

Tumor Cell Specific Function of IL-1 Signaling in the Pathogenesis of K-ras Mutant Lung Cancer

Milind Mutala¹, Avantika Krishna^{1,2}, Bo Yuan¹, Michael J. Clowers^{1,2}, Seyed Javad Moghaddam^{1,2}

¹Department of Pulmonary Medicine, UT MD Anderson Cancer Center, ²The University of Texas M.D. Anderson Cancer Center UTHealth Graduate School of Biomedical Sciences, Houston, TX, USA.

THE UNIVERSITY OF TEXAS
MD Anderson
Cancer Center

Making Cancer History®

Objective

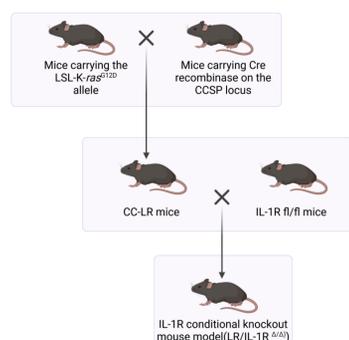
Conditional knockout of the IL-1R receptor in KM-LUAD mouse models will allow for a greater understanding of the involvement of the IL-1 cytokine family in tumor promotion and potentially support the idea of a new target for treatment.

Introduction

- As of last year, 23% of cancer-related deaths have been attributed to lung cancer. It is the third most common cause of cancer in males and females and its low survival rate has also been an area of concern with only 26% of diagnosed patients surviving up to 5 years.
- Non-small cell lung cancer (NSCLC) is the most common type of lung cancer, in which mutations of the *K-ras* oncogene are the driving factor leading to the promotion of lung adenocarcinoma (LUAD), a histological subtype of NSCLC.
- Mutations of the *K-ras* oncogene have been shown to cause increased activation of the NF- κ B pathway and inflammatory response, promoting a pro-tumor microenvironment.
- The pro-inflammatory cytokine IL-1 β plays a major role in the activation of the NF- κ B pathway and has been found in abundance at tumor sites.

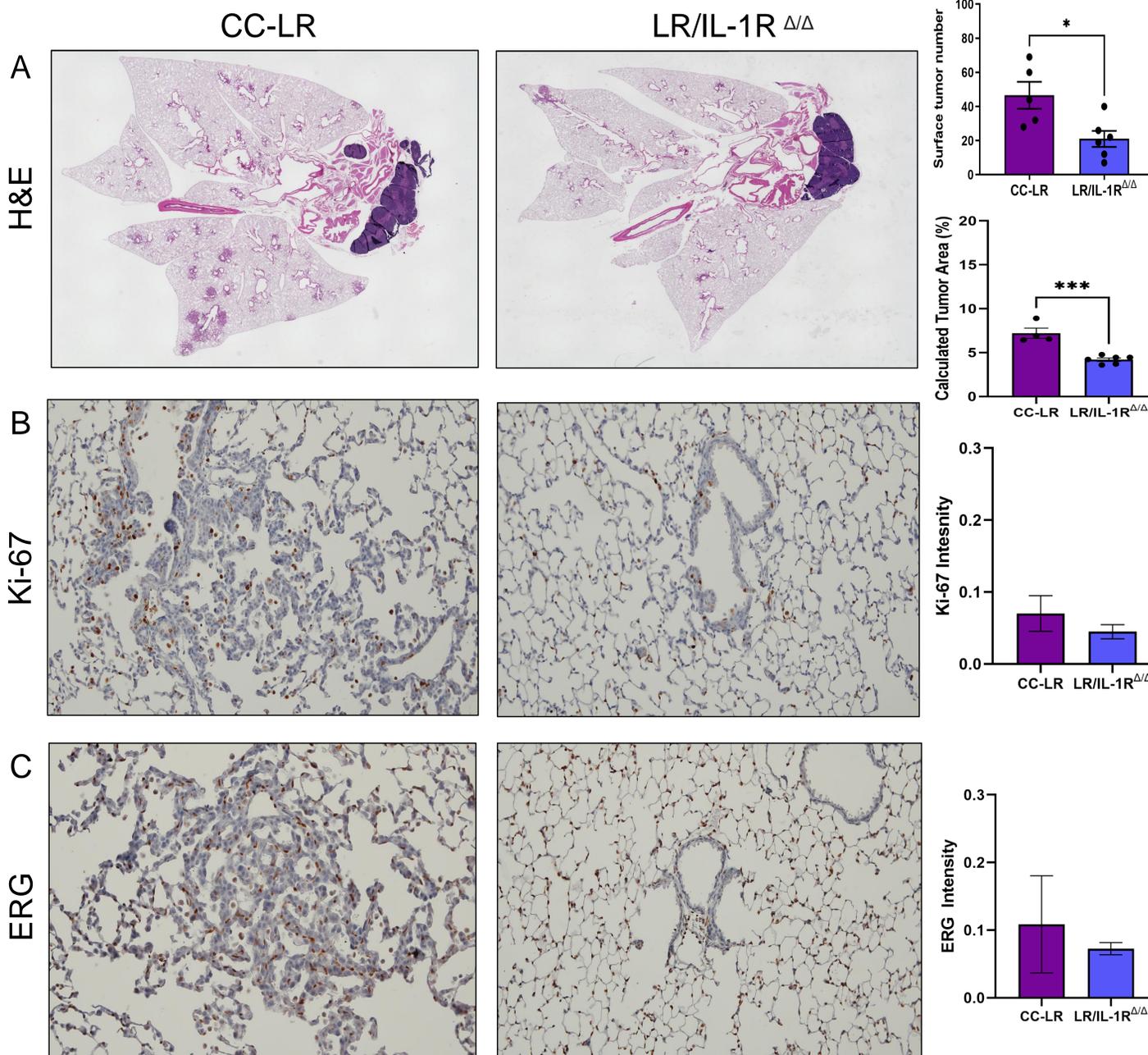
Methods

Animals:



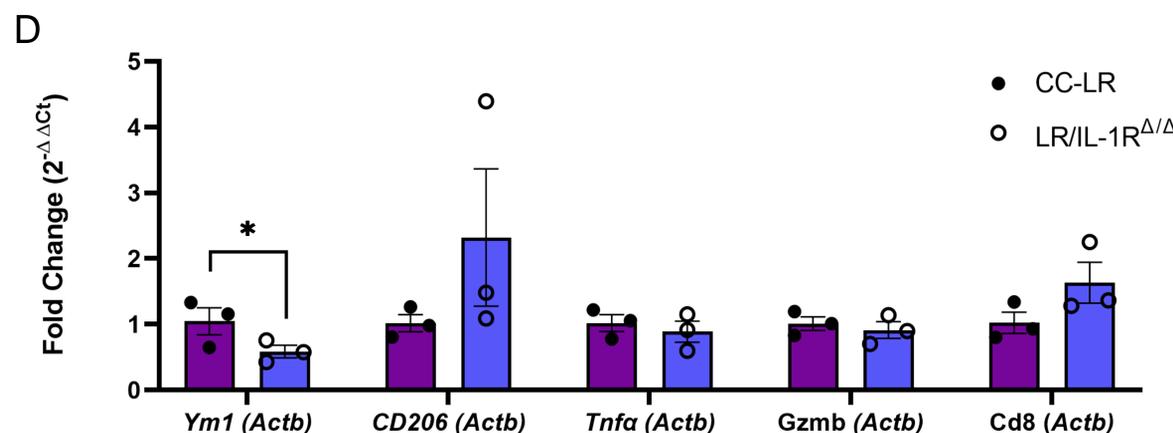
Step 1

Results



Conditional Knockout of the IL-1 receptor leads to a significant decrease in tumor area. (A) Representative photomicrographs of the H&E stained whole lung sections from 14 week old CC-LR mice as well as LR/IL-1R $\Delta\Delta$ mice (4X). Representative photomicrographs of the stained IHC sections (20x) and corresponding quantification of (B) proliferation marker Ki-67 and (C) angiogenesis marker ERG. Data represent mean \pm SEM; unpaired t-test, * p <0.05.

Step 2



Conditional Knockout of the IL-1 receptor led to a decrease in M2 phenotype. Relative mRNA expression of Ym1, CD206, TNF- α , Gzmb, and CD8 in the whole lung. Data represent mean \pm SEM; unpaired t-test, * p <0.05.

Conclusion

- IL-1R conditional knockout results in a significant decrease in tumor burden as shown by the H&E staining.
- IHC staining indicated a negative trend in the expression of Ki67 along with ERG, which could mean the IL-1R conditional knockout suppressed tumor proliferation and angiogenesis.
- qRT-PCR analysis of RNA extracted from the lung revealed an increase in Cd8a expression along with a significant decrease in Ym1 expression, which supports the notion of increased antitumor cytotoxic activity.

Future Directions

- Repeat qPCR analysis to confirm results and add additional markers to gain a more complete understanding of the tumor microenvironment.
- Stain samples with p65 marker to evaluate the effects the conditional knock-out of IL-1R will have on the NF- κ B pathway.

References

- Aredo, Jacqueline V et al. "Impact of KRAS mutation subtype and concurrent pathogenic mutations on non-small cell lung cancer outcomes." *Lung cancer (Amsterdam, Netherlands)* vol. 133 (2019): 144-150. doi:10.1016/j.lungcan.2019.05.015
- Dinarelli CA. Overview of the IL-1 family in innate inflammation and acquired immunity. *Immunol Rev.* 2018 Jan;281(1):8-27. doi: 10.1111/imr.12621. PMID: 29247995; PMCID: PMC5756628.
- <https://doi.org/10.1016/j.lungcan.2019.05.015>.
- Yuan B, Clowers MJ, Velasco WV, Peng S, Peng Q, Shi Y, Ramos-Castaneda M, Zarghooni M, Yang S, Babcock RL, Chang SH, Heymach JV, Zhang J, Ostrin EJ, Watowich SS, Kadara H, Moghaddam SJ. Targeting IL-1 β as an immunopreventive and therapeutic modality for K-ras-mutant lung cancer. *JCI Insight.* 2022 Jun 8;7(11):e157788. doi: 10.1172/jci.insight.157788. PMID: 35471938; PMCID: PMC9220853

Acknowledgments

R01 grant from NIH/NCI (R01CA225977) Lung Cancer Discovery Award from the American Lung Association (grant LCD- 821433)

Step 1(H&E and IHC):

H&E staining was used to calculate tumor area. IHC staining for Ki-67 and ERG markers were performed to measure proliferation and angiogenesis, respectively.

Step 2(qPCR):

Used to find the concentration of desired DNA in our samples of interest. Fold changes between test groups and controls were calculated using the 2 $^{-\Delta\Delta Ct}$ method.