Some cancer prognoses have been radically improved in recent years, but little headway has been made with others. One of these diseases, small-cell lung cancer (SCLC), has a five-year survival rate of less than 7% and a standard of care that has been essentially unchanged for forty years. One promising avenue to improve SCLC outcomes is to understand the cancer’s underlying genetic alterations that drive its formation and growth. Functional inactivation of the Rb gene is seen in a number of cancers and is a genetic hallmark of SCLC. Normally Rb promotes differentiation by regulating lineage-specific transcription factors, including pluripotency factors such as Sox2. However, there is evidence that when certain tissues lose Rb, Sox2 becomes upregulated and promotes oncogenesis. To understand this relationship in the pursuit to uncover new treatments for SCLC, we have studied the role of Sox2 in Rb loss initiated tumors by investigating both the tumor initiation in a SCLC genetically engineered mouse model, as well as tumor maintenance in SCLC cell lines and organoid culture.

B27
IH7 Acts as a Tumor Suppressor of Lung Adenocarcinoma by Repressing Reactive Oxygen Species
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Background: Aberrant activation of the Hedgehog (Hh) signaling pathway, a crucial developmental pathway, drives the tumor growth of basal cell carcinoma, medulloblastoma, and rhabdomyosarcoma. However, recent data suggest that paracrine activation of the pathway is tumor suppressive rather than oncogenic in sporadic epithelial cancers. The recent data suggest that paracrine activation of the pathway is tumor suppressive rather than oncogenic in sporadic epithelial cancers. The role of the pathway in the non-small lung cancer is poorly understood. Thus, we explored the role of stromal Hh pathway activation in growth of lung tumor epithelia. Methods: Human and murine lung adenocarcinoma cell lines and murine fibroblasts were used to probe SHH mRNA and protein expression and to verify paracrine activation of the Hh signaling pathway. The role of paracrine SHH was tested in vivo using KraLSL-G12D/fl, Trp53G6/10 (KP) and LSL-KraG12D/fl, Trp53G6/10, Shhfl/fl (KPS) autochthonous murine lung cancer models. The role of IH7 was examined in vivo using the pSCEC CRISPR system in KP;Rosa26LSL-Fluc/+ and tumor growth monitored by bioluminescence imaging. Results: In human lung adenocarcinoma (LAD) patients, higher expression of SHH mRNA correlated with poor overall and progression free survival. Co-culture of high SHH-expressing tumor epithelial cells and Shh-Light2 reporter fibroblasts demonstrated that SHH activated the Hh pathway in the fibroblasts in a paracrine manner. Surprisingly, genetic loss of SHH in an autochthonous mouse model, KPS, did not affect overall survival compared to KP mice. However, early inhibition of stromal Hh pathway by 5E1, an anti-SHH/HH pathway, in KP mice resulted in significantly worse survival with increased metastatic burden. We tested the loss of IH7 in vivo with the pSCEC CRISPR system. IH7-loss in airway epithelia led to more aggressive tumor growth, suggesting that IH7, not SHH, activates the pathway in stroma to drive its tumor suppressive effects—a novel role for IH7 in the lung. Tumors from mice treated with 5E1 had decreased blood vessel density and increased reactive oxygen species (ROS). Treatment of KP mice with 5E1 and N-acetylcysteine, as a ROS scavenger, decreased tumor ROS levels, inhibited tumor growth, and prolonged mouse survival, suggesting that increased ROS levels from stromal Hh pathway inhibition accelerated lung tumor growth. Conclusions: IH7 activates the Hh signaling pathway in lung stroma in a paracrine manner to suppress tumor growth and metastases, in part, by limiting ROS production.

B28
Intertumoral Hypoxia Exacerbates Tumor Progression in a Mouse Model of Lung Cancer

Background: Obstructive sleep apnea (OSA) is a very prevalent disorder characterized by chronic intermittent hypoxia (CIH), and some reports suggested that OSA is related to increased incidence of cancer as well as cancer progression. The purpose of this study was to evaluate whether obstructive sleep apnea (OSA)-related chronic intermittent hypoxia (CIH) influences lung cancer progression and to elucidate the associated mechanisms in a mouse model of lung cancer. Methods: C57/B6 mice in a CIH group were exposed to intermittent hypoxia for two weeks after tumor induction and compared with control mice (room air). Hypoxia inducible factor 1α (HIF-1α), vascular endothelial growth factor (VEGF), and metastasis-related matrix metalloproteinases (MMP) were measured. The expression levels of several hypoxia-related pathway proteins including HIF-1α, VEGF, and mammalian target of rapamycin-ERK were measured by Western blot. Results: The number (P < 0.01) and volume (P < 0.05) of tumors were increased in the CIH group. The activity of MMP-2 was enhanced after CIH treatment. The level of VEGF was increased significantly in the CIH group (p < 0.05), β-catenin, and Nrf2 were translocated to the nucleus and the levels of downstream effectors of Wnt/β-catenin signaling increased after H exposure. Conclusions: CIH enhanced proliferative and migratory properties of tumors in a mouse model of lung cancer. β-Catenin and Nrf2 appeared to be crucial mediators of tumor growth. These results suggest evidence for the causal link between OSA and lung cancer progression.

B30
The Role of SMARCA4 as an EGFR-Independent Mechanism of Resistance to Osimertinib

Targeted therapies have replaced conventional chemotherapy as first-line treatment for patients with non-small cell lung cancers harboring epidermal growth factor receptor (EGFR) alterations. Although tyrosine-kinase inhibitors (TKI) targeting these proteins lead to responses in ~70% of cases, tumors almost inevitably become resistant. Acquired resistance is commonly caused by secondary mutations in the target oncogene, activation of bypass signaling pathways, histologic transformation of the tumor, or unknown mechanisms (~20-40%). Epigenetic mechanisms are responsible for regulating genes involved in cell lineage specificity, and they are known to modulate tumorigenesis. In recent years, several epigenetic modifiers have also been implicated in processes related to drug resistance. We hypothesized that dysfunction of epigenetic processes plays a role in mediating resistance to TKIs. To examine this possibility, we generated three isogenic osimertinib-sensitive/resistant cell line pairs and mined whole-exome and RNA sequencing (RNA-seq) data. Distinct alterations and phenotypes were identified in the different models, highlighting the importance of the baseline biologic context for the type of osimertinib resistance