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Transcriptional Coactivator Cited2 Induces Bmi1 and Mel18 and Controls Fibroblast Proliferation via Ink4a/ARF

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Cited2 (CBP/p300 interacting transactivator with ED-rich tail 2) is required for embryonic development, coactivation of transcription factor AP-2, and inhibition of hypoxia-inducible factor 1 transactivation. Cited2 is induced by multiple growth factors and cytokines and oncogenically transforms cells. Here, we show that the proliferation of Cited2−/− mouse embryonic fibroblasts ceases prematurely. This is associated with a reduction in growth fraction, senescent cellular morphology, and increased expression of the cell proliferation inhibitors p16INK4a, p19ARF, and p15INK4b. Deletion of INK4a/ARF (encoding p16INK4a and p19ARF) completely rescued the defective proliferation of Cited2−/− fibroblasts. However, the deletion of INK4a/ARF did not rescue the embryonic malformations observed in Cited2−/− mice, indicating that INK4a/ARF-independent pathways are likely to be involved here. We found that Cited2−/− fibroblasts had reduced expression of the polycomb-group genes Bmi1 and Mel18, which function as INK4a/ARF and Hox repressors. Complementation with CITED2-expressing retrovirus enhanced proliferation, induced Bmi1/Mel18 expression, and decreased INK4a/ARF expression. Bmi1- and Mel18-expressing retroviruses enhanced the proliferation of Cited2−/− fibroblasts, indicating that they function downstream of Cited2. Our results provide genetic evidence that Cited2 controls the expression of INK4a/ARF and fibroblast proliferation, at least in part via the polycomb-group genes Bmi1 and Mel18.

p300 and its paralog, CBP (CREB-binding protein), are ubiquitously expressed nuclear proteins that function as transcriptional coactivators and histone acetyl transferases, connecting DNA-bound transcription factors to the core transcription machinery (reviewed in references 19 and 50). p300 and CBP are essential for normal cardiovascular, neural, and hematopoietic development (31, 39, 60). p300 and CBP also play a fundamental role in cellular growth control (reviewed in reference 19). Genetic evidence indicates that CBP is a tumor suppressor. Patients with CBP mutations (Rubinstein-Taybi syndrome) (41) have a high incidence of neural and developmental tumors (38), and mice lacking a single CBP allele develop hematologic malignancies (31). Consistent with this finding, p300 and CBP function as coactivators of the tumor suppressor p53 (8, 34). Paradoxically, p300 and CBP are also required for cell proliferation. Embryonic fibroblasts lacking p300 proliferate poorly in culture (60), and neutralization of p300/CBP by antibody injection inhibits progression through the G1/S transition (1). In keeping with this finding, many oncogenic transcription factors require either p300 or CBP for transactivation (reviewed in reference 19).

p300 and CBP also interact with members of the CBP/p300 interacting transactivator with ED-rich tail (CITED) family. These include CITED1/MSG1 (58); CITED2, splice isoforms of which are known as p35srj/Mrg1 (12, 33); and CITED4 (15, 59). Loss of Cited2 in mice results in embryonic lethality as a consequence of cardiac malformations, neural tube defects, and adrenal gland agenesis (9, 10, 43, 57, 61). At a biochemical level, CITED2 physically interacts with and coactivates all transcription factor AP-2 (TFAP2) isoforms and is necessary for TFAP2 function (9, 14). CITED2 also inhibits hypoxia-inducible factor 1 alpha (HIF-1α) transactivation by disrupting the HIF-1α–p300 interaction (12, 61). These molecular mechanisms are thought to underlie the embryonic malformations observed in mice lacking Cited2. CITED2 is induced by multiple growth factors and cytokines (e.g., interleukin-1α [IL-1α], IL-2, IL-4, IL-6, IL-9, IL-11, granulocyte-macrophage colony-stimulating factor, platelet-derived growth factor, and insulin), and overexpression of CITED2 results in oncogenic cell transformation (51). The response of CITED2 to mitogenic stimuli and its ability to transform cells suggest that it may function in cell growth control (51). To understand the genetic pathways by which CITED2 may control cell proliferation, we studied mouse embryonic fibroblasts lacking Cited2.

MATERIALS AND METHODS

Mice. Cited2−/− and Cited2+/+ embryos were on a 129Ola/C57BL/6J mixed background or a C57BL/6J background as indicated and were generated as previously described (9). Cited2−/−:INK4a/ARF−/− (INK4, inhibitor of cyclin-dependent kinase; ARF, alternative reading frame) embryos and the relevant controls (see Fig. 5 and 6) were on an FVB/129Ola/C57BL/6J mixed background and were generated by intercrossing Cited2−/−:INK4a/ARF−/− mice. INK4a/ARF−/− mice (45) were kind gifts from Ronald DePinho (DFCI, Boston, Mass.). Mice, embryos, and fibroblasts were genotyped by using PCR with allele-specific primers (9).
CITED2 CONTROLS FIBROBLAST PROLIFERATION

Vol. 23, 2003

Fibroblast isolation, passage, growth curves, and colony formation assays. Murine embryonic fibroblasts were prepared from littermate embryos at 13.5 or 15.5 days postcoitus (dpc) as previously described (36). Adherent fibroblasts were harvested the following day and plated at a density of 1.5 × 10^5 cells per cm^2 (passage 0) and passed every 3 days thereafter at the same density. Population doubling per passage was calculated as log(ni/ni+1)/log2, where ni is the initial and ni+1 the final number of cells at each passage (13). When ni+1 was greater than ni, the population doubling was defined as 0. Cumulative population doubling (CDP) at each passage was calculated by adding population doubling per passage (13). Senescence-associated β-galactosidase was detected as previously described (17). For growth curves, the indicated number of cells per well of a 12-well plate were plated and harvested at the indicated time points and relative cell numbers were measured with crystal violet as previously described (46). Values were normalized to day 1 for the indicated culture, and each point was determined in quadruplicate. For colony formation, fibroblasts were plated at the densities indicated and the colonies were visualized with Giemsa stain as previously described (13).

Growth fraction. Cells were plated at a density of 1.5 × 10^4/cm^2 on glass coverslips at the indicated passages, and after 48 h, 10 μM bromodeoxyuridine (BrdU) was added for 24 h. The coverslips were fixed, incubated in HCl, and then stained with anti-BrdU monoclonal antibody (Becton-Dickinson) followed by secondary sheep anti-mouse fluorescein isothiocyanate-conjugated antibody. Cells were stained with propidium iodide (PI) and mounted in Vectashield containing PI (Vector). Nuclear uptake of BrdU and PI was quantitated on a WinCyte software according to the manufacturer’s instructions. Data from the results for 3,000 to 5,000 cells were acquired for each individual experiment. The growth fraction was calculated as the percentage of BrdU-positive cells in the culture.

Retroviruses. CITED2, CITED2Δ2, Mel18, Bmi1, and control retroviral supernatants were generated by using the bicistronic pLZR5-IRE-RES-GFP plasmid and Phoenix producer cells (gifts from Garry Nolan, Stanford, Calif.). Fibroblasts were infected as previously described (25). The Mel18 retrovirus has been described previously (25). The Mel18 retrovirus was generated from pg55-Mel18, which contains a mouse Mel18 cDNA insert (gift from M. Kanno, Hiroshima, Japan) with a modified translation start site (GCCACCATTG) that changes the second amino acid from H to D. We used PCR to convert the Mel18 translation start site back to the wild-type form (GGCATCATGC) (GenBank accession no. D99088) and subcloned an EcoRI-SalI fragment (containing the open reading frame) into pLZR5-IRE-RES-GFP. Induction efficiencies typically exceeded 80%. Plasmid vectors were generated by using standard molecular cloning techniques (7).

Blotting. Northern blotting was performed as previously described (7) by using 6 to 10 μg of total RNA (NEasy Mini-kits; QIAGEN) transferred to Hybond N+ membranes (Amersham). 28S and 18S RNA species were visualized by staining with 0.05% methylene blue. Mouse p19ARF, p16INK4a, and p15INK4b cDNA plasmids were gifts from Charles Sherr (HHML, Memphis, Tenn.) and Gordon Peters (Cancer Research UK, London, United Kingdom). p16INK4a, and p19ARF-specific probes (INK4a/ARF1s and INK4a/ARF1β) were generated from respective cDNA templates by PCR with exon-specific primers. Northern blotting for Mel18, Bmi1, and Mph1 was performed from early passage, non-confluent mouse embryonic fibroblasts by using the respective murine probes. The Mph1 probe (IMAGE:3512187) was obtained from MRC-HGMP, Cambridge, United Kingdom. The relative signal intensity was measured by using NIH Image software on scanned autoradiograms. Western blotting with anti-Bmi1 monoclonal antibody (229F6; Upstate Biotechnology, Lake Placid, N.Y.), p16INK4a (M156; Santa Cruz) and anti-p19ARF (Ab80; Abcam, Cambridge, United Kingdom) polyclonal antibodies, and antitubulin antibody (T-2029; Sigma) was performed according to the instructions of the manufacturer.

Histology. Embryos were fixed in 4% paraformaldehyde, dehydrated in ethanol, and embedded in paraffin wax. Sections (7 μm thick) were stained with hematoxylin and eosin.

RESULTS

Premature proliferation arrest of CITED2−/− fibroblasts. We investigated the proliferative capacity of CITED2+/+ and CITED2−/− primary mouse embryonic fibroblasts. Fibroblasts were derived from littermate embryos and passed every 3 days in parallel and under identical conditions at a density of 1.5 × 10^5 cells/cm^2, equivalent to a 3T3 protocol (36). CITED2−/− fibroblasts proliferated rapidly and then slowly transiently at passages 5 and 6, following which proliferation was rapid and continuous (Fig. 1A). CITED2−/− fibroblasts proliferated normally in the first three passages but slowed dramatically and then arrested permanently. This premature proliferation arrest was confirmed in three independent experiments by using independently isolated fibroblasts (data not shown) as well as fibroblasts isolated from cosogenic mice generated by back-crossing the CITED2 mutation to the C57BL/6j background for more than nine generations (Fig. 1B). These results were also corroborated by plating fibroblasts at passage 3 and assaying cell growth over the next 14 days without replating (Fig. 1C). By passage 3, the CITED2−/− fibroblasts had a decreased ability to proliferate in comparison to that of the CITED2+/+ fibroblasts. Cultures of either genotype had indistinguishable spindle-shaped cells at initial plating. With passage, CITED2−/− fibroblasts cultures rapidly accumulated cells that had a flattened appearance and cytoplasmic enlargement and expressed senescence-associated β-galactosidase (17) (Fig. 1D and E).

Reduction in growth fraction. To determine the mechanism of impaired proliferation in CITED2−/− fibroblasts, the fraction of actively proliferating cells (11) was measured by labeling parallel cultures with BrdU for 24 h. Bivariate analysis for BrdU uptake and PI staining showed that the growth fraction of CITED2+/+ and CITED2−/− fibroblasts was close to 100% when initially plated in culture (Fig. 2A and B). With serial passage, the growth fraction of CITED2−/− fibroblasts declined more rapidly than that of CITED2+/+ fibroblasts. The fall in growth fraction in the CITED2−/− cultures was evident as early as the first passage. All cultures were initially predominantly diploid, as determined by PI staining (Fig. 2A). With serial passage, cultures became tetraploid (Fig. 2A, middle and lower panels). Although the wild-type culture shown in the top panel of Fig. 2A was still predominantly diploid at this passage, it became predominantly tetraploid at the next passage (data not shown).

CITED2−/− fibroblasts have increased expression of INK4a/ARF. The above results showed that CITED2−/− fibroblasts cease to proliferate prematurely when cultured, do not spontaneously immortalize, and express morphological features of cellular senescence. Senescence in mouse embryonic fibroblasts is associated with increased levels of the alternatively spliced products of the INK4a/ARF locus, p16INK4a, and p19ARF (Fig. 3A) (reviewed in references 48 and 49). Members of the INK4 family are known to interfere with cyclin-dependent kinases 4 and 6, whereas p19ARF functions to inhibit MDM2, a repressor of p53. We examined the expression of INK4a/ARF in early passage CITED2+/+ and CITED2−/− fibroblasts (Fig. 3B through F). Northern blots were probed with a p19ARF cDNA, which detects both p16INK4a and p19ARF (Fig. 3A and B) as comigrating products. This probe showed that INK4a/ARF expression was clearly increased (2.7-fold) in CITED2−/− fibroblasts (Fig. 3B). Specific probes that discriminate between the alternatively spliced p16INK4a and p19ARF transcripts showed that the expression of both transcripts was increased (4.2- and 2.7-fold, respectively) in fibroblasts lacking Cited2 (Fig. 3C and D). The expression of p15INK4b, a member of the INK4 family (21) (Fig. 3E), was also increased (2.5-fold) in CITED2−/− fibroblasts (Fig. 3A). We observed no change in the expression of p19INK4a, another INK4 family member (23) (Fig. 3F). Consistent with these observations, the levels of p16INK4a and...
p19\textsuperscript{ARF} proteins were also increased (2.5- and 2.7-fold, respectively) in \textit{Cited2}\textsuperscript{−/−} fibroblasts at passage 1 (Fig. 3G and H).

**Complementation with CITED2 enhances proliferation.** To determine if these changes were specific for loss of \textit{Cited2}, we infected \textit{Cited2}\textsuperscript{−/−} fibroblasts with a bicistronic retrovirus expressing human CITED2 (which is 94\% conserved with mouse \textit{Cited2}) and GFP (green fluorescent protein) driven by a retroviral long terminal repeat (Fig. 4A). These fibroblasts showed a marked enhancement of proliferation that was maintained for the period studied (i.e., greater than 30 days) and retained their spindle shape (data not shown). In comparison, parallel infection with a control GFP-expressing retrovirus (LZRS) or with retrovirus expressing CITED2\textsubscript{Δ}, a mutant lacking the overlapping CBP/p300 and TFIAP2 binding domains (9, 12), did not enhance proliferation. These results were confirmed in two further experiments using independently isolated \textit{Cited2}\textsuperscript{−/−} fibroblasts. Infection of \textit{Cited2}\textsuperscript{−/−} fibroblasts with the CITED2-expressing retrovirus also led to a modest reduction in \textit{INK4a/ARF} and \textit{INK4b} expression (1.5- and 1.6-fold, respectively) (Fig. 4B and C) in comparison to control retrovirus.

**An intact \textit{INK4a/ARF} gene is essential for proliferation arrest in \textit{Cited2}\textsuperscript{−/−} fibroblasts.** The above results suggested that \textit{Cited2} enhances fibroblast proliferation by repressing \textit{INK4a/ARF} and/or \textit{INK4b}. To definitively establish the role of \textit{INK4a/ARF} in mediating the premature proliferation arrest of \textit{Cited2}\textsuperscript{−/−} fibroblasts, we generated fibroblasts lacking both \textit{Cited2} and \textit{INK4a/ARF} (\textit{Cited2}\textsuperscript{−/−}: \textit{INK4a/ARF}\textsuperscript{−/−}) and compared their proliferation with that of \textit{Cited2}\textsuperscript{−/−}: \textit{INK4a/ARF}\textsuperscript{+/−}, \textit{Cited2}\textsuperscript{−/−}: \textit{INK4a/ARF}\textsuperscript{+/-}, and wild-type fibroblasts (Fig. 5A and B). Consistent with previous observations (45), fibroblasts lacking \textit{INK4a/ARF} proliferated more rapidly than wild-type fibroblasts during serial passage in culture, with no slowing of proliferation for the duration of the experiment (48 days). Consistent with the observations in Fig. 1, \textit{Cited2}\textsuperscript{−/−} fibroblasts arrested prematurely and permanently. Fibroblasts lacking both \textit{Cited2} and \textit{INK4a/ARF} proliferated almost as rapidly as those lacking \textit{INK4a/ARF}, with no slowing of prolif-
that Cited2 controls other pathways that are relevant for embryonic development.

Cells lacking Cited2 have reduced Bmi1 and Mel18 expression. The above data indicated that Cited2 enhances cell proliferation by repressing INK4a/ARF. Genetic evidence indicates that in primary mouse fibroblasts the polycomb-group gene Bmi1 represses p16INK4a and p19ARF and that Mel18 (a Bmi1 paralog) represses p16INK4a (25). We therefore examined the expression of these genes in early passage Cited2+/+ and Cited2−/− fibroblasts derived from littermate embryos. We found that both Mel18 and Bmi1 expression was reduced (2.3- and 2.2-fold, respectively) in Cited2−/− fibroblasts (Fig. 7A and B). There was no significant change in the expression of Mph1, another polycomb group gene. We also examined the fibroblasts for TBX2, another INK4a/ARF repressor (24), but expression of this gene was not detected in wild-type or Cited2−/− fibroblasts. These results indicate that Cited2 controls other pathways that are relevant for embryonic development.

Elimination of INK4a/ARF does not rescue embryonic malformations in Cited2−/− mice. To determine if INK4a/ARF plays a role in the genesis of embryonic malformations in mice lacking Cited2, we examined embryos lacking both Cited2 and INK4a/ARF. Like embryos lacking only Cited2 (9), those lacking both Cited2 and INK4a/ARF had cardiac malformations (Fig. 6B), adrenal agenesis (Fig. 6D), and exencephaly (Fig. 6F). In these experiments, exencephaly was observed in 4 of 8 embryos lacking Cited2 and 6 of 13 embryos lacking both Cited2 and INK4a/ARF. Control embryos that were wild type for Cited2 but lacked INK4a/ARF had normal heart, adrenal, and neural development (Fig. 6A, C, and E). These results indicate that Cited2 controls other pathways that are relevant for embryonic development.

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ed2\(^{-/-}\) fibroblasts (data not shown). Infection of Cited2\(^{-/-}\) fibroblasts with CITED2-expressing retrovirus resulted in a modest increase in expression of Mel18 (1.5-fold) and Bmi1 (1.4-fold) (Fig. 7C and D).

**Bmi1 and Mel18 enhance proliferation of cells lacking Cited2.** The above data indicated that Cited2 is required for normal Bmi1 and Mel18 expression. To determine if the observed deficiency of Bmi1 and Mel18 expression in Cited2\(^{-/-}\) fibroblasts would explain their premature proliferation arrest, we infected Cited2\(^{-/-}\) and Cited2\(^{-/-}\) fibroblasts with bicistronic retroviruses expressing either Bmi1 and GFP, Mel18 and GFP, or GFP alone (as the control) and performed cell proliferation and colony formation assays (Fig. 8A and B). Infection with Bmi1- or Mel18-expressing retroviruses led to enhanced proliferation compared to that for the control retrovirus, regardless of the Cited2 genotype (Fig. 8A). The Bmi1 retrovirus enhanced proliferation better than the Mel18 retrovirus, again regardless of Cited2 genotype. We next examined the ability of these fibroblasts to form colonies when plated at low density (Fig. 8B). Consistent with the above results, infection with Bmi1- or Mel18-expressing retroviruses led to enhanced colony formation (with Bmi1 being more efficient than Mel18) compared to that for the control retrovirus, regardless of the Cited2 genotype. This enhancement of proliferation by Bmi1 and by Mel18 was reproducibly observed in Cited2\(^{-/-}\) fibroblasts obtained from independently harvested embryos. These results indicate that Bmi1 or Mel18 can enhance the proliferation of fibroblasts lacking Cited2 and that Cited2 function is not required for proliferation enhancement by these polycistron-expressing proteins.

**DISCUSSION**

Normal primary mouse embryonic fibroblasts proliferate rapidly when explanted but soon slow down in response to the stress of culture (reviewed in reference 48). This is associated with cell cycle exit, cytoplasmic enlargement, and expression of senescence-associated \(\beta\)-galactosidase, which are characteristics of cellular senescence (17). At a molecular level in murine cells, this senescent phenotype is associated with increased levels of the cell proliferation inhibitors p16\(^{INK4a}\) and p19\(^{ARF}\) and activation of the p19\(^{ARF}\) target p53 and the p53 target gene p21\(^{WAF1}\) (reviewed in references 48 and 49). Genetic evidence indicates that intact p19\(^{ARF}\) and p53 are necessary for the senescent phenotype of cultured mouse embryonic fibroblasts (45; reviewed in reference 48). Deletion of p19\(^{ARF}\) alone abolishes the senescent phenotype, whereas deletion of p16\(^{INK4a}\) alone does not have this effect, indicating that p19\(^{ARF}\) plays a critical role (28, 30, 47). Notably, the spontaneous immortalization of wild-type fibroblasts observed when they are serially passaged in culture typically results from spontaneous mutations in p19\(^{ARF}\) or p53 (28, 62). Premature senescence of primary fibroblasts is associated with mutations in genes such as Bmi1, JunD, Atm, and Lig4 that function upstream of p19\(^{ARF}\) (18, 25, 27, 56). It is also associated with mutations in Mel18 (25) and Id1 (5), which repress p16\(^{INK4a}\) expression. Overexpression of activated cellular oncogenes...
FIG. 6. Embryonic malformations in embryos lacking both Cited2 and Ink4a/Arf. (A through D) Hematoxylin- and cosin-stained transverse sections from Cited2−/−;Ink4a/Arf−/− and Cited2−/−;Ink4a/Arf−/− embryos at 14.5 dpc. Scale bars, 0.1 mm. Axes: d, dorsal; v, ventral; r, right; l, left. (A) Transverse section through the thorax of a Cited2−/−;Ink4a/Arf−/− embryo shows normal cardiac anatomy. The interatrial septum (ias) separates the right (ra) and left (la) atria. The right (rv) and left (lv) ventricles are separated by a normal interventricular septum (ivs) (B) Corresponding section of a Cited2−/−;Ink4a/Arf−/− embryo showing normal right (rad) and left (lad) adrenal glands and left kidney (lk). (C) Transverse section through the abdomen of a Cited2−/−;Ink4a/Arf−/− embryo showing ventricular septal defect (VSD). (D) Corresponding section of a Cited2−/−;Ink4a/Arf−/− embryo showing normal cranial development. (F) Cited2−/−;Ink4a/Arf−/− embryo showing exencephaly.

FIG. 7. Expression of Mel18 and Bmi1 in fibroblasts lacking Cited2. (A) Northern blots of total RNA obtained from Cited2−/− (w) and Cited2−/− (n) fibroblasts derived from littermate embryos. The top panels show blots probed for Mel18, Bmi1, or Mph1, as indicated. Two Mel18 isoforms are noted (52). The bottom panels show 28S and 18S RNA species to show equal loading. (B) Western blot of total cell lysate probed with an anti-Bmi1 monoclonal antibody (top) and re-probed with an antitubulin monoclonal antibody (bottom) to show equal loading. (C) Northern blot of total RNA obtained from Cited2−/− fibroblasts infected in parallel with retroviruses. LZ indicates infection with LZRS (control) retrovirus, and C2 indicates infection with CITED2-expressing retrovirus. The top panel shows a blot probed with a Mel18 cDNA. The bottom panel shows 28S and 18S RNA species to show equal loading. (D) Western blot of total cell lysate from Cited2−/− fibroblasts. LZ indicates infection with LZRS (control) retrovirus and C2 indicates infection with CITED2-expressing retrovirus. The experiment was performed in duplicate using independently isolated Cited2−/− fibroblast lines (nos. 1 and 2). The top panel shows a blot probed with an anti-Bmi1 antibody. The bottom panel shows a blot probed with an antitubulin antibody to show equal loading.

such as Ras (37, 46) and MEK (35) in primary cells results in senescence by activating the p19ARF-p53 pathway.

The results presented here provide genetic evidence that Cited2, a growth factor and cytokine-inducible gene with oncogenic potential (51), is necessary for fibroblast proliferation in culture. Fibroblasts lacking Cited2 stop dividing prematurely, display typical senescent morphology, and express senescence-associated β-galactosidase (Fig. 1 and 2). These results suggested that they are hypersensitive to culture-induced stress. The expression of p16INK4a, p19ARF, and p15INK4b but not p16INK4a was markedly increased in Cited2−/− fibroblasts (Fig. 3). These results indicated that Cited2 is required for the coordinated repression of the physically linked Ink4a/Arf and Ink4b genes. Ink4a/Arf and Ink4b were also repressed by complementation with CITED2, which also enhanced cellular proliferation, suggesting a causal mechanism (Fig. 4). Although the reduction of Ink4a/Arf and Ink4b by retroviral transduced CITED2 was modest, it was reproducible and supports the idea that Cited2 represses these genes.

Embryos lacking Cited2 invariably have heart and adrenal gland defects, and ~50% of embryos have exencephaly (9, 42–44). Premature fibroblast senescence occurred regardless of the presence or absence of exencephaly (e.g., the mutant embryos in Fig. 1A and B had normal neural development). Nevertheless, it was possible that the embryonic heart or adrenal malformation itself affects (through secondary changes) the growth of fibroblasts. To address this issue, we performed the complementation experiment with CITED2 and showed, reproducibly, that complementation with retroviral CITED2 markedly enhances proliferation of Cited2−/− fibroblasts (Fig. 4A). This effect was specific, as it was not seen with a CITED2 mutant lacking residues 215 to 270. Successful complementation indicates that no secondary change, e.g., one induced by the heart, adrenal, or neural defect, was responsible for the fibroblast growth defect. Thus, the premature senescence observed in fibroblasts lacking Cited2 is indeed specific and is unlikely to be due to the preexisting embryonic malformation.
INK4a 
the elevated levels of complementation with CITED2, these experiments show that 
as assay.

Complementation factors: The proliferation of retrovirally complemented 
4 (referred to as P0), and passaged in parallel every 3 days at identical 
harvested, infected with retroviruses the following day, replated on day 

Amount of DNA derived from littermate embryos on a C57BL/6J background at 13.5 dpc) 
senescence phenotype and that hurricanes play a major causal role in generating the premature 
stream target of Cited2 

Blasts have two independent functions: first, a role in embryonic 
development, and second, a role in fibroblast proliferation under conditions of culture-induced stress. One possible mechanism is that Cited2 positively regulates genes that not only repress INK4a/ARF but also have independent roles in development. Members of the polycomb family (e.g., Bmi1 and Mel18) are known to play these dual roles (2, 25, 55).

Genetic evidence indicates that in primary mouse fibroblasts Bmi1 represses p16INK4a and p19ARF and Mel18 represses p16INK4a (25). Deletion of either Bmi1 or its paralog Mel18 leads to reduced lymphocyte precursor proliferation and premature proliferation arrest of primary mouse embryonic fibroblasts (3, 25). Bmi1 is also necessary for self-renewal of hematopoietic stem cells (32, 40). Deletion of INK4a/ARF in mice lacking Bmi1 rescues premature fibroblast senescence and postnatal cerebellar and lymphoid defects (25). However, Mel18 has more complex functions, as evidenced by the fact that it can also function as a cell proliferation inhibitor in other cell types (29, 53). Differences in Bmi1 and Mel18 function are also suggested by distinct phenotypes observed in mutant mice: for instance, cerebellar defects are observed in mice lacking Bmi1, and colonic smooth muscle defects are seen in mice lacking Mel18 (2, 55).

Bmi1 and Mel18 function as transcriptional repressors that interact with a similar set of polycomb-group proteins (6, 20, 22, 54). They function during development to repress Hox genes, and deletion of either gene leads to defects in anteroposterior axis formation (2, 55). Mel18 and Bmi1 act synergetically in a dose-dependent manner during development to maintain Hox gene expression and cell survival (4). Importantly, we have observed anteroposterior patterning defects in Cited2 mutant embryos. These include fusion of cranial ganglia (9) and of the cervical vertebrae (S. D. Bamforth and S. Bhattacharya, unpublished observations). These observations prompted us to examine Bmi1 and Mel18 expression in fibroblasts lacking Cited2.

We found that fibroblasts lacking Cited2 have a marked reduction in levels of Bmi1 and Mel18 transcripts (Fig. 7). Complementation of Cited2−/− fibroblasts with CITED2 led to a modest increase in the expression of Bmi1 and Mel18. We also found that both Bmi1 and Mel18 enhanced the proliferation of fibroblasts regardless of the Cited2 genotype (Fig. 8), indicating that Cited2 is not necessary for proliferation enhancement by these polycomb-group genes and supporting the idea that Bmi1 and Mel18 function downstream of Cited2. However, after infection with Bmi1- and Mel18-expressing retroviruses, Cited2+/- fibroblasts proliferated faster than Cited2−/− fibroblasts, implying that Cited2 deficiency cannot be completely rescued by overexpression of Bmi1 or Mel18 indi-

Notably, residues 215 to 270 of CITED2 contain the overlapping TFAP2 and CBP/p300 binding domains (9, 12), suggesting that binding of CITED2 to CBP/p300 and/or TFAP2 is required for enhancement of cell proliferation.

We also found that deletion of INK4a/ARF completely rescued the proliferation defect in fibroblasts lacking Cited2 (Fig. 5). This finding was observed reproducibly in fibroblasts obtained from independently isolated embryos. As the proliferation defect in Cited2-deficient fibroblasts is reproducibly observed on both mixed (Fig. 1A and Fig. 5A and C) and pure (Fig. 1B) genetic backgrounds, the reproducible rescue of senescence by deletion of INK4a/ARF in Cited2−/− fibroblasts indicates that random segregation of genetic modifiers in these experiments does not likely play an important role. Taken together with the increased levels of INK4a/ARF observed in Cited2−/− fibroblasts and the suppression of INK4a/ARF by complementation with CITED2, these experiments show that the elevated levels of INK4a/ARF observed in Cited2−/− fibroblasts play a major causal role in generating the premature senescence phenotype and that INK4a/ARF is a critical downstream target of Cited2 in fibroblasts. The complete rescue in cell proliferation that we observed also indicates that no other downstream mechanism (e.g., the activation of p53 or p27 by a different mechanism, such as HIF-1 activation in cells lacking Cited2) is likely to be involved. These results are also supported by experiments which show that fibroblasts lacking Cited2 are efficiently immortalized by overexpression of the p19ARF repressor TBX2 (24) and by antisense p19ARF retrovirus (16) (K. R. Kranc and S. Bhattacharya, unpublished observations).

The above data indicated that Cited2 enhances cell proliferation by repressing INK4a/ARF. However, deletion of the INK4a/ARF locus did not rescue the embryonic malformations (cardiac, adrenal, and neural) associated with mutation in Cited2 (Fig. 6). Thus, the Cited2-mediated repression of INK4a/ARF observed in fibroblasts does not appear to play a significant role in embryonic development. This finding indicates that Cited2 has two independent functions: first, a role in embryonic development, and second, a role in fibroblast proliferation under conditions of culture-induced stress. One possible mechanism is that Cited2 positively regulates genes that not only repress INK4a/ARF but also have independent roles in development. Members of the polycomb family (e.g., Bmi1 and Mel18) are known to play these dual roles (2, 25, 55).

Genetic evidence indicates that in primary mouse fibroblasts Bmi1 represses p16INK4a and p19ARF and Mel18 represses p16INK4a (25). Deletion of either Bmi1 or its paralog Mel18 leads to reduced lymphocyte precursor proliferation and premature proliferation arrest of primary mouse embryonic fibroblasts (3, 25). Bmi1 is also necessary for self-renewal of hematopoietic stem cells (32, 40). Deletion of INK4a/ARF in mice lacking Bmi1 rescues premature fibroblast senescence and postnatal cerebellar and lymphoid defects (25). However, Mel18 has more complex functions, as evidenced by the fact that it can also function as a cell proliferation inhibitor in other cell types (29, 53). Differences in Bmi1 and Mel18 function are also suggested by distinct phenotypes observed in mutant mice: for instance, cerebellar defects are observed in mice lacking Bmi1, and colonic smooth muscle defects are seen in mice lacking Mel18 (2, 55). Bmi1 and Mel18 function as transcriptional repressors that interact with a similar set of polycomb-group proteins (6, 20, 22, 54). They function during development to repress Hox genes, and deletion of either gene leads to defects in anteroposterior axis formation (2, 55). Mel18 and Bmi1 act synergistically in a dose-dependent manner during development to maintain Hox gene expression and cell survival (4). Importantly, we have observed anteroposterior patterning defects in Cited2 mutant embryos. These include fusion of cranial ganglia (9) and of the cervical vertebrae (S. D. Bamforth and S. Bhattacharya, unpublished observations). These observations prompted us to examine Bmi1 and Mel18 expression in fibroblasts lacking Cited2.

We found that fibroblasts lacking Cited2 have a marked reduction in levels of Bmi1 and Mel18 transcripts (Fig. 7). Complementation of Cited2−/− fibroblasts with CITED2 led to a modest increase in the expression of Bmi1 and Mel18. We also found that both Bmi1 and Mel18 enhanced the proliferation of fibroblasts regardless of the Cited2 genotype (Fig. 8), indicating that Cited2 is not necessary for proliferation enhancement by these polycomb-group genes and supporting the idea that Bmi1 and Mel18 function downstream of Cited2. However, after infection with Bmi1- and Mel18-expressing retroviruses, Cited2+/- fibroblasts proliferated faster than Cited2−/− fibroblasts, implying that Cited2 deficiency cannot be completely rescued by overexpression of Bmi1 or Mel18 indi-

FIG. 8. Growth properties of Cited2−/− and Cited2+/- embryonic fibroblasts infected with Bmi1- and Mel18-expressing retroviruses. (A) Proliferation of Cited2−/− (n) and Cited2+/- (w) fibroblasts (derived from littermate embryos on a C57BL/6J background at 13.5 dpc) infected with Bmi1, Mel18, or control retroviruses. Fibroblasts were harvested, infected with retroviruses the following day, replated on day 4 (referred to as P0), and passaged in parallel every 3 days at identical conditions. The proliferation of retrovirally complemented fibroblasts is shown as plots of CPD versus passage number. (B) Colony formation assay. Cited2−/− and Cited2+/- fibroblasts (at passage 6) infected with the indicated retroviruses were plated at a density of 4,000 cells per 9-cm plate, and colonies were visualized with Giemsa stain after 16 days.
vidually. One possibility is that other Cited2 functions that are independent of Bmi1 or Mel18 may be important. Another possibility is that Cited2 is required for the coordinated induction of Bmi1 and Mel18, which is not mimicked by the forced expression of either Bmi1 or Mel18 alone.

In summary, these data indicate that Cited2 is required for normal Bmi1 and Mel18 expression in primary mouse embryonic fibroblasts and that Bmi1 and Mel18 function downstream of Cited2. The mechanism by which Cited2 induces Bmi1 and Mel18 is not understood at present. One possibility is that a coactivation function of Cited2 is required for Bmi1/Mel18 expression. Alternatively, Cited2 may function several steps away, even perhaps via nonautonomous cell mechanisms, to control Bmi1/Mel18 expression. These possibilities require further investigation. Taken together, our results provide genetic evidence that Cited2 controls the expression of INK4a/ARF and fibroblast proliferation at least in part via the polycomb-group genes Bmi1 and Mel18 and provide a mechanism by which Cited2 may function as an oncogene.

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