iScience

Article

HIB/SPOP inhibits Ci/Gli-mediated tumorigenesis by modulating the RNA Polymerase II components stabilities

Gao et al., iScience 26, 107334 August 18, 2023 © 2023 The Author(s). [https://doi.org/10.1016/](https://doi.org/10.1016/j.isci.2023.107334) [j.isci.2023.107334](https://doi.org/10.1016/j.isci.2023.107334)

iScience

Article

HIB/SPOP inhibits Ci/Gli-mediated tumorigenesis by modulating the RNA Polymerase II components stabilities

Yuxue Gao,[1,](#page-1-0)[3](#page-1-1) Zhaoliang Shan,[1](#page-1-0)[,3](#page-1-1) Chunhua Jian,[1](#page-1-0) Ying Wang,[1](#page-1-0) Xia Yao,[1](#page-1-0) Shengnan Li,[1](#page-1-0) Xiuxiu Ti[,1](#page-1-0) Guochun Zhao,[1](#page-1-0) Chen Liu,^{[2](#page-1-2),[*](#page-1-3)} and Qing Zhang^{[1](#page-1-0)[,4](#page-1-4),*}

SUMMARY

Hedgehog (Hh) signaling mediated by transcription factor Ci/Gli plays a vital role in embryonic development and adult tissue homeostasis in invertebrates and vertebrates, whose dysregulation leads to many human disorders, including cancer. However, till now, cofactors of Ci/Gli which can affect tumorigenesis are not well known. Here, through genetic screen, we find overexpression of active Ci alone is not sufficient to generate tumor-like eye phenotype in Drosophila, however, its overexpression combined with knockdown of hib causes a striking tumor-like big eye phenotype. Mechanistically, HIB/SPOP inhibits Ci/Gli-mediated tumorigenesis by modulating the RNA polymerase II (RNAPII) components Rpb3/ Rpb7 stabilities in E3 ligase dependent manner. In addition, Ci/Gli can promote HIB/SPOP-mediated Rpb7/Rpb3 degradation. Taken together, our results indicate Ci/Gli needs to hook up with suitable RNAPII together to achieve the tumor-like eye phenotype and HIB/SPOP plays dual roles through controlling Ci/ Gli and Rpb3/Rpb7 protein stabilities to temper Ci/Gli/RNAPII-mediated tumorigenesis.

INTRODUCTION

First identified in Drosophila, later shown to be largely conserved in mammals, the Hedgehog (Hh) family of secreted proteins plays a vital role in embryonic development and adult tissue homeostasis from inverte-brates to vertebrates.^{[1–4](#page-11-0)} Dysregulation of Hh signaling leads to many human disorders including birth defects such as holoprosencephaly, polydactyly, skeletal malformations, and kinds of human cancer, for example, basal cell carcinoma (BCC) and medulloblastoma (MB).^{[4–11](#page-11-1)}

In Drosophila, Hh pathway mainly consist of Hh ligand, twelve-span transmembrane protein Patched (Ptc), GPCR family of seven-span transmembrane protein Smoothened (Smo), kinesin family protein Costal-2 (Cos2), serine-threonine kinase Fused (Fu), the Suppressor of fused (Sufu), and zinc finger transcription fac-tor Cubitus interruptus (Ci).^{[12–20](#page-12-0)} In the absence of Hh, Ptc switches off the signaling by inhibiting Smo plasma membrane accumulation and activity.^{[21–30](#page-12-1)} In this situation, Cos2 forms a complex with Fu, Ci, and Sufu, allowing its recruited protein kinase A (PKA), glycogen synthase kinase 3 (GSK3), and casein kinase 1 (CK1) to sequentially phosphorylate Ci. This hyperphosphorylated Ci further recruits F box protein Slimb of SCF (Skp1-Cullin1-F-box) ubiquitin ligase complex to proteolytically process the full-length Ci into a truncated form (CiR) that acts as a transcriptional repressor of Hh target gene decapentaplegic (dpp). $31-37$ In the presence of Hh, it dissociates Cos2/Ci complex, leading to blocking Ci phosphorylation, processing, and at the end turning Ci to be an active form (CiA) to promote Hh target gene dpp, patched(ptc) and engrailed (en) expression.^{[8](#page-11-2)[,11,](#page-11-3)[22](#page-12-3)}

As mentioned previously, Hh signaling-triggered human cancers are mediated by transcription factor Gli, including Gli1, 2, 3. Among them, Gli3 mainly functions as a repressor, while Gli1 and Gli2 as promoters.[4](#page-11-1),[11](#page-11-3)[,38](#page-12-4) However, in some cases, only Gli is not sufficient to initiate tumorigenesis, the underlying mechanism and what cofactors of Ci/Gli affect tumorigenesis are not well known.

hib encodes Hh-induced BTB protein, which is also a Hh target gene in Drosophila wing discs. HIB together with Cul3 functions as an E3 ubiquitin ligase to modulate Ci degradation, forming a negative feedback loop

¹State Key Laboratory of Pharmaceutical Biotechnology and MOE Key Laboratory of Model Animals for Disease Study, Jiangsu Key Laboratory of Molecular Medicine, Model Animal Research Center, School of Medicine, Nanjing University, Nanjing 210061, China

ll OPEN ACCESS

2Department of Medical Genetics, Nanjing Medical University, Nanjing 211166, China

3These authors contributed equally

4Lead contact

*Correspondence: liuchen@njmu.edu.cn (C.L.), zhangqing@nju.edu.cn (Q.Z.) [https://doi.org/10.1016/j.isci.](https://doi.org/10.1016/j.isci.2023.107334) [2023.107334](https://doi.org/10.1016/j.isci.2023.107334)

Figure 1. Co-overexpression of Ci^{-3P} and hib RNAi causes tumor-like eye phenotype in Drosophila $(A-C)$ Co-overexpression of hib RNAi and Ci^{-3P}, which is an active form of Ci with three PKA kinase sites mutated, led to tumor-like Drosophila eyes.

(D and E) Knockdown of ci or co-expression SPOP could totally rescue the tumor-like eye phenotype.

(F) Overexpressing Ci^{-3P} in hib clones also caused tumor-like eye phenotype. Scale bars: 2 mm.

to fine-tune Hh pathway activity.^{[39](#page-12-5)[,40](#page-12-6)} Speckle-type POZ protein (SPOP), which is a mammalian homologue of HIB, also modulates Gli2, Gli3 but not Gli1 protein levels.[41](#page-12-7),[42](#page-12-8) In humans, studies have shown that loss of function of SPOP mutation has closely related to many kinds of cancer, including glioma, prostate, breast cancer, etc.^{[43–47](#page-12-9)} Why it can affect so wide spectrum tumorigenesis, the underlying mechanism is also not well known.

Here, we find overexpression of active Ci alone is not sufficient to generate tumor-like eye phenotype in Drosophila, however, its overexpression combined with hib RNAi causes a striking tumor-like big eye phenotype manifested by enlarged and protruded outward eye. Interestingly, in our Ci overexpression case, the occurrence of tumor-like eye is not due to hib RNAi-mediated upregulation of Ci level. Instead, it is related to the relative high RNA polymerase II (RNAPII) components Rpb3/Rpb7 levels. Our study demonstrates that except inhibiting Ci, HIB also synergistically inhibits RNAPII complex to prevent the occurrence of tumorigenesis. Further, we demonstrate SPOP is functionally conserved which can substitute HIB to inhibit Ci/Gli plus hib RNAi-mediated tumor-like eye phenotype.

RESULTS

Genetic screen demonstrates that HIB inhibits ci-mediated tumor-like eye phenotype in Drosophila

Hh signaling-related tumorigenesis is mediated by transcription factor Ci/Gli, however, in some cases, only Ci/Gli is not sufficient to initiate tumorigenesis, the underlying mechanism and what cofactors of Ci/Gli affect tumorigenesis are not clear. To address this question, in Drosophila, overexpressing Ci^{-3P}, which is an active form of Ci with three PKA kinase sites mutated, 48 48 48 we found that Ci alone was insufficient to cause eye tumorigenesis ([Figure 1A](#page-2-0)). Next, we overexpressed Ci^{-3P} and simultaneously knocked down around 7000 conserved genes between fly and human to screen Ci/Gli cofactors by checking the eye phenotype. In this first round of genetic screen, we got two hits, hib, and warts. Since knockdown of warts alone showed the similar phenotype with co-expression of Ci and warts RNAi (data not shown), we later gave up this hit. For hib, we found its knockdown alone showed normal eye phenotype, however, its knockdown together with overexpression of Ci^{-3P} caused very striking tumor-like eye phenotype [\(Figures 1](#page-2-0)A-1C). In this condition, either knockdown of ci or co-expression of mammalian homologue of HIB, SPOP, could completely inhibit the tumor-like eye phenotype ([Figures 1D](#page-2-0) and 1E), indicating expression of Ci and inhibition of HIB are both necessary for making the tumor-like eye phenotype. Similarly, in Drosophila eye, we found that overexpressing Ci^{-3P} in hib^{-/-} mutant clones generated with FLP/FRT-mediated mitotic recombina-tion also resulted in tumor-like phenotype inside hib^{-/-} clones ([Figure 1](#page-2-0)F). Taken together, these results

Figure 2. Inhibition of Ci-mediated tumor-like eye phenotype is dependent on HIB/SPOP E3 ligase activity (A–D) Lacking of MATH (C) or BTB (D) domain could not rescue the tumor-like eye phenotype caused by cooverexpression of Ci^{-3P} and hib RNAi with GMR Gal4.

(E) Overexpression of SPOPD3box, which lacks of 299aa to 330aa could not rescue the tumor-like eye phenotype. Scale bars: 2 mm.

support the notion that Ci synergizing with hib RNAi causes the eye tumorigenesis. Of note, overexpression of HIB could not efficiently rescue the big eye phenotype owing it was eliminated by its powerful RNAi. To overcome this drawback, we overexpressed SPOP to substitute HIB to do the rescue experiment.

Inhibition of Ci-mediated tumorigenesis is dependent on HIB/SPOP E3 ligase activity

HIB/SPOP usually binds with Cul3 through its BTB domain (residues 190 aa–297 aa) to form an E3 ligase complex and recognizes substrates via its MATH domain (residues 28 aa-166 aa).^{[49,](#page-13-1)[50](#page-13-2)} To test whether HIB/SPOP E3 ligase activity is needed to inhibit the tumor-like eye phenotype, in co-overexpression of Ci^{-3P} and hib RNAi background, we first check the rescue effects of two SPOP E3 ligase-dead truncated forms, SPOP \triangle MATH and SPOP \triangle BTB, respectively. As shown in [Figure 2](#page-3-0), we found SPOP but not SPOP \triangle MATH and SPOPABTB could rescue the tumor-like eye phenotype, SPOPAMATH and SPOPABTB even manifested the dominant negative effect shown by boosting the tumor-like eye phenotype ([Figures 2C](#page-3-0) and 2D), indicating that both MATH and BTB domain are needed to inhibit the eye tumorigenesis. Next, we further test whether another SPOP E3 ligase mutant, SPOP $\Delta 3$ box, can inhibit the tumor-like eye phenotype. SPOPA3box lacks of 299 aa to 330 aa, which decreases the association with Cul3 and weakens func-tional SPOP-Cul3 E3 ligase.^{[51](#page-13-3)} We found that unlike SPOP, overexpression of SPOPA3box no longer rescued the tumor-like eye phenotype [\(Figure 2](#page-3-0)E). Taken together, these results suggest that HIB/SPOP inhibits the eye tumorigenesis dependent on its E3 ligase activity.

The accumulation of ci alone is not sufficient to cause eye tumorigenesis

Given HIB/SPOP modulates the tumor-like eye phenotype dependent on its E3 ligase activity, we next test whether the previous tumor-like eye phenotype caused by co-expression of Ci and hib RNAi is only due to hib RNAi-mediated upregulation of Ci level. Through modulating different copies of both Ci^{-3P} and GMR Gal4 to increase the amount of Ci^{-3P} , we found that expression of one and two copies of Ci^{-3P} with one ([Figures 3A](#page-4-0) and 3C) or two copies of GMR Gal4 [\(Figures 3B](#page-4-0) and 3D), none of them could cause tumorlike eye phenotype. Taken together, these results suggest that Ci^{-3P} alone is not sufficient to generate the big eye phenotype, which needs hib RNAi together to produce the tumor-like eye phenotype.

Ci together with suitable Rpb3/Rpb7 of RNAPII complex mediates the tumorigenesis

To unveil hib RNAi-affected factors which together with Ci mediate tumorigenesis, we did the secondround genetic screen in previous co-overexpression of Ci^{-3P} and hib RNAi tumor-like eye background. By crossing with the RNAi lines of around 7000 conserved genes between fly and human, we got several hits, including rpb3, rpb7, and su(var)3-9. Because HIB did not bind SuVar3-9 and affect its stability

Figure 3. Ci^{-3P} alone is not sufficient to generate the tumor-like eye phenotype

 $(A$ and B) One copy of Ci^{-3P} driven by one copy of GMR (A), or by two copies of GMR Gal4 (B) was not sufficient to generate the tumor-like eye phenotype.

(C and D) Two copies of Ci^{-3P} driven by one copy of GMR (C), or by two copies of GMR Gal4 (D) were also not sufficient to generate the tumor-like eye phenotype. Scale bars: 2 mm.

(data not shown), we ruled out the su(var)3–9. For rpb3 and rpb7, we found knockdown of rpb3 or rpb7, two subunits of the RNAPII complex, could quench the tumor-like eye phenotype ([Figures 4A](#page-5-0)-4E"), suggesting that RNAPII machinery may be involved in mediating the tumor-like eye phenotype. Consistently, in S2 cells, we found HIB downregulated both Rpb3 and Rpb7 protein levels through proteasome ([Figures 4](#page-5-0)F–4G'). Further IP experiments showed that HIB bound Rpb3 but not Rpb7, while Rpb3 bound Rpb7, suggesting that Rpb3 may recruit HIB to mediate itself and promote Rpb7 degradation ([Figures 4H](#page-5-0) and 4I). Supporting this idea, we found co-transfection of Rpb3 indeed increased Rpb7 degradation ([Figure 4I](#page-5-0)). Not surprisingly, SPOP also mediated Rpb3/Rpb7 degradation in its E3 ligase depen-dent manner ([Figures 4J](#page-5-0) and 4J'). Given Ci binds HIB and also forms a transcription complex with RNAPII to mediate its target gene expression, we speculate that it may modulate HIB/Rpb3/Rpb7 complex, affecting HIB-mediated Rpb3/Rpb7 degradation. To test that, co-transfecting Rpb3/Rpb7 and HIB with or without Ci, we found that Ci bound Rpb3, Rpb7 and promoted their degradation by HIB [\(Figures 4K](#page-5-0)–4L'). Consistently, in S2 cells or in vivo, in overexpression of Ci background, knockdown of hib inhibited Rpb3/ Rpb7 degradation ([Figures 4](#page-5-0)M–4N′). Taken together, HIB may inhibit Ci-mediated tumorigenesis by downregulating Rpb3/Rpb7, suggesting both Ci and RNAPII complex together enable the occurrence of tumorlike eye phenotype.

The tumor-like eye phenotype is related to excess interommatidial cells at pupal stage

The tumor-like eye phenotype may be generated from over proliferation at the larval stage or dampened apoptosis during the pupal stage. To determine which stage is important for formation of tumor-like eye, we use Gal80 to block the GMR-driven co-overexpression of Ci^{-3P} and hib RNAi at different stages, the results showed that the tumor-like eye phenotype was generated at the pupal stage ([Figures 5](#page-6-0)A and 5B). Our subsequent study revealed that compared with wild type ([Figure 5](#page-6-0)C), Ci^{-3P} ([Figure 5D](#page-6-0)) and hib RNAi ([Fig](#page-6-0)[ure 5](#page-6-0)E) alone, co-overexpression of Ci^{-3P} and hib RNAi ([Figure 5F](#page-6-0)) presented excessive interommatidial cells, which are usually produced at the larval stage and then eliminated through apoptosis at the pupal stage for giving rise to the adult lattice. Collectively, the earlier results suggest that impaired apoptosis may be related to the tumor-like eye phenotype. Since Grim/Hid/Rpr-mediated apoptosis is involved in elimination of excessive interommatidial cells, next we check whether their mediated apoptosis was blocked upon co-overexpression of Ci^{-3P} and hib RNAi. As shown in [Figures 5G](#page-6-0)-5I^{'''}, co-overexpression of Ci^{-3P} and hib RNAi blocked Grim/Rpr but not Hid-mediated apoptosis, implying that the tumor-like eye may be related to disability of apoptosis mediated by Grim/Rpr but not Hid in our case. Of note, overexpression of Ci alone already blocked Grim-mediated apoptosis, but that did not produce any eye phenotype, indicating that Grim is also not responsible for the tumor-like eye phenotype mediated by the cooperation of both Ci and hib RNAi. For Rpr, we surprisingly found that overexpression of Ci or hib RNAi alone could not, but they together completely blocked Rpr-mediated apoptosis, suggesting Ci and hib RNAi synergistically achieve the tumor-like eye phenotype possibly partially through inhibiting Rpr-mediated apoptosis ([Figures 5G](#page-6-0)–51^m). Consistently, TUNEL assay showed compared with Ci^{-3P} and hib RNAi alone, co-expression of them inhibited apoptosis ([Figures 5J](#page-6-0)-5J"').

SPOP and Gli2 can substitute HIB and ci to mediate the tumor-like eye phenotype in Drosophila

As mammalian homologue of HIB, SPOP is involved in various human cancers. In our previous study, we showed that SPOP function is evolutionally conserved in mediating Ci/Gli degradation. Consistently, using

Figure 4. Ci and RNAPII complex together contribute to the tumor-like eye phenotype

(A–E") Knockdown of rpb3 and rpb7 could rescue the tumor-like eye phenotype. Scale bars: 2 mm.

(F and F') HIB degraded Rpb3/Rpb7 through proteasome.

(G and G') HIB could mediate Rpb3/Rpb7 ubiquitination.

(H and H') HIB bound Rpb3 but not Rpb7.

(I) Rpb7 bound Rpb3.

(J and J') The degradation of Rpb3/Rpb7 by SPOP was dependent on its E3 ligase activity.

(K-L') Ci^{-3P} could bind Rpb3/Rpb7 and enhanced their degradation by HIB.

(M–M') In overexpression of Ci^{-3P} background, knockdown of *hib* dramatically upregulated Rpb3/7 levels in S2 cells.

(N–N') In overexpression of Ci^{-3P} background, knockdown of hib dramatically upregulated Rpb3/Rpb7 levels in vivo.

SPOP to replace HIB, it totally rescued the eye phenotype, confirming SPOP function is also conserved in mediating the tumor-like eye phenotype. In addition, we generated transgenic flies of several human can-cer-related mutants of SPOP, [46,](#page-12-10)[52](#page-13-4) including SPOPY87N, SPOPF102C, SPOPS119N, SPOPF125L, SPOPW131G, which all prevented SPOP E3 ligase activity as shown by loss of the abilities to degrade both Ci ([Figures 6](#page-7-0)G–6K‴) and Rpb3/Rpb7 [\(Figures 7](#page-9-0)A and 7A'). They no longer inhibited, instead even enhanced the tumor-like phenotype possibly due to their dominant negative effects ([Figures 6](#page-7-0)G–6K), further indicating SPOP E3 ligase activity is necessary for the rescue of tumor-like eye phenotype. Similarly, Gli2 could

Larva 29°C, Pupa 25°C **GMR UAS-Ci UAS-hib RNAi** Gal80 Pupa 29°C

B

A

G G' G'' G'''

GMR

GMR

GMR

UAS-Rpr

UAS-Hid

S-Ci

180

S-hib RNAi

GMR GMR

UAS-Grim

UAS-Grim UAS-Ci-3P

UAS-Hid

UAS-Ci-3P

GMR

UAS-Rpr

UAS-Ci-^{3P}

UAS-hib RNAi

C

D

GMR

UAS-Ci-3P

w

H \sim M \sim H' \sim H' \sim H' \sim H''. **GMR**

 $\mathbf{I}^{\prime\prime}$ is a set of $\mathbf{I}^{\prime\prime}$ in the set of $\mathbf{I}^{\prime\prime}$

 J' J''

GMR **UAS-Grim UAS-hib RNAi**

GMR

UAS-Rpr

UAS-Ci^{3P}

UAS-hib RNAi

E

F-actin

F-actin

GMR

GMR UAS-Rpr

UAS-Ci^{3P}

UAS-hib RNAi

UAS-hib RNAi

F-actin

F-actin

F

GMR

UAS-Ci^{3P}

UAS-hib RNAi

GMR

UAS-hib RNA

UAS-Grim UAS-Ci^{3P} **UAS-hib RNAi GMR GMR UAS-Hid UAS-Hid** UAS-Ci-3P **UAS-hib RNAi UAS-hib RNAi**

GMR GMR GMR UAS Ci^{3P}

Figure 5. The tumor-like eye phenotype is related to excess interommatidial cells at pupal stage

(A and B) Gal80 blocks Gal4 function at 25°C but not 29°C. Keeping 29°C at the larva stage then switching temperature to 25° C in the pupal stage (A), or keeping 25 °C at the larva stage then switching temperature to 29°C in the pupal stage (B), only in case of (B), overexpression of Ci^{-3P} plus knockdown of hib with GMR Gal4 caused the tumor-like eye phenotype. Scale bars: 2 mm.

(C–F) Compared with yw (C), Ci^{-3P} (D) or hib RNAi (E) alone, overexpression of Ci^{-3P}and hib RNAi together showed excess interommatidial cells in pupal retinae stained with TRITC-labeled phalloidin that preferentially labels filamentous actin (F-actin) (F). Scale bars: $10 \mu m$.

(G-I"') Ci^{-3P} synergizing with hib RNAi resisted the Grim/Rpr-mediated but not Hid-mediated apoptosis. Scale bars: 2 mm.

(J–J%) TUNEL staining (Green) was used to assess the status of apoptosis in a whole-mounted 25-h-APF retina. Compared with GMR (A), hib RNAi (B) or Ci^{-3P} (C) alone, co-overexpression of Ci^{-3P} and hib RNAi together (D) showed fewer apoptosis signals. Scale bars: 100 µm.

Figure 6. SPOP and Gli2 can substitute HIB and Ci to mediate the tumor-like eye phenotype in Drosophila (A) A big eye as the control.

(B) Gli2 could substitute Ci to form tumor-like eye.

(C) Overexpressing SPOP also could completely rescue Gli2-mediated the tumor-like eye phenotype.

(D–E') Knockdown *rpb3* or *rpb7 could rescue the Gli-mediated tumor-like eye phenotype. (A-E' scale bars: 2 mm).* (F–K^m) Among SPOP mutants, overexpression of SPOP^{Y87N} (G), SPOP^{F102C} (H), SPOP^{S119N} (I), SPOP^{F125L} (J) and SPOPW131G (K), could not rescue the tumor-like eye phenotype. These SPOP mutants lacked of E3 ligase activities as shown by loss of the abilities to degrade Ci. (F'–F‴) Wing discs expressing Flag-SPOP with MS1096 were immunestained with Ci^{FL} (red) antibody. The results showed that wild type SPOP degraded Ci. (F,G,H,I,J, and K scale bars: 2 mm). (G'–K‴) Overexpression of SPOP mutants such as SPOP^{Y87N} (G–G‴), SPOP^{F102C} (H–H‴), SPOP^{S119N} (I'–I‴), SPOP^{F125L} (J'– J‴) and SPOP^{W131G} (K'–K‴) could not degrade Ci. Scale bars: 20 µm.

substitute Ci together with knockdown of hib to cause the tumor-like eye phenotype ([Figures 6](#page-7-0)A–6C). And knockdown rpb3/rpb7 could also totally rescue the Gli2-mediated tumor-like eye phenotype ([Figures 6](#page-7-0)D–6E').

Next, in mammalian 293T cells, we found SPOP could bind both hRpb3 and hRpb7, but only obviously degraded hRpb7 in its E3 ligase dependent manner ([Figures 7B](#page-9-0) and 7C). Contrastingly, cancer-related SPOP mutants no longer modulated hRpb7 stability ([Figure 7](#page-9-0)D). Except 293T cells, in A375, DaoY tumor cells, taking SPOP^{S119N} as an example, it also inhibited hRpb7 degradation [\(Figure 7](#page-9-0)E). Like Ci, Gli2 also bound hRpb7 and promoted hRpb7 degradation by SPOP [\(Figures 7F](#page-9-0) and 7G). Consistently, in overexpression of Gli2 background, knockdown of spop dramatically upregulated hRpb7 levels in 293T cells and A375, DaoY tumor cells ([Figure 7](#page-9-0)H). Taken together, all these results suggest that in Drosophila, Gli2 and SPOP can substitute Ci and HIB to mediate the same tumor-like eye phenotype, in mammalian cells, hRpb7 stability is similarly regulated by Gli2 and SPOP, indicating that our model and the identified rescue genes are useful for future study of Hh- and SPOP-related tumorigenesis.

DISCUSSION

Hh signaling transduced by transcription factor Ci/Gli is essential for embryonic development and adult tissue homeostasis across species. Its dysregulation links with many kinds of human disorders, including various types of cancer, however, till now, cofactors of Ci/Gli which affect tumorigenesis are not well known. In this study, we find overexpression of active Ci alone is not sufficient to generate a tumor-like eye phenotype in Drosophila, however, its overexpression combined with knockdown of hib causes the striking tumor-like eye phenotype. Subsequent results indicate HIB regulates this phenotype dependent on its E3 ligase activity. Considering HIB can function as an E3 ligase of Ci, we first test whether hib RNAi mediates the phenotype only by boosting Ci accumulation. The results indicate it is not the case, since only increasing Ci is not sufficient to initiate the tumorigenesis. Taking advantage of our tumor-like eye fly model, we finally find HIB inhibits Rpb3/Rpb7 to prevent the tumor-like eye phenotype. Rpb3 and Rpb7, as two subunits of RNAPII complex, function in the same manner, suggesting HIB regulates the phenotype through modulating RNAPII machinery and the levels of RNAPII subunits are important for together with Ci to generate the tumorigenesis.

Why overexpression of Ci alone is not sufficient to mediate tumorigenesis but it plus knockdown of hib can? No matter HIB is expressed as Ci target gene (in Drosophila wing disc) or not (in the posterior region of Drosophila eye disc), in the presence of HIB, on the one hand, it can downregulate Rpb3/Rpb7 to inhibit RNAPII function independent on Ci. On the other hand, Ci promotes HIB-mediated Rpb3/Rpb7 degradation, resulting in RNAPII machinery working inefficiently. Therefore, only Ci overexpression case, overall high Ci with relatively low RNAPII machinery does not work effectively to generate tumor-like phenotype. However, co-overexpression of Ci and hib RNAi not only further increases Ci level but also lifts the burden on RNAPII machinery by stabilizing its subunits, therefore, leading to achieving the tumor-like eye phenotype more easily and effectively.

As mentioned, SPOP is involved in kinds of human cancer. The reason why it is related to a wide spectrum of tumors is not clear. Based on our study, dysfunction of SPOP will upregulate transcription machine RNAPII levels, leading to relatively easily reaching the thresholds of specific tumorigenesis. Additionally, except Ci/Gli, defective SPOP may also upregulate the levels of some other specific transcription factors which are usually degraded by SPOP. Taken these two aspects together, loss of function of SPOP may upregulate both specific transcription factors and general RNAPII machinery levels, leading to reaching the thresholds of various SPOP-regulated transcription factors-driven cancers easily.

Figure 7. SPOP function is conserved in the regulation of RNAPII component hRpb7 stability in mammalian cells

(A and A') SPOP mutants which lack of E3 ligase activities could not degrade Rpb3/Rpb7 in S2 cells.

(B and B') SPOP could bind both hRpb3 and hRpb7 but only obviously degraded hRpb7 in mammalian 293T cells.

(C) The degradation of hRpb7 by SPOP was dependent on its E3 ligase activity.

(D) In 293T cells, SPOP mutants (SPOP^{Y87N}, SPOP^{F102C}, SPOP^{S119N}, SPOP^{F125L}, and SPOP^{W131G}) did not degrade hRpb7.

iScience 26, 107334, August 18, 2023 9

Figure 7. Continued

(E) In 293T cells and DaoY, A375 tumor cells, SPOPS119N inhibited hRpb7 degradation. (F and G) Gli2 also bound hRpb7 and promoted hRpb7 degradation by SPOP. (H) In overexpression of Gli2 background, knockdown of spop upregulated hRpb7 levels in 293T cells and A375, DaoY tumor cells.

As mammalian homologue of Ci, Gli includes Gli1, Gli2, and Gli3. Among them, Gli1 itself is also an Hh target gene, which and Gli2 mainly function as activators, while Gli3 as a repressor. It is reported that in some cases, loss of function of Ptch1 (Ptch1^{-/-}) and gain of function of Smo (SmoM2) effectively induce MB, but overexpression of active Gli2 alone is not sufficient, the underlying reason is not clear.^{[53](#page-13-5)} Based on our study, any essential subunit dysfunction will impair the function of the whole RNAPII complex. We think in the context with SPOP, SPOP degrades the subunit hRpb7 of RNAPII, leading to forming low level functional RNAPII complex. Additionally, overexpressed Gli2 further promotes degradation of hRpb7, leading to forming low level functional Gli2/RNAPII complex according to limited Rpb7 level but not the overexpressed Gli2 level, subsequently causing correspondent low level Gli1/RNAPII complex. Therefore, high level overexpressed Gli2 with non-matched low level RNAPII generates low level functional Gli2/RNAPII and Gli1/RNAPII complexes, which is not easy to reach the needed threshold to effectively trigger MB. However, even Ptch1^{-/-} and SmoM2 activate Gli2, whose level is lower than overexpressed Gli2, therefore, it mediates relatively weak inhibition on RNAPII. Consequently, this leads to a little higher RNAPII, therefore, resulting in forming correspondent a little higher functional Gli2/RNAPII and Gli1/RNA-PII complexes. But such little increase may be still not enough to effectively trigger MB. Except this, Ptch1 $^{-/-}$ and SmoM2 also inhibit Gli3's repressor function, this may decrease MB threshold or further tip the balance to trigger MB. Taken together, these results imply that Gli1/2/3, all streams may function together to effectively fulfill induction of MB. In contrast to previous case, in the context without SPOP or SPOP dysfunction, RNAPII maintains relative high level, leading to forming relative high level Gli2/RNA-PII complex, which consequently causes correspondent high level Gli1/RNAPII. Therefore, they together may already reach the threshold of tumorigenesis. In this case, we predict Ptch1^{-/-}, SmoM2 and overexpression of active Gli2 alone all may effectively cause tumorigenesis, including MB.

As for why overexpression of Ci together with knockdown of hib antagonizes the apoptosis mediated by Rpr, but either overexpression of Ci or knockdown of hib alone not, the underlying mechanism is elusive. The specificity is possibly due to that overexpression of Ci plus knockdown of hib forms high level Ci/RNAPII complex which surpasses the specific transcription threshold needed for antagonizing Rprmediated apoptosis or high Ci/RNAPII may selectively express Ci target genes needed to fulfill the distinct roles compared with their alone. Of note, simultaneous overexpression of Ci and DIAP1 or P35 which blocks apoptosis did not cause tumor-like eye phenotype ([Figure S1\)](#page-11-4), suggesting Ci and hib RNAi together-mediated tumor-like eye phenotype may be only partially through inhibiting apoptosis, except that, other unidentified process must be involved in driving the big eye phenotype.

Finally, SPOP and Gli2 can substitute HIB and Ci to mediate the same tumor-like eye phenotype in Drosophila, in mammalian cells, hRpb7 stability is similarly regulated by Gli2 and SPOP, indicating that they are functionally conserved. Our study demonstrates SPOP is a very important cofactor of Ci/Gli, which plays a dual role in regulating both Ci/Gli and RNAPII machinery for suitable Hh signaling outcome, its loss of function will unleash Ci/Gli and RNAP II machinery to cause abnormal high Hh signaling or parallel signaling, leading to various Hh-related tumorigeneses. Accordingly, this mechanism is also applied to SPOP-regulated other transcription factors-related kinds of cancer.

Limitations of study

As mentioned previously, HIB/SPOP inhibits Ci/Gli-mediated tumorigenesis by modulating the RNAPII components Rpb3/Rpb7 stabilities and Ci/Gli can promote HIB/SPOP-mediated Rpb7/Rpb3 degradation. However, the detail underlying mechanism needs to be further addressed, for examples, except apoptosis inhibition, what is other unidentified process involved in driving the big eye phenotype? Except Rpb3/Rpb7, whether other RNAPII complex components are regulated by HIB/SPOP needs to be checked. In addition, we indicate SPOP is a very important cofactor of Ci/Gli, which plays a dual role in regulating both Ci/Gli and RNAPII machinery, how HIB/SPOP-RNAPII axis affects Ci/Gli-mediated tumorigeneses needs to be further investigated.

STAR★METHODS

Detailed methods are provided in the online version of this paper and include the following:

Cell_{ress} OPEN ACCESS

- **C** [KEY RESOURCES TABLE](#page-14-0)
- **[RESOURCE AVAILABILITY](#page-14-1)**
	- O Lead contact
	- \circ Materials availability
	- O Data and code availability
- **[EXPERIMENTAL MODEL AND STUDY PARTICIPANT DETAILS](#page-14-2)**
	- B Cell lines
	- \cap Animal model
- \bullet [METHOD DETAILS](#page-15-0)
	- O Genetic screen
	- O Plasmid constructs
	- O Immunostaining
	- O TUNEL assay
	- O Cell-based ubiquitination assay
	- O Generating clones of mutant cells
	- O Cell transfection, Western blot assays
- **.** [QUANTIFICATION AND STATISTICAL ANALYSIS](#page-16-0)

SUPPLEMENTAL INFORMATION

Supplemental information can be found online at <https://doi.org/10.1016/j.isci.2023.107334>.

ACKNOWLEDGMENTS

We thank Fly Stocks of National Institute of Genetics of Japan (NIG-Fly), Vienna Drosophila RNAi Center (VDRC), the Bloomington Stock Center and Developmental Studies Hybridoma Bank at the University of Iowa for providing fly stocks and reagents. This work was supported by grants from the National Natural Science Foundation of China (32070801, 31771615, 31271531, 31071264 and 30971679) and the Fundamental Research Funds for the Central Universities (021414380533 and 090314380019).

AUTHOR CONTRIBUTIONS

Conceptualization, Q.Z. and C.L.; Methodology, Y.G., Z.S., C.J., Y.W., X.Y., S.L., X.T. G.Z. and C.L.; Writing - Review & Editing, Y.G. and Q.Z.

DECLARATION OF INTERESTS

The authors declare no competing interests.

Received: March 8, 2023 Revised: June 9, 2023 Accepted: July 5, 2023 Published: July 10, 2023

REFERENCES

- 1. [Ingham, P.W., and McMahon, A.P. \(2001\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref1) [Hedgehog signaling in animal development:](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref1) [paradigms and principles. Genes Dev.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref1) 15, [3059–3087.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref1)
- 2. Lum, L., and Beachy, P.A. (2004). The Hedgehog response network: sensors, switches, and routers. Science 304, 1755–1759. <https://doi.org/10.1126/science.1098020>.
- 3. Jia, J., and Jiang, J. (2006). Decoding the Hedgehog signal in animal development. Cell. Mol. Life Sci. 63, 1249–1265. [https://doi.](https://doi.org/10.1007/s00018-005-5519-z) [org/10.1007/s00018-005-5519-z](https://doi.org/10.1007/s00018-005-5519-z).
- 4. [Jiang, J., and Hui, C.C. \(2008\). Hedgehog](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref4) [Signaling in Development and Cancer. Dev.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref4) Cell 15[, 801–812.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref4)
- 5. [Villavicencio, E.H., Walterhouse, D.O.,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref5) [Iannaccone, P.M., and Philip, M. \(2000\). The](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref5) [Sonic Hedgehog–Patched–Gli Pathway in](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref5) [Human Development and Disease. Am. J.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref5) [Hum. Genet.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref5) 67, 1047–1054.
- 6. Taipale, J., and Beachy, P.A. (2001). The Hedgehog and Wnt signalling pathways in cancer. Nature 411, 349–354. [https://doi.org/](https://doi.org/10.1038/35077219) [10.1038/35077219.](https://doi.org/10.1038/35077219)
- 7. [Nieuwenhuis, E., and Hui, C.C. \(2005\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref7) [Hedgehog signaling and congenital](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref7) [malformations. Clin. Genet.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref7) 67, 193–208.
- 8. [Bonifas, J.M., Pennypacker, S., Chuang, P.T.,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref8) [Mcmahon, A.P., Williams, M., Rosenthal, A.,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref8) [De Sauvage, F.J., and Epstein, E.H., Jr. \(2001\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref8) [Activation of Expression of Hedgehog Target](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref8)

[Genes in Basal Cell Carcinomas. J. Invest.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref8) Dermatol. 116[, 739–742.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref8)

- 9. [Yauch, R.L., Dijkgraaf, G.J.P., Alicke, B.,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref9) [Januario, T., Ahn, C.P., Holcomb, T., Pujara,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref9) [K., Stinson, J., Callahan, C.A., Tang, T., et al.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref9) [\(2009\). Smoothened Mutation Confers](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref9) [Resistance to a Hedgehog Pathway Inhibitor](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref9) [in Medulloblastoma. Science](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref9) 326, 572–574.
- 10. Pasca di Magliano, M., and Hebrok, M. (2003). Hedgehog signalling in cancer formation and maintenance. Nat. Rev. Cancer 3, 903–911. <https://doi.org/10.1038/nrc1229>.
- 11. Ingham, P.W., Nakano, Y., and Seger, C. (2011). Mechanisms and functions of Hedgehog signalling across the metazoa.

Nat. Rev. Genet. 12, 393–406. [https://doi.org/](https://doi.org/10.1038/nrg2984) [10.1038/nrg2984](https://doi.org/10.1038/nrg2984).

- 12. [Hooper, J.E., and Scott, M.P. \(1989\). The](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref12) [Drosophila patched gene encodes a putative](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref12) [membrane protein required for segmental](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref12) [patterning. Cell](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref12) 59, 751–765.
- 13. [Yue, S., Tang, L.Y., Tang, Y., Tang, Y., Shen,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref13) [Q.H., Ding, J., Chen, Y., Zhang, Z., Yu, T.T.,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref13) [Zhang, Y.E., and Cheng, S.Y. \(2014\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref13) [Requirement of Smurf-mediated endocytosis](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref13) [of Patched1 in sonic hedgehog signal](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref13) [reception. Elife](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref13) 3, e02555.
- 14. [Alcedo, J., Ayzenzon, M., Ohlen, T.V., Noll,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref14) [M., and Hooper, J.E. \(1996\). The Drosophila](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref14) [Smoothened Gene Encodes a Seven-Pass](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref14) [Membrane Protein, a Putative Receptor for](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref14) [the Hedgehog Signal. Cell](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref14) 86, 221–232.
- 15. Riobo, N.A., Saucy, B., Dilizio, C., and Manning, D.R. (2006). Activation of heterotrimeric G proteins by Smoothened. Proc. Natl. Acad. Sci. USA 103, 12607–12612. <https://doi.org/10.1073/pnas.0600880103>.
- 16. [Ogden, S.K., Fei, D.L., Schilling, N.S., Ahmed,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref16) [Y.F., Hwa, J., and Robbins, D.J. \(2008\). G](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref16) [protein Galphai functions immediately](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref16) [downstream of Smoothened in Hedgehog](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref16) [signalling. Nature](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref16) 456, 967–970.
- 17. [Sisson, J.C., Ho, K.S., Suyama, K., and Scott,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref17) [M.P. \(1997\). Costal2, a novel kinesin-related](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref17) [protein in the Hedgehog signaling pathway.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref17) .
Cell 90[, 235–245.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref17)
- 18. [Jia, J., Tong, C., and Jiang, J. \(2003\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref18) [Smoothened transduces Hedgehog signal by](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref18) [physically interacting with Costal2/Fused](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref18) [complex through its C-terminal tail. Genes](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref18) Dev. 17[, 2709–2720.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref18)
- 19. Motzny, C.K., and Holmgren, R. (1995). The Drosophila cubitus interruptus protein and its role in the wingless and hedgehog signal transduction pathways. Mech. Dev. 52, 137–150. [https://doi.org/10.1016/0925-](https://doi.org/10.1016/0925-4773(95)00397-j) [4773\(95\)00397-j](https://doi.org/10.1016/0925-4773(95)00397-j).
- 20. Chen, C.H., von Kessler, D.P., Park, W., Wang, B., Ma, Y., and Beachy, P.A. (1999). Nuclear trafficking of Cubitus interruptus in the transcriptional regulation of Hedgehog target gene expression. Cell 98, 305–316. [https://doi.org/10.1016/s0092-8674\(00\)](https://doi.org/10.1016/s0092-8674(00)81960-1) [81960-1.](https://doi.org/10.1016/s0092-8674(00)81960-1)
- 21. [Berman, D.M., Karhadkar, S.S., Hallahan,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref21) [A.R., Pritchard, J.I., Eberhart, C.G., Watkins,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref21) [D.N., Chen, J.K., Cooper, M.K., Taipale, J.,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref21) [Olson, J.M., and Beachy, P.A. \(2002\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref21) [Medulloblastoma growth inhibition by](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref21) [hedgehog pathway blockade. Science](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref21) 297, [1559–1561.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref21)
- 22. Qi, X., Schmiege, P., Coutavas, E., and Li, X. (2018). Two Patched molecules engage distinct sites on Hedgehog yielding a signaling-competent complex. Science 362, eaas8843. [https://doi.org/10.1126/science.](https://doi.org/10.1126/science.aas8843) [aas8843.](https://doi.org/10.1126/science.aas8843)
- 23. Qi, X., Schmiege, P., Coutavas, E., Wang, J., and Li, X. (2018). Structures of human Patched and its complex with native palmitoylated sonic hedgehog. Nature 560, 128–132. <https://doi.org/10.1038/s41586-018-0308-7>.
- 24. Gong, X., Qian, H., Cao, P., Zhao, X., Zhou, Q., Lei, J., and Yan, N. (2018). Structural basis for the recognition of Sonic Hedgehog by human Patched1. Science 361, eaas8935. <https://doi.org/10.1126/science.aas8935>.
- 25. Zhang, Y., Bulkley, D.P., Xin, Y., Roberts, K.J., Asarnow, D.E., Sharma, A., Myers, B.R., Cho, W., Cheng, Y., and Beachy, P.A. (2018). Structural Basis for Cholesterol Transport-like Activity of the Hedgehog Receptor Patched. Cell 175, 1352–1364.e14. [https://doi.org/10.](https://doi.org/10.1016/j.cell.2018.10.026) [1016/j.cell.2018.10.026.](https://doi.org/10.1016/j.cell.2018.10.026)
- 26. Qian, H., Cao, P., Hu, M., Gao, S., Yan, N., and Gong, X. (2019). Inhibition of tetrameric Patched1 by Sonic Hedgehog through an asymmetric paradigm. Nat. Commun. 10, 2320. [https://doi.org/10.1038/s41467-019-](https://doi.org/10.1038/s41467-019-10234-9) [10234-9](https://doi.org/10.1038/s41467-019-10234-9).
- 27. Myers, B.R., Neahring, L., Zhang, Y., Roberts, K.J., and Beachy, P.A. (2017). Rapid, direct activity assays for Smoothened reveal Hedgehog pathway regulation by membrane cholesterol and extracellular sodium. Proc. Natl. Acad. Sci. USA 114, E11141–E11150. [https://doi.org/10.1073/pnas.1717891115.](https://doi.org/10.1073/pnas.1717891115)
- 28. Rudolf, A.F., Kinnebrew, M., Kowatsch, C., Ansell, T.B., El Omari, K., Bishop, B., Pardon, E., Schwab, R.A., Malinauskas, T., Qian, M., et al. (2019). The morphogen Sonic hedgehog inhibits its receptor Patched by a pincer grasp mechanism. Nat. Chem. Biol. 15, 975–982. [https://doi.org/10.1038/s41589-](https://doi.org/10.1038/s41589-019-0370-y) [019-0370-y](https://doi.org/10.1038/s41589-019-0370-y).
- 29. Petrov, K., Wierbowski, B.M., Liu, J., and Salic, A. (2020). Distinct Cation Gradients Power Cholesterol Transport at Different Key Points in the Hedgehog Signaling Pathway. Dev. Cell 55, 314–327.e7. [https://doi.org/10.1016/](https://doi.org/10.1016/j.devcel.2020.08.002) [j.devcel.2020.08.002](https://doi.org/10.1016/j.devcel.2020.08.002).
- 30. Qi, C., Di Minin, G., Vercellino, I., Wutz, A., and Korkhov, V.M. (2019). Structural basis of sterol recognition by human hedgehog receptor PTCH1. Sci. Adv. 5, eaaw6490. <https://doi.org/10.1126/sciadv.aaw6490>.
- 31. Jiang, J., and Struhl, G. (1995). Protein kinase A and hedgehog signaling in drosophila limb development. Cell 80, 563–572. [https://doi.](https://doi.org/10.1016/0092-8674(95)90510-3) [org/10.1016/0092-8674\(95\)90510-3](https://doi.org/10.1016/0092-8674(95)90510-3).
- 32. Jia, J., Amanai, K., Wang, G., Tang, J., Wang, B., and Jiang, J. (2002). Shaggy/GSK3 antagonizes Hedgehog signalling by regulating Cubitus interruptus. Nature 416, 548–552. <https://doi.org/10.1038/nature733>.
- 33. Jia, J., Zhang, L., Zhang, Q., Tong, C., Wang, B., Hou, F., Amanai, K., and Jiang, J. (2005). Phosphorylation by double-time/CKIepsilon and CKIalpha targets cubitus interruptus for Slimb/beta-TRCP-mediated proteolytic
processing. Dev. Cell 9, 819–830. [https://doi.](https://doi.org/10.1016/j.devcel.2005.10.006) [org/10.1016/j.devcel.2005.10.006.](https://doi.org/10.1016/j.devcel.2005.10.006)
- 34. Price, M.A., and Kalderon, D. (2002). Proteolysis of the Hedgehog signaling effector Cubitus interruptus requires phosphorylation by Glycogen Synthase Kinase 3 and Casein Kinase 1. Cell 108, 823–835. [https://doi.org/10.1016/s0092-](https://doi.org/10.1016/s0092-8674(02)00664-5) [8674\(02\)00664-5.](https://doi.org/10.1016/s0092-8674(02)00664-5)
- 35. Zhang, W., Zhao, Y., Tong, C., Wang, G., Wang, B., Jia, J., and Jiang, J. (2005). Hedgehog-regulated Costal2-kinase complexes control phosphorylation and proteolytic processing of Cubitus interruptus. Dev. Cell 8, 267–278. [https://doi.org/10.1016/](https://doi.org/10.1016/j.devcel.2005.01.001) [j.devcel.2005.01.001](https://doi.org/10.1016/j.devcel.2005.01.001).
- 36. Smelkinson, M.G., and Kalderon, D. (2006). Processing of the Drosophila hedgehog signaling effector Ci-155 to the repressor Ci-75 is mediated by direct binding to the SCF component Slimb. Curr. Biol. 16, 110–116. <https://doi.org/10.1016/j.cub.2005.12.012>.
- 37. Jiang, J., and Struhl, G. (1998). Regulation of the Hedgehog and Wingless signalling pathways by the F-box/WD40-repeat protein Slimb. Nature 391, 493–496. [https://doi.org/](https://doi.org/10.1038/35154) [10.1038/35154.](https://doi.org/10.1038/35154)
- 38. [Wilson, C.W., and Chuang, P.T. \(2010\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref38) [Mechanism and evolution of cytosolic](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref38) [Hedgehog signal transduction. Development](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref38) 137[, 2079–2094.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref38)
- 39. Kent, D., Bush, E.W., and Hooper, J.E. (2006). Roadkill attenuates Hedgehog responses through degradation of Cubitus interruptus. Development 133, 2001–2010. [https://doi.](https://doi.org/10.1242/dev.02370) [org/10.1242/dev.02370.](https://doi.org/10.1242/dev.02370)
- 40. [Zhang, Q., Zhang, L., Wang, B., Ou, C.Y.,](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref40) [Chien, C.T., and Jiang, J. \(2006\). A](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref40) [hedgehog-induced BTB protein modulates](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref40) [hedgehog signaling by degrading Ci/Gli](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref40) [transcription factor. Dev. Cell](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref40) 10, 719–729.
- 41. [Wang, C., Pan, Y., and Wang, B. \(2010\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref41) [Suppressor of fused and Spop regulate the](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref41) [stability, processing and function of Gli2 and](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref41) [Gli3 full-length activators but not their](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref41) [repressors. Development](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref41) 137, 2001–2009.
- 42. [Cai, H., and Liu, A. \(2017\). Spop regulates Gli3](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref42) [activity and Shh signaling in dorsoventral](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref42) [patterning of the mouse spinal cord. Dev.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref42) Biol. 432[, 72–85.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref42)
- 43. Liu, Y., Melin, B.S., Rajaraman, P., Wang, Z., Linet, M., Shete, S., Amos, C.I., Lau, C.C., Scheurer, M.E., Tsavachidis, S., et al. (2012). Insight in glioma susceptibility through an analysis of 6p22.3, 12p13.33-12.1, 17q22-23.2 and 18q23 SNP genotypes in familial and non-familial glioma. Hum. Genet. 131, 1507– 1517. [https://doi.org/10.1007/s00439-012-](https://doi.org/10.1007/s00439-012-1187-x) [1187-x.](https://doi.org/10.1007/s00439-012-1187-x)
- 44. Berger, M.F., Lawrence, M.S., Demichelis, F., Drier, Y., Cibulskis, K., Sivachenko, A.Y., Sboner, A., Esgueva, R., Pflueger, D., Sougnez, C., et al. (2011). The genomic complexity of primary human prostate cancer. Nature 470, 214–220. [https://doi.org/](https://doi.org/10.1038/nature09744) [10.1038/nature09744.](https://doi.org/10.1038/nature09744)
- 45. Kan, Z., Jaiswal, B.S., Stinson, J., Janakiraman, V., Bhatt, D., Stern, H.M., Yue, P., Haverty, P.M., Bourgon, R., Zheng, J., et al. (2010). Diverse somatic mutation patterns and pathway alterations in human cancers. Nature 466, 869–873. [https://doi.org/10.1038/](https://doi.org/10.1038/nature09208) [nature09208.](https://doi.org/10.1038/nature09208)
- 46. Barbieri, C.E., Baca, S.C., Lawrence, M.S., Demichelis, F., Blattner, M., Theurillat, J.P., White, T.A., Stojanov, P., Van Allen, E., Stransky, N., et al. (2012). Exome sequencing

identifies recurrent SPOP, FOXA1 and MED12 mutations in prostate cancer. Nat. Genet. 44, 685–689. [https://doi.org/10.1038/](https://doi.org/10.1038/ng.2279) [ng.2279](https://doi.org/10.1038/ng.2279).

- 47. Li, C., Ao, J., Fu, J., Lee, D.F., Xu, J., Lonard, D., and O'Malley, B.W. (2011). Tumor-suppressor role for the SPOP ubiquitin ligase in signal-dependent proteolysis of the oncogenic co-activator SRC-3/AIB1. Oncogene 30, 4350–4364. [https://doi.org/10.](https://doi.org/10.1038/onc.2011.151) [1038/onc.2011.151](https://doi.org/10.1038/onc.2011.151).
- 48. Wang, G., Wang, B., and Jiang, J. (1999). Protein kinase A antagonizes Hedgehog signaling by regulating both the activator and repressor forms of Cubitus interruptus. Genes Dev. 13, 2828–2837. [https://doi.org/](https://doi.org/10.1101/gad.13.21.2828) [10.1101/gad.13.21.2828.](https://doi.org/10.1101/gad.13.21.2828)
- 49. Xu, L., Wei, Y., Reboul, J., Vaglio, P., Shin, T.H., Vidal, M., Elledge, S.J., and Harper, J.W. (2003). BTB proteins are substrate-specific adaptors in an SCF-like modular ubiquitin ligase containing CUL-3. Nature 425, 316–321. [https://doi.org/10.1038/](https://doi.org/10.1038/nature01985) [nature01985.](https://doi.org/10.1038/nature01985)
- 50. Pintard, L., Willems, A., and Peter, M. (2004). Cullin-based ubiquitin ligases: Cul3-BTB complexes join the family. EMBO J. 23, 1681– 1687. [https://doi.org/10.1038/sj.emboj.](https://doi.org/10.1038/sj.emboj.7600186) [7600186.](https://doi.org/10.1038/sj.emboj.7600186)
- 51. Liu, C., Zhou, Z., Yao, X., Chen, P., Sun, M., Su, M., Chang, C., Yan, J., Jiang, J., and Zhang, Q. (2014). Hedgehog signaling downregulates suppressor of fused through the HIB/SPOP-Crn axis in Drosophila. Cell Res. 24, 595–609. [https://doi.org/10.1038/cr.2014.29.](https://doi.org/10.1038/cr.2014.29)
- 52. Theurillat, J.-P.P., Udeshi, N.D., Errington, W.J., Svinkina, T., Baca, S.C., Pop, M., Wild, P.J., Blattner, M., Groner, A.C., Rubin, M.A., et al. (2014). Ubiquitylome analysis identifies dysregulation of effector substrates in SPOPmutant prostate cancer. Science 346, 85–89. [https://doi.org/10.1126/science.1250255.](https://doi.org/10.1126/science.1250255)
- 53. Yin, W.-C., Satkunendran, T., Mo, R., Morrissy, S., Zhang, X., Huang, E.S., Uusküla-Reimand, L., Hou, H., Son, J.E., Liu, W., et al. (2019). Dual Regulatory Functions of SUFU and Targetome of GLI2 in SHH Subgroup Medulloblastoma. Dev. Cell 48, 167–183.e5. [https://doi.org/10.](https://doi.org/10.1016/j.devcel.2018.11.015) [1016/j.devcel.2018.11.015](https://doi.org/10.1016/j.devcel.2018.11.015).
- 54. Aza-Blanc, P., Lin, H.Y., Ruiz i Altaba, A., and Kornberg, T.B. (2000). Expression of the vertebrate Gli proteins in Drosophila reveals a distribution of activator and repressor activities. Development 127, 4293–4301. [https://doi.org/10.1242/dev.127.](https://doi.org/10.1242/dev.127.19.4293) [19.4293](https://doi.org/10.1242/dev.127.19.4293).
- 55. Rubin, G.M., and Spradling, A.C. (1982). Genetic transformation of Drosophila with transposable element vectors. Science 218, 348–353. [https://doi.org/10.1126/science.](https://doi.org/10.1126/science.6289436) [6289436.](https://doi.org/10.1126/science.6289436)
- 56. [Wang, G., Wang, B., and Jiang, J. \(1999\).](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref56) [Protein kinase A antagonizes Hedgehog](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref56) [signaling by regulating both the activator](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref56) [and repressor forms of Cubitus interruptus.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref56) Genes Dev. 13[, 2828–2837.](http://refhub.elsevier.com/S2589-0042(23)01411-6/sref56)
- 57. Zhou, Z., Yao, X., Li, S., Xiong, Y., Dong, X., Zhao, Y., Jiang, J., and Zhang, Q. (2015). Deubiquitination of Ci/Gli by Usp7/HAUSP Regulates Hedgehog Signaling. Dev. Cell 34, 58–72. [https://doi.org/10.1016/j.devcel.2015.](https://doi.org/10.1016/j.devcel.2015.05.016) [05.016.](https://doi.org/10.1016/j.devcel.2015.05.016)

STAR+METHODS

KEY RESOURCES TABLE

RESOURCE AVAILABILITY

Lead contact

Further information and requests for resources should be directed to and will be fulfilled by the lead contact, Qing Zhang [\(zhangqing@nju.edu.cn](mailto:zhangqing@nju.edu.cn)).

Materials availability

This study did not generate new unique reagents.

Data and code availability

- The data reported in this paper will be shared upon request to the lead corresponding author (zhangqing@nju.edu.cn).
- This paper does not report original code.
- Any additional information required to reanalyze the data reported in this paper is available from the [lead contact](#page-14-3) upon request.

EXPERIMENTAL MODEL AND STUDY PARTICIPANT DETAILS

Cell lines

The Drosophila embryonic cell line S2 were obtained from Yun Zhao lab (CAS Center for Excellence in Molecular Cell Science, Shanghai, China) and was cultured in the Schneider's Drosophila Medium (Invitrogen) with 10% fetal bovine serum (FBS, Gibco), 100 U/ml of penicillin and 100 µg/mL of Streptomycin (P/S, Gibco) in a humidified incubator at 28°C. The human embryonic kidney cell line 293T and human melanoma cell line A375 were obtained from Ying Cao lab (Nanjing university, Nanjing, China). The human medulloblastoma cell line DaoY was obtained from Chen Liu lab (Nanjing Medical University, Nanjing, China). They were cultured in Dulbecco's modified Eagle's medium (DMEM, HyClone), with 10% fetal bovine serum (FBS, Gibco),100 U/ml of penicillin and 100 µg/ml of Streptomycin (P/S, Gibco) in a humidified incubator with 5% $CO₂$ at 37°C.

Animal model

The RNAi lines that targeted ci (2125R-1), hib (9924R-1), rpb3 (7885R-1) and rpb7 (v100309) used in genetic screen were obtained from the National Institute of Genetics Stock Center (NIG), Japan and the Vienna Drosophila RNAi Center (VDRC), Austria. eyflp, hsflp and FRT82 flies were obtained from the Bloomington

Stock Center. Flies of MS1096, GMR-Gal4, Gal80, HA-Ci^{-3P}, Fq/Myc-SPOP, Fq-SPOPA3box, Fq-SPOPA MATH, Fg-SPOP Δ BTB, Myc-mouse-Gli2 have been described.^{[40,](#page-12-6)[54](#page-13-6)} hib⁴⁶ is a hib mutant allele, whose coding sequence is replaced by the white gene. The transgenic flies of Fg-Rpb3, Fg-Rpb7, HA-Grim, HA-Hid, HA-Rpr, Fg-SPOP^{Y87N}, Fg-SPOP^{F102C}, Fg-SPOP^{S119N}, Fg-SPOP^{F125L}, Fg-SPOP^{K129E}, Fg-SPOP^{W131G} and Fg-SPOPF133V were generated by injection of corresponding constructs into Drosophila embryos according to the methods described previously.^{[55](#page-13-7)} The parental strain for all germline transformations is w1118. All stocks used in this study were maintained and raised under standard conditions.

METHOD DETAILS

Genetic screen

The RNAi lines were obtained from the National Institute of Genetics Stock Center (NIG), Japan and the Vienna Drosophila RNAi Center (VDRC), Austria. We totally screened around 7000 RNAi lines which target to the conserved genes between Drosophila and Mammals. We did the first and the second round of genetic screening in GMR Ci^{-3P} and in GMR Ci^{-3P}/hib RNAi background, respectively.

Plasmid constructs

The constructs for S2 cell transfection experiments are as follows: HA/Myc-Rpb3, HA/Myc-Rpb7, Myc/Fg-HIB, HA-Grim, HA-Hid, HA-Rpr and Myc/Fg-SPOP, their corresponding cDNA fragments were amplified and cloned into the pUAST vectors. Similarly, the constructs of Fg-hRpb3, Fg-hRpb7, Myc-hGli2, Myc-SPOP and Myc-SPOPA3box used in mammalian cells were cloned into the pcDNA3.1 vectors. The mutants of Myc/Fg-SPOP, including Fg/Myc-SPOP^{Y87N}, Fg/Myc-SPOP^{F102C}, Fg/Myc-SPOP^{5119N}, Fg/Myc-SPOP^{F125L}, Fg/Myc-SPOP^{K129E}, Fg/Myc-SPOP^{W131G} and Fg/Myc-SPOP^{F133V}, were made through the PCR-based site-directed mutagenesis and cloned into the pUAST and pcDNA3.1 vectors, respectively.

Immunostaining

Immunostaining of imaginal discs were performed with standard protocols.^{[56,](#page-13-8)[57](#page-13-9)} To immunostain the pupal Drosophila visual system, samples should be dissected after 48h when third-instar larvae begin to enter pupation. Then the whole visual system is held in fixative and primary antibody solution with shaking in overnight at 4°C. After adding the secondary antibody, the Drosophila pupal visual system can be mounted for confocal microscopy (TCS SP5II, Leica Company, Germany). Antibodies were used in this study: rat anti-Ci (2A) (1:50; DSHB); mouse anti-Flag (M2) (1:200; Sigma); mouse anti-HA (F7) (1:200; Santa Cruz); TRITClabeled phalloidin (1:250; Sigma).

TUNEL assay

Pupal retinas were dissected in PBS and fixed for 30 min in 4% paraformaldehyde. The apoptotic cells were determined by TUNEL staining using a One Step TUNEL Apoptosis Assay Kit (Beyotime, Jiangsu, China). Tissues were incubated at 37°C for 1 h in the mixture of enzyme and fluorescent labeling solution then rinsed in PBS. Images were captured using a confocal microscopy (TCS SP5II, Leica Company, Germany).

Cell-based ubiquitination assay

For cell-based ubiquitination assays, cells were lysed with denaturing buffer (1% SDS, 50 mM Tris-base, pH 7.5, 0.5 mM EDTA, and 1 mM DTT) and incubated at 100°C for 5 min. The lysates were then diluted 10-fold with regular lysis buffer (50 mM Tris pH8.0, 0.1 M NaCl, 10 mM NaF, 1 mM Na3VO4, 0.5% NP-40, 10% Glycerol and 1 mM EDTA pH8.0) and subject to IP and IB analysis.

Generating clones of mutant cells

Clones of mutant cells were generated by FLP/FRT-mediated mitotic recombination as described.[31](#page-12-2),[40](#page-12-6) Genotypes for generating clones are as follows: hib clones in eye with expressing UAS transgenes: ey-flp; $(GMR-Gal4/UAS-Ci^{-3P})$; FRT82 hib/FRT82 Gal80.

Cell transfection, Western blot assays

Transfection of S2 cells were carried out using calcium phosphate transfection method. Usually, S2 cells are transfected in 10 cm plates with no more than 20 µg of total DNA for a ubiquitin-Gal4 construct and other co-transfected pUAST expression vectors. 36–48 h after transfection, cells are harvested for Western blot analysis with standard protocols as previously described. 293T cells were transfected using lipofectamine

2000 (Invitrogen) according to the manufacturer's instructions. 48 h after transfection, cells were harvested for immunoprecipitation and Western blot analysis with standard protocols as described.

The primary antibodies used were mouse anti-HA (F7) (1:5000; Santa Cruz); mouse anti-Myc (9E10) (1:2500; Santa Cruz); mouse anti-Flag (1:2500; Sigma) and mouse anti-Actin (A00702) (1:5000: Genscript). After incubation with HRP-coupled secondary antibodies (goat anti-mouse diluted 1:10000, Jackson ImmunoResearch), the blots were visualized using a chemiluminescent detection kit (GE healthcare).

QUANTIFICATION AND STATISTICAL ANALYSIS

The data shown in all figures were representative of three or more independent experiments.