DEEP TREATMENT RESPONSE ASSESSMENT AND PREDICTION OF COLORECTAL CANCER LIVER METASTASES

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Introduction

Colorectal cancer:

- second leading cause of cancer death

- \approx half of patients develop a distant recurrence

Liver, through the development of **liver metastases**, is the most common spread site. When surgery is not an option due to the tumor burden, **palliative** oncological treatments are administrated.

Assessing treatment response involves CT image analysis and RECIST1.1 criterion [1]. These tasks are largely performed manually by clinicians, hence time-consuming and prone to expert variability.

Network architecture

- **Siamese network** built upon a modified 3D ResNeXt-26 architecture. Each **residual** block is fitted with a 3D concurrent spatial and channel **squeeze-and-excitation** layer [3].
- Introduced long skip-connections that transfer features arising from different network depths to the very end of the pipeline by aggregating them with the final output of the ResNeXt blocks.



- To predict the early treatment response (TRP), we adopted the TRA pipeline by modifying it to a single 3D ResNeXt branch instead of a Siamese style architecture.
- [1] E. A Eisenhauer et al. New response evaluation criteria in solid tumours: revised RECIST guideline (version 1.1). European Journal of Cancer, 2009.
- [2] T. Aparicio et al. Bevacizumab+chemotherapy versus chemotherapy alone in elderly patients with untreated metastatic colorectal cancer: a randomized phase II trial—PRODIGE 20 study results. Annals of Oncology, 2018.
- [3] A. G Roy et al. Concurrent spatial and channel squeeze & excitation in fully convolutional networks. In International Conference on Medical Image Computing and Computer-Assisted Intervention, 2018.

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Methodology

We propose a deep learning based treatment response assessment pipeline and its extension for prediction purposes based on a newly designed 3D Siamese classification network.

Treatment response assessment (TRA) exploits pair of volumetric CT scans from consecutive pretreatment and post-treatment examinations, acquired at times t and t + 1.

Treatment response prediction (TRP) at time t + 1 given only the CT scan acquired at time t.

A liver metastasis segmentation task has also been tackled as it helps improving both TRA and TRP tasks.

Results

TRA/TRP pipelines were evaluated with overall accuracy (acc), sensitivity (sens), specificity (spec), F1 score (F1) and area under the curve (AUC). An **ablation study** was performed to compare the TRA pipeline with 3 striped versions: Model A (w/o GRUbased feature fusion), Model B (w/o long skipconnections and transition blocks) and Model C (w/o segmentation masks as inputs).

TRA TRP Mode Mode

Our pipeline for TRA achieved 94.94% overall accuracy with high F1-score and TRP pipeline achieved 86.86%. **GradCAM** was used to see what the model is focusing on to make the decision.





	acc	sens	spec	F1	AUC
TRA	94.94	94.89	97.22	94.72	95.56
TRP	86.86	83.22	94.17	83.02	89.25
Model A	92.42	93.17	97.22	92.30	94.17
Model B	89.39	86.89	96.46	82.23	90.39
Model C	87.87	89.17	95.51	84.66	92.34



Data

PRODIGE20 dataset [2] collected from a phase-II multi-center clinical trial in colorectal cancer with liver metastasis that evaluated chemotherapy alone or combined with Bevacizumab during follow-up.



- A total of 400 consecutive CT scan pairs from 102 patients were considered, divided into 4 re**sponse groups**: Complete Response (CR), Partial Response (PR), Progression (P) and Stable (S) in accordance with RECIST1.1 [1].

Conclusion

Both treatment response assessment and prediction issues can be effectively solved with **deep learning**.

▷ Our contributions **alleviate** manual liver metastasis segmentation and RECIST1.1 evaluation **burdens**.

▷ A future possible extension is to build a **complete** chemotherapy regimen recommendation system able to predict the best treatment for each patient and which would include other organ assessment such as lungs, lymph nodes, peritoneal cavity and bones.



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