

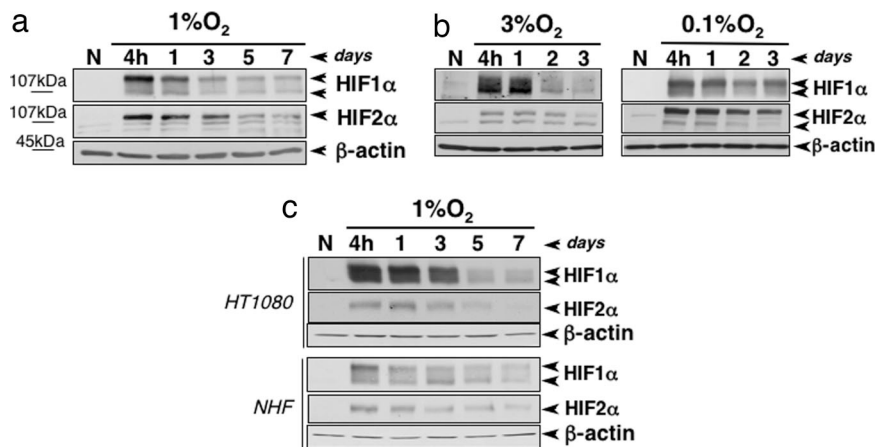


Fig. 1. Levels of HIF α proteins decline during chronic hypoxia. Cells were incubated in hypoxic conditions for different periods of time, and the levels of HIF1 α , HIF2 α , and β -actin (loading control) were analyzed by Western blotting. (a) HeLa cells at 1% O $_2$. (b) HeLa cells at 3% and 0.1% O $_2$. (c) HT1080 and NHF cells at 1% O $_2$.

deficient cell line, RCC4, under hypoxia at 1% O₂ for 4 h to 4 days and looked for the HIF α proteins. In these cells, HIF α level never declined, whereas in the stable clone RCC4-pVHL (RCC4 in which pVHL had been reintroduced) HIF α dramatically decreased after 1 day at 1% O₂ (Fig. 2b). Finally, we investigated HIF α protein hydroxylation as the signal allowing pVHL binding, and we observed that the addition of the hydroxylase inhibitor, dimethylallyl glycine (DMOG), restored HIF α accumulation across hypoxic stress (Fig. 2c). Accordingly, the HIF1 α construct mutated on the two proline residues (Pro⁴⁰² and Pro⁵⁶⁴) remained stable upon chronic hypoxia (Fig. 2d). Therefore, during chronic hypoxia, HIF α levels decline because of increased protein instability. Indeed, HIF α proteins appear to be hydroxylated, ubiquitinated, and targeted through the proteasome, suggesting that PHDs are “active” despite the hypoxic conditions.

Chronic Hypoxia Overactivates the Three PHDs. To evaluate PHD activity during chronic hypoxia, we performed *in vitro* pVHL capture assays. HeLa cells were exposed to hypoxia at 1% O₂ for 4 h up to 7 days. GST-HIF1 α constructs were incubated with the cell lysates and thereafter with the radio-labeled pVHL protein. Because pVHL binds to HIF α only when the relevant proline residues have been previously hydroxylated by the PHDs, the binding of *in vitro*-translated [³⁵S]pVHL to GST-HIF1 α is accepted as a reliable measure of PHD activity (Fig. 3*a Inset*). Compared with normoxia, as soon as 24 h of 1% O₂, PHDs appeared overactivated (Fig. 3*a*, lane 3). Furthermore, from day 1 to day 7, the assay highlighted a stronger and progressive activation of the PHDs (Fig.

3a). PHD activation was indeed observed with the recombinant GST-HIF1 α P402A mutant constructs but not with P564A or P402/564A, meaning that P564 hydroxylation is absolutely required and limits HIF1 α hydroxylation (data not shown). Because PHD activity could be caused by an increase in the pool of these proteins, we quantified the level of the hypoxia up-regulated PHD2 and PHD3 isoforms. As expected, both proteins are induced under hypoxia but their maximal induction were reached after 24 h and the proteins were no longer increased in subsequent days (Fig. 3b). This result strongly suggests that in addition to an increase in the amount of PHD proteins, chronic hypoxia overactivates these enzymes.

We investigated the specific contribution of each PHD isoform on the HIF α degradation pathway by using the siRNA approach. During hypoxic kinetics at 1% O $_2$ and at two different points (on days 1 and 3 according to our PHD activity assay), PHDs isoforms were invalidated alone or in combination. Fig. 3c shows that at day 1 and consistent with our previous work (8), only PHD2 silencing leads to HIF1 α stabilization (lane 3). For HIF2 α , in addition to PHD2, we observed a slight contribution of PHD1 (Fig. 3c, lanes 2 and 3). Surprisingly, at day 3, not only PHD2 but also PHD3 silencing rescued HIF α protein accumulation (Fig. 3c, lanes 3 and 4). Moreover, cosilencing of PHDs suggests that, in combination with the other isoforms, PHD1 acts on HIF α regulation (Fig. 3c, lanes 5, 6, and 8). We conclude that, in contrast to normoxia, during chronic hypoxia, the three PHDs contribute to HIF α degradation.

Chronic Hypoxia Increases O₂ Availability for PHDs. Hagen *et al.* (9) showed that NO and other chemical inhibitors of mitochondrial

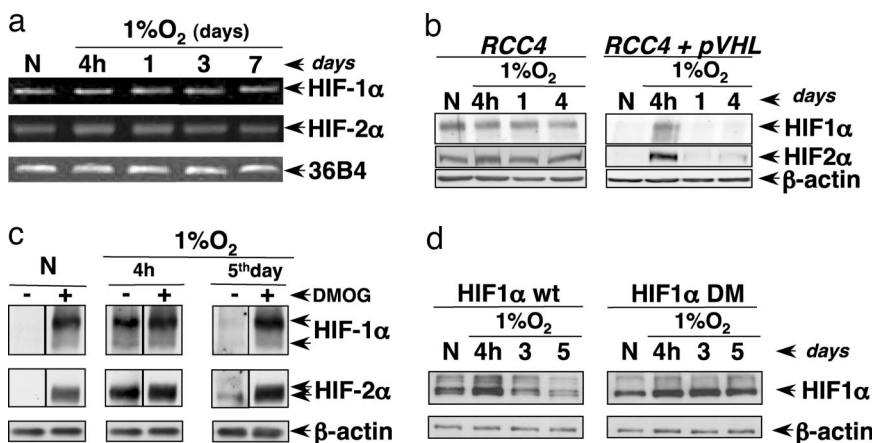


Fig. 2. Chronic hypoxia promotes HIF α protein degradation depending on the hydroxylation–pVHL–proteasome pathway. (a) Hypoxic kinetics in HeLa cells at 1% O₂. HIF-1 α , HIF-2 α , and 36B4 mRNA levels were analyzed by RT-PCR. (b) RCC4 cells (pVHL-deficient) or the stable clone RCC4-pVHL (which overexpresses pVHL) were incubated in normoxia (20% O₂) or hypoxia (1% O₂) for different periods of time. (c) HeLa cells were incubated in normoxic (20% O₂) or hypoxic conditions (1% O₂) for 4 h or 5 days, and 4 h before the lysis, cells were incubated in the presence or absence of DMOG (1 mM). The levels of HIF1 α , HIF2 α , and β -actin (loading control) were analyzed by Western blotting. (d) 293 cells were incubated in normoxia (20% O₂) or hypoxia (1% O₂) for different periods of time, and 48 h before the lysis, cells were transfected with the HIF1 α WT or DM constructs. The levels of HIF1 α were analyzed by Western blotting using the Myc-tag antibody.

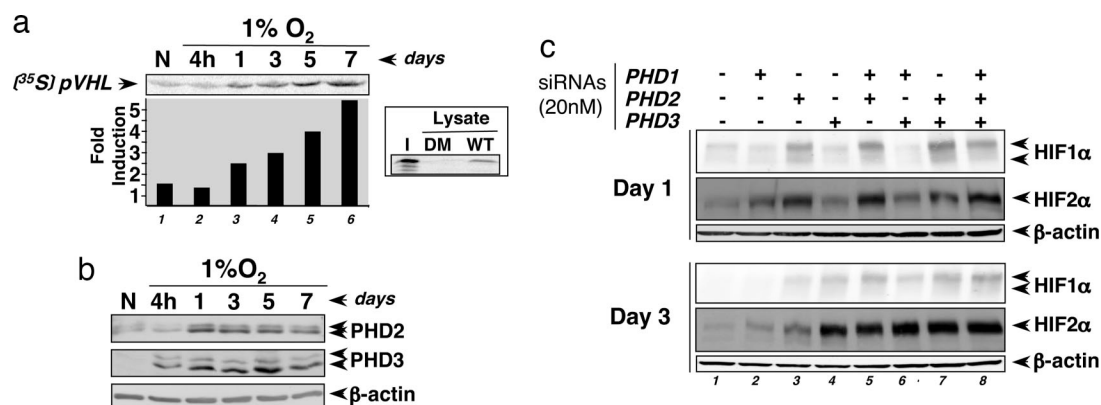


Fig. 3. Chronic hypoxia overactivates the three PHDs. (a) Autoradiogram showing PHD activity during hypoxic kinetics in HeLa cells at 1% O₂. (Inset) [³⁵S]pVHL input (I) and pVHL binding using the P402/564A mutant (DM) or the WT GST-HIF1 α constructs. (b) Western blot analysis showing PHD2 and PHD3 protein levels during hypoxic kinetics. (c) To evaluate the specific contribution of each PHD isoform, 48 h before extraction, cells were transfected with siRNAs as indicated. The levels of HIF1 α , HIF2 α , and β -actin were analyzed by Western blotting.

respiration prevent hypoxia-induced HIF1 α stabilization. Indeed, because mitochondrial respiration pumps most of the intracellular O₂, its inhibition increases intracellular O₂ availability. Moreover, PDK1 (pyruvate dehydrogenase kinase), which is a HIF1 α dependent gene product, has been recently reported as a natural inhibitor of mitochondrial activity in hypoxia (10, 11). Based on these results and because we observed that HIF α desensitization did not occur in drastic hypoxic conditions (0.1% O₂; Fig. 1*b*), we assumed that chronic hypoxia by inhibiting mitochondrial activity increases intracellular O₂ availability. We further decided to use GSK3 cells, a respiratory-deficient cell line (12). Indeed, we hypothesized that in GSK3 cells, in which O₂ cannot be redistributed because of the constitutive inhibition of mitochondrial inhibition, we would prevent PHDs reactivation and thus chronic hypoxia-induced HIF α desensitization. According to our hypothesis, we observed that HIF1 α protein, which accumulates after acute hypoxia, was not degraded during chronic hypoxia in GSK3 cells unlike in the control cells (Fig. 4). Therefore, we conclude that chronic hypoxia, by inhibiting mitochondrial respiration, progressively increases O₂ availability for PHDs, leading to their overactivation and thus triggering HIF α desensitization.

Chronic Hypoxia-Induced HIF α Desensitization Protects Cells from Necrosis. To elucidate the physiological relevance of the intracellular O₂ redistribution during chronic hypoxia, we exposed GSK3 cells to acute or chronic 1% O₂ hypoxia and measured cell death by determining the percentage of permeabilized (trypan blue-positive) cells. As shown in Fig. 5*a Left*, in chronic hypoxia the GSK3 cells (deficient for HIF α desensitization) showed a drastic increase in cell death compared with the control cells fully competent to degrade HIF α . Indeed, almost 100% of GSK3 cells died

after 7 days, suggesting that HIF α degradation is essential for cell survival in chronic hypoxia.

To support our hypothesis and confirm the role of PHDs, we performed hypoxic kinetics in HeLa cells and forced sustained HIF1/2 α expression by silencing the PHDs. As we expected, PHDs silencing-mediated HIF α stabilization during chronic hypoxia led to massive cell death (Fig. 5a *Right*). To definitively prove that HIF α subunits mediate this process, we cosilenced the PHDs and HIF α subunits (HIF1 α and HIF2 α) at the same time upon chronic hypoxia. Interestingly, silencing of HIF α subunits completely blocks PHD silencing-mediated cell death (Fig. 5b *Right*). Similar results were obtained when we silenced HIF1 α expression in GSK3 cells upon chronic hypoxia (Fig. 5b *Left*). Therefore, we conclude that HIF α desensitization induced by chronic hypoxia-mediated PHD overactivation is required to prevent cells from cell death.

To characterize this cell death, we analyzed by flow cytometry the proportion of apoptotic cells represented by the sub-G₁ population stained for propidium iodide (PI). We measured a slight and nonsignificant increase in the number of apoptotic cells in GSK3 cells or PHD-inactivated cells after chronic hypoxia (Fig. 6 and data not shown). Furthermore, Western blot analysis showed no cleavage of the apoptotic marker, poly(ADP-ribose)polymerase (PARP), and LC3 lipidation was not involved in the autophagic process during the chronic hypoxia (Fig. 6 and data not shown). Taken together, we conclude that during chronic hypoxia, HIF α desensitization is required to prevent cells from a massive cell death that is largely independent of the apoptotic and/or autophagic pathways. Moreover, as a feedback survival mechanism, chronic hypoxia-induced PHD overactivation, by ensuring HIF α desensitization, escapes cells from necrosis.

Mice Adapt to Chronic Hypoxia by Overactivating PHDs to Desensitize HIF α . To confirm our results *in vivo*, we exposed mice to hypoxia at 8% O₂ for 6 h (acute) and 24 h (chronic). HIF1 α protein level in the kidney (Fig. 7a), the brain, and thymus (data not shown) of mice was then analyzed by immunohistochemistry. We observed an HIF1 α increase in the tissues of mice exposed to 6 h hypoxia but none in the 24 h group. Nevertheless, when these mice were exposed for an additional 2 h at more drastic hypoxia (6% O₂), we observed a reaccumulation of HIF1 α protein (Fig. 7a). Therefore, we hypothesized that mice, as cells, “adapt” to chronic hypoxia by increasing the rate of HIF1 α degradation. To test whether the HIF1 α desensitization we observed in mice tissues was caused by PHD overactivation, we performed a pVHL capture assay. Fig. 7b shows that after 6 h of hypoxia PHDs are less active compared with the control group. However, after chronic hypoxia of 24 h, the assay

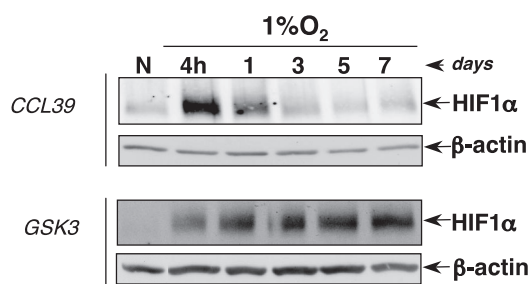


Fig. 4. Intracellular O₂ redistribution is required for PHD overactivation. Hypoxic kinetics in CCL39 and GSK3 cells at 1% O₂ are shown. HIF1 α and β -actin levels were analyzed by Western blotting.

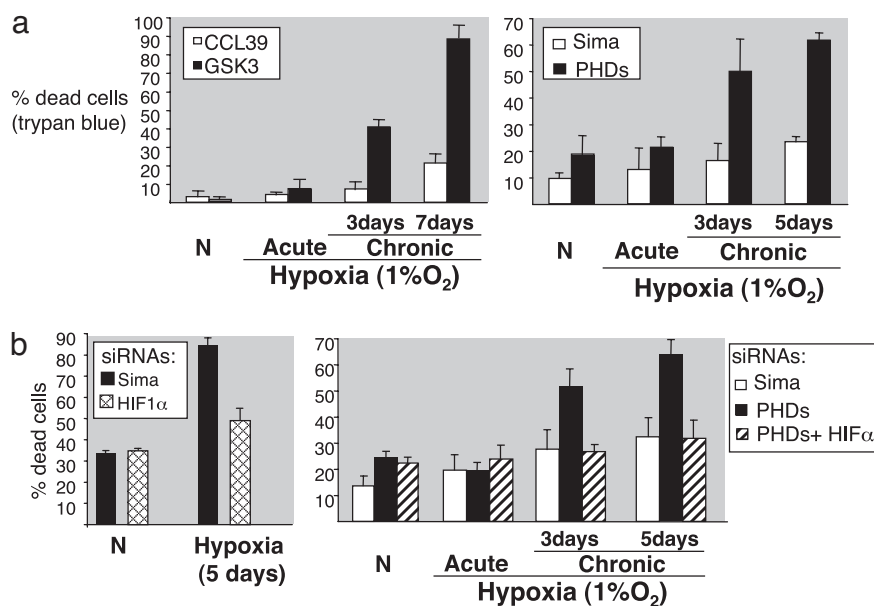


Fig. 5. Chronic hypoxia-induced HIF α desensitization protects cells from necrosis. Cell death was measured in cells incubated in hypoxia at 1% O₂ for different periods of time by a trypan blue incorporation assay. (a) CCL39 and GSK3 cells (Left) or HeLa cells transfected with siRNA targeting the three PHDs isoforms or an irrelevant sequence (Right) for 5 days and incubated either in normoxia (N) or upon hypoxia for different periods of time (4 h, 3 days, or 5 days). (b) GSK3 cells (Left) and HeLa cells transfected as described in a (Right) in which HIF α expression has been silenced. The siHIF α transfections started 16 h after the cells were incubated upon hypoxia and were carried on every day until the end of the experiment for days 3 and 5. In the case of the N and H (4 h) conditions, siHIF α were transfected during the last 24 h.

reveals a reactivation of the hydroxylases. Moreover, the autoradiogram shows less pVHL binding, reflecting a new decrease in the activity of the PHDs, in mice subjected to an additional and more severe hypoxic exposure (2 h at 6% O₂). These results showed a perfect correlation between the expression of HIF1 α and the inhibition of PHDs activity *in vivo*. Finally, to confirm the implication of PHDs in this mechanism, we used the siRNA approach *in vivo*. Mice were injected with siRNAs targeting the three PHDs or an irrelevant sequence before the hypoxic exposure, and HIF1 α protein expression was analyzed (Fig. 7c). As previously observed in the control group, we did not detect HIF1 α protein after 24 h at 8% O₂, whereas in mice injected with siRNAs silencing the three PHDs isoforms, we observed a clear stabilization of HIF1 α protein

similar to the desferrioxamine (DFO)-treated animals (Fig. 7c). These data reinforced our *in cellulo* experiments, showing that mice, which accumulate HIF α protein during acute hypoxia, also adapt to chronic hypoxia by overactivating PHDs to desensitize HIF α .

Discussion

A perfect balance between O₂ delivery and demand in tissues (O₂ homeostasis) is essential for survival. When O₂ balance is disrupted, organisms have developed adaptive mechanisms to restore the equilibrium and go out of danger. The transcription factor HIF is at the heart of these mechanisms implicated in several physiopathological processes such as cancer or ischemia.

HIF α subunits are tightly regulated by O₂. When O₂ is restricted, the high turnover rate of the HIF α subunits allows their rapid stabilization and activation. Nevertheless, we show here that chronic hypoxic stress desensitizes HIF α proteins. Our results agree with previous reports showing that HIF1 α proteins vanish with prolonged hypoxia (13) and extend this regulatory pathway to HIF2 α . Indeed, in contrast with previous data (14, 15), HIF2 α behavior appears very close to that of HIF1 α . Furthermore, we demonstrate that HIF1/2 α decline is observed in a large range of O₂ levels and cell types (Fig. 1 and data not shown), supporting HIF α desensitization during chronic hypoxia as a general adaptive mechanism.

Analysis of the HIF α mRNAs did not show any variation in acute or chronic hypoxia, ruling out the putative contribution of the antisense HIF1 α previously described (16). However, we show that the bona fide hydroxylation–ubiquitination–proteasome pathway mediates HIF α protein decrease as recently reported by Stiehl *et al.* (13). Furthermore, we demonstrate an unexpected and gradual PHD overactivation across chronic hypoxia despite low global O₂ availability (Fig. 3a). P⁵⁶⁴ remains the first hydroxylated residue, and mutation on this proline results in the complete loss of P⁴⁰² hydroxylation (data not shown). Moreover, siRNA experiments reveal that, in contrast to normoxia, the three PHD isoforms contribute to HIF α degradation and hence to the HIF α desensitization mechanism set up during chronic hypoxia (Fig. 3c). However, these results point out differences in the affinity of each PHD isoform for HIF1 α and/or HIF2 α that need further work to be clarified.

Accumulation of PHD2 and PHD3, could account, at least partially, for the increase in the overall PHD activity. However, PHD2 and PHD3 analysis revealed no additional increase in the

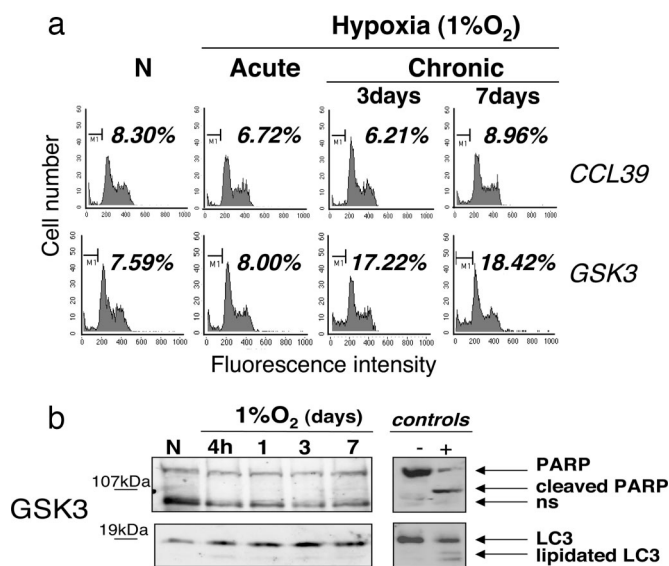


Fig. 6. Chronic hypoxia-induced cell death is independent of the apoptotic and/or autophagic pathways. Cells were incubated in hypoxia (1% O₂) for different periods of time. (a) Measurement by flow cytometry of the sub-G₁ PI-positive population in CCL39 and GSK3 cells. (b) PARP and LC3 proteins were analyzed by Western blotting on GSK3 extracts. Staurosporin (1 μ M) was used as an apoptotic cell death activator. Nutrient-starved cells were used as a positive control for autophagy.

favours or, in contrast, suppresses tumor progression, wound healing, or ischemic diseases may yield insights into the suitable use of PHDs as an attractive therapeutic strategy for those pathologies in which hypoxia is implicated.

Methods

Cell Culture. HeLa, HT1080, NHF, LS174, CAL51, 293, CCL39, and GSK3 (12) cells were grown in DMEM with 7.5% FBS. RCC4 cells and the stable clone RCC4+pVHL were generated were grown in DMEM/Ham's F10 (1:1) medium with 10% FBS. NHK cells were grown in a keratinocyte serum-free medium (Invitrogen) supplemented in L-glutamine, recombinant EGF (0.2 μ g/ml) and bovine pituitary extracts (30 μ g/ml).

Hypoxic conditions were performed by incubation of cells in a "bug box" anaerobic workstation (Ruskin Technologies). The oxygen concentrations were fixed with a gas mixture containing 5% carbon dioxide and balance nitrogen. For hypoxic kinetics, cells were split sparsely (to reach 60% of confluency at the end of the experiment) 3 days before lysis, outside or inside the bug box depending on the timing, and cell medium was changed every day. Cell extraction was performed inside the bug box to avoid reoxygenation.

Transfections were carried out as described (8). siRNAs were purchased from Eurogentec. Sequences of the irrelevant siRNA and the human HIF1 α and PHDs have been reported (8). The same regions were targeted by the mouse siPHDs sequences. siRNA sequence targeting HIF2 α (GenBank accession no. NM.001430) corresponds to regions 1590–1609 relative to the start codon. The HIF1 α expression vector mutated on Pro⁴⁰² and Pro⁵⁶⁴ [HIF1 α double mutant (DM)] was generated by mutagenesis using the QuikChange Site-Directed Mutagenesis kit (Stratagene).

Animal Studies. C57BL/6 mice (8-week-old females) were subjected to systemic hypoxia in the anaerobic workstation at 25°C and in an humidified atmosphere (95%) supplied with the gas mixture allowing it to adjust to the hypoxic environment by gradually decreasing the O₂ level from 20.9% to 8% and 92% N₂ during an adaptation time of 2 h 30 min. The mice were provided with food and water ad libitum. For the siRNA experiment, mice were i.p.-injected for 7 days every day with 3 μ g of each siRNA by using jetPEI (Polyplus Transfection). The animal study protocols were conducted according to institutional guidelines for animal use. Three mice were used for each experimental condition and experiments were reproduced twice. Mice were killed inside the bug box. Half of the dissected organs were frozen and stored at –80°C (to measure PHDs activity), and the rest were fixed in 4% formaldehyde.

RT-PCR, Western Blotting, and Immunohistochemistry. RNA was purified with the RNeasy Mini Kit (Qiagen). The oligonucleotide sequences for RT-PCR are available on request.

For Western blot analysis, cells were lysed in Laemmli buffer. Protein concentration was determined with Lowry assay, and 50 μ g of cell extracts was resolved

by SDS/PAGE and electrophoretically transferred onto a PVDF (Millipore) or nitrocellulose membrane (Amersham Biosciences). Immunoreactive bands were visualized with the Amersham Biosciences ECL system.

HIF1 α immunohistochemistry was performed as described (21). Immunohistochemical studies were analyzed with a Leica Zeiss inverted microscope (Axiovert 200 M, $\times 40$ objective) equipped with a CoolSnap HQ cooled CCD camera (Roeper Scientifique).

Antibodies. The anti-HIF1 α has been characterized (22). The Ab to PHD2 (anti-serum 923) was raised in rabbits immunised against the first 15 N-terminal amino acids of human PHD2. Antibodies recognizing β -actin (Sigma), PHD3 (Interchim and Novus Biologicals), HIF2 α (Novus Biologicals), Myc (9B11; Cell Signaling), PARP (Biomol), and LC3 (MBL) were used.

pVHL Capture Assay. GST-HIF1 α WT was obtained by subcloning the 0.714-kb BamHI/XmaI PCR fragment of the pcDNA3-HA-HIF1 α plasmid (corresponding to the 1032–1746 nucleotides of the human HIF1 α sequence) into the BamHI/SmaI site of PGEX-6P-3. GST-HIF1 α P402A, P564A, and P402AP564A mutants were obtained by mutagenesis of the GST-HIF1 α construct with the QuikChange Site-Directed Mutagenesis Kit (Stratagene).

pVHL was *in vitro*-translated by using a TNT Transcription/Translation Kit (Promega) supplemented with Redivue L- [³⁵S]methionine (Amersham Biosciences) to produce [³⁵S]pVHL. The protocol used to cell extracts preparation and hydroxylation reaction has been described (23). It is important to note that cell lysis and hydroxylation assay of hypoxic extracts were performed inside the bug box. The same procedure was applied for tissue hydroxylation assay.

Cell Death and Apoptosis. Cell death was analyzed by using the trypan blue staining procedure (24).

For PI staining, total cell pellets were washed once with PBS and finally fixed in ice-cold 70% ethanol. The proportion of apoptotic cells was determined as described (25).

All experiments were repeated at least three times (excepted for the mice studies) with similar results. Representative immunostainings, blots, and autoradiograms are shown.

ACKNOWLEDGMENTS. We thank Dr. R. Buscà (Faculty of Medicine, University of Nice) for providing NHK and NHF cells and critically reading the manuscript, Dr. A. O. Hueber and her team for help with the death experiments, and Drs. P. Lenormand and E. Vial for their comments. Financial support was from the Centre National de la Recherche Scientifique, Ministère de l'Éducation, de la Recherche et de la Technologie, Ligue Nationale Contre le Cancer (Equipe Labellisée), Cancéropôle Provence-Alpes-Côte d'Azur, and Association pour la Recherche sur le Cancer (subvention 3693). E.B. is supported by Ministerio de Educación y Ciencia Grant SAF2007-64597, ETORTEK Research Program 05-07, and the Bizkaia Xede Program from Bizkaia County.

- Semenza GL (1998) *Curr Opin Genet Dev* 8:588–594.
- Ivan M, Kondo K, Yang H, Kim W, Valiando J, Ohn M, Salic A, Asara JM, Lane WS, Kaelin WG, Jr (2001) *Science* 292:464–468.
- Jaakkola P, Mole DR, Tian YM, Wilson MI, Gielbert J, Gaskell SJ, Kriegsheim A, Hebestreit HF, Mukherji M, Schofield CJ, et al. (2001) *Science* 292:468–472.
- Maxwell PH, Wiesener MS, Chang GW, Clifford SC, Vaux EC, Cockman ME, Wykoff CC, Pugh CW, Maher ER, Ratcliffe PJ (1999) *Nature* 399:271–275.
- Lando D, Peet DJ, Gorman JJ, Whelan DA, Whitelaw ML, Bruck RK (2002) *Genes Dev* 16:1466–1471.
- Berra E, Ginouves A, Pouyssegur J (2006) *EMBO Rep* 7:41–45.
- Appelhoff RJ, Tian YM, Raval RR, Turley H, Harris AL, Pugh CW, Ratcliffe PJ, Gleadle JM (2004) *J Biol Chem* 279:38458–38465.
- Berra E, Benizri E, Ginouves A, Volmat V, Roux D, Pouyssegur J (2003) *EMBO J* 22:4082–4090.
- Hagen T, Taylor CT, Lam F, Moncada S (2003) *Science* 302:1975–1978.
- Kim JW, Tchernyshyov I, Semenza GL, Dang CV (2006) *Cell Metab* 3:177–185.
- Papandreou I, Cairns RA, Fontana L, Lim AL, Denko NC (2006) *Cell Metab* 3:187–197.
- Pouyssegur J, Franchi A, Pages G (2001) *Novartis Found Symp* 240:186–196; discussion 196–188.
- Stiehl DP, Wirthner R, Koditz J, Spielmann P, Camenisch G, Wenger RH (2006) *J Biol Chem* 281:23482–23491.
- Holmquist-Mengelbier L, Fredlund E, Lofstedt T, Noguera R, Navarro S, Nilsson H, Pietras A, Vallon-Christersson J, Borg A, Gradin K, et al. (2006) *Cancer Cell* 10:413–423.
- Uchida T, Rossignol F, Matthey MA, Mounier R, Couette S, Clottes E, Clerici C (2004) *J Biol Chem* 279:14871–14878.
- Thrash-Bingham CA, Tartof KD (1999) *J Natl Cancer Inst* 91:143–151.
- Schofield CJ, Ratcliffe PJ (2004) *Nat Rev Mol Cell Biol* 5:343–354.
- Schofield CJ, Zhang Z (1999) *Curr Opin Struct Biol* 9:722–731.
- Gerald D, Berra E, Frapart YM, Chan DA, Giaccia AJ, Mansuy D, Pouyssegur J, Yaniv M, Mehta-Grigoriou F (2004) *Cell* 118:781–794.
- Tuckerman JR, Zhao Y, Hewitson KS, Tian YM, Pugh CW, Ratcliffe PJ, Mole DR (2004) *FEBS Lett* 576:145–150.
- Trastour C, Benizri E, Ettore F, Ramaoli A, Chamorey E, Pouyssegur J, Berra E (2007) *Int J Cancer* 120:1443–1450.
- Richard DE, Berra E, Gothie E, Roux D, Pouyssegur J (1999) *J Biol Chem* 274:32631–32637.
- Huang J, Zhao Q, Mooney SM, Lee FS (2002) *J Biol Chem* 277:39792–39800.
- Forcet C, Ye X, Granger L, Corset V, Shin H, Bredesen DE, Mehlen P (2001) *Proc Natl Acad Sci USA* 98:3416–3421.
- Cahuzac N, Baum W, Kirkin V, Conchonaud F, Wawrzyniec L, Marguet D, Janssen O, Zornig M, Hueber AO (2006) *Blood* 107:2384–2391.