells suggests a novel therapeutic approach of targeting HIF2α as an alternative therapy for VHL-associated RHs.

# **Disclosure/Conflict of Interest**

The National Eye Institute Intramural Research Program supported the study.

The authors declare that they have no conflict of interest.

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#### **Fig. 1.**

Clinical photos and photomicrographs in Patient 1 with von Hippel-Lindau (VHL) associated retinal hemangioblastoma (RH). Fundus photographs of VHL-associated RH in the right eye taken in January 2005 (**a**) and March 2010 (**b**). **a** The baseline fundus photograph before the intravitreal ranibizumab trial showed a prominent optic nerve tumor (arrow), focal hemorrhages and yellowish retinal hard exudates. **b** Ranibizumab and radiation therapy shows multiple vitreous hemorrhages, enlarged optic nerve tumor, serous retinal detachment, lipid accumulation and multiple pigmentary/yellowish chorioretinal lesions in the retina. Macroscopic photographs of VHL-associated RH in the right eye after enucleation in January 2012 (**c**, **d**). **c** The inferior calotte of the globe showed several patches of vitreous hemorrhages, yellowish chorioretinal lesions and retinal detachment. **d**. A whitish mass with focal hemorrhages around the optic nerve head (juxtapapillary RH) and several smaller lesions are observed in the peripapillary retina. **e** A large juxtapapillary RH was noted at the optic nerve head with a huge cyst (C) (hematoxylin and eosin (H&E) stain, original magnification,  $\times$ 100). Avidin-biotin-complex immunohistochemistry staining corresponding to the same section with panel shows CD68 positive macrophages (**f**) and CD45RO positive T lymphocyte (**g**) infiltration in the juxtapapillary RH (original magnification, ×100). **h** Marked gliosis (glial fibrillary acidic protein positive) occurs in the whole areas of hemangioblastoma and retina (Avidin-biotin-complex immunohistochemistry, original magnification, ×200)

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# **Fig. 2.**

HIF1α expression in von Hippel-Lindau-associated retinal hemangioblastoma (RH) in 3 patients. **a** HIF1A transcript is comparably upregulated in the RH of 3 patients. Photomicrographs of HIF1α protein expression in the RH cells in Patients 1(**b**), 2 (**c**) and 3 (**d**) (Avidin-biotin-complex immunohistochemistry, original magnification, ×200)

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# **Fig. 3.**

HIF2α expression in von Hippel-Lindau-associated retinal hemangioblastoma (RH). **a**  HIF2A transcript expression is higher in the RH of Patients 1 and 2 than Patient 3. Photomicrographs of HIF2α protein expression in the RH cells in Patient 1 (**b**) and Patient 2 (**c**), but not in Patient 3 (**d**) (Avidin-biotin-complex immunohistochemistry, original magnification, ×200)



#### **Fig. 4.**

Transmission electron microscopy photomicrograph of von Hippel-Lindau-associated retinal capillary hemangioblastoma (RH) cells of Patient 1 (original magnification, ×2500). Ultrathin sections were prepared and double-stained with uranyl acetate and lead citrate. **a** A typical RH cell has degenerated cytoplasm and a small nucleus (N). **b** Degenerative cytoplasm autophagosomes (arrows) and damaged mitochondria (asterisks) in a RH cell (N, nucleus). **c** Glycogen aggregations (asterisks) and autophagosomes arrows) are noted in RH

cells (N, nucleus). **d** Lipid droplets (asterisks) and nuclear chromatin disintegration are noted in a RH cell (N, nucleus)

# **Table 1**

# Brief summary of the clinical features in three VHL patients



 $a<sub>T</sub>$ The enucleated eye presented in this study.