Carcinoma of unknown primary (CUP): The role of tumor genomic profiling

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Background

- CUP is a heterogenous group of cancers characterized by early metastatic dissemination from an unknown site of origin.¹
- Overall survival is a dismal 6-12 months and untreated CUP is associated with a 4 week life expectancy.^{2,3}
 - A 2014 review of the molecular profile of 1806 cases of CUP within the Caris Life Sciences database identified biomarkers with potential therapeutic benefits in over 96% of cases ⁴
- CUP continues to be a diagnostic and treatment challenge and comprehensive genomic profiling may provide therapeutic insight.

Study Methods

- Molecular profiles of tumors noted as 'unknown' for tumor primary site within the CARIS Life Sciences database were analyzed utilizing CODEai, a platform that integrates real-world clinical information obtained from insurance claims and medical records with genomic data.
- This real-world cohort consisted of 3,841 tumors
 - 2,137: Adenocarcinoma (ADC)
 - 385: Squamous cell carcinoma (SQ)
 - 1,319: Carcinoma not otherwise specified (NOS).
- CUP-ALL: CUP-ADC + CUP-SQ + CUP-NOS
- Overall survival (OS) was calculated from time of tissue collection to last contact assessed by Kaplan-Meier estimates.



Figure 1.

Within CUP-ALL, the targeted therapy cohort had a longer mOS of 638 days compared to 374 days in the chemotherapy cohort

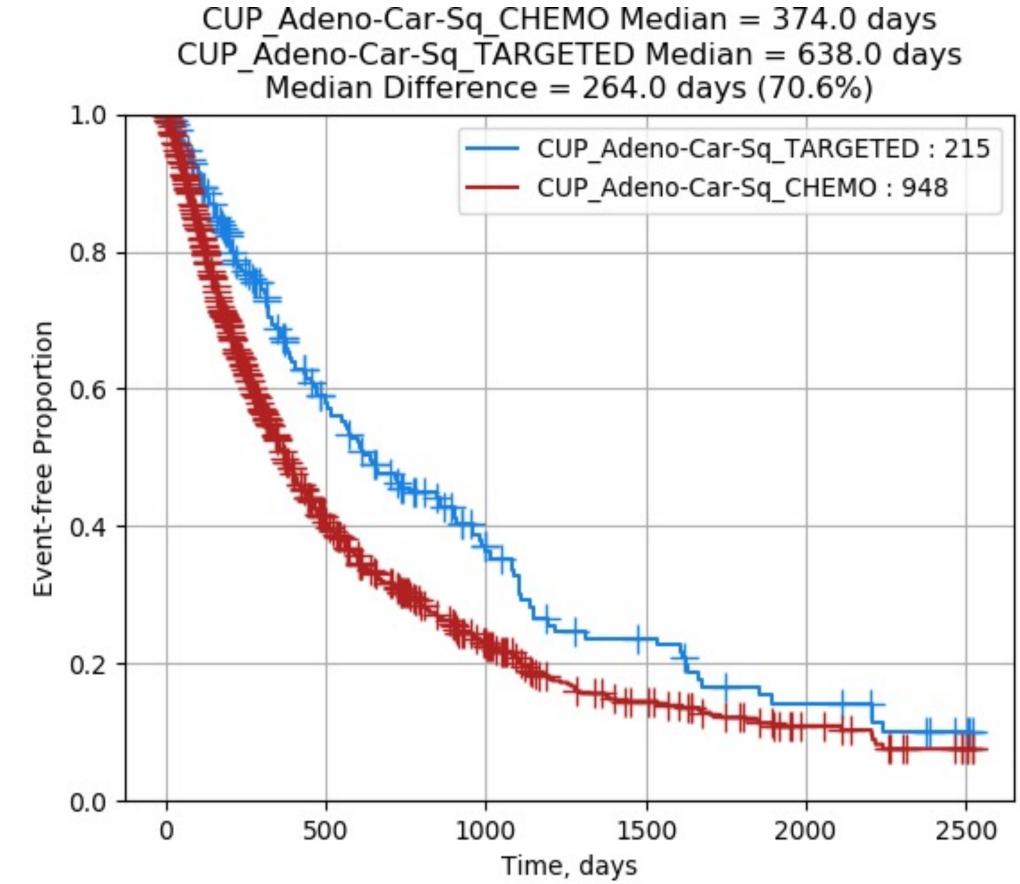
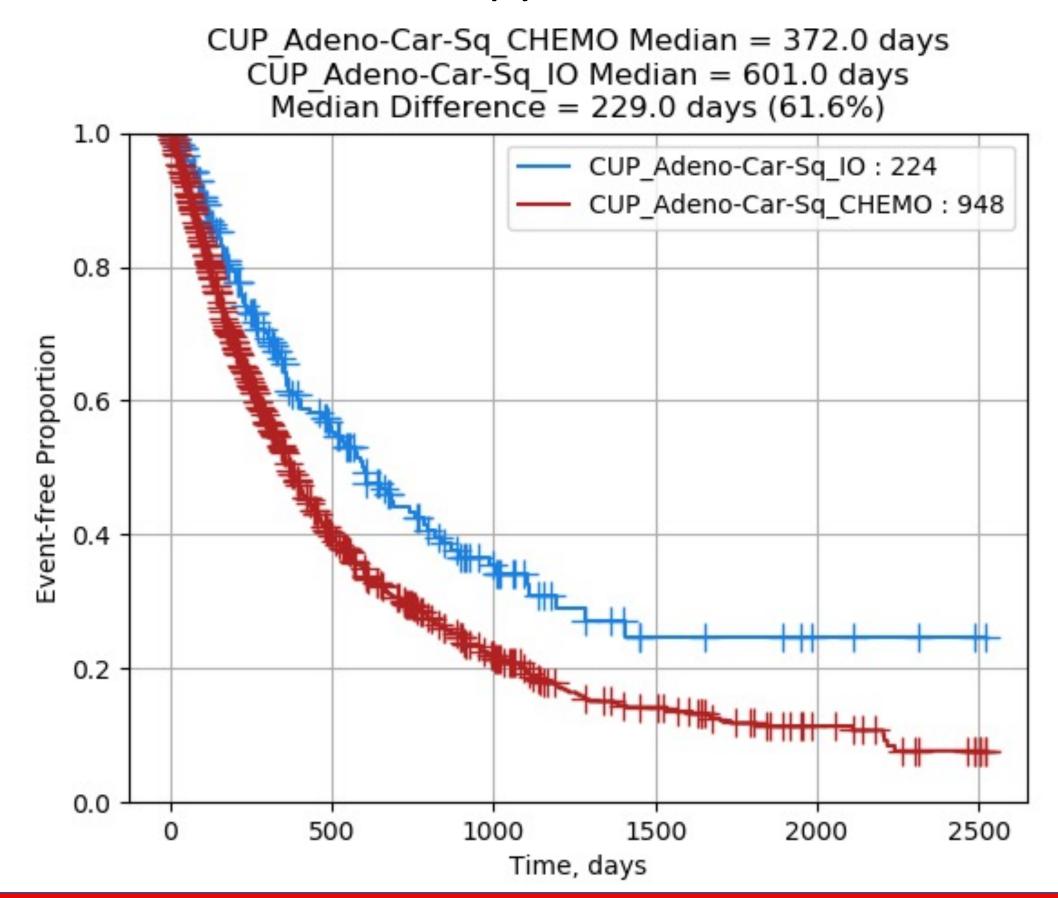


Figure 2.
Within CUP-ALL, the immunotherapy cohort had a longer mOS of 601 days compared to the chemotherapy cohort with a mOS of 372 days



Results

Figure 3.

In CUP-ADC, tumors with KRAS wild type had a longer mOS of 397 days compared to 202 days in tumors with a KRAS mutant variant

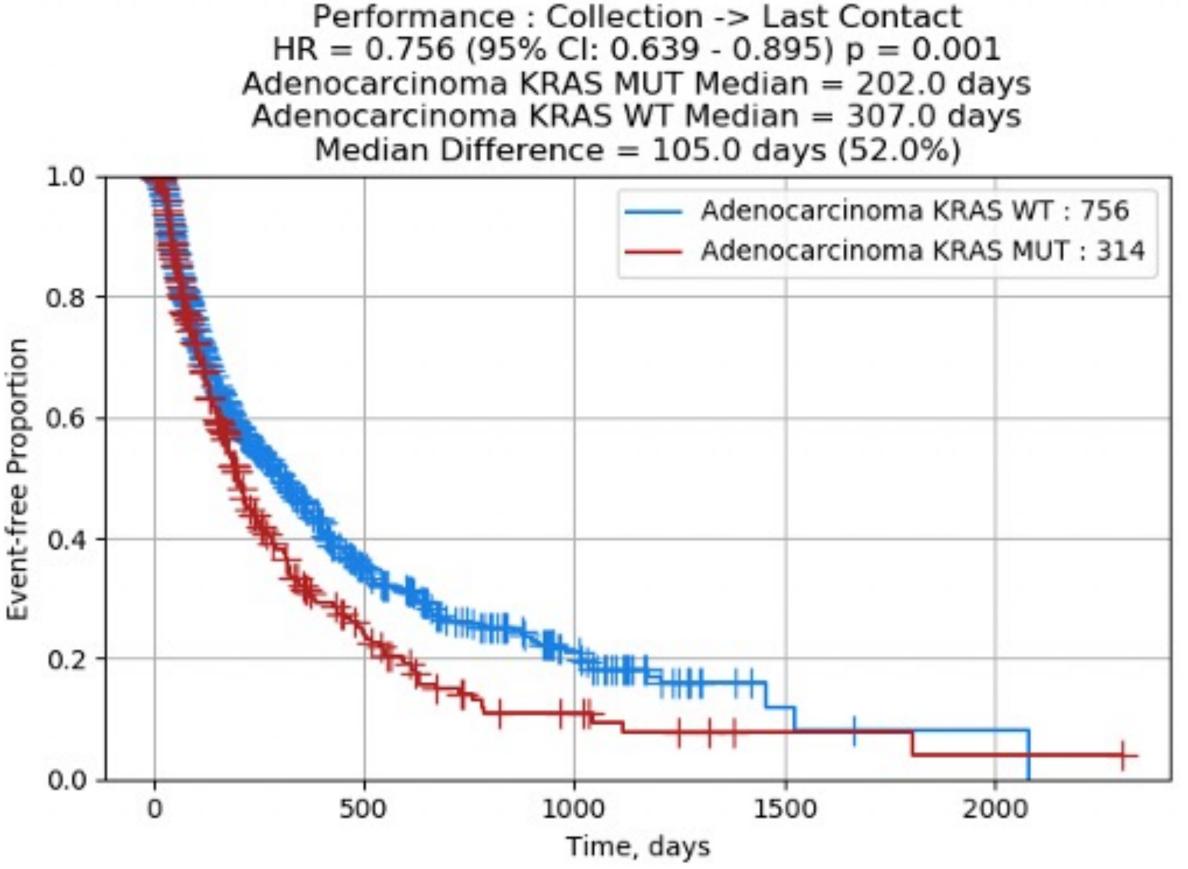
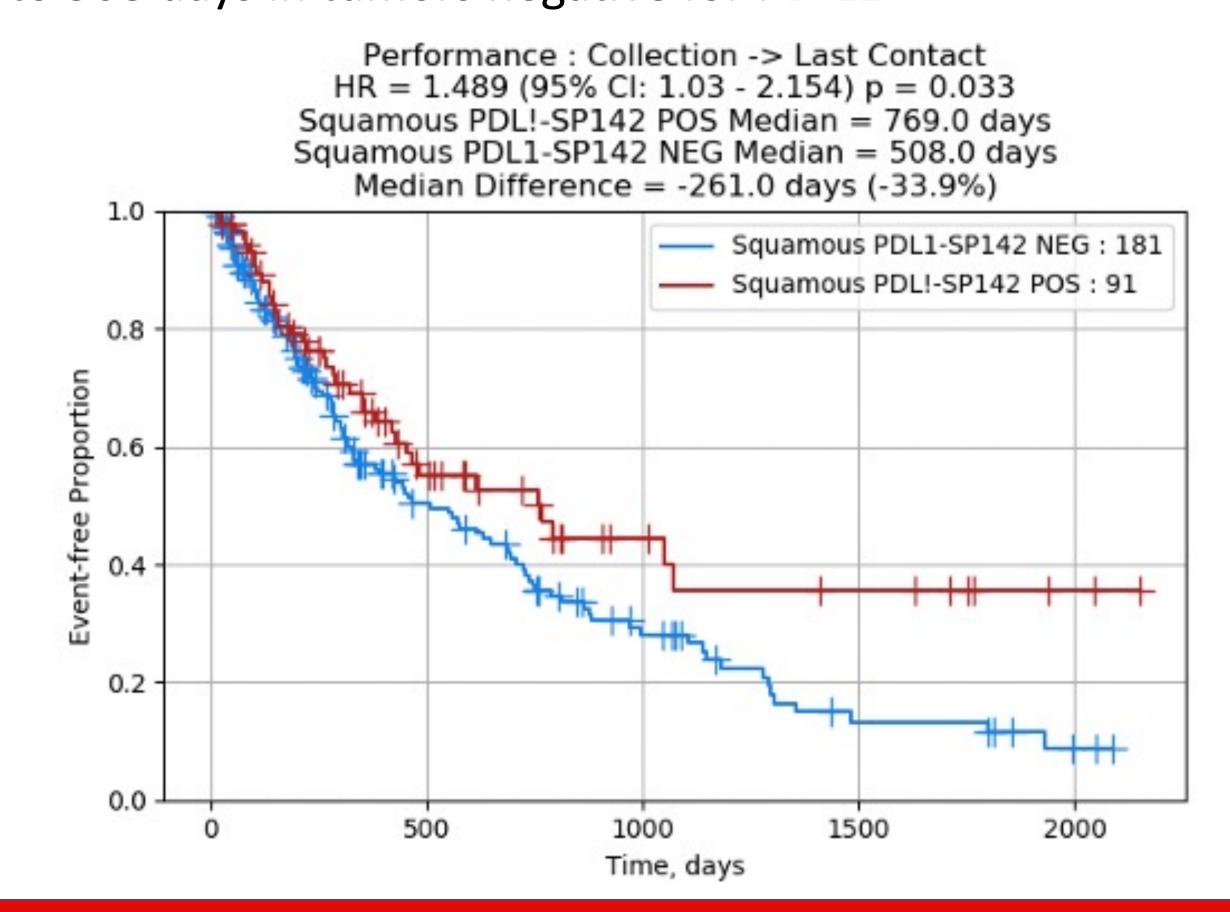


Figure 4.
In CUP-SQ, tumors positive for PD-L1 had a longer mOS of 769 days compared to 508 days in tumors negative for PD-L1



Conclusion

- The findings from this large real-world cohort demonstrate that key molecular alterations have prognostic and predictive roles in CUP.
- To maximize clinical benefit, prospective studies with various therapeutic classes of cancer treatments exploiting these differences are warranted.

References

- . Varadhachary GR, Lenzi R, Raber MN, Abbruzzese JL. Carcinoma of Unknown Primary. In: Abeloff's Clinical Oncology: Fifth Edition.; 2014. doi:10.1016/B978-1-4557-2865-7.00094-1
- 2. Massard C, Voigt JJ, Laplanche A, et al. Carcinoma of an unknown primary: Are EGF receptor, Her-2/neu, and c-Kit tyrosine kinases potential targets for therapy? Br J Cancer. Published online 2007. doi:10.1038/sj.bjc.6603942
- 3. Conway AM, Mitchell C, Kilgour E, Brady G, Dive C, Cook N. Molecular characterisation and liquid biomarkers in Carcinoma of Unknown Primary (CUP): taking the 'U' out of 'CUP.' Br J Cancer. Published online 2019. doi:10.1038/s41416-018-0332-2
 4. Gatalica Z, Millis SZ, Vranic S, et al. Comprehensive tumor profiling identifies numerous biomarkers of drug response in cancers of unknown primary site: Analysis of 1806 cases. Oncotarget. Published online 2014. doi:10.18632/oncotarget.2574