



Artificial Intelligence in Genomic Surveillance: Mitigating Lung Cancer Risk and Metastasis in Aviation and Space Warfighters

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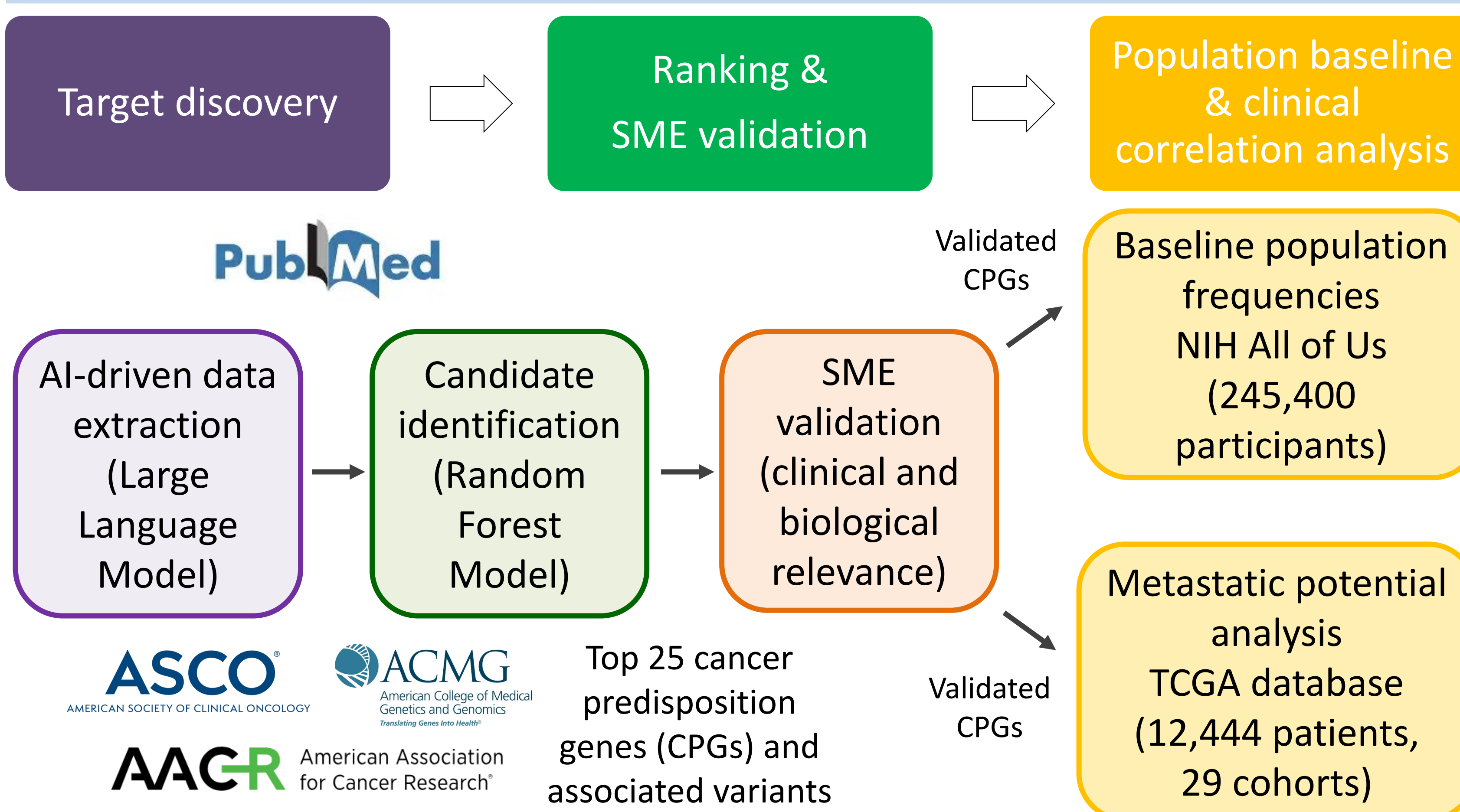
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Background

- Ionizing radiation from space is a major cancer risk for DoW pilots and aircrew, who frequently operate at high altitudes and/or high latitudes, and a primary threat to USSF Guardians and astronauts during their extended or deep space missions.
- Lung cancer, a leading cause of cancer-related death in the U.S., is not only common in DoW personnel but also highly sensitive to radiation.
- Currently, there is a significant gap in understanding how genetic predispositions interact with occupational radiation exposure, particularly concerning the development and metastasis of lung cancer.
- This lack of knowledge limits the effectiveness of current strategies for cancer prevention and early detection among high-risk groups like pilots, astronauts, and aircrew.

Methods



Results

- Prevalent pathogenic CPG mutations in the US population include *TP53*, *TSC2*, *TERT*, *PALB2*, *ERCC2*, *BRCA2*, *CHEK2*, *CDKN2A*, *BLM*, and *ATM*.
- Our AI/ML-powered big data analysis revealed significant links between specific mutations in genes like *EGFR*, *ALK*, *TP53*, *STK11*, *ATM*, *PALB2*, *CDKN2A*, *TSC2*, and *TERT* and higher rates of lung cancer metastasis.
- Direct impact on risk stratification, providing a genomic basis for identifying personnel who would benefit from enhanced surveillance and proactive risk mitigation.

The views expressed in this material are those of the authors and do not necessarily reflect the official policy or position of the Defense Health Agency, Department of Defense, nor the U.S. Government. This study was conducted under Defense Health Agency-approved Clinical Investigation Numbers FDG20240051R and FDG20240062R.

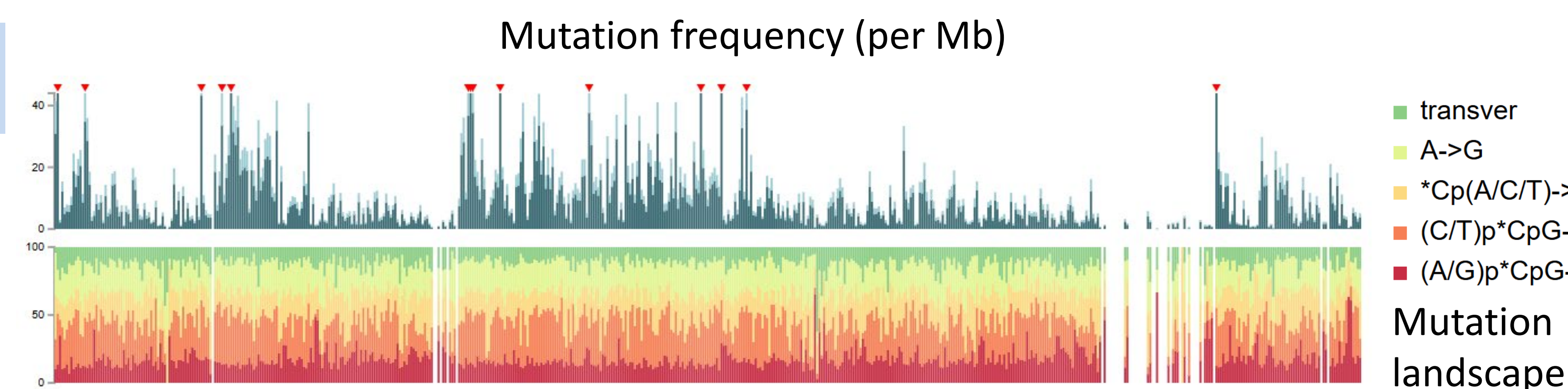


Fig. 1: Mutation signature of lung cancer patients in TCGA database

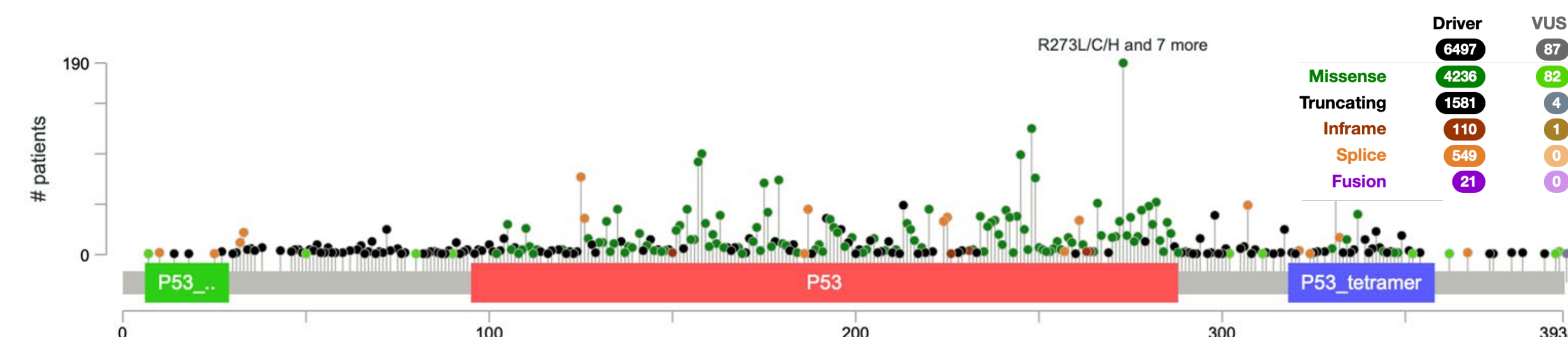


Fig. 2: Comprehensive mapping of *TP53* gene mutation landscape in lung tumors

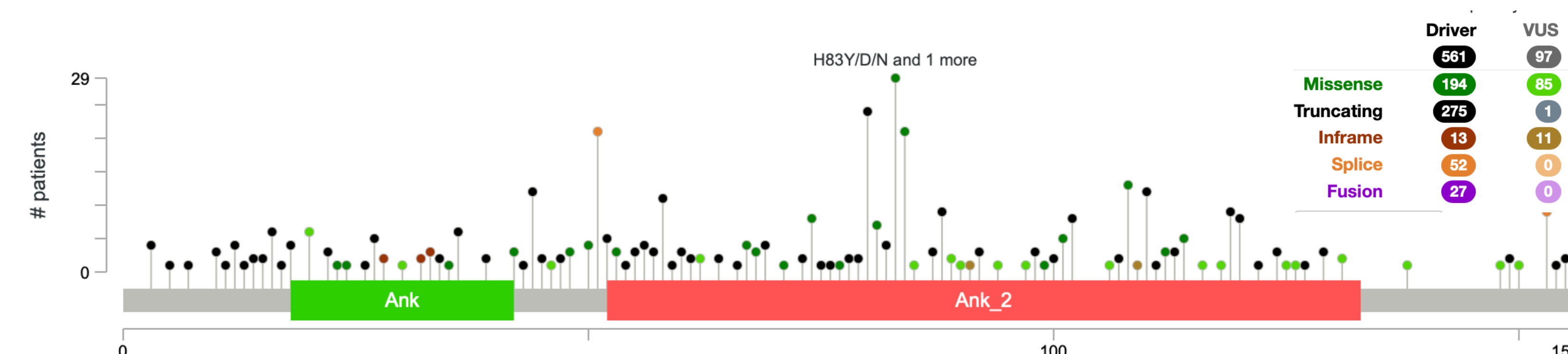


Fig. 3: Comprehensive mapping of *CDKN2A* gene mutation landscape in lung tumors

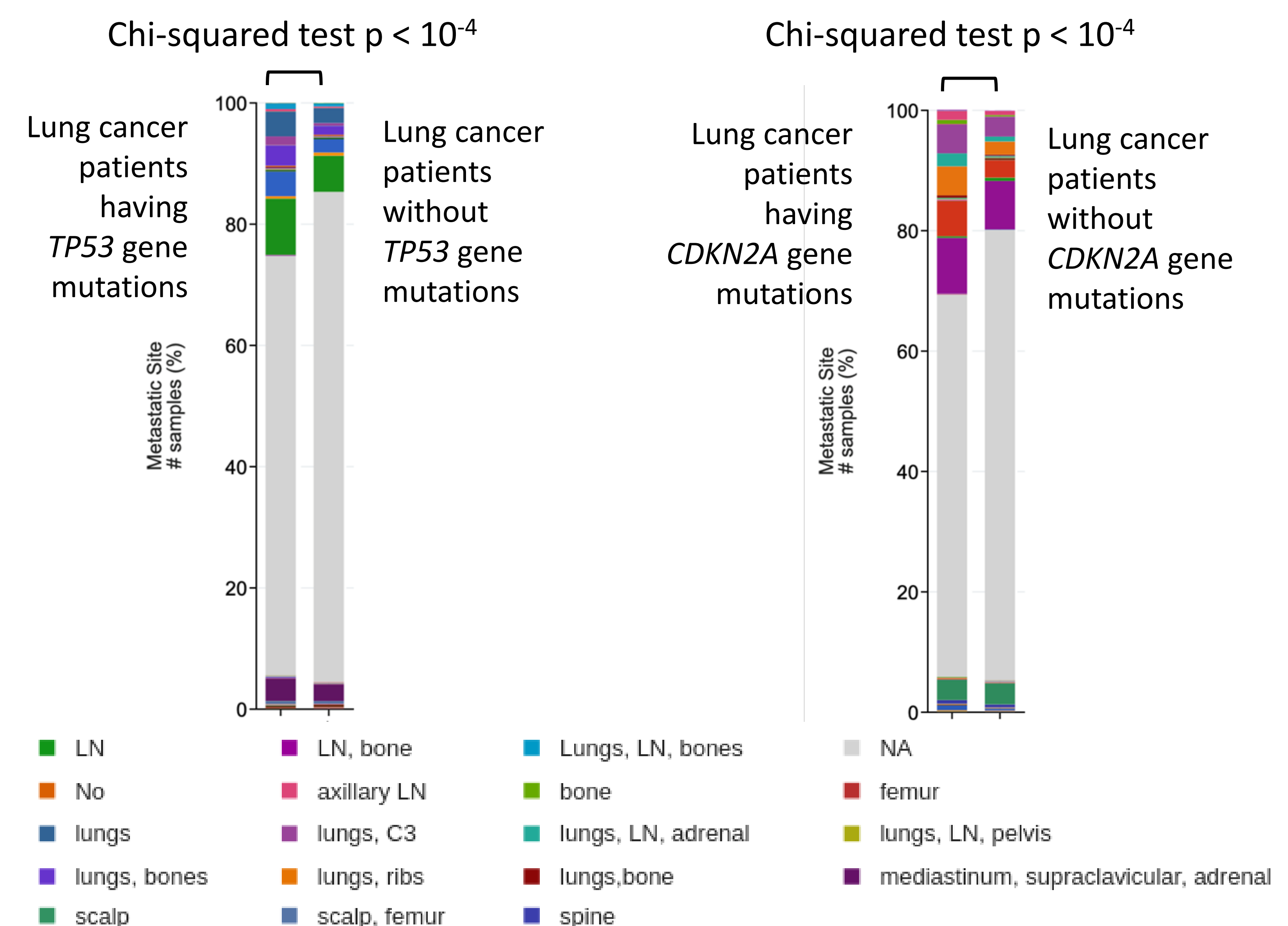


Fig. 4: Significant increase in lung cancer metastasis in the presence of *TP53* and *CDKN2A* cancer predisposition gene mutations

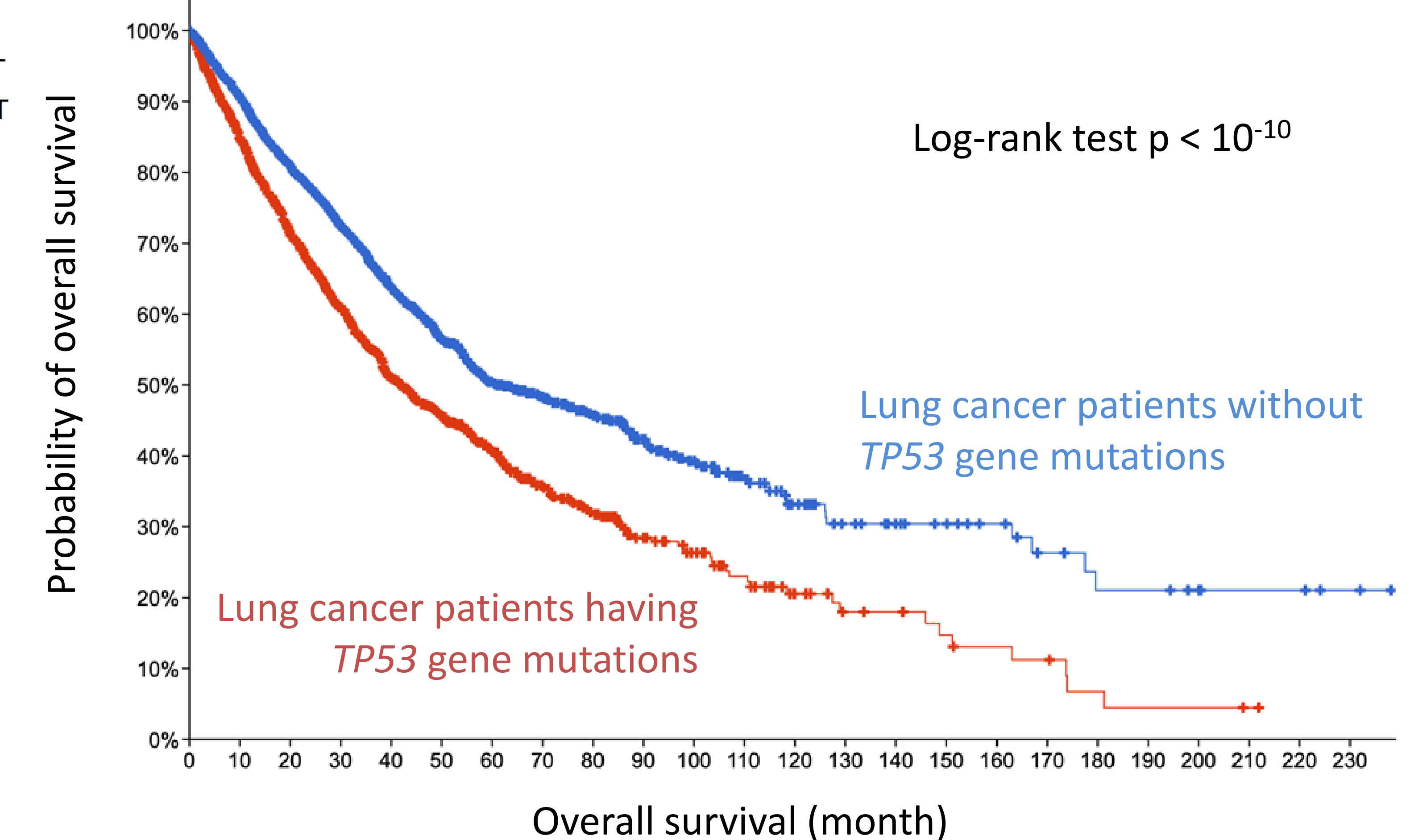


Fig.5: *TP53* cancer predisposition gene mutations are associated with significantly shortened overall survival in lung cancer patients

Conclusion

- This research underscores the role of CPGs in personalized healthcare for high-risk occupational groups.
- Individuals with specific CPG mutations linked to lung cancer metastasis may benefit from intensive screening and proactive interventions.
- Further investigation into the prevalence of these mutations in DoW/USAF/USSF flying forces is necessary.
- Future work includes refining our AI/ML analysis with data from these cohorts to enhance the precision of lung cancer prevention and treatment strategies in at-risk DoW personnel populations.

References

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