Contrasting Expression of Canonical Wnt Signaling Reporters TOPGAL, BATGAL and Axin2LacZ during Murine Lung Development and Repair

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Abstract

Canonical Wnt signaling plays multiple roles in lung organogenesis and repair by regulating early progenitor cell fates: investigation has been enhanced by canonical Wnt reporter mice, TOPGAL, BATGAL and Axin2LacZ. Although widely used, it remains unclear whether these reporters convey the same information about canonical Wnt signaling. We therefore compared beta-galactosidase expression patterns in canonical Wnt signaling of these reporter mice in whole embryo versus isolated prenatal lungs. To determine if expression varied further during repair, we analyzed comparative pulmonary expression of beta-galactosidase after naphthalene injury. Our data show important differences between reporter mice. While TOPGAL and BATGAL lines demonstrate Wnt signaling well in early lung epithelium, BATGAL expression is markedly reduced in late embryonic and adult lungs. By contrast, Axin2LacZ expression is sustained in embryonic lung mesenchyme as well as epithelium. Three days into repair after naphthalene, BATGAL expression is induced in bronchial epithelium as well as TOPGAL expression (already strongly expressed without injury). Axin2LacZ expression is increased in bronchial epithelium of injured lungs. Interestingly, both TOPGAL and Axin2LacZ are up regulated in parabronchial smooth muscle cells during repair. Therefore the optimal choice of Wnt reporter line depends on whether up- or down-regulation of canonical Wnt signal reporting in either lung epithelium or mesenchyme is being compared.

Introduction

Canonical Wnt signaling plays multiple roles during lung organogenesis and repair by controlling survival, proliferation and differentiation of early progenitor cells in epithelium and mesenchyme [1,2]. Canonical Wnt signaling is mediated mainly by the multifunctional beta-catenin protein which is a potent co-activator of transcription factors such as Lymphoid Enhancer Factor (LEF) and T-Cell Factor (TCF) [3]. Canonical s-catenin activation requires binding of secreted lipoglycoproteins termed Wnts to the Frizzled receptor, thereby raising cytoplasmic levels of activated s-catenin, and ultimately nuclear translocation of s-catenin. Without Wnt activation, s-catenin is mainly located at epithelial junctions where it acts as a cell adhesion molecule by interacting with E-cadherin and alpha-catenin [4]. s-catenin has a rapid turnover; excess s-catenin binds to a APC/Axin/GSK3β complex that mediates its phosphorylation, ubiquitination and degradation [5]. Upon frizzled/LRP5/6 activation, the destruction complex is dismantled by release of Axin such that s-catenin is now released, stabilized and translocated into the nucleus (reviewed in [6,7,8]).

Several reporter mice have been designed to track Wnt signaling in vivo: two allow monitoring of formation of the s-catenin/TCF transcription complex. The Tcf optimal promoter (TOP)-beta-galactosidase (TOPGAL) transgenic mice were made fusing three LEF/TCF binding sites to c-fos minimal promoter [9]. These mice were originally reported to follow activation of LEF/TCF transcription complexes during hair development and differentiation. The second reporter line, the s-catenin activated transgene (BAT) driving the expression of nuclear beta-galactosidase, was designed by fusing seven TCF/LEF binding sites upstream of a 0.13 kb fragment containing the minimal promoter of the Siamois gene [10]. However, transgenic mice relying upon random insertion of an expression cassette may be unstable with increased number of generations.

In contrast to the first 2 reporter lines, a stable knock in of LacZ in frame with the endogenous start codon of the Axin2 gene has recently been generated [11]. Axin2 induces s-catenin degradation...
However, Axin2, also known as conductin, is also a target of the canonical Wnt signaling pathway and its expression can therefore be used to report the activation of this pathway [11].

Herein, we have directly compared reporter expression in the BATGAL and TOPGAL transgenics, seven and eleven years respectively after their first publication, using the stable Axin2 for additional comparison. We have tested each reporter not only during development but also during repair after naphthalene-induced airway injury. We show the optimal choice of canonical Wnt reporter line depends on whether an up- or down-regulation of canonical Wnt signaling in either lung epithelium or mesenchyme is being evaluated.

**Results**

Embryonic LacZ expression differs significantly between reporter mice

We examined LacZ activity in the whole embryo at E11.5 and E12.5 with a focus on ectoderm-derived organs controlled by Wnt signaling such as whiskers, mammary placodes and limbs. At E11.5, TOPGAL is expressed diffusely in the forebrain, nasal process and inner ear. Further specific expression is seen in the apical ectodermal ridge (AER) of the limb, and epithelium of the mammary placode (Fig. 1A). TOPGAL expression is also found in the dorsal and ventral somites as well as the tip of the tail. At E12.5 (Fig. 1B), the expression in ectodermal appendages is maintained and enhanced in the whisker placodes in the nasal region as well as in the AER and the mammary buds. LacZ expression is also detected in discrete mesenchymal condensations within the limbs. In contrast, at E11.5, BATGAL expression is found throughout the embryo with a “salt and pepper” pattern (Fig. 1C). LacZ expression is found in the mammary buds but significant staining was also found in the surrounding tissue. At E12.5, the BATGAL embryo still shows “salt and pepper” expression throughout the embryo. LacZ expression is found in the mammary buds and the AER are clearly positive for Axin2LacZ. In addition, strong LacZ expression is found in the developing ear and in the somites.

**Figure 1.** LacZ expression in whole embryos of TOPGAL, BATGAL and Axin2LacZ mice. (A) E11.5 TOPGAL embryo shows staining in the forebrain, the nasal process, the inner ear, the apical ectodermal ridge (AER) of the limb, the epithelium of the mammary placode, the somites and the tip of the tail. (B) E12.5 TOPGAL embryo shows LacZ expression in ectodermal appendages, the whisker placodes in the nasal region as well as the AER, the mammary buds in between the limbs and discrete mesenchymal condensations within the limbs. (C) E11.5 BATGAL embryo shows expression throughout the embryo with a “salt and pepper” pattern with higher signals in ectodermal domains such as the nasal process, the forebrain, the AER and the tip of the tail. (D) At E12.5, the BATGAL embryo shows LacZ expression in the mammary buds but significant staining was also found in the surrounding tissue. (E) E11.5 Axin2LacZ embryo shows homogenous LacZ expression throughout the embryo with higher expression in the AER and the developing mammary placode. (F) At E12.5, individualized whisker placodes as well as mammary buds and the AER are clearly positive for Axin2LacZ. In addition, strong LacZ expression is found in the developing ear and in the somites.

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Figure 2. LacZ expression in TOPGAL, BATGAL and Axin2LacZ whole lungs during prenatal development. TOPGAL is expressed in the epithelium of the trachea and the lung at E11.5 (A), E13.5 (B) and E16.5 (C); whereas at E18.5 (D) TOPGAL expression is also detected in the terminal bronchioles and the surrounding parenchyma. BATGAL expression is detected in a “salt and pepper” manner in the lung epithelium at E11.5 (E) and E13.5 (F). This expression is drastically reduced at E16.5 (G) and E18.5 (H). Axin2LacZ expression is found in both the epithelium and mesenchyme at E11.5 (I), E13.5 (J), E16.5 (K) and e18.5 (L).

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found in the developing vascular network underneath the skin. We next examined the expression of LacZ in the developing lung. Sustained TOPGAL and Axin2LacZ expression contrasts with decreased BATGAL expression in developing lung Mouse embryonic lungs at early, mid and late pseudoglandular stages (E11.5, E13.5 and E16.5) as well as at the saccular stage (E18.5) were isolated and examined for LacZ expression. At E11.5, E13.5 and E16.5 TOPGAL expression is mostly detected in the epithelium of the proximal and distal lung as well as in the trachea (Fig. 2A–C). At E18.5, LacZ expression is detected in the terminal bronchioles as well as the surrounding parenchyma (Fig. 2D). In contrast, at E11.5, BATGAL expression is detected in a “salt and pepper” fashion in the lung epithelium (Fig. 2E). This expression is drastically reduced at the other stages examined. Only patches of LacZ expression are observed (Fig. 2F–G). At E18.5, there is almost no LacZ expression detectable (Fig. 2H). We finally examined the expression of Axin2LacZ (Fig. 2I–L). At E11.5, LacZ expression is found in the lung epithelium with lower levels in the mesenchyme. In the trachea and the primary bronchi, it appears that the expression is in both the epithelium and the adjacent mesenchyme. At E13.5, this expression pattern is maintained with the exception of the primary bronchi where LacZ is now expressed only in the mesenchyme as a ring-like structure. At E16.5, LacZ expression is found in both the epithelium and mesenchyme. Such expression is maintained at E18.5. Vibratome sections of the E13.5 TOPGAL, BATGAL and Axin2LacZ lungs. TOPGAL is expressed in both epithelium and mesenchyme at the level of the bronchi (A, B) and restricted to the epithelium in the distal lung (C). BATGAL “salt and pepper” expression is found in a heterogeneous fashion in the lung (D). The expression is restricted to the mesenchyme adjacent to the bronchial epithelium (E). In the distal lung, BATGAL is sporadically expressed in the epithelium (F). Axin2LacZ is found exclusively in the mesenchyme adjacent to the epithelium at the bronchial level (G, H). While in the distal lung, LacZ expression is found mainly in the epithelium and at lower level in the mesenchyme (I).

![Figure 3. Vibratome sections of E13.5 TOPGAL, BATGAL and Axin2LacZ lungs.](https://example.com/figure3.png)

**Axin2LacZ** is a superior line to assess the response to naphthalene injury We next examined the response of the bronchial airway epithelium of the different Wnt reporter lines to naphthalene, a simple and robust model of lung injury. Corn oil was used as control. Only females were used for the naphthalene injury since females are more susceptible to this type of injury than males [13]. In the conducting airway, naphthalene caused significant epithelial cell death within 12 hours followed by re-epithelization of the airways, presumably by the P450neg variant of Clara cells [14,15]. Three days after injury onset, the epithelium is in part re-populated, with complete re-epithelialisation 10–14 days after injury [14,16]. We first visualized the impact of the injury on whole left lung of the Wnt reporter mice studied herein. TOPGAL expression in corn oil control shows a high baseline of LacZ expression in the bronchial epithelium (Fig. 4A, B). Since the epithelium was injured and sloughed off, this expression in the epithelium is significantly decreased in the naphthalene-treated lungs (Fig. 4C, D). LacZ expression is now detected in a stripe-like pattern around the bronchi in the parabronchial smooth muscle cells surrounding the damaged conducting airways (inset in Fig. 4D), suggesting a possible role of these cells in the repair process. In contrast, BATGAL expression is not detected in the control lung (Fig. 4E, F) and a slight but significant increase in LacZ expression is observed in the bronchial epithelium of...
naphthalene-treated lung (Fig. 4G, H). \(Axin2^{\text{LacZ}}\) expression is detected homogeneously throughout the corn oil control lung in both the conducting and respiratory airways (Fig. 4I, J). In the naphthalene-treated lung, LacZ expression is sharply upregulated in both areas of the lung (Fig. 4K, L). To confirm that the naphthalene injury occurred in our samples, we carried out CC10 staining to follow the Clara cells in control and experimental lungs. Typical robust CC10 expression was present in the corn oil treated lungs of \(TOPGAL\) (Fig. 5A, B and Fig. 6A, B), \(BATGAL\) (Fig. 5E, F and Fig. 6E, F) and \(Axin2^{\text{LacZ}}\) (Fig. 5I, J and Fig. 6I, J) in both proximal (Fig. 5) and distal airways (Fig. 6). Whereas, Clara cells in naphthalene-treated lungs were severely damaged in the three reporter lines as indicated by the localized expression of CC10 in the proximal (Fig. 5C, D; G, H; K, L) and distal airways (Fig. 6C, D, G, H, K, L). Consistent with the whole mount data, \(TOPGAL\) was expressed in proximal (Fig. 5A, B) and distal epithelium (Fig. 6A–B) in control lung on LacZ/CC10 staining, as well as in the parabronchial smooth muscle cells (PBSMCs) at both proximal (Fig. 7A, B) and distal (Fig. 8A, B) levels as indicated by SMA staining. While CC10 staining decreased drastically in the naphthalene-treated \(TOPGAL\) lungs, \(TOPGAL\) expression is present in the CC10-positive cells re-populating the airways mainly in the proximal compartment (Fig. 5C, D) as compared to the distal compartment (Fig. 6C, D). Moreover, \(TOPGAL\) expression was still present in the PBSMCs surrounding the proximal (Fig. 7C, D) and distal airways (Fig. 8C, D). Moreover, \(BATGAL\) expression is not detected in control lungs at the proximal (Fig. 5E, F) or distal level (Fig. 6E, F). However, \(BATGAL\) is found at discrete spots in the bronchial epithelium of naphthalene-treated lungs (Fig. 5G, H) and is still completely absent in the distal airways (Fig. 6G, H) and in the PBSMCs (Fig. 7G, H and Fig. 8G, H). Finally, \(Axin2^{\text{LacZ}}\) expression is found at low level throughout the lung in both the epithelium (Fig. 5I, J and Fig. 6I, J) and the PBSMCs (Fig. 7I, J and 8I, J) in proximal and distal airways. An increase of LacZ expression is observed in both proximal (Fig. 5K, L) and distal epithelium (Fig. 6K, L) of naphthalene-

Figure 4. Corn oil and Naphthalene-treated whole mount left lungs cleared with Benzyl benzoate. In the adult lung, \(TOPGAL\) is strongly expressed in the epithelium (A, B) and decreases slightly after naphthalene injury (C, D). \(BATGAL\) expression is totally absent in the adult lung (E, F) and a discrete expression appears after injury (G, H). While a homogenous \(Axin2^{\text{LacZ}}\) expression is present throughout the adult lung (I, J), a marked increase surrounding the bronchial epithelium is observed after injury (K, L).

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treated lungs as well as in the PBSMCs surrounding proximal (Fig. 7K,L) and distal airways (Fig. 8K,L). The arrows displayed in Fig. 7 and 8 show co-localization of SMA and LacZ after naphthalene injury in TOPGAL and Axin2LacZ mice. qRT-PCR analyses showed a 10-fold increase in beta-galactosidase expression in the Axin2LacZ naphthalene-treated lungs as compared to controls accompanied with a 8-fold decrease in CC10 expression (data not shown).

Discussion

Our aim was to provide a systematic comparison of the expression patterns of three different but classical Wnt reporter lines that are in common use and on whose output depend a growing number of published research findings. We exploited the stable Axin2LacZ line to provide an internal control for the older TOPGAL and BATGAL lines where random insertion may alter expression from the original reports (both first made 6 years ago). We chose to use the developing lung and the repairing lung as model systems in which to test such expression since both scenarios require canonical Wnt signaling. Our findings can be a significant resource to not only the lung field but also the many other bioscience research areas in which Wnt function is being investigated. Crosses to introduce one of these reporter alleles in already complex combinations of driver and responder lines is resource and time intensive but our findings can guide the critical selection of the appropriate Wnt reporter line.

Our data indicate that Axin2LacZ mice are the Wnt reporter line of choice for the specific detection of increased Wnt signaling in the epithelium. Surprisingly, only one paper so far makes use of this line to follow Wnt signaling in vivo in the lung [17]. Most of the papers published to date make extensive use of TOPGAL mice. The usefulness of the BATGAL mice to detect an increase in Wnt signaling is due to the progressive disappearance of LacZ signal as the embryo develops with almost complete absence of LacZ expression in the E18.5 lungs. Similar results are observed in the lungs of 2 month-old BATGAL mice. In harmony with these
results, these mice have been used to demonstrate that a Gata6-Wnt pathway is required for epithelial stem cell development and airway regeneration [18]. Gata6 is a transcription factor, which negatively regulates the canonical Wnt pathway. Inactivation of Gata6 in the lung epithelium in the background of the BATGAL allele leads to a drastic increase in LacZ expression in the epithelium at E13.5 and E16.5 demonstrating the negative role played by Gata6 on the activation of Wnt signaling. The same authors also reported the progressive appearance of BATGAL positive cells, starting at 2 days post-naphthalene injury, in the broncho-alveolar duct junction of the airways. Our results concerning the use of BATGAL are therefore in agreement with this report.

The TOPGAL reporter has been used most extensively, probably because it was the first published line made available to monitor Wnt signaling in vivo. During development, TOPGAL was present in epithelium of the developing lung from E11.5 to E18.5, with higher expression in proximal compared to distal epithelium [19]. This line has mainly been used to demonstrate reduced canonical Wnt signaling. For example this line was used to validate the biological activity of recombinant Dickkopf-1, a canonical Wnt inhibitor [20], to show that deletion of R-Spondin leads to reduction in TOPGAL reporter activity [21] and to demonstrate reduction of Wnt signaling in elastase and cigarette smoke-induced lung emphysema [22]. Moreover, an increase in TOPGAL expression was reported in hyperoxia injury model in the lungs of neonate mice [23].

Our results with Axin2lacZ mice indicate that this line is the most robust and faithful line to detect Wnt signaling. As expected for such a critical pathway, this line indicates that Wnt signaling occurs throughout lung development as well as in the adult lung, in both the epithelium and the mesenchyme, in both the conducting and respiratory airways. Harris-Johnson et al. showed that Axin2lacZ expression is expressed early on (E9.5) in the prospective respiratory region and it is restricted to the ventral foregut that will form the lung and trachea [24]. This line has also been used by another group reporting that
adult alveolar type II cells do not exhibit constitutive beta-catenin signaling in vivo. However, after bleomycin injury, a significant increase in the number of LacZ/SPC double positive cells is observed [17]. We also showed increased expression of Axin2LacZ in response to naphthalene injury in adult mice, and in response to hyperoxia injury in neonates (data not shown). Although, this line will be used more extensively in the future, an important inherent limitation of the Axin2LacZ line is the corresponding up-regulation of endogenous Wnt signaling, since the Axin-related protein, AXIN2, modulates beta-catenin stability. Deregulation of beta-catenin is important in the genesis of several malignancies and this aspect will therefore have to be taken into consideration. The Axin2lacZ mice therefore represent more than just stable reporters for Wnt signaling; they are also a gain of function of canonical beta-catenin signaling, both in the epithelium and mesenchyme. This line has been used with success to demonstrate that Wnt signaling allows self-renewal of mammary stem cells and promotes their long-term propagation in culture [25].

In conclusion, our data indicate that the choice of the appropriate Wnt reporter line should be tailored to the need to detect either up- or down-regulation of the canonical Wnt signaling pathway in the lung epithelium versus the mesenchyme. Until better tools to follow Wnt signaling are available (e.g. a transgenic construct with Axin2 regulatory sequences upstream of LacZ), this conclusion is highly relevant to the wide range of research fields studying Wnt signaling. Moreover, our findings represent a reference resource for researchers pursuing such work within the pulmonary field and beyond.

Materials and Methods

Mice

The 3 reporter mice used in this study were obtained from the Jackson Lab. TOPGAL mice (Tg(Fos-lacZ)34Efu/J, stock number 004623) were generated by Das Gupta and Fuchs in 1999. BATGAL mice (B6.Cg-Tg(BAT-lacZ)3Picc/J, 005317) were
generated by Maretto et al. in 2003. \(\text{Axin}^{2\text{lac}Z}\) (B6.129P2-Axin\textsuperscript{2\text{lac}Z/Wbm}/J, stock number 009120) were generated by Lustig et al. in 2002 [11]. BATGAL and \(\text{Axin}^{2\text{lac}Z}\) lines purchased from the Jackson lab are on a C57 black (C57BL) background while \(\text{TOPGAL}\) mice are on CD1 background. To eliminate any background variabilities, we backcrossed the \(\text{TOPGAL}\) line with C57BL for more than 6 generations to obtain a pure C57BL background. Animal experiments were performed under the research protocol approved by the Animal Research Committee at Children’s Hospital Los Angeles and in strict accordance with the recommendations in the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. The approval identification for Children’s Hospital Los Angeles is AAALAC A3276-01. These experiments were done under the protocol 31-08.

**X-gal staining**

Mouse embryos were isolated at E11.5 and E12.5 in Hank’s solution, washed briefly in PBS, pre-fixed for 10 min with 4% PFA and washed twice in LacZ buffer solution. Embryos were incubated overnight at 37°C with the LacZ buffer solution containing 40 mg/mL of X-gal (rpi research products). Embryonic lungs were dissected out in DMEM from embryos at E11.5, E13.5, E16.5 and E18.5. Lungs were washed briefly in PBS and pre-fixed for 10 min in 4% PFA before incubation overnight with the X-gal solution. Vibratome lung sections 20 \(\mu\)m thickness were carried out at E13.5. Three independent litters for each time point were collected. Littermates that do not carry the beta-galactosidase were used as controls and did not show any LacZ staining.

For adult lungs, transcardiac perfusion with PBS was performed to remove the red blood cells in the lung. The lungs were inflated trans-tracheally at 25 cm of water pressure and submerged in 4% PFA for 5 min, washed 5 minutes in PBS and 5 minutes in LacZ buffer. The lungs were then inflated again with the LacZ buffer containing 40 mg/mL of X-gal. The trachea was ligated using sutures to maintain the staining solution inside the lung. The whole lung was then incubated in 10 mL of staining solution overnight at 37°C. The

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**Figure 8. SMA and LacZ staining in the distal lung compartment after naphthalene injury.** SMA and LacZ co-staining in \(\text{TOPGAL}\) control lungs at low (A) and high (B) magnification showed co-localization in the PBSMCs of control (A, B) and naphthalene-treated lungs (C, D). \(\text{BATGAL}\) expression is not detected in the PBSMCs of adult control lungs (E, F) and naphthalene-treated lungs (G, H). \(\text{Axin}^{2\text{lac}Z}\) sections showed low level staining in the PBSMCs surrounding the distal bronchioles in control adult lungs (I, J) and an increase after injury (K, L). The arrows show co-localisation of SMA and LacZ. Dotted lines show the basal membrane separating the epithelium from the PSMC layer. Scale bars are 100 \(\mu\)m. doi:10.1371/journal.pone.0023139.g008
lungs were subsequently washed with PBS and fixed again in 4% PFA in PBS at room temperature overnight. For better visualization of the staining inside the lung, lungs were dehydrated and cleared with BABB (1:2 Benzyl Alcohol and Benzyl Benzoate). Dehydrated lungs were transferred to 1:2 BABB and ethanol for 20 min, 2:1 BABB and ethanol for 20 min, and 100% BABB for 20 min.

Naphthalene injury

Naphthalene, NA (Fisher, Aschaffenburg, Germany) was dissolved in corn oil at 30 mg/mL. TOPGAL, BATGAL and Axin2LacZ 2-month-old ice were injected IP with 250 mg/kg body weight of either naphthalene or the same volume of a vial control of corn oil alone and the mice were sacrificed 3 days later. For TOPGAL and BATGAL, 4 adult females were injected with corn oil and 4 adult females were injected with naphthalene. For Axin2LacZ 8 adult females were used for control group and 8 for experimental group. The time-point of 3 days was chosen because experiments studying the kinetics of naphthalene-induced acute airway injury revealed that epithelial cells in the conducting airways are proliferating to repair the damaged bronchial epithelium between 2–4 days. It has been reported that complete exfoliation was observed 24 h after naphthalene injection.

Immunohistochemistry

For microtome sections, after 4% PFA fixation, lungs were washed in PBS, dehydrated, and embedded in paraffin. Sections were performed at 5 microns. The sections were cleared with 2 changes of xylene and hydrated in successive graded Ethanol solutions, equilibrated in water then washed in 3% H2O2 for 20 min at room temperature. The sections were incubated with primary antibodies anti-CC10 (santa Cruz, 1:200 dilution) and anti-alpha smooth muscle actin (Dako cytomation, 1:200 dilution) at 4°C overnight. Immunohistochemistry was performed using Dako EnVision Kit following the manufacturer’s instructions. Slides were mounted using xylene-based mounting media. Brightfield images were acquired on an Axio Observer.Z1 microscope equipped with an AxioCam MRc color CCD camera (Carl Zeiss Microimaging, Thornwood, NY). Microscope control and image processing were done with AxioVision 4.8.1.0 software (Carl Zeiss). Images at different magnifications were acquired with the following objective lenses: 20x/0.8 Plan- APOCHROMAT, 40x/1.3 Plan-NEOFLUAR oil immersion, and 63x/1.4 Plan-APOCHROMAT oil immersion (Carl Zeiss). For some fields of view several focus planes were acquired and an Extended Focus projection is shown. Where larger areas were needed four adjacent fields of view were acquired and stitched together with the MosaicX module of the software. CC10 staining was performed on each adult animal included in this study to verify that the injury did occur.

Real time PCR

RNA was extracted from Axin2LacZ lungs treated with naphthalene or corn oil for 3 days. RNA was reverse-transcribed into cDNA using Transcriptor High Fidelity cDNA Synthesis Kit (Roche Applied Science) according to the manufacturer’s instructions. cDNA was used for dual color Hydrolysis Probe – Universal probe library based real time PCR, using the LightCycler 480 from Roche Applied Science. GAPDH assay commercially available from Roche Applied Science was used as reference gene. The primers and probes used for CC10 and beta-galactosidase are as follows: CC10 Left 5’-gatcgccatcacaatg3’; Right 5’-catccagtcggaaagaatgctg-3’; probe #56 (Roche Applied Science UPL); beta-galactosidase: Left 5’-aattgatgagcagacgatgg-3’; Right 5’-cgcccctgaattttgtgc-3’ with probe #18 (Roche Applied Science).

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Author Contributions

Conceived and designed the experiments: SB EJ RV DW. Performed the experiments: DA MG RJ SP JB FS EA SD. Analyzed the data: DA SB CT FS. Contributed reagents/materials/analysis tools: DA SB FS. Wrote the paper: DA SB EJ DW CT.

References

1. De Langhe SP, Reynolds SD (2008) Wnt signaling in lung organogenesis. Organogenesis 4: 100–108.
2. Warburton D, El-Haddad A, Carraro G, Tiozzo C, Sala F, et al. Lung organogenesis. Curr Top Dev Biol 90: 73–138.
3. Nusse R (1999) WNT targets. Repression and activation. Trends Genet 15: 1–3.
4. Bremsbeck FH, Schwarz-Romond T, Bakkers J, Wilhelm S, Hammerschmidt M, et al. (2004) Essential role of BCL-2 in the switch between beta-catenin’s adhesive and transcriptional functions. Genes Dev 18: 2225–2230.
5. Price MA (2006) CKI, there’s more than one: casein kinase I family members in development. Development 133: 3205–3214.
6. Gordon MD, Nusse R (2006) Wnt signaling: multiple pathways, multiple receptors, and multiple transcription factors. J Biol Chem 281: 22429–22433.
7. van Amerongen R, Nusse R (2009) Towards an integrated view of Wnt signaling. Trends in Developmental Biology 22: 316–331.
8. Flozac AS, Lam AP, Russell S, Jain M, Peled ON, et al. (2010) Beta-catenin/T-cell factor signaling is activated during lung injury and promotes the survival and migration of alveolar epithelial cells. J Biol Chem 285: 3157–3167.
9. Zhang Y, Goss AM, Cohen ED, Kardos R, Lepori JJ, et al. (2008) A Gat6-Wnt pathway required for epithelial stem cell development and airway regeneration. Nat Genet 40: 482–487.
10. Okubo T, Hogan BL (2004) Hyperactive Wnt signaling changes the developmental potential of embryonic lung endoderm. J Biol 3: 11.
11. De Langhe SP, Sala FG, Del Moral PM, Fairbanks TJ, Yamada KM, et al. (2005) Dickkopf-1 (DKK1) reveals that fibronectin is a major target of Wnt signaling in branching morphogenesis of the mouse embryonic lung. Dev Biol 277: 316–331.
12. Bell SM, Schreiner CM, Wett SE, Muencens ML, Scott WJ, et al. (2008) R-spondin 2 is required for normal larval lung-branchial, lung and limb morphogenesis. Development 135: 1049–1058.
13. Van Winkle LS, Gunderson AD, Shizuma JA, Baker GJ, Brown CD (2002) Gender differences in naphthalene metabolism and naphthalene-induced acute lung injury. Am J Physiol Lung Cell Mol Physiol 282: L1122–L1134.
14. Stripp BR, Masson K, Mera R, Singh G (1995) Plasticity of airway cell proliferation and gene expression after acute naphthalene injury. Am J Physiol 269: L791–799.
15. Mahvi D, Bank H, Harley R (1977) Morphology of a naphthalene-induced bronchiolar lesion. Am J Pathol 86: 558–572.
16. Van Winkle LS, Buckpitt AR, Nishio SJ, Isaac JM, Plopper CG (1995) Cellular response in naphthalene-induced Clara cell injury and bronchiolar epithelial repair in mice. Am J Physiol 269: L810–818.
17. Flozac AS, Lam AP, Russell S, Jain M, Peled ON, et al. (2010) Beta-catenin/T-cell factor signaling is activated during lung injury and promotes the survival and migration of alveolar epithelial cells. J Biol Chem 283: 3157–3167.
18. Zhang Y, Goss AM, Cohen ED, Kardos R, Lepori JJ, et al. (2008) A Gat6-Wnt pathway required for epithelial stem cell development and airway regeneration. Nat Genet 40: 482–487.
19. Okubo T, Hogan BL (2004) Hyperactive Wnt signaling changes the developmental potential of embryonic lung endoderm. J Biol 3: 11.
20. De Langhe SP, Sala FG, Del Moral PM, Fairbanks TJ, Yamada KM, et al. (2005) Dickkopf-1 (DKK1) reveals that fibronectin is a major target of Wnt signaling in branching morphogenesis of the mouse embryonic lung. Dev Biol 277: 316–331.
21. Bell SM, Schreiner CM, Wett SE, Muencens ML, Scott WJ, et al. (2008) R-spondin 2 is required for normal larval lung-branchial, lung and limb morphogenesis. Development 135: 1049–1058.
22. Kurcinski N, Yildirim AO, Cellegari J, Takenaka S, Stein MM, et al. (2010) Activation of the WNT/ﬁbeta-Catenin Pathway Attenuates Experimental Emphysema. Am J Respir Crit Care Med.
23. Dasgupta C, Sakurai R, Wang Y, Guo P, Ambalavanan N, et al. (2009) Hyperoxia-induced neonatal rat lung injury involves activation of TGF-{beta} and Wnt signaling and is protected by rosiglitazone. Am J Physiol Lung Cell Mol Physiol 296: L1031–1041.

24. Harris-Johnson KS, Domyan ET, Vezina CM, Sun X (2009) beta-Catenin promotes respiratory progenitor identity in mouse foregut. Proc Natl Acad Sci U S A 106: 16287–16292.

25. Zeng YA, Nusse R (2010) Wnt proteins are self-renewal factors for mammary stem cells and promote their long-term expansion in culture. Cell Stem Cell 6: 568–577.