

Analysis and Prediction of Patient Survival After Radiotherapy For Liver Cancer **Based On Volumetric Segmental Response and Clinically Relevant Factors**

Aditya Prasad¹, Aashish C. Gupta¹, Mais M. Al Taie¹, Iwan Paolucci², Austin H. Castelo¹, Tien T. Tang¹, Eugene J. Koay³, Kristy K. Brock¹

¹Department of Imaging Physics, The University of Texas MD Anderson Cancer Center, Houston, TX, USA. ²Department of Interventional Radiology, The University of Texas MD Anderson Cancer Center, Houston, TX, USA. ³Department of GI Radiation Oncology, The University of Texas MD Anderson Cancer Center, Houston, TX, USA.

THE UNIVERSITY OF TEXAS MDAnderson **Cancer** Center

Making Cancer History[®]

Introduction

Primary liver cancer is one of the leading causes of cancer fatalities globally, with over 905,000 people diagnosed and over 830,000 deaths in 2020^1 . The three major types of cholangiocarcinoma (CC), colorectal metastasis and (CRM).

The liver's unique capability to regenerate functional tissue allows for significant recovery following loss of liver function due to cancer-associated damage. Namely, marked increases in normal tissue volume following external beam radiotherapy² (RT) and contralateral hypertrophy after radioembolization³ have been documented. However, **there** is limited literature documenting liver segment-specific hypertrophy due to RT and its hypothesized connection with liver cancer patient survival.

Results: Survival Analysis



Key Points: Figure 2 and Table 1

1. Liver cancer patients in the overall patient cohort with hypertrophy in segments 2 and 3 have a 56.67% greater probability of surviving relative to those who do not at any given time point.

2. Patients in all cohorts with PVT, with cirrhosis in the overall cohort, and with PHT in the HCC patient cohort, respectively, are approximately 2 times as likely to die at any given time point [compared to the

negative group].

Results: Predictive Modeling



AUPRC

0.71

0.75

0.75

0.71

0.61





. Liver hypertrophy in segments 2 and 3 significantly correlates with better survival, with PVT, PHT, and cirrhosis significantly associated with greater mortality

in certain liver cancer patient cohorts.

2. We developed binary risk prediction ML models to predict patient survival 15 months following the end of

Created with Biorender.com

Sensitivity, Specificity, and **ROC and PR curves.**

Fig 1. Flowchart of study, including clinical analysis, predictive modeling, and model validation.

comparatively higher probability of death in the positive group relative to the negative group (or between comparative subgroups) at any given time point and HR lesser than 1 indicates comparatively lesser probability of death [at any given time point]. Log-rank test results ($\alpha =$ 0.05) indicate whether there is a significant difference in the effect on patient survival between positive and negative groups, or subgroups, for each predictor.

RT treatment. In the future, we aim to expand the prediction model to predict survival as a continuous outcome, in days survived after the end of radiotherapy. 3. All model accuracies, AUROC, and sensitivities were equal to or above 0.72, 0.79, and 0.90, respectively, demonstrating strong performance. Further steps include to integrate additional clinical and radiomic factors and increase the number of patients included in the survival analysis and predictive modeling.

4. Lasso regression seems to be the most robust model due its relatively greater AUROC and comparatively similar accuracy, specificity, and sensitivity, although Random Forest and Support Vector Machine also exhibit relatively high predictive power.

References

. Rumgay, Harriet, et al. Journal of Hepatology . 2022.https://doi.org/10.1016/j.jhep.2022.08.021 . Su, Ting-Shi, et al. Frontiers in Oncology. 2021. 10.3389/fonc.2021.680303 Vouche, Michael, et al. Journal of Hepatology. 2013. https://doi.org/10.1016%2Fj.jhep.2013.06.015

Acknowledgements

thank the members of Morfeus lab, collaborators, and medical illustrator Kelly Kage for their support and guidance. Research reported was supported in part by the National Cancer Institute of the National Institutes of Health under award number R01CA221971. This work was also supported by the Image Guided Cancer Therapy Research Program at The University of Texas MD Anderson Cancer Center through a generous gift from the Apache Corporation, by the Helen Black Image Guided Fund, and by the Tumor Measurement Initiative through the MD Anderson Strategic Initiative Development Program (STRIDE). AP was supported by the CPRIT Research Training Award CPRIT Training Program (RP210028).