suggesting down-regulation of DNA replication and providing evidence for a role of human Orc6 in DNA replication. BrdU incorporation studies revealed 60 to 70% cells positive for BrdU after 60 hours of transfection, and decreased to 20 to 30% cells after 108 hours (Fig. 5A). In the majority of multinucleate cells, one or two nuclei showed much lower than normal BrdU staining and the remaining nuclei had no BrdU incorporation over a 24-hour period (Fig. 5B). Some cells transfected with Orc6 siRNA duplexes showed an overall intensity of BrdU staining considerably lower than that observed in control cells (see figs. S5 and S6), suggesting infrequent replication origin-firing. Because the origins of DNA replication are redundant, it is likely that most of the Orc6 had to be depleted to reveal a complete defect in DNA replication. BrdU incorporation followed by colcemid-induced metaphase spreads following 108 hours of Orc6 siRNA showed incomplete and lower levels of BrdU incorporation during a 24-hour label (Fig. 5C) (fig. S7). However, the control cells (Gl3) showed complete staining of duplicated chromosomes. The punctate incorporation of BrdU suggests that fewer origins fire, resulting in a slower S phase. Similarly, Drosophila larvae homozygous for Orc2 and Orc5 mutants progressed slowly through S phase and showed vastly reduced BrdU incorporation; these defects may have been caused by fewer active origins (24).

In human cells, Orc6 functions in multiple aspects of the cell division cycle, including DNA replication, chromosome segregation, and cytokinesis. It may be that Orc6 participates independently in the mechanics of DNA replication, centromere function, and cytokinesis. Alternatively, a significant defect in DNA replication caused by silencing of Orc6 might trigger aberrant mitosis (or other downstream effects) by interfering with centrosome duplication (25). Although this scenario could explain the spindle defects and polyploidy, it would not explain the cytokinesis defects in cells lacking Orc6. A reduction in DNA replication after Orc2 siRNA treatment did not yield similar phenotypes. Also, the Orc6 siRNA phenotypes are consistent with Orc6 distribution to the kinetochores and midbody. A third possibility is consistent with the Orc6 localization pattern during the cell cycle: Orc6 could be essential for inducing or signaling to multiple cell cycle checkpoints that coordinate the many processes of the cell division cycle (26). In this model, chromosome duplication, segregation, and cytokinesis would be coordinated by a common participant. A protein such as Orc6 could cycle from origins of DNA replication to centromeres and the cytokinesis apparatus to ensure the correct order and coordination of events.

## **References and Notes**

- S. Waga, B. Stillman, Annu. Rev. Biochem. 67, 721 (1998).
- 2. S. P. Bell, Genes Dev. 16, 659 (2002).

- REPORTS
- 3. \_\_\_\_\_, B. Stillman, *Nature* **357**, 128 (1992).
- T. J. Kelly, G. W. Brown, Annu. Rev. Biochem. 69, 829 (2000).
- I. Chesnokov, M. Gossen, D. Remus, M. Botchan, Genes Dev 13, 1289 (1999).
   I. Chesnokov, D. Remus, M. Botchan, Proc. Natl. Acad.
- Sci. U.S.A. 98, 11997 (2001).
- 7. J. Mendez et al., Mol. Cell 9, 481 (2002).
- A. A. Van Hooser, M. A. Mancini, C. D. Allis, K. F. Sullivan, B. R. Brinkley, *FASEB J.* **13** (suppl. 2), S216 (1999).
- 9. S. G. Prasanth, B. Stillman, unpublished data.
- D. A. Skoufias, P. R. Andreassen, F. B. Lacroix, L. Wilson, R. L. Margolis, Proc. Natl. Acad. Sci. U.S.A. 98, 4492 (2001).
- 11. M. Glotzer, Annu. Rev. Cell Dev. Biol. 17, 351 (2001).
- 12. S. G. Zeitlin, K. F. Sullivan, Curr. Biol. 11, R514 (2001).
- 13. P. Fortugno et al., J. Cell Sci. 115, 575 (2002).
- J. M. Mullins, J. R. McIntosh, J. Cell Biol. 94, 654 (1982).
- W. C. Earnshaw, R. L. Bernat, *Chromosoma* **100**, 139 (1991).
- D. G. Lee, S. P. Bell, *Mol. Cell. Biol.* **17**, 7159 (1997).
  M. Weinreich, C. Liang, H. H. Chen, B. Stillman, *Proc. Natl. Acad. Sci. U.S.A.* **98**, 11211 (2001).
- Natl. Acad. Sci. U.S.A. 98, 11211 (2001). 18. P. J. Gillespie, A. Li, J. J. Blow, BioMed Central Bio-
- chem. 2, 15 (2001).
- 19. S. M. Elbashir et al., Nature 411, 494 (2001).
- Orc6 siRNA oligos 6.13 (5'-AAUGGUAGCCACAUC-CGGUGU-3') and 6.17 (5'-AAGATTGGACAGCAG-

- GUCGAC-3') were obtained from Dharmacon Research Inc.
- 21. R. R. Adams, H. Maiato, W. C. Earnshaw, M. Carmena, J. Cell Biol. 153, 865 (2001).
- 22. A. G. Uren et al., Curr. Biol. 10, 1319 (2000).
- J. Harborth, S. M. Elbashir, K. Bechert, T. Tuschl, K. Weber, J. Cell Sci. 114, 4557 (2001).
  M. F. Pflurm, M. R. Botchan, Development 128, 1697
- (2001).
  25. O. C. Sibon, A. Kelkar, W. Lemstra, W. E. Theurkauf, Nature Cell Biol. 2, 90 (2000).
- 26. H. Murakami, P. Nurse, Biochem. J. 349, 1 (2000).
- S. A. Jablonski, G. K. Chan, C. A. Cooke, W. C. Earnshaw, T. J. Yen, *Chromosoma* 107, 386 (1998).
- S. We thank all members of the Stillman laboratory for stimulating discussions; H. Zou-Yang and S. Dike for anti-ORC antibodies; T. J. Yen, B. R. Brinkley, K. H. A. Choo, and F. McKeon for kinetochore antibodies; A. Krainer for SF2/ASF antibody; D. L. Spector and D. Helfman for suggestions; J. Mendez, A. Stenlund, and G. Hannon for reading the manuscript; and J. Duffy for artwork and photography. Supported by National Cancer Institute grant CA13106.

### **Supporting Online Material**

www.sciencemag.org/cgi/content/full/297/5583/1026/ DC1

Figs. S1 to S7

11 April 2002; accepted 22 May 2002

# Impaired B and T Cell Antigen Receptor Signaling in p110 $\delta$ PI 3-Kinase Mutant Mice

Klaus Okkenhaug,<sup>1</sup> Antonio Bilancio,<sup>1\*</sup> Géraldine Farjot,<sup>1\*</sup> Helen Priddle,<sup>2\*†</sup> Sara Sancho,<sup>3</sup> Emma Peskett,<sup>1</sup> Wayne Pearce,<sup>1</sup> Stephen E. Meek,<sup>2</sup> Ashreena Salpekar,<sup>1</sup> Michael D. Waterfield,<sup>1,4</sup> Andrew J. H. Smith,<sup>2</sup> Bart Vanhaesebroeck<sup>1,4</sup>‡

Class IA phosphoinositide 3-kinases (PI3Ks) are a family of p85/p110 heterodimeric lipid kinases that generate second messenger signals downstream of tyrosine kinases, thereby controlling cell metabolism, growth, proliferation, differentiation, motility, and survival. Mammals express three class IA catalytic subunits: p110 $\alpha$ , p110 $\beta$ , and p110 $\delta$ . It is unclear to what extent these p110 isoforms have overlapping or distinct biological roles. Mice expressing a catalytically inactive form of p110 $\delta$  (p110 $\delta^{D910A}$ ) were generated by gene targeting. Antigen receptor signaling in B and T cells was impaired and immune responses in vivo were attenuated in p110 $\delta$  mutant mice. They also developed inflammatory bowel disease. These results reveal a selective role for p110 $\delta$  in immunity.

There is increasing evidence for an important role for class IA PI3Ks in regulation of the immune system (1-4). Mice lacking the

<sup>1</sup>Ludwig Institute for Cancer Research, 91 Riding House Street, London W1W 7BS, UK. <sup>2</sup>Gene Targeting Laboratory, Center for Genome Research, University of Edinburgh, West Mains Road, Edinburgh, EH9 3JQ, UK. <sup>3</sup>Frimorfo, Rue du Musée 12, CH-1705 Fribourg, Switzerland. <sup>4</sup>Department of Biochemistry and Molecular Biology, University College London, Gower Street, London WC1E 6BT, UK.

\*These authors contributed equally to this report. †Present address: Roslin Institute, Roslin, Midlothian, EH25 9PS, UK.

‡To whom correspondence should be addressed. Email: bartvanh@ludwig.ucl.ac.uk p85 $\alpha$  regulatory subunit show impaired B cell development and activation but normal T cell activation, whereas T cell-restricted deletion of the gene for the phosphoinositide 3-phosphatase PTEN, a negative regulator of PI3K signaling, results in a lethal lymphoproliferative disease (5–7). To date, the specific role for each of the three class IA PI3K catalytic subunits in lymphocyte signaling has not been determined. p110 $\delta$  is expressed predominantly in leukocytes (8, 9), which indicates that it may play a unique role in immune signaling.

The study of class IA PI3Ks is complicated by the heterodimeric nature of these proteins, and altering the expression of one subunit can affect the expression of the others (10). In cells from p85a knockout mice, p85ß expression was up-regulated, and expression of  $p110\alpha$ , p110B, and p110b was down-regulated (5, 6, 11). In contrast,  $p85\alpha$  was overexpressed in p110a knockout embryos, which died after 9.5 days of gestation (12). In this study, p1108 was inactivated by point mutation instead of by deletion to prevent changes in the expression levels of the other PI3K catalytic and regulatory subunits (see fig. S1 for details of the targeting strategy). Homozygous mutant mice ( $p110\delta^{D910A/D910A}$ ) were born at normal Mendelian ratios, were fertile, and did not show gross anatomical or behavioral abnormalities (13).

Expression of the mutated p110 $\delta$  protein was equivalent to that of the wild-type (WT) protein, as was the expression of the other PI3K subunits (Fig. 1A). p110 $\delta$  lipid kinase activity was completely abrogated in p110 $\delta^{D910A/D910A}$  mice, with no alteration in the kinase activities of p110 $\alpha$  and p110 $\beta$ (Fig. 1B). Total class IA PI3K activity was reduced 30 to 50% in thymocytes as well as in mature B and T cells (*13*).

Antibodies to immunoglobulin M and to CD3 (anti-IgM and anti-CD3) were used to stimulate the B and T cell antigen receptors (BcR and TcR), respectively, and phosphorylation of the PI3K target Akt/protein kinase B (PKB) was used as an indirect measure of PI3K activity. In both cases, the induction of phosphoserine-473 [Ser(P)<sup>473</sup>] Akt/PKB was compromised in p1108<sup>D910A/D910A</sup> cells (Fig. 1C). Phosphorylation of the mitogen-activated protein kinase Erk was also reduced in p1108<sup>D910A/D910A</sup> B and T cells (Fig. 1C), which is consistent with the observation that, depending on signal strength, Erk activation can be regulated by PI3Ks (14).  $Ca^{2+}$  flux in response to anti-IgM crosslinking and to anti-CD3 crosslinking was attenuated in p1108<sup>D910A/D910A</sup> mice (Fig. 1D), in accord with an important role for PI3Ks upstream of Tec family kinase-mediated activation of phospholipase  $C\gamma$  (PLC $\gamma$ ) and  $Ca^{2+}$  signaling by antigen receptors (3, 15). These results implicate p1108 as an important mediator of antigen receptor signals in B and T cells.

The bone marrow of p1108<sup>D910A/D910A</sup> mice had reduced numbers of B220<sup>+</sup>IgM<sup>+</sup> B cell progenitors (Fig. 2A). More specifically, the ratios of pre- to pro-B cells were altered: in WT mice there were almost four times as many CD43<sup>-</sup>IgM<sup>+</sup> pre-B cells as CD43<sup>+</sup>IgM<sup>-</sup> pro-B cells, whereas the p1108<sup>D910A/D910A</sup> mice had equivalent numbers of pre- and pro-B cells (Fig. 2A). The spleens of p1108<sup>D910A/D910A</sup> mice had, on average, 50% fewer cells than the spleens of WT mice, with a more severe reduction of B cells than T cells (13). Analysis of the  $B220^+$ population in the spleen indicated similar levels of IgM<sup>+</sup>IgD<sup>+</sup> follicular B cells; however, the fraction of CD23<sup>-</sup>CD21<sup>+</sup>IgM<sup>+</sup> marginal zone B cells was reduced in p110 $\delta^{WT/D910A}$ spleens and was undetectable in the p110 $\delta^{D910A/D910A}$  spleens (Fig. 2B). CD5<sup>+</sup>IgM<sup>+</sup> peritoneal B-1 cells were readily detected in WT mice, were detected to a lesser extent in p110 $\delta^{WT/D910A}$  littermates, and were almost undetectable in p110 $\delta^{D910A/D910A}$  mice (Fig. 2C). These results show that p110 $\delta$  plays an important role in the development and differentiation of B cells.

T cell development in the thymus appeared to be normal as judged by CD4 versus CD8 profiles (Fig. 2D). The lymph nodes contained normal ratios of CD4<sup>+</sup> and CD8<sup>+</sup> T cells, but the expression of CD44, an activation marker that is also used to distinguish CD44<sup>lo</sup> naïve T cells from CD44<sup>hi</sup> memory T cells (*16*), was considerably reduced in T cells from p110 $\delta^{D910A/D910A}$  mice (Fig. 2E). These results indicate that p110 $\delta$  plays a role in the differentiation or survival of effector and, possibly, memory T cells. Peripheral blood cell counts appeared normal, with the exception of a twofold increase in the numbers of neutrophils (table S1).

Proliferation of purified B cells by anti-IgM cross-linking was almost completely abrogated (Fig. 3A). In contrast, proliferation in response to anti-CD40, interleukin 4 (IL-4), or lipopolysaccharide (LPS) was impaired but not abrogated (Fig. 3A). Stimulation with anti-CD40 in combination with IL-4 was also impaired (13). Proliferation of purified CD4<sup>+</sup> T cells in response to anti-CD3 was reduced

Fig. 1. Impaired PI3K signaling in B and T cells. immunoblots of (A) class IA PI3K subunits expressed in thymocytes. As a control for equal protein loading, the filter was probed with anti- $\beta$ -actin. (B) p110 subunits were immunoprecipitated from thymocyte lysates with isoform-specific antibodies, and the associated lipid kinase activity was assayed with phosphatidylinositol-4,5-bisphosphate as a substrate. Average results from duplicate measurements are represented as arbitrary units (a.u.). No PI3K activity was detected in p110δ immunoprecipitates from

in p1108<sup>D910A/D910A</sup> mice (Fig. 3B). In contrast, proliferation in response to costimulation with anti-CD3 and anti-CD28 was normal, or even enhanced, in  $p110\delta^{D910A/D910A}$ T cells (Fig. 3B). Stimulation of IL-2 production by anti-CD3 and anti-CD28 was normal (Fig. 3C), as was the capacity of phorbol 12,13-dibutyrate (PdBu)-stimulated T cells to proliferate in response to IL-2 (Fig. 3B). To examine the response of T cells to a physiological ligand, we crossed the p1108D910A/D910A mice with DO11-10 mice, which express a TcR specific for an ovalbumin peptide [Ova(323-339)] (17). T cells from p1108<sup>D910A/D910A</sup> DO11-10 mice exhibited diminished proliferative responses to the Ova peptide presented by WT antigen presenting cells (APCs) (Fig. 3D). IL-2 production was also attenuated in response to peptide stimulation, especially at elevated peptide concentrations (Fig. 3E). These results indicate that p1108 plays an important role in T cell activation but that the requirement for p110 $\delta$  can be overcome by stimulation through CD28, a receptor that can enhance proliferation and IL-2 production independently of PI3K (18). The functional interaction between B and T cells depends on integrins that stabilize cellcell adhesion. However, anti-CD3-induced cell adhesion to fibronectin via  $\beta 1$  integrins and to ICAM-1 via B2 integrins was unaffected by the p1108<sup>D910A</sup> mutation (Fig. 3F). TcR signaling is thought to be amplified by CD28-dependent accumulation of lipid rafts, membrane structures enriched in signaling proteins, at the interface between the T cell and the APC (19). The recruitment of lipid rafts, as induced by



p110 $\delta^{D910A/D910A}$  mice. Solid bars, WT; open bars, p110 $\delta^{D910A/D910A}$ . (**C**) Purified B cells were treated with or without soluble anti-lgM ( $\alpha$ IgM) F(ab')<sub>2</sub> (10 µg/ml). Purified T cells were treated with or without anti-CD3 ( $\alpha$ CD3) (10 µg/ml) bound to polystyrene beads. Immunoblots of total cell lysates with antibodies to Akt/PKB [total or Ser(*P*)<sup>473</sup>] or to Erk [total or Tyr(*P*)<sup>204</sup>] are shown. (**D**) Ca<sup>2+</sup> flux was measured in B cells stimulated with  $\alpha$ IgM F(ab')<sub>2</sub> (10 µg/ml) and in T cells stimulated with  $\alpha$ CD3 (5 µg/ml) and anti-hamster immunoglobulin (10 µg/ml) (24). Black, WT; gray, p110 $\delta^{D910A/D910A}$ .

В

stimulating T cells with beads coated with anti-CD3 and anti-CD28 (19, 20), was about 60% less frequent in p1108<sup>D910A/D910A</sup> T cells than in WT T cells (Fig. 3, G and H). Therefore, although raft recruitment appears to be dispensable for T cell activation by antibodies, it may play a critical role during T cell activation by peptide-major histocompatibility complex ligands presented by APCs.

The p110 $\delta^{D910A/D910A}$  mice had somewhat lower serum immunoglobulin levels than WT littermates, the IgM and IgA isotypes being most consistently reduced (Fig. 4A). To examine in vivo humoral immune responses, we immunized mice with TNP-KLH [(2,4,6-



trinitrophenyl)-keyhole limpet hemocyanin] and measured relative TNP-specific antibody titers in the serum on day 12. The total titer of TNP-specific antibodies made by the  $p110\delta^{D910A/D910A}$  mice was about 4% of that from the WT mice, with a more profound reduction of the IgM, IgG1, and IgG3 isotypes than IgG2a and IgG2b (Fig. 4B). Consistent with these findings, the spleens of TNP-KLHimmunized WT mice contained numerous germinal centers (GCs), whereas formation of these was largely absent in the spleens of TNP-KLH-immunized p1108<sup>D910A/D910A</sup> mice (Fig. 4C). These results demonstrate that T celldependent humoral immune responses are de-

Α

cpm (x 10²)

jm/gr

L-2 (ng/ml)

0

10-1

10-2 µmol peptide

10-3

0

ficient in p1108<sup>D910A/D910A</sup> mice. To measure T cell-independent antibody responses, we immunized mice with TNP-Ficoll and measured relative TNP-specific antibody titers on day 7. Consistent with the impaired capacity of p1108<sup>D910A/D910A</sup> B cells to signal through the BcR, T cell-independent humoral responses were strongly impaired (Fig. 4D).

Histological examination revealed lymphoid hypoplasia and lack of GCs in the spleen, lymph nodes, and Peyer's patches (PPs) in  $p110\delta^{D910A/D910A}$  mice (Fig. 4E, i to iv) (13). All other organs appeared to be normal, with the interesting exception of the large intestine. The  $p110\delta^{D910A/D910A}$  mice developed a mild

fibronectin

Fig. 3. Impaired B and T cell responses in vitro. (A) Purified spleen B cells were stimulated with anti-IgM ( $\alpha$ IgM) F(ab')<sub>2</sub> (10  $\mu$ g/ml),  $\alpha$ CD40 (10 µg/ml), IL-4 (20 ng/ ml), or LPS (10 µg/ml). Proliferation was measured by [3H]thymidine incorporation during the last 6 hours of a 48-hour culture. Similar results were obtained after 24 and 72 hours (13). Bars are as in Fig. 1. (B) Purified lymph node CD4<sup>+</sup> T cells were stimulated with polystyrene beads coated with  $\alpha$ CD3 (0.1, 1, or 10 µg/ml) with or withαCD28 (10 out µg/ml), and proliferation was measured as described in (A). Proliferation was also measured in response to PdBu (50 ng/ml) and IL-2 (10 ng/ml)



and in response to PdBu (50 ng/ml) and ionomycin (iono) (1  $\mu$ g/ml). (C) CD4<sup>+</sup> T cells were stimulated with polystyrene beads coated with  $\alpha$ CD3 (1  $\mu$ g/ml) with or without αCD28 (10 µg/ml), and supernatant was harvested after 24 hours for analysis of IL-2 production by enzyme-linked immunosorbent assay (ELISA). (D) CD4<sup>+</sup> T cells from WT and  $p110\delta^{D910A/D910A}$  mice that had been crossed onto the DO11-10 background were stimulated with mitomycin C-treated Balb/c spleen B cells and the indicated concentrations of Ova<sub>323-339</sub> peptide. Proliferation was measured by [<sup>3</sup>H]thymidine incorpo-

αCD28

1h

25 min

Fig. 2. Impaired B and T cell maturation. Cells from bone marrow (A), spleen (B), peritoneal washes (C), thymus (D), and lymph nodes (E) were analyzed by flow cytometry with the indicated antibodies (24). The percentage of gated cells in particular quadrant is indicated in (A to D). The percentage of CD44<sup>lo</sup> cells is indicated in (E).

ration during the last 6 hours of a 48-hour culture. Solid symbols, WT; open symbols, p110 $\delta^{D910A/D910A}$  (E) CD4<sup>+</sup> DO11-10 WT and p110 $\delta^{D910A/D910A}$  T cells were stimulated as in (D). Supernatants were harvested at 40 hours and assayed for IL-2 content by ELISA. (F) Adhesion of T cells stimulated with anti-CD3 to plate-bound fibronectin and ICAM-1 is expressed as fold increase in adhesion of anti-CD3-stimulated versus nonstimulated T cells. (G) Purified spleen and lymph node T cells were stimulated for 25 min with polystyrene beads coated with  $\alpha$ CD3,  $\alpha$ CD28, or  $\alpha$ CD3 and  $\alpha$ CD28. (Left) Bead (dotted circle) with a WT T cell, scored as positive (presenting raft aggregation toward the bead). (Right) p110 $\delta^{D910A/D910A}$  T cell/bead complex, scored as negative. Graph represents the percentage of cells presenting raft aggregation toward the bead. The values presented for stimulation by  $\alpha$ CD3 and  $\alpha$ CD28 are the average of six independent experiments and of two experiments for the other conditions. (H) Cells were incubated with  $\alpha$ CD3 and  $\alpha$ CD28coated beads, fixed after the indicated incubation times, and analyzed as described in (G). All results presented are from three or four measurements  $\pm$  SD, except where noted (24).







stained sections of mesenteric lymph nodes (i and ii), Peyer's patches (PPs) (iii and iv), and cecum (v to viii) of WT and p1108<sup>D910A/D910A</sup> mice. The mesenteric lymph node (i) and the PPs (iii) of WT mice have secondary follicles with prominent GCs (star). GCs are conspicuously absent from the follicles (star) of the mesenteric lymph node (ii) and PPs (iv) of p1108<sup>D910A/D910A</sup> mice. (v and vii) Normal intestinal mucosa of the cecum of a control mouse with short crypts (arrow) and sparsely cellular lamina propria. (vi) Cecum of p1108<sup>D910A/D910A</sup> mice have thickened mucosa with hyperplasic glands and mixed inflammatory infiltration of the lamina propria. (vii) Higher magnification of the cecum of a p1108<sup>D910A/D910A</sup> mouse showing focal erosion of mucosa with neutrophilic inflammation (arrowhead) and a regenerating gland (arrow). (i to vi) Bar, 300 µm; (vii and viii) bar, 50 µm (24).

inflammatory bowel disease (IBD), which was segmental and mostly limited to the cecal and rectal sections of the large intestine. The lesions were characterized by mucosal hyperplasia, crypt abscesses, and mixed leukocyte infiltrates, associated with regenerative changes of the glandular epithelium (Fig. 4E, v to viii). Other gene-targeted mice with T cell defects develop IBD, which may reflect an important role for regulatory T cells in maintaining tolerance to gut flora by secreting anti-inflammatory cytokines such as IL-10 and transforming growth factor- $\beta$  (21). IBD in humans, which includes Crohn's disease and ulcerative colitis, can result from inherited mutations in disease susceptibility genes interacting with environmental factors such as the intestinal bacterial flora. The human p1108 gene (PIK3CD) maps to the IBD7 susceptibility locus on chromosome 1p36 (22, 23). Given the IBD phenotype of p110 $\delta^{D910A/D910A}$  mice, *PIK3CD* should be further investigated as a candidate human IBD susceptibility gene.

p110 $\delta$  plays a unique role in antigen receptor signaling that is not readily compensated for by p110 $\alpha$  or p110 $\beta$ . The selective attenuation of immune function in p110 $\delta^{D910A/D910A}$  mice suggests that a specific inhibitor of  $p110\delta$  could effectively suppress B and T cell-mediated autoimmunity and possibly B and T cell transformation.

# **References and Notes**

- 1. B. Vanhaesebroeck et al., Annu. Rev. Biochem. 70, 535 (2001).
- R. Katso et al., Annu. Rev. Cell Dev. Biol. 17, 615 (2001).
- D. A. Fruman, L. C. Cantley, Semin. Immunol. 14, 7 (2002).
- T. Sasaki, A. Suzuki, J. Sasaki, J. M. Penninger, J. Biochem. (Tokyo) 131, 495 (2002).
- 5. D. A. Fruman et al., Science 283, 393 (1999).
- 6. H. Suzuki et al., Science 283, 390 (1999).
- 7. A. Suzuki et al., Immunity 14, 523 (2001).
- B. Vanhaesebroeck et al., Proc. Natl. Acad. Sci. U.S.A. 94, 4330 (1997).
- D. Chantry et al., J. Biol. Chem. 272, 19236 (1997).
  K. Okkenhaug, B. Vanhaesebroeck, Science's STKE 16 January 2001. http://stke.sciencemag.org/cgi/ content/full/OC\_sigtrans;2001/65/pe1
- J. M. Lu-Kuo, D. A. Fruman, D. M. Joyal, L. C. Cantley, H. R. Katz, J. Biol. Chem. 275, 6022 (2000).
- L. Bi, I. Okabe, D. J. Bernard, A. Wynshaw-Boris, R. L. Nussbaum, J. Biol. Chem. 274, 10963 (1999).
- 13. K. Okkenhaug et al., data not shown.
- B. C. Duckworth, L. C. Cantley, J. Biol. Chem. 272, 27665 (1997).
- 15. A. M. Scharenberg, J. P. Kinet, Cell 94, 5 (1998).
- J. Sprent, C. D. Surh, Annu. Rev. Immunol. 20, 551 (2002).

- K. M. Murphy, A. B. Heimberger, D. Y. Loh, Science 250, 1720 (1990).
- K. Okkenhaug et al., Nature Immunol. 2, 325 (2001).
  A. Viola, S. Schroeder, Y. Sakakibara, A. Lanzavecchia,
- Science 283, 680 (1999). 20. P. J. Ebert, J. F. Baker, J. A. Punt, J. Immunol. 165, 5435
- (2000). 21. K. J. Maloy, F. Powrie, *Nature Immunol.* **2**, 816 (2001).
- 22. N. Seki et al., DNA Res. 4, 355 (1997).
- 23. J. H. Cho et al., Hum. Mol. Genet. 9, 1425 (2000).
- 24. Supplementary Methods are available as supporting material on *Science* Online.
- 25. We thank L. Dobbie, M. Gomez, G. Wame, J. Punt, D. Cantrell, F. Powrie, S. Leevers, D. van Heel, A. Ridley, C. Sawyer, K. Ahmadi, and H. Okkenhaug for help and advice. Supported by the Ludwig Institute for Cancer Research, the UK Biotechnology and Biological Sciences Research Council (B.V., A.J.H.S., and W.P.), Diabetes UK (B.V., A.J.H.S., and A.S.), European Union Fifth Framework Programme QLG1-2001-02171 (B.V. and K.O.), l'Association pour la Recherche sur le Cancer and European Union Marie Curie (G.F.), Fondazione Italiana per la Ricerca sul Cancro and Università di Napoli Federico II (A.B.), and the Flanders Fund for Scientific Research, Belgium (B.V.).

# Supporting Online Material

www.sciencemag.org/cgi/content/full/1073560/DC1 Materials and Methods

- Fig. S1 Table S1
- Table 31

3 May 2002; accepted 3 July 2002

- Published online 18 July 2002;
- 10.1126/science.1073560

Include this information when citing this paper.