

The dependence receptor Ret induces apoptosis in somatotrophs through a Pit-1/p53 pathway, preventing tumor growth

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Somatotrophs are the only pituitary cells that express Ret, GFR α 1 and GDNF. This study investigated the effects of Ret in a somatotroph cell line, in primary pituitary cultures and in Ret KO mice. Ret regulates somatotroph numbers by inducing Pit-1 overexpression, leading to increased p53 expression and apoptosis, both of which can be prevented with Ret or Pit-1 siRNA. The Pit-1 overexpression is mediated by sustained activation of PKC δ , JNK, c/EBP α and CREB induced by a complex of Ret, caspase 3 and PKC δ . In the presence of GDNF, Akt is activated, and the Pit-1 overexpression and resulting apoptosis are blocked. The adenopituitary of Ret KO mice is larger than normal, showing Pit-1 and somatotroph hyperplasia. In normal animals, activation of the Ret/Pit-1/p53 pathway by retroviral introduction of Ret blocked tumor growth *in vivo*. Thus, somatotrophs have an intrinsic mechanism for controlling Pit-1/GH production through an apoptotic/survival pathway. Ret might be of value for treatment of pituitary adenomas.

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Introduction

Final adult height and body composition are maintained in a constant interval for each given species. Somatotrophs, the pituitary cells secreting growth hormone (GH), are specialized secretory cells. However, they are not terminally differentiated: the pituitary retains some plasticity throughout life,

enabling appropriate responses to events such as puberty and pregnancy (Melmed, 2003). Somatotrophs play a key role in the control of growth during infancy and puberty, although they continue to secrete GH throughout adult life. Somatotroph number and function are tightly regulated, as GH hypersecretion leads to excessive growth in the form of either gigantism or acromegaly. Genetic factors leading to somatotroph hypoplasia are well known, but little is known about the mechanisms controlling physiological adjustments in somatotroph number. In this connection, the incidence of pituitary somatotroph adenomas is surprisingly low (about 3–4 cases per 10⁶ per year) compared with adenomas of other endocrine glands (Holdaway and Rajasoorya, 1999). More puzzling still is the consistent benignity of these tumors, which never metastasize (Melmed, 2003; Kaltsas *et al*, 2005).

We have recently demonstrated that somatotrophs are the only pituitary secretory cells expressing Ret, GDNF and GFR α 1, both in humans (Japon *et al*, 2002) and in rats (Urbano *et al*, 2000). This expression pattern is maintained in all somatotroph adenomas. Ret is a transmembrane receptor well known for its differentiation- and survival-controlling functions in epithelial cells, neurons and neuroendocrine cells. Alternative splicing produces two main isoforms: Ret long (Ret-L or Ret51), 1072 amino acids (aa) long, and Ret short (Ret-S or Ret9), 1114 aa long, with different sequences in the C-terminal region (51 aa long in Ret-L and 9 aa long in Ret-S) (Tahira *et al*, 1990). Ret has four coreceptors (GFR α 1, 2, 3 and 4), with ligands GDNF, NTN, ART and PSP, respectively. Extensive research on Ret function in mammals indicates that it has two main functions. First, it actively promotes survival, growth and extension/migration of cells such as neurons (neurite and axonal sprouting) and epithelial renal cells (branching). In these roles, Ret binds to its ligand GDNF and its GPI-anchored extracellular coreceptor GFR α 1, and this binding induces the tyrosine kinase activity of Ret, triggering various signal transduction pathways (Bredesen *et al*, 2004; Porter and Dhakshinamoorthy, 2004; Arighi *et al*, 2005). Second, Ret appears to induce apoptosis, but only in the absence of GDNF; to date, however, this effect has only been demonstrated in transient transfection of HEK293T and neuroblastoma cell lines (Bordeaux *et al*, 2000). In line with this, Ret has been included in the family of ‘dependence receptors’, together with p75NTR, DCC, UNC5H, PTC1 and the androgen receptor. In humans, mutations that activate Ret lead to multiple endocrine neoplasia type 2 (MEN 2), whereas inactivating mutations lead to Hirschprung’s disease (Santoro *et al*, 2004; Arighi *et al*, 2005).

Here, we report a study of Ret functions in a somatotroph cell line and in primary pituitary cultures. We uncover a pathway in which Ret, in the absence of GDNF, induces Pit-1 overexpression, leading to increased p53 expression and apoptosis. We identify the kinases involved in the induction

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of Pit-1 promoter overexpression by Ret. The cytoplasmic Ret portion (IC-Ret) complexes with caspase 3 and the kinase PKC delta, all becoming cleaved during the process. We also present results indicating that newborn Ret KO mice pituitaries are larger than normal, showing Pit-1-expressing cell and somatotroph hyperplasia. Finally, the Ret/Pit-1/p53 apoptotic pathway acts in an *in vivo* model of pituitary hyperplasia. In this model, retroviral Ret delivery suppressed estrogen-induced hyperplasia but did not alter normal pituitary function.

Results

Ret-induced apoptosis is associated with Pit-1 expression

To investigate the role of Ret in somatotrophs, we used a pituitary cell line, GH4C1, that does not express some of the characteristic receptors of normal somatotrophs (e.g. GHRH-R), but that nonetheless expresses the ghrelin receptor (NRL unpublished) and small amounts of Pit-1, and secretes some GH. This cell line does not express Ret but expresses both GFR α 1 and GDNF mRNA (data not shown) and the corresponding proteins (Figure 1A, left). We tried to produce transfectants stably expressing human Ret-L or Ret-S; colonies were obtained in various transfections, but in no case was Ret expression detected (Figure 1A). While transient transfection of GH4C1 or human embryonic kidney (HEK293) cells led to Ret expression (Figure 1A), Ret stable transfection had a deleterious effect on colony number after selection with neomycin (G418) (Supplementary Figure 1). In a previous study using the HEK293T cell line, transient transfection with Ret induced apoptosis that could be blocked by GDNF (Bordeaux *et al*, 2000). Similarly, in our somatotroph cells, transfection with Ret (either isoform) potently induced apoptosis within 24 h, whereas addition of GDNF (50 ng/ml) completely blocked this response (Figure 1B, left). zVAD, a broad caspase inhibitor, likewise blocked the apoptosis, suggesting an involvement of caspase activity (Figure 1B, right).

We next investigated the effects of Ret transfection and GDNF on the expression of specific pituitary somatotroph genes. In the absence of Ret, GDNF had no consistent effect on the hormones GH or PRL (data not shown), but had a small dose-dependent inhibitory effect on Pit-1 mRNA expression (Figure 1C). Transfection with Ret induced a marked increase in Pit-1 mRNA, which was blocked by incubation with GDNF (Figure 1D). The caspase inhibitor zVAD, which blocked Ret-induced apoptosis, also blocked the effect of Ret on Pit-1 mRNA expression (Figure 1E). Ret likewise induced an increase in Pit-1 protein levels, which were again reduced by GDNF and zVAD (Figure 1F and G). Ret has been classed as a 'dependence receptor': in HEK and neuroblastoma cell lines, it actively induces apoptosis in the absence of its ligand GDNF, whereas the presence of the ligand allows cell survival (Bordeaux *et al*, 2000; Bredesen *et al*, 2004; Porter and Dhakshinamoorthy, 2004). The Ret protein is processed by caspase 3 at two intracytoplasmic sites, releasing a fragment of about 40–50 kDa (IC-Ret) that has been proposed to induce apoptosis. IC-Ret was readily detected by Western blotting in our GH4C1 cells after transient transfection with either Ret-L or Ret-S. IC-Ret was not detectable in cells treated with zVAD, and levels were significantly reduced by GDNF (Figure 1F and

G), in contrast with the previous findings in HEK cells. Basal levels of activated caspase 3 were present in the GH4C1 cells, and caspase-3 activity was strongly induced by Ret transfection and reduced by GDNF or zVAD; that is, the effects on caspase 3 paralleled the effects on apoptosis, IC-Ret presence and Pit-1 level (Figure 1F, G and B, E).

In the presence of GDNF, Ret forms a heterodimeric complex with GDNF and GFR α 1, activating Ret's tyrosine kinase activity (so that it phosphorylates susceptible cytoplasmic substrates). We therefore investigated whether the tyrosine kinase activity of Ret is necessary for induction of apoptosis and/or of Pit-1 expression. The kinase-dead mutants of Ret-L (Ret-L_{KR}, K758R substitution) and Ret-S (Ret-S_{KR}, K758R substitution) were transiently transfected into GH4C1 cells. The transfected cells did not show increased tyrosine kinase activity in the presence of GDNF (unlike cells transfected with Ret-L or Ret-S) (Figure 2B), although Ret was expressed to similar levels (Figure 2C). While all Ret isoforms had apoptotic effects, the protective effects of GDNF were not seen in the cells transfected with the kinase-dead mutants (Figure 2D). Interestingly, both kinase-dead Ret isoforms strongly induced Pit-1 mRNA and protein expression (like the normal isoforms), but, unlike in cells transfected with the normal isoforms, GDNF did not inhibit this response (Figure 2E and F); again, this is in line with a relationship between Ret's apoptotic activity and its induction of Pit-1 expression.

Given our results showing that Ret's apoptotic activity is associated with caspase-3 processing and the appearance of the intracytoplasmic fragment IC-Ret, we transfected cells with Ret-S_{DN}, a Ret-S mutant with a D707N substitution affecting the first caspase-3 consensus site in the cytoplasmic tail of the receptor (Figure 2A). As previously demonstrated (Bordeaux *et al*, 2000), transfected Ret-S_{DN} could not be processed by caspase 3 *in vivo*, and no IC-Ret band was detected in Western blotting (Figure 2G). Ret-S_{DN} did not induce either apoptosis (Figure 2H) or Pit-1 mRNA or protein expression (Figure 2I and J).

Ret-induced apoptosis is mediated by Pit-1, and Pit-1 overexpression increases p53 levels

The data shown in Figures 1 and 2 suggested a direct relationship between caspase-3-processed intracytoplasmic Ret, Pit-1 expression and apoptosis. The time course of caspase-3 activity in extracts of transfected cells showed that the maximal activity was reached immediately after initiating the Ret transfection (2 h) (Figure 3A). Coincidentally, the induction of Pit-1 expression in somatotrophs was a very early event, seen as soon as 2 h after Ret transfection (Figure 3B), and maintained throughout the experiment (Figure 3C). Ret also induced a marked increase in p53 at 24 h after transfection (Figure 3D) before the induction of apoptosis, which peaked around 48 h after transfection (see Figures 1 and 2). Incubation with zVAD (which blocked IC-Ret generation, induction of Pit-1 expression and apoptosis; Figures 1B, E and G) also blocked the p53 increase (Figure 3E).

This sequence of events suggested that Ret-induced apoptosis in somatotrophs is mediated by Pit-1 induction. Transfection of a Pit-1 expression construct induced similar levels of apoptosis to Ret transfection, and the two treatments were not additive (Figure 3F). However, Pit-1-induced apoptosis was blocked by GDNF (Figure 3G), simultaneously

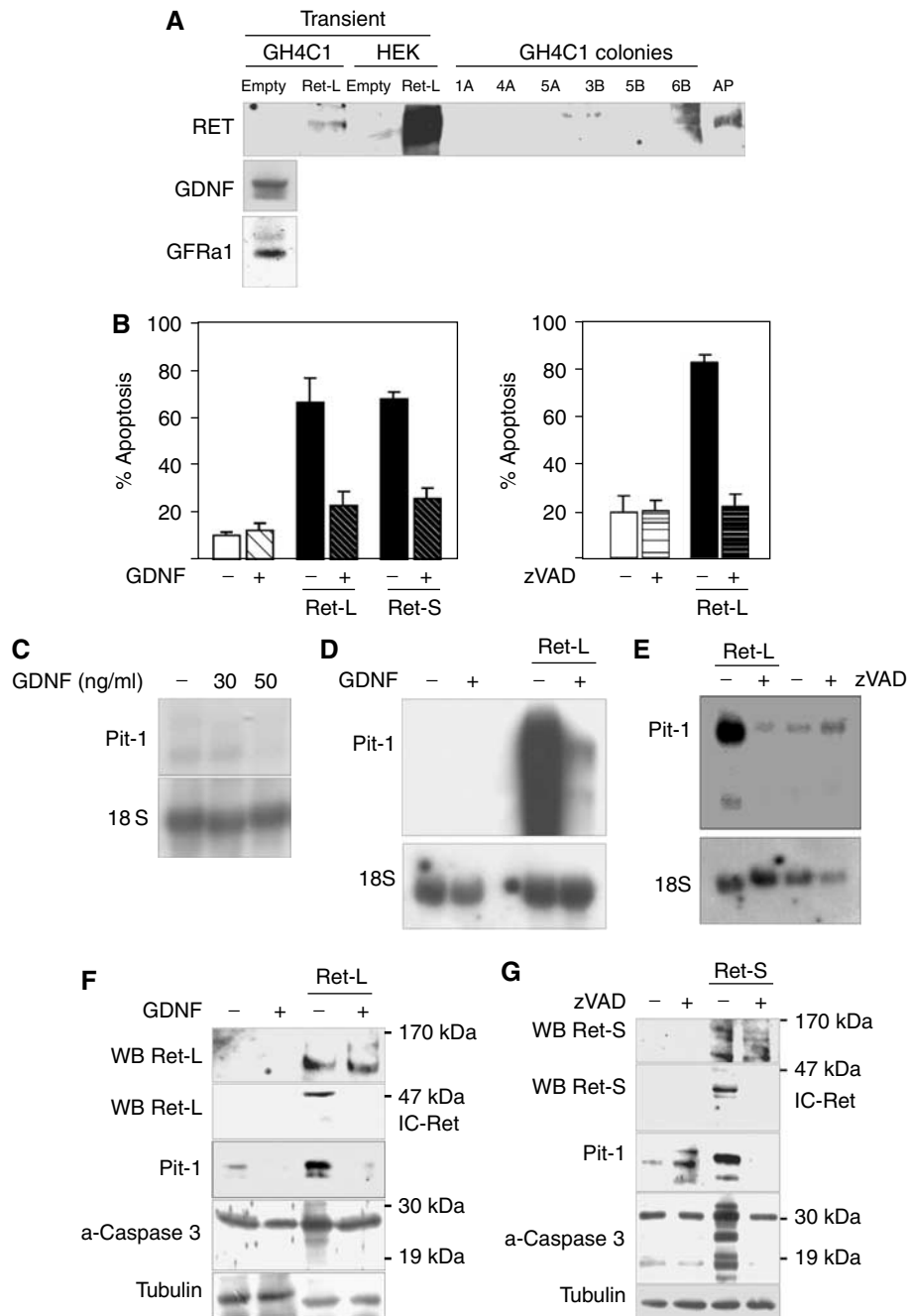


Figure 1 Both the Ret-L and Ret-S isoforms strongly induced apoptosis and Pit-1 expression in the GH4C1 somatotroph line. **(A)** GH4C1 cells expressed GDNF and GFR α 1 but not Ret. No stable Ret-expressing transfectants could be obtained (results shown only for Ret-L): the few colonies obtained were all Ret-negative, except colony 6B, which died after the second passage. In transient transfections, full-length Ret (150–170 kDa) was expressed strongly in HEK293 cells and weakly in GH4C1 cells. **(B)** Forty-eight hours after transient transfection of GH4C1 cells with either Ret-L or Ret-S, marked apoptosis was detected. Apoptosis could be prevented by treatment with GDNF or the broad caspase inhibitor zVAD. **(C)** In the absence of Ret, GDNF had a weak inhibitory effect on Pit-1 mRNA expression in GH4C1 cells. **(D)** Transfection with Ret induced a marked increase in Pit-1 mRNA expression, which was blocked by GDNF. **(E)** The broad caspase inhibitor zVAD blocked the effect of Ret on Pit-1 mRNA expression. **(F)** Ret transfection likewise induced a marked increase in Pit-1 protein expression, again blocked by GDNF (results shown only for Ret-L). Note that detection of full-length Ret required immunoprecipitation or loading more than 150 μ g proteins/lane, whereas the processed intracellular fragment (IC-Ret) could be detected easily with 50 μ g proteins/lane. Both IC-Ret and activated caspase 3 (19 kDa) were induced by transfection with Ret when GDNF was absent but not when GDNF was present. **(G)** zVAD also reduced the Ret-induced increase in Pit-1 protein expression (results shown only for Ret-S) and blocked IC-Ret formation and activation of caspase 3.

with the inhibition of transfected Pit-1 expression (Supplementary Figure 2A). To confirm that Pit-1 expression is essential for Ret-induced apoptosis, we used Pit-1 siRNA to block basal and Ret-induced Pit-1 expression (Figure 3H). The presence of Pit-1 siRNA entirely blocked the apoptosis

induced by transfection with Ret-L, Ret-S or Pit-1 (Figure 3I), and also blocked p53 induction (Figure 3J). These results suggest that caspase-3-processed IC-Ret induces Pit-1 expression, which in turn increases p53 expression and causes apoptosis.

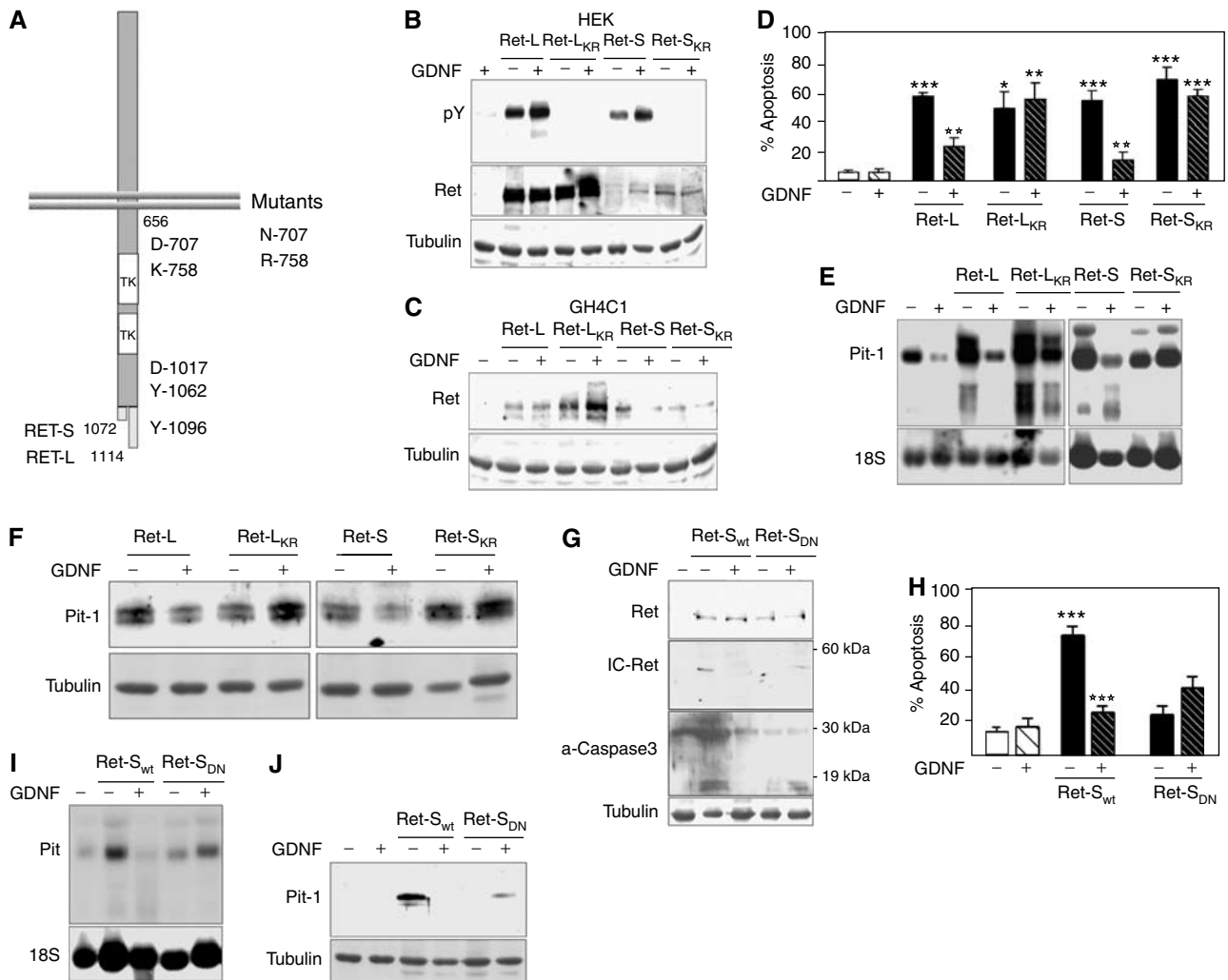


Figure 2 Ret-induced apoptosis and Pit-1 expression are independent of Ret's tyrosine kinase activity, but dependent on activation of caspase 3. (A) Cartoon showing the wild-type and mutant receptors used in these experiments. Ret-L_{KR} is a kinase-dead mutant (K758R) of Ret-L; Ret-S_{KR} is a kinase-dead mutant (K758R) of Ret-S; Ret-S_{DN}, cloned in the pBR expression vector, has a substitution in the proximal caspase-3-processing consensus site of the Ret-S isoform (control: Ret-S_{wt} in the same vector). (B) When expressed in HEK293 cells, neither kinase-dead mutant showed tyrosine kinase activity. As HEK293 cells secrete GDNF, a basal phosphorylation of the Ret receptor is seen in the untreated lanes. (C) The kinase-dead mutants were expressed in GH4C1 somatotrophs and (D) strongly induced apoptosis that could not be blocked by GDNF, like the wild-type isoforms. The kinase-dead mutants likewise markedly increased Pit-1 mRNA (E) and protein (F) expression; however, these increases were not blocked by GDNF. (G) In cells transfected with the Ret-S_{DN} mutant, neither IC-Ret (the processed intracellular form) nor activated caspase 3 could be detected. In line with this, Ret-S_{DN} did not induce either apoptosis (H) or Pit-1 mRNA (I), or protein (J) expression (**P* < 0.05; ***P* < 0.01; ****P* < 0.001).

Ret-induced c/EBP α binding to the Pit-1 promoter depends on PKC δ , JNK and CREB

Next, we performed experiments designed to identify kinases involved in the induction of Pit-1 expression by Ret. Sustained p-JNK and p-CREB expressions were detected in somatotrophs transfected with all Ret isoforms and mutants that induced apoptosis, but not in somatotrophs transfected with the non-apoptosis-inducing mutant Ret-S_{DN} (Figure 4A). As previously described (Chiariello *et al*, 1998; Feng *et al*, 1999; Poteryaev *et al*, 1999; Trupp *et al*, 1999; Hayashi *et al*, 2000; Pezeshki *et al*, 2001), short-term (15 min) treatment with GDNF also induced p-JNK or p-CREB activation (Figure 4B) that returned to basal levels after 30 min (Supplementary Figure 2B) and was not altered by Ret expression. However, long-term (24 h) treatment with GDNF, which blocks Ret-induced apoptosis, also blocked

the Ret-induced p-JNK and p-CREB sustained expression (Figure 4C). Likewise, we investigated whether PKC δ is activated by Ret. PKC δ has been implicated in apoptosis induced by DNA-damaging agents or Fas ligand: after its induction, PKC δ is phosphorylated (p-PKC δ) and processed, and the free catalytic domain migrates to the nucleus inducing apoptosis (DeVries *et al*, 2002; Mecklenbrauker *et al*, 2002, 2004), although the mechanism remains unknown. p-PKC δ was detected in somatotrophs transfected with the apoptosis-inducing Ret forms, but not in cells transfected with Ret-S_{DN} (Figure 4D). In contrast to the effects on JNK or CREB, short-term treatment (15 min) with GDNF reduced p-PKC δ levels in cells transfected with Ret-L and Ret-S but not in cells transfected with the kinase-dead mutant Ret-L_{KR} (Figure 4D). This reduction in p-PKC δ levels by GDNF was maintained after 24 h of treatment, coinciding with the

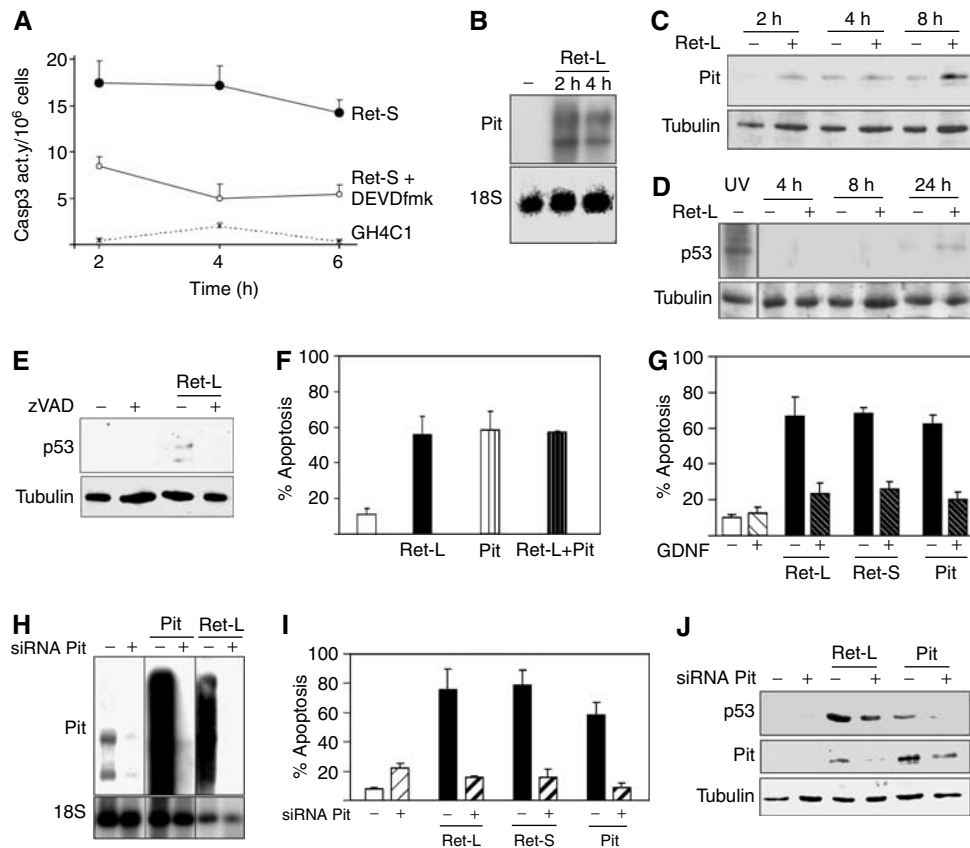


Figure 3 Ret-induced caspase-3 activation and Pit-1 overexpression in somatotrophs are early events causing a later p53 increase and apoptosis. (A) Time-course assay of caspase-3 activity after initiation of Ret-S transfection compared with empty-vector transfected cells or in the presence of the caspase-3 inhibitor DEVDfmk. The highest activity was seen at the first time point, only 2 h after the initiation of Ret transfection. (B) The Ret-induced increase in Pit-1 mRNA expression was already detectable within 2 h of transfection, suggesting a direct effect on the Pit-1 promoter that increased Pit-1 protein (C) with time after transfection. (D) An increase in p53 protein expression was only detectable 24 h after transfection. (E) Caspase inhibition with zVAD (which inhibited caspase-3 activity, Ret processing, apoptosis and Pit-1 overexpression; Figures 1 and 2) blocked the Ret-induced p53 increase. (F) Transfection with Pit-1 induced similar levels of apoptosis in somatotrophs as transfection with Ret; the two effects were not additive. (G) Intriguingly, Pit-1-induced apoptosis was also inhibited by GDNF (Supplementary Figure 2A). (H) Pit-1 siRNA blocked Pit-1 mRNA expression, whether basal or induced by Ret or Pit-1 transfection. For each Pit-1 siRNA treatment (+) and its control (-, GFP siRNA), lanes are shown with different exposure times. Pit-1 siRNA blocked apoptosis induced by Ret-L, Ret-S and Pit-1 (I) and the Ret-induced increase in Pit-1 and p53 protein expression (J).

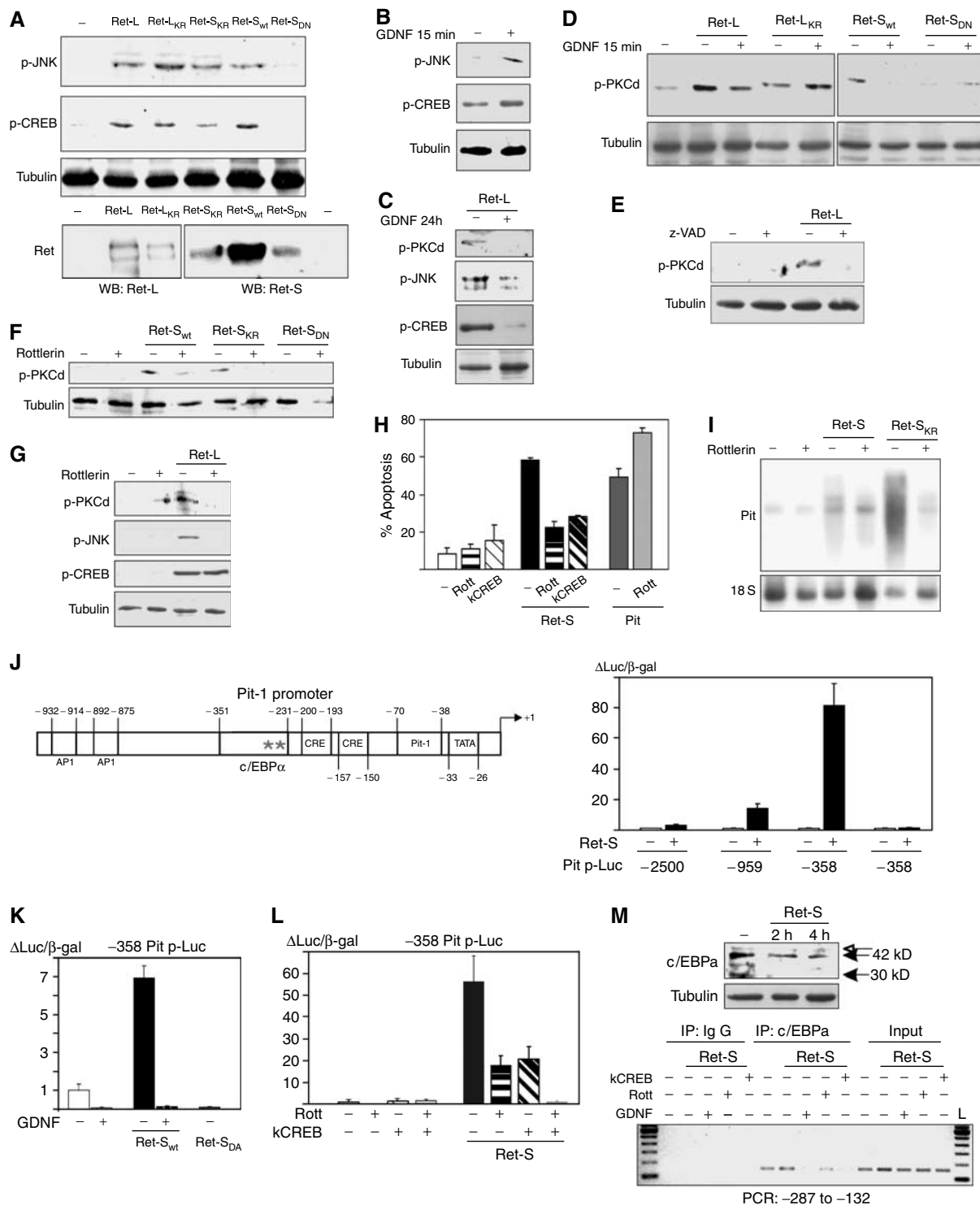
reduction in p-CREB and p-JNK levels (Figure 4C). Ret-induced p-PKC δ production was also blocked by the caspase inhibitor zVAD (Figure 4E). These results suggested that Ret may induce Pit-1 and apoptosis through activation of PKC δ . Rottlerin, a PKC δ inhibitor, blocked PKC δ phosphorylation by all the apoptosis-inducing Ret forms (Figure 4F). Rottlerin also blocked Ret-induced p-JNK expression but not Ret-induced p-CREB expression (Figure 4G). Rottlerin strongly reduced Ret-dependent Pit-1 expression, and in line with this reduced Ret-induced apoptosis; however, it did not affect Pit-induced apoptosis (Figure 4H and I). Finally, killer CREB, a dominant-negative form of CREB, significantly reduced Ret-induced apoptosis (Figure 4H). These data suggest a two-branched pathway for Ret-induced Pit-1 expression, one branch dependent on activation of PKC δ /JNK and the other dependent on activation of CREB. Pit-1 mRNA induction by this pathway was very rapid, suggesting a transcriptional effect. The most important consensus binding sites on the rat Pit-1 promoter are shown in Figure 4J. Both CRE elements are essential for GHRH- and ghrelin-induced Pit-1 expression (Soto *et al*, 1995; Garcia *et al*, 2001). In somatotrophs cotransfected with Ret and promoter/Luc con-

structs with decreasing promoter length, luciferase activity was detected with the -2500, -959 and -358 promoter fragments (strongest activity with the -358 fragment), whereas further shortening to -231 abolished the Ret-dependent activity (Figure 4J). Luciferase activity was also completely abolished by treatment with GDNF or by cotransfection with Ret-S_{DN} instead of Ret-S (Figure 4K). Luciferase activity in cells cotransfected with Ret-S and the -358 promoter fragment was markedly reduced by rottlerin and by killer CREB, and completely abolished by adding both together, confirming the additive effects of the PKC δ /JNK and CREB pathways (Figure 4L). These results indicate that the Ret-responsive region of the rat Pit-1 promoter lies between this protein's positions -358 and -231. A consensus binding site for the Pit-1 transcription factor *c/EBP α* appears near the first CRE, just before -231. Interestingly, the human Pit-1 promoter has a similar conserved site located upstream to the TATA region (Pfaffle *et al*, 1992), and initial results corroborate a Ret-dependent regulation of the hPit-1 promoter (data not shown). To confirm that Ret induces binding of *c/EBP α* to the rat Pit-1 promoter, we used chromatin immunoprecipitation (ChIP) assay with *c/EBP α* -specific

antibodies, followed by PCR using primers for a small region of the Pit-1 promoter, including the c/EBP α binding site. A basal level of c/EBP α binding was detected in untransfected cells, and an increase in this binding in Ret-transfected cells was detected within only 4 h of transfection. The increase in c/EBP α binding was completely abolished by GDNF and significantly reduced by rottlerin or cotransfection with killer CREB (Figure 4M).

The intracytoplasmic portion of Ret forms a complex with caspase 3 and PKC δ

Our data suggested a strong relationship between the intracellular Ret portion, caspase 3, PKC δ , Pit-1 expression and apoptosis. It seemed that blockage of Ret or caspase-3 processing also inhibited Pit-1 expression. In fact, caspase-3 activity was detected as early as 2 h after transfection of either Ret-S, the kinase-dead mutant Ret-S_{KR} or the intracellular portion of



Ret (IC-Ret₇₀₇₋₁₀₁₇). However, there was no caspase-3 activity after transfection with Ret-S_{DN}, a mutant that could not be processed by caspase-3 (Figure 5A, upper panel). GDNF was able to block Ret-induced caspase-3 activity, Pit-1 and apoptosis, but could not stop the apoptosis induced by IC-Ret₇₀₇₋₁₀₁₇, which correspondingly induced Pit-1 expression (Figure 5A, lower panels). We have found another caspase-3-processed protein, PKC δ (Voss *et al*, 2005), implicated in the pathway from Ret to the Pit-1 promoter. As both the catalytic fragment of PKC δ and caspase 3 have been located in the nucleus, we investigated the location of the processed intracellular fragment of Ret. Both the full-length Ret-S and the fragment IC-Ret₇₀₇₋₁₀₁₇ transfected in GH4C1 showed cytoplasmic location (Figure 5B and Supplementary Figure 2D).

As it has been demonstrated that PKC δ binds to and phosphorylates caspase 3 in monocytes (Voss *et al*, 2005), we investigated whether IC-Ret binds either of these proteins. Initial experiments showed the presence of PKC δ in Ret immunoprecipitates, with a weak band around 80 kDa and a stronger band around 40 kDa (data not shown); straight Western blotting against PKC δ showed the 80-kDa band and a shorter (unphosphorylated) band around 60 kDa (Figure 5C, right). We thought that the smaller 40-kDa band that appeared in the Ret immunoprecipitates could be the processed/phosphorylated catalytic fragment of PKC δ , absent in the 60-kDa band. Therefore, we performed IP-Ret IP followed by Western blotting against phosphorylated PKC δ (Figure 5C, upper left). Full-length Ret-S and IC-Ret₇₀₇₋₁₀₁₇ both co-immunoprecipitated with the p-PKC δ ; the strongest band was the processed 40-kDa band of p-PKC δ and the weakest band was the full-length 80-kDa p-PKC δ . Formation of this Ret/p-PKC δ complex was prevented in the presence of the ligand GDNF. The Ret-S_{DN} mutant was not able to associate with p-PKC δ , in line with its inability to induce Pit-1 expression or apoptosis. In the absence of GDNF, full-length Ret-S and IC-Ret₇₀₇₋₁₀₁₇ also co-immunoprecipitated with unprocessed (32 kDa) or processed (20/19 kDa) caspase 3 (Figure 5C, lower left). However, Ret-S_{DN} associated very weakly with unprocessed caspase 3, in keeping with its inability to induce caspase-3 processing/activation. To further study this complex, we immunoprecipitated PKC δ and assessed its association with either caspase 3 or the intracellular portion of Ret. The presence of any Ret construct, even Ret-S_{DN}, induced the formation of the PKC δ /

full-length caspase-3 complex, albeit in residual amounts in the presence of GDNF (Figure 5D, lower left). However, IC-Ret was found to be associated with PKC δ only in the presence of Ret-S or IC-Ret₇₀₇₋₁₀₁₇ (Figure 5D, upper left), that is, the same samples in which PKC δ was associated with Ret (compare upper panels of Figure 5C and D). These results indicated that the intracellular portion of Ret induces the formation of a complex with both full-length caspase 3 and PKC δ that results in cleavage (and activation) of the trio, and suggested that the complex was unstable when Ret could not be processed (Ret S_{DN}), or when GDNF induced Ret dimerization and tyrosine kinase activation.

GDNF prevents Ret-induced Pit-1 expression and apoptosis through Akt activation in both the somatotroph line and in primary pituitary cultures

Many pathways induced by GDNF have been described, both in the presence (Chiariello *et al*, 1998; Feng *et al*, 1999; Hayashi *et al*, 2000; Santoro *et al*, 2004; Arighi *et al*, 2005) and absence (Poteryaev *et al*, 1999; Trupp *et al*, 1999; Pezeshki *et al*, 2001) of Ret. In neurons, the blocking of Ret-induced apoptosis by GDNF is mediated by the kinase Akt: specifically, a GDNF/GFR α 1/Ret complex induces PI3K activation, leading to Akt phosphorylation and activation (Besset *et al*, 2000; De Vita *et al*, 2000; Hayashi *et al*, 2000). We therefore investigated whether this same pathway acts in somatotrophs. GDNF strongly induced Akt phosphorylation in cells transfected with wild-type Ret, but not in cells transfected with the kinase-dead mutants Ret-L_{KR} or Ret-S_{KR} (Supplementary Figure 2C). The PI3K inhibitor LY294002 partially reduced the GDNF-induced phosphorylation of Akt (Figure 6A). Likewise, the blocking of Ret-induced apoptosis (Figure 6B) and Pit-1 expression (Figure 6C) by GDNF was abolished by coincubation with LY294002.

We next investigated whether the results obtained in the GH4C1 somatotroph line were reproduced in primary pituitary cultures expressing endogenous Ret receptors. Pituitary cultures were grown in collagen-IV-coated dishes. Around 30% of cells were somatotrophs, in line with the high proportion of somatotrophs seen *in vivo*, and 99.9% were GH and Ret positive (Figure 6D); note that somatotrophs are the only cells expressing Ret and GFR α 1 (Urbano *et al*, 2000; Japon *et al*, 2002; and Supplementary Figure 3) in the rat and human pituitary. After 2 days, the original medium (containing 10% FCS) was changed to serum-deprived medium (only

Figure 4 Ret-induced sustained activation of PKC δ , JNK and CREB leads to binding of c/EBP α to the Pit-1 promoter. (A) Twenty-four hours after transfection of GH4C1 cells with apoptosis-inducing Ret forms, but not after transfection with Ret-S_{DN}, p-JNK and p-CREB levels showed a sustained increase. (B) We could not study the short-term effects of GDNF on p-JNK or p-CREB levels, as GDNF induced short-term phosphorylation of both kinases in the absence of Ret (peak at 15 min, return to basal at 30 min; Supplementary Figure 2B). (C) Long-term treatment with GDNF (24 h) blocked the Ret-induced phosphorylation of both kinases. (D) Transfection of GH4C1 cells with the apoptosis-inducing Ret forms, but not Ret-S_{DN}, induced a sustained increase in levels of p-PKC δ . Brief treatment with GDNF (15 min) inhibited this PKC δ phosphorylation, which was maintained after long-term (24 h) GDNF treatment (C). (E) zVAD treatment blocked the Ret-induced phosphorylation of PKC δ . (F) Rottlerin (Rott), a specific PKC δ inhibitor, effectively blocked the Ret-induced phosphorylation of PKC δ and (G) JNK, but not of CREB, suggesting that PKC δ /JNK and CREB form part of two parallel and additive branches of the pathway. (H) Both rottlerin and the dominant-negative form of CREB (killer CREB, kCREB) blocked Ret- but not Pit-induced apoptosis. (I) Rottlerin blocked Ret-induced Pit-1 mRNA expression. (J) Schematic representation of the rat Pit-1 promoter, with known (Carneiro *et al*, 1998; Garcia *et al*, 2001) (inside boxes) and putative (outside boxes) response elements. Cells were cotransfected with Ret plus Pit-1 promoter-luciferase constructs with increasing deletions of the promoter. In the presence of Ret-S, Pit-1 promoter deleted to -2500 showed a three-fold induction in luciferase activity that was enhanced with further deletions, especially to -358, but that was lost with further deletion to -231. (K) The -358 bp region appears to be very important for the response to Ret, as the response was blocked by GDNF treatment and could not be induced by the Ret-S_{DN}. (L) Both rottlerin and kCREB reduced Pit-1 promoter induction by Ret, although complete abolishment could only be achieved with both inhibitors together. (M) A consensus c/EBP α site was present in this small region of the promoter. Specific binding of c/EBP α to the Pit-1 promoter could be demonstrated by ChIP only 4 h after transfection with Ret. This was abolished by GDNF treatment and reduced by treatment with rottlerin or kCREB. c/EBP α Western blotting showed both the 42 and 30-kDa isoforms present in pituitary cell lines (Schaufele, 1996).

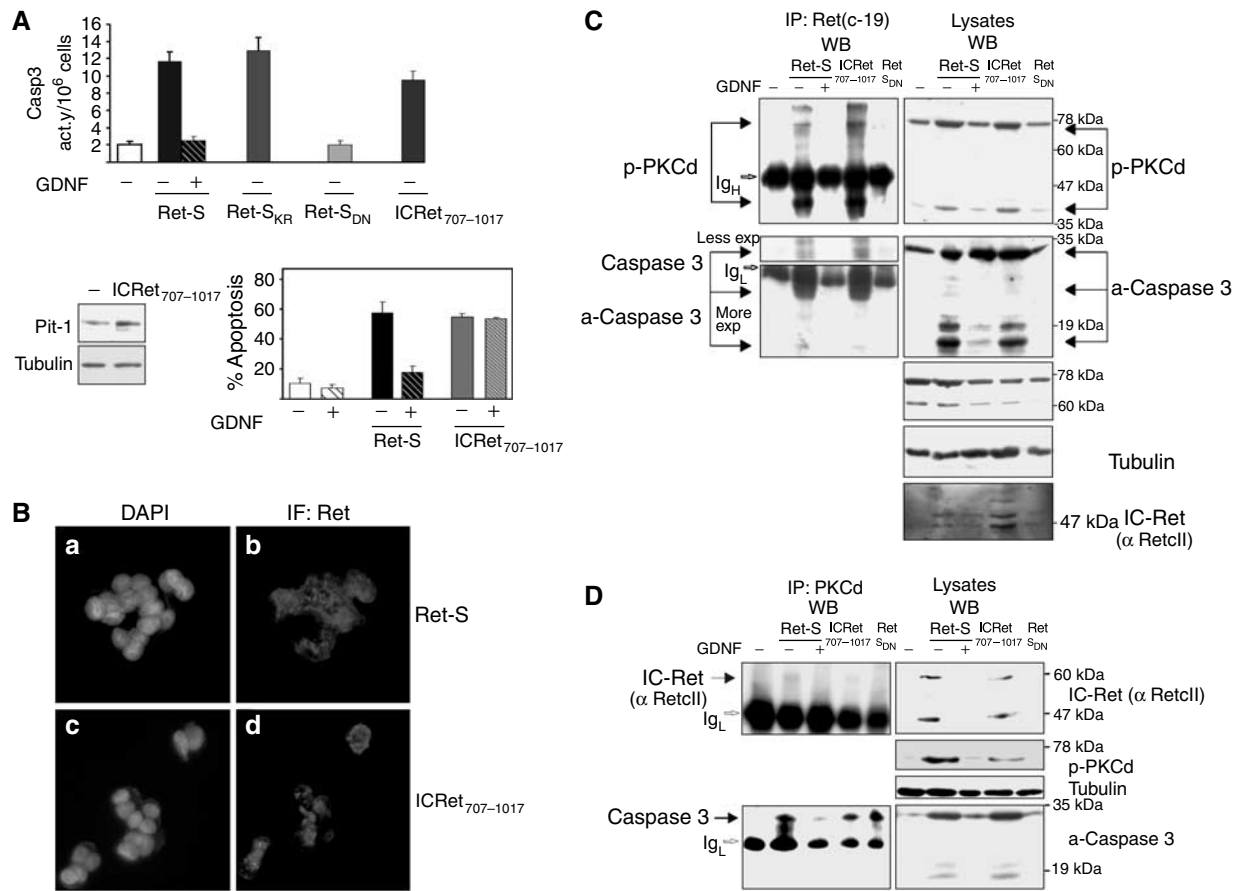


Figure 5 The intracellular portion of Ret forms a complex with caspase 3 and PKC δ in which all three become cleaved and activated. (A) Transfection of the intracellular portion of Ret, IC-Ret₇₀₇₋₁₀₁₇, induced an early (2 h) caspase-3 activity similar to full-length Ret-S or the kinase-dead Ret-S_{KR} but not Ret-S_{DN}. IC-Ret₇₀₇₋₁₀₁₇ induced Pit-1 expression and apoptosis not inhibited by GDNF. (B) Ret immunofluorescence in cells transfected with full-length Ret-S (b) or ICRet₇₀₇₋₁₀₁₇ (d) showed peripheral cytoplasmic location. DAPI staining was used as a nuclear marker (a, c). As it is a transient transfection, not all the cells expressed Ret (see Supplementary Figure 2E). (C) Left: immunoblotting with anti-p-PKC δ and anti-caspase 3 antibodies of anti-Ret immunoprecipitates indicated specific binding of Ret either to the cleaved 40 kDa p-PKC δ (more) and 25–19 kDa caspase 3 (less), or to the full-length PKC δ (70 kDa, less) and caspase 3 (30 kDa, more). The complex was prevented by GDNF treatment and was not present after transfection of the Ret-S_{DN} mutant. Right: levels of proteins in the lysates. (D) Similar anti-PKC δ immunoprecipitation revealed binding of PKC δ to full-length caspase 3 in the presence of any isoform of the Ret receptor, even the inactive Ret-S_{DN}; thus the binding was independent of caspase-3 activation. However, PKC δ bound to IC-Ret exclusively in the presence of Ret-S or IC-Ret₇₀₇₋₁₀₁₇, but not in the presence of Ret-S_{DN}. Again, GDNF treatment prevented the formation of both the IC-Ret–PKC δ complex and the Ret-induced PKC δ –caspase-3 complex.

0.5% FCS). Serum deprivation induced massive apoptosis and c/EBP α , Pit-1 and p53 overexpression that could be blocked by addition of GDNF, which blocked also IC-Ret expression (Figure 6D). In line with our findings in GH4C1, short-term GDNF treatment induced phosphorylation of Akt and CREB, but not of other kinases such as ERK. Treatment with the somatotroph-specific factors GHRH or ghrelin (both known to induce a short-term increase in Pit-1 expression; Soto *et al*, 1995; Garcia *et al*, 2001) likewise induced phosphorylation of Akt (Figure 6E). These results indicate that Akt is a key mediator in Pit-1 induction to prevent apoptosis, controlling its time course. Long-term treatment of the cultures with GDNF reduced p-CREB, p-JNK and p-PKC δ expression (Figure 6E). Either Pit-1 or Ret siRNA transfection of the cultures completely blocked the apoptosis or the Pit-1 expression (Figure 6F), indicating a role of Ret in maintenance of the somatotroph phenotype in adult pituitary.

Activation of the Ret/Pit-1/p53/apoptosis pathway in vivo prevents tumor growth

The above results indicated that the apoptotic pathway in primary pituitary cultures is the same as in the GH4C1 line. To investigate whether this pathway acts *in vivo*, we analyzed the pituitaries of 20 d.p.c. Ret KO mice in comparison with their wild-type littermates. Ret KO mice die around the first day of life owing to the absence of kidney (Schuchardt *et al*, 1994). Coronal and sagittal (data not shown) sections of the heads were obtained every 5 μ m. The first section in which the pituitary appeared was called level 0. From level 0 to the end, 5- μ m sections were obtained every 25 μ m, H&E-stained and used to measure the three pituitary lobes, that is, the adenopituitary (AP), intermediate lobe (IL) and neuropituitary (NP), and to reconstruct their volume (μ m³) using ImageJ software (NIH) (Figure 7A, B and D). Areas of the pituitary lobes (μ m²) in the three largest sections were also calculated (Figure 7A, E and F). AP volume in Ret KO mice

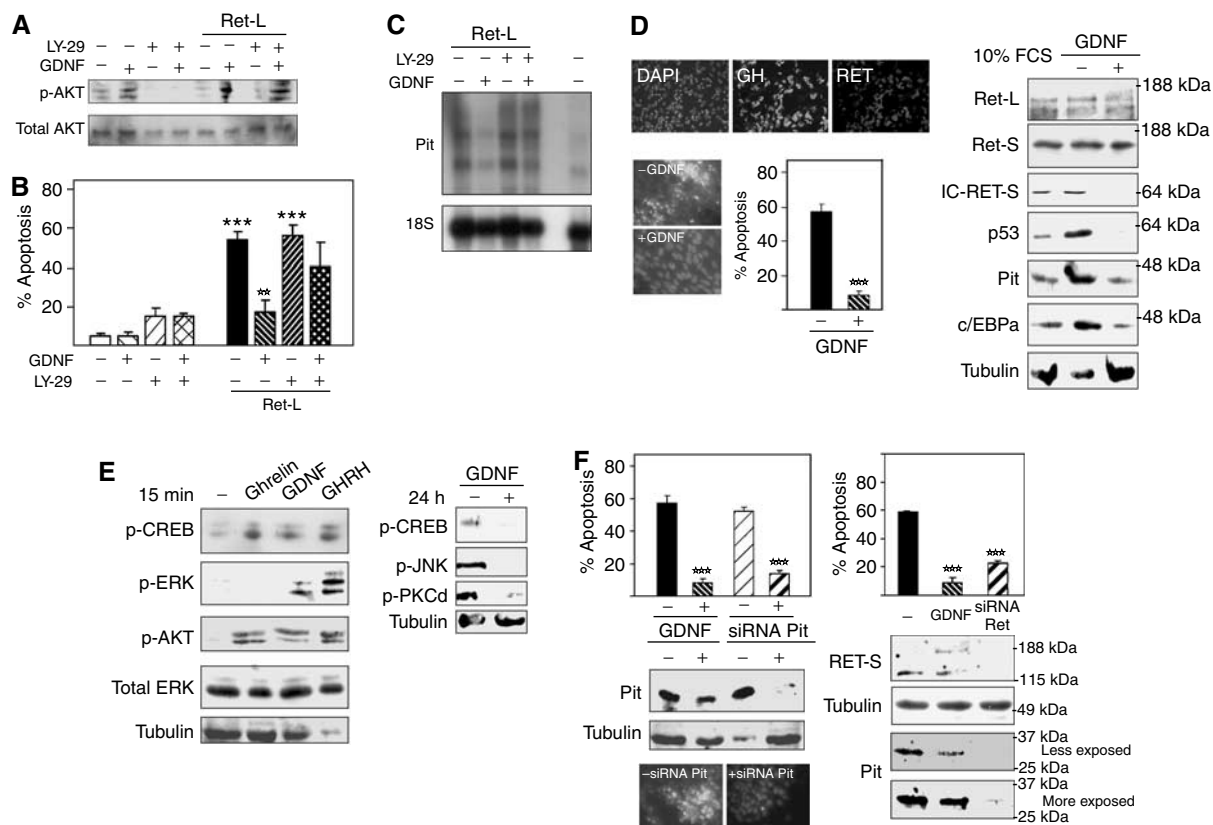


Figure 6 GDNF prevents Ret-induced p-PKC δ , p-JNK and apoptosis in both GH4C1 cells and primary pituitary cultures by phosphorylating Akt and thus blocking the Ret-induced increase in Pit-1. Both Pit-1 and Ret siRNA prevent apoptosis and Pit-1 expression. (A) GDNF induced Akt phosphorylation in Ret-transfected cells, and the PI3K inhibitor LY294002 (LY-29), Ret induced apoptosis (B) and Pit-1 mRNA overexpression (C) despite the presence of GDNF. (D) The percentage of GH+/Ret+ cells in primary cultures correlated with that in the whole pituitary (more than 40%). In these cultures, change from DMEM + 10% FCS to serum-deprived medium (DMEM + 0.5% FCS) induced marked apoptosis and IC-Ret, c/EBPa, Pit-1 and p53 expression. All of them could be rescued by GDNF. (E) Short-term treatment (15 min) of primary cultures with Pit-1 regulators (ghrelin, GDNF or GHRH) induced phosphorylation of CREB and Akt, whereas phosphorylation of ERK was induced only by GHRH (Pombo *et al*, 2000). Ghrelin and GHRH (Soto *et al*, 1995; Garcia *et al*, 2001) induced increased Pit-1 mRNA expression over a period of about 2 h, whereas GDNF reduced Ret-induced Pit-1 mRNA expression. As in the GH4C1 cell line, prolonged treatment with GDNF in primary cultures blocked the expression of p-CREB, p-JNK and p-PKC δ . (F) Both Pit-1 and Ret siRNA abolished Ret-induced Pit-1 protein expression after serodeprivation in primary cultures, in parallel with prevention of apoptosis (** $P < 0.01$; *** $P < 0.001$).

was significantly larger than in the wild type ($P = 0.007$; Figure 7D), as was AP area ($P = 2.4 \times 10^{-5}$; Figure 7E), whereas IL and NP volumes and areas were similar to those of the wild type. To rule out bias in the different sections, we calculated the ratios of AP to IL area and NP to IL area for each section. The AP/IL ratio was 1.4 times greater in the Ret KO mice ($P = 0.009$), whereas the NP/IL ratio showed no difference from the wild type ($P = 0.38$; Figure 7F).

There are five different types of secretory cell in the adenopituitary: somatotrophs (producing GH), lactotrophs (PRL) and thyrotrophs (TSH) are Pit-1-expressing cells derived from a common precursor, whereas gonadotrophs (LH and FSH) and corticotrophs (ACTH) are Pit-1 negative. Gonadotrophs (LH, FSH) also express a specific nuclear transcription factor called SF-1 (Morohashi and Omura, 1996). Whereas Pit-1-positive nuclei showed increased abundance in Ret KO mice ($P = 0.0097$), the number of SF-1-positive nuclei was similar in the two groups of mice ($P = 0.98$; Figure 7G). We counted the different Pit-1-positive lineages. Ret KO mice presented more GH-producing cells per hemipituitary ($P = 9.8 \times 10^{-6}$), whereas the number of TSH-produ-

cing cells was slightly, although not significantly, decreased ($P = 0.09$), and the number of PRL-producing cells was similar to that in wild type ($P = 0.92$). The total number of GH-, TSH- and PRL-producing cells in the wild-type pituitary equalled the number of Pit-1-positive cells. In the Ret KO mice, however, a subset of Pit-1-positive cells were not stained for any hormone marker, suggesting a delay in differentiation. Therefore, as GH cells are the only cells expressing Ret in the normal pituitary, somatotroph hyperplasia seems to have been the main cause of the larger size of the Ret KO pituitaries. There was no difference between the two groups in proliferative index in the pituitary, as measured by BrdU staining (Figure 7C and H); similar results were found by Ki-67 staining (data not shown). We conclude that somatotroph hyperplasia in Ret KO mice is likely due to enhanced survival through the differentiation process.

To investigate whether the most important feature of the pathway *in vivo* is Pit-1 expression or GH phenotype, we engineered the overexpression of Ret in Ret-negative/Pit-1-expressing pituitary lactotrophs, and then looked for increased Pit-1 expression, increased p53 expression and apoptosis. We used a retrovirus bearing the human Ret-S

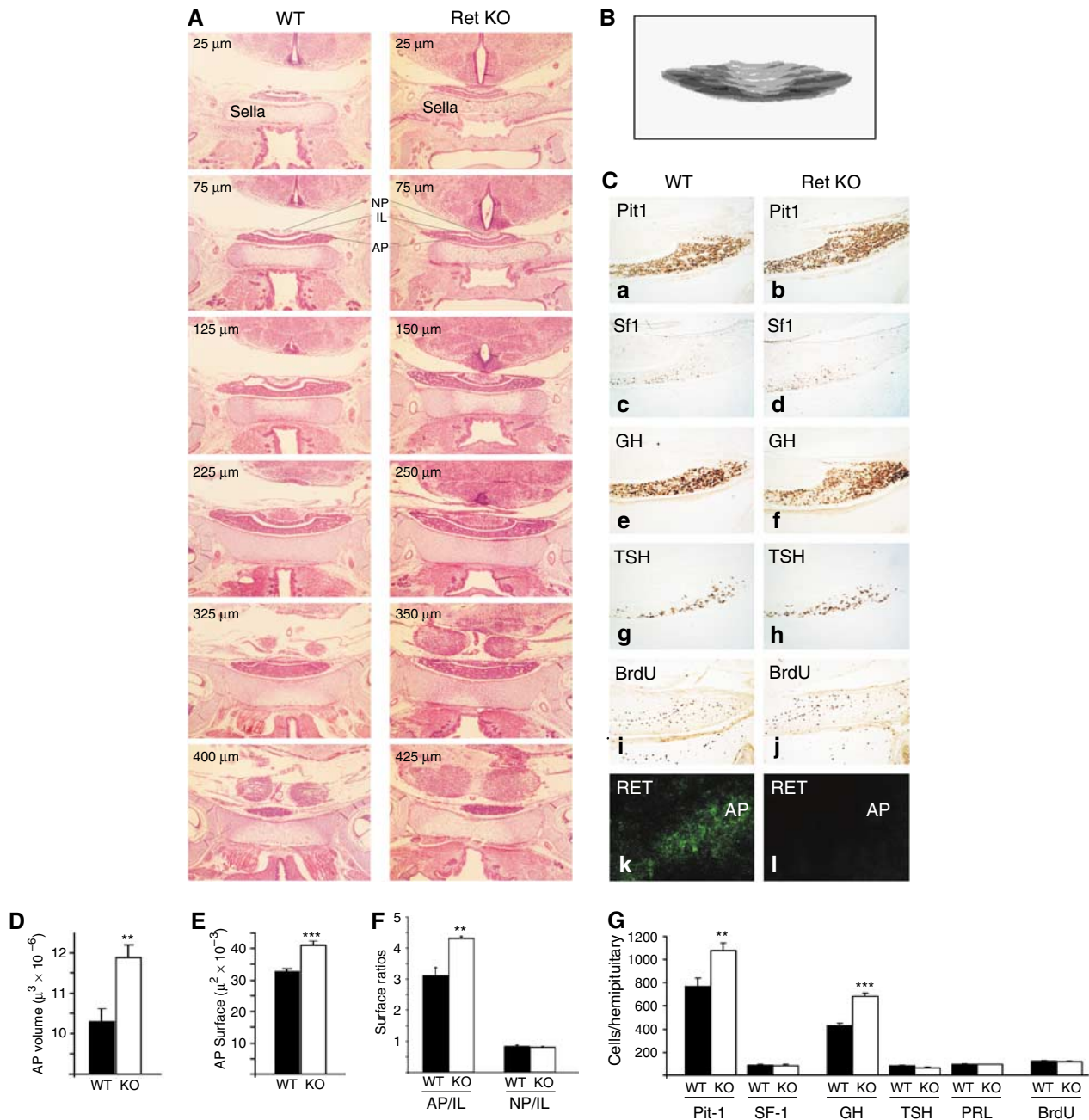


Figure 7 The Ret/Pit-1/p53 pathway acts *in vivo*: Ret KO mice present somatotroph hyperplasia. (A) H&E pituitary series of wild-type and KO mice (selected sections, $\times 40$). The first section in which the pituitary appeared on top of the sella turcica (Sella) was labelled as $0 \mu\text{m}$. The Ret KO mice showed a longer pituitary. (B) From level $0 \mu\text{m}$ to the end, $5\text{-}\mu\text{m}$ sections were taken every $25 \mu\text{m}$ and used to measure the three pituitary lobes, adenopituitary (AP), intermediate lobe (IL) and neuropituitary (NP), and to reconstruct their volume. (C) The central sections were used to study Pit-1-, Sf-1-, GH-, TSH-, PRL- (not shown) and BrdU-positive cells (hemipituitaries, $\times 200$). As a control, Ret immunofluorescence of pituitary sections is shown in the bottom panel. (D) Ret KO pituitaries had significantly higher volume and AP area (E) than wild-type littermates. (F) Area ratios between the three pituitary lobes in each section were calculated. The AP/IL ratio was higher in the Ret KO mice, although no difference could be detected in the NP/IL ratio in the same sections. (G) Ret KO mice showed Pit-1 hyperplasia but no alteration in SF-1 cells (gonadotrophs). Of the three Pit-1-positive cell types, only GH were more abundant; no difference was detected in TSH or PRL abundance. Likewise, we did not detect any difference in the abundance of BrdU-positive cells (** $P < 0.01$; *** $P < 0.001$).

cDNA (LPC-Ret) or the empty construct (LPC). For retroviral expression, we needed actively dividing cells, and so we used estrogen-induced lactotroph hyperplasia (Tsukahara *et al*, 1994). The LPC or LPC-Ret retrovirus was delivered to estrogen-injected female rats by stereotaxic injection. One week later, the pituitaries were dissected out, weighed and processed. Estrogens induced a strongly significant increase in pituitary weight in LPC-infected rats by comparison with non-

treated controls ($P = 0.00019$). Infection with the Ret-expressing retrovirus prevented this estrogen-induced hyperplasia (LPC + E2 versus LPC-Ret + E2 $P = 0.00077$, LPC-Ret + E2 versus control $P = 0.34$) (Figure 8A and Supplementary Figure 4C). In line with this, PRL immunostaining was much weaker in LPC-Ret-infected pituitaries than in LPC-infected pituitaries; however, the small areas that showed detectable PRL staining in LPC-Ret-infected pituitaries also

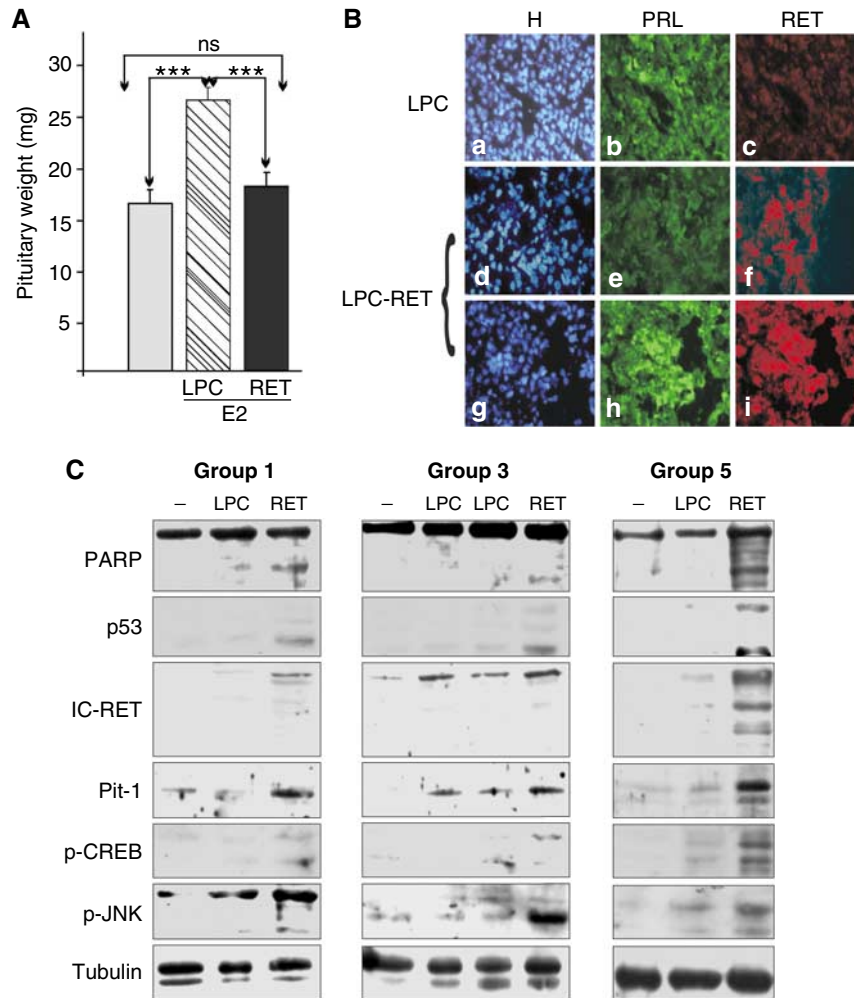


Figure 8 The Ret/Pit-1/p53 pathway acts *in vivo*: retroviral RET delivery in rat pituitary prevents tumor growth. (A) Mean (\pm s.e.m.) pituitary weights in estrogen- (E2) and vehicle-treated (C) rats, 7 days after the stereotaxic delivery of empty (LPC) or Ret-containing (LPC-RET) retrovirus. LPC-RET significantly reduced the E2-induced pituitary hyperplasia, returning pituitary weight to basal levels (for individual pituitary weights, see Supplementary Figure 3C). (B) The LPC-RET-infected pituitaries showed a marked decrease in PRL immunostaining. Areas still positive for PRL were also Ret-positive, in contrast with the absence of any PRL+/Ret+ areas in LPC-transfected pituitaries. (C) LPC-RET-infected pituitaries showed increased IC-Ret, Pit-1, p53, p-CREB and p-JNK expression 7 days after transfection. Cleaved PARP (80 kDa) was only present in LPC-RET-infected pituitaries. Data for groups 1, 3 and 5 are shown. Similar increases were already detectable 24 h after transfection (Supplementary Figure 4) (** $P < 0.001$).

showed strong Ret costaining (Figure 8B). The estrogen-treated LPC-Ret-infected pituitaries contained detectable levels of IC-Ret and increased amounts of Pit-1, p53, p-CREB and p-JNK by comparison with both estrogen-treated LPC-infected pituitaries and untreated pituitaries. Cleaved PARP (80 kDa) was only present in LPC-RET-infected pituitaries (Figure 8C). Some pituitaries were dissected out just 24 h after retroviral delivery, when no marked alteration in lactotroph numbers was expected. The proportion of Ret-expressing cells was $37.8 \pm 9.2\%$ in LPC-infected pituitary sections versus $61.8 \pm 7.3\%$ in LPC-Ret-infected pituitary, whereas the opposite pattern was seen in PRL-expressing cells ($62.2 \pm 9.2\%$ in LPC-infected pituitary versus $38.1 \pm 7.4\%$ in LPC-Ret-infected pituitary). More interestingly, as in normal pituitary (Urbano *et al*, 2000), the proportion of double labelled Ret/PRL cells was only $0.66 \pm 0.73\%$ in LPC-infected pituitary, versus $10.7 \pm 1.8\%$ in LPC-Ret-infected pituitary ($P = 0.0011$) (see Supplementary Figure 3A). p-PKC δ , IC-Ret, Pit-1 and p53 levels were increased within 24 h of LPC-RET infection.

Phosphorylated c/EBP α levels were increased in LPC-Ret-infected pituitary, whereas active caspase-3 levels, reduced in LPC-infected pituitary, returned to control levels (Supplementary Figure 3B). Taken together, these results demonstrate that Ret expression blocks estrogen-induced lactotroph hyperplasia *in vivo* by the same mechanism as revealed in the GH4C1 somatotroph cell line and in primary pituitary cultures, that is, via Pit-1 overexpression leading to increased p53 expression and thus apoptosis.

Discussion

Dependence receptors like Ret only induce apoptosis in the absence of a particular ligand, so that cell survival depends on the presence of the ligand. Besides, it seems likely that these proteins also play key roles in cell differentiation (Bredesen *et al*, 2004). Our present results rule out the possibility that Ret is necessary for differentiation to the somatotroph fate, and establish that Ret's role in the pituitary is to regulate Pit-1 expression and control survival of differ-

entiated somatotrophs (Figures 6F and 7A–C). We might speculate that this acts as a mechanism to prevent excessive body growth.

For apoptotic effect, dependence receptors need to be intracellularly processed by proteases, releasing a fragment with apoptotic properties. It remains unclear which specific protease processes each receptor (Bredesen *et al*, 2004). We found processing of the Ret receptor to be caspase-dependent, both in cell cultures and *in vivo*. It has been postulated that initial processing of Ret takes place under low levels of caspase-3 activation, which induces greater caspase activation in an activating loop (Bredesen *et al*, 2004). In somatotrophs, the maximal caspase-3 activity was detected immediately after Ret transfection (Figures 3 and 5), with the caspase-3 processing Ret intracellularly (to IC-Ret) in the absence of GDNF, and inducing sustained activation of PKC δ , JNK, c/EBP α and CREB, leading to strong and sustained overexpression of Pit-1, which in turn led to increased p53 expression and apoptosis. By contrast, GDNF activated Ret's tyrosine kinase activity, leading to Akt activation, and thus reduced Pit-1 expression, promoting survival (Figure 6B). Long-term phosphorylation of either CREB or JNK seems to be the key to the Ret-induced process, as GDNF inducing short-term phosphorylation of both has opposite effects (Chiariello *et al*, 1998; Feng *et al*, 1999; Poteryaev *et al*, 1999; Trupp *et al*, 1999; Hayashi *et al*, 2000; Pezeshki *et al*, 2001). Recently, the UNC5H2 dependence receptor has been found to bind and activate DAP kinase to induce apoptosis (Llambi *et al*, 2005). We provide evidence that the intracellular portion of Ret forms a complex with PKC δ and caspase 3 (Figure 5) and binding of caspase 3 and PKC δ has been described previously (Voss *et al*, 2005). Ternary complex formation was dependent solely on the presence of Ret, even an inactive form (Figure 5D), but was not enough to initiate the activation of the complex. Our data suggest that after binding to the complex, IC-Ret initiates the processing and activation of both PKC δ and caspase 3. A certain basal level of activated caspase 3 could initiate Ret cleavage and thus trigger an activating loop (Llambi *et al*, 2005); alternatively, the presence of the monomeric Ret cytoplasmic tail may elicit formation of the ternary complex in which caspase 3 would become stabilized and activated. We favor this latter model, as total caspase-3 levels are increased by Ret transfection. In contrast to previous findings in HEK cells (Bordeaux *et al*, 2000), in somatotrophs, GDNF was able to suppress Ret processing by dimerization of the Ret receptors and Akt recruiting, disrupting the IC-Ret/PKC δ /caspase-3 complex.

The PKC δ catalytic fragment migrates to the nucleus (DeVries *et al*, 2002), but its targets remain unknown. In many systems, PKC δ is implicated in apoptosis downstream of p53 but upstream of JNK (Humphries *et al*, 2006; Liu *et al*, 2006; Miyamoto *et al*, 2002). PKC δ -dependent CREB phosphorylation has been described for the B-cell receptor (Blois *et al*, 2004). However, in our cells, inhibition of PKC δ , although it blocked JNK activation, did not abolish CREB phosphorylation. Our results indicate a cooperative action of c/EBP α and CREB on the Pit-1 promoter. Similar cooperation has been described in human monocyte differentiation (Brenner *et al*, 2003). Taken together, our data suggest a model in which, once the ternary complex (IC-Ret/PKC δ /caspase 3) is formed and processed, PKC δ either alone or bound to caspase-3 activates JNK. It remains to be deter-

mined whether JNK activation occurs in the cytosol or in the nucleus, and how CREB phosphorylation is initiated and maintained independent of PKC δ activation.

Previous reports have already suggested a role of c/EBP α in the pituitary (Schaufele, 1996). Pit-1 establishes somatotroph (GH), lactotroph (PRL) and thyrotroph (TSH) phenotypes during pituitary development. Based on *in vitro* experiments in lactotroph cell lines, it is currently accepted that Pit-1 has two functions, regulating both differentiation and proliferation (Castrillo *et al*, 1991). However, there is no substantial *in vivo* evidence in rodents or humans to support this latter possibility: the Pit-1 gene is not altered in adenomas, and Pit-1 expression appears to be more closely related to GH secretion than to tumorigenesis (Delhase *et al*, 1993; Hoggard *et al*, 1993; Pellegrini *et al*, 1994). Moreover, Pit-1 mutations have been found in human hypopituitarism without pituitary hypoplasia (Pfaffle *et al*, 1992), whereas no increase in Pit-1 expression is required for *in vivo* estrogen-induced lactotroph hyperplasia in rats (Tsukahara *et al*, 1994). Our data show that too much Pit-1 is deleterious for the somatotroph; conversely, knocking down Pit-1 prevents apoptosis. The apoptosis induced by dependence receptors appears to be independent of both the intrinsic (caspase 9/Apaf1) and the extrinsic (FADD/caspase 8) pathways (Porter and Dhakshinamoorthy, 2004). The induction of p53 expression by Pit-1 links Ret-induced apoptosis with p53. To date, the only reported relationship of dependence receptors with p53 is via netrin-1, a ligand that blocks p53-dependent apoptosis by binding to the UNC5B dependence receptor (Tanikawa *et al*, 2003).

In the presence of the PI3K inhibitor LY-294002, Ret induced Pit-1 overexpression and apoptosis despite the presence of GDNF, although Akt phosphorylation was not completely abolished. GDNF induces Akt activation through binding of a Gab-p85PI3K complex to the phosphorylated Y1062 of activated Ret, thus preventing apoptosis (Besset *et al*, 2000; De Vita *et al*, 2000; Hayashi *et al*, 2000). In somatotrophs, the PI3K–Akt pathway seems to be essential for Pit-1 downregulation and cell survival. This pathway could act at various levels: by downregulating Ret processing by caspase inhibition, by decreasing c/EBP α phosphorylation through PP2A activation (Wang *et al*, 2004) or by regulating p53-dependent apoptosis. The fact that GDNF is not able to block Pit-1 induction/apoptosis induced by transfection of the IC-Ret_{707–1017} fragment, but decreases Pit-1 induction of exogenous transfected Pit-1 and consequent apoptosis, suggests that the GDNF action is before the Pit-1 action, and rules out any direct effect on p53; it seems probable that IC-Ret_{707–1017} sequesters PI3K/Akt, preventing its activation near the membrane. GDNF (via Akt induction) and siRNA Pit-1 were also able to prevent apoptosis in the primary pituitary culture. GHRH, ghrelin and GDNF share the ability to activate Akt and regulate Pit-1 transcription. GHRH and ghrelin cause very short-lived Pit-1 induction (2 h) (Soto *et al*, 1995; Garcia *et al*, 2001), whereas GDNF reduces Pit-1 transcription. We propose that Akt activation by GHRH and ghrelin switches off the Pit-1 promoter, confining the induction to this short period. In fact, GDNF is able to decrease Pit-1 levels after transfection of exogenous Pit-1 expression vector; in this case, the positive feedback of exogenous Pit-1 activating the endogenous Pit-1 promoter is disrupted by Akt. A role for Akt in pituitary tumorigenesis is

suggested by the finding of increased p-Akt levels in pituitary tumors (Musat *et al*, 2005).

Our results support the notion of dependence receptors being tumor suppressors (Mazelin *et al*, 2004), where Ret, like the other dependence receptors, would be a two-sided tumor regulator: a tumor suppressor because of its dependence receptor activity and an oncogene because of its tyrosine kinase activity. We further show that activation of the Ret/Pit-1/p53 pathway was able to block tumor growth *in vivo*, probably by the same mechanisms described in cultures. These results raise the possibility of Ret-based therapy as a treatment for conventional Pit-1-expressing chemotherapy-resistant adenomas.

The Ret/Pit-1/p53 pathway, conserved in mouse, rat and human somatotrophs, appears to be a mechanism to prevent oversized individuals, exerting a strong pressure on somatotrophs to stay in the pituitary, and limiting their numbers (Supplementary Figure 5). Disrupting Ret expression during development provokes somatotroph hyperplasia (Figure 7); the final result in the adult animal might perhaps be gigantism. If a precursor cell is to become a somatotroph, it needs Pit-1, and to maintain sufficient Pit-1 expression in a somatotroph after birth, Ret expression is required (Urbano *et al*, 2000; Japon *et al*, 2002). But if a somatotroph produces too much Pit-1 or migrates to areas in which GDNF is present at much lower concentrations, Ret-induced Pit-1 overexpression will kill it. This would further explain why all GH-secreting adenomas maintain normal Ret/GFR α 1 and GDNF expression, and why GH-secreting adenomas are unable to metastasize.

Materials and methods

A more detailed list of methods is given as Supplementary data.

Cell culture and apoptosis detection

Rat GH4C1 pituitary cells and HEK293 cells were cultured in DMEM plus 10% FCS and apoptosis was detected by Hoechst staining (Bravo *et al*, 2003). zVAD (Sigma) was dissolved in DMSO and used at 50 μ M. Rottlerin (Sigma) was dissolved in DMSO and used at 20 nM. LY294002 (Sigma) was dissolved in DMSO and used at 10 μ M.

Constructs and transfections

Cells were transfected with Eugene (Roche) or JetSi (PolyPlus). A detailed list of the constructs and siRNA sequences used in this work is provided as Supplementary data.

Primary adenopituitary culture

Pituitary cells were cultured by a modification of our previously described procedure (Coya *et al*, 1999; Garcia *et al*, 2001; Supplementary data). Cells were plated in Ikawi culture dishes coated with type-IV collagen (Sigma) and incubated for 4 days. To induce apoptosis, the medium was changed to DMEM plus 0.5% FCS for 36 h. Apoptosis was detected by Hoechst staining.

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Northern blotting

RNA was extracted with RNAeasy (Qiagen).

Caspase-3 activity

We used the Caspase 3 Apoalert kit (Clontech) following the manufacturer's instructions.

Immunodetection

Cell extracts were obtained with Triton lysis buffer as described (Carneiro *et al*, 1998; Garcia *et al*, 2001). A detailed list of the antibodies used is provided as Supplementary data.

Immunocytochemistry

To estimate the proportion of secretory cells in pituitary cultures and cryosections, we used a double immunofluorescence procedure, with the two immunostainings performed sequentially. See Supplementary data for the list of antibodies and procedures.

Chromatin immunoprecipitation assay

ChIP assay was performed with a kit from Upstate, using c/EBP α (Santa Cruz Biotechnology) and rabbit IgG as control. PCR was performed at 59°C for 35 cycles with forward primer 5'-CCAAAAGA CGCCTATTTTTCA-3' and reverse primer 5'-GGAGAAACCTGGTGTG GACGTA-3'.

Mouse pituitary analysis

Full-term pregnant mothers (20 days) were injected with 100 μ g/g BrdU (Sigma) i.p. 1 h before killing. Pups' heads were fixed in formalin for 24 h followed by immersion in 70% ethanol until analysis by standard paraffin inclusion. The heads were completely sectioned (5 μ m) sagittally (initial group, $n=4$ /genotype) or coronally (group 2, $n=3$ /genotype; and group 3, $n=4$ /genotype), and every 25 μ m a hematoxylin–eosin staining was performed for estimation of pituitary lobe volume ($n=4$ /genotype) and area ($n=7$ /genotype) (ImageJ, <http://rsb.info.nih.gov/ij/>). Pit-1, Sf-1 ($n=4$ /genotype), and GH, TSH and PRL ($n=7$ /genotype) immunohistochemistry in the pituitaries from littermates were performed by standard techniques. A detailed procedure is provided as Supplementary data. ImageJ was also used to count the stained cells.

In vivo stereotaxic retroviral delivery

Ret-expressing or empty retrovirus were delivered after stereotaxic injection to estrogen-treated rats. See Supplementary data for a more detailed description.

Statistical analysis

Results were compared by non-parametric *t*-tests. (* $P<0.05$; ** $P<0.01$; *** $P<0.001$).

Supplementary data

Supplementary data are available at *The EMBO Journal* Online (<http://www.embojournal.org>).

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