

# #423650: Association of <sup>18</sup>F-FDG PET characteristics and survival outcomes using whole body tumor analysis in patients (pts) with metastatic genitourinary (GU) malignancies

Nicholas I. Simon, Rod Carlo Columbres, Elias Chandran, Scot Anthony Niglio, Lisa M. Cordes, Lisa Ley, Tzu-fang Wang, Amir Mortazavi, Sumanta Kumar Pal, Rajkumar Munian-Govindan, Timothy G. Perk, Esther Mena Gonzalez, Liza Lindenberg, Andrea B. Apolo

## Background:

- <sup>18</sup>F-FDG PET scans are widely used for staging and monitoring pts with metastatic GU malignancies
- <sup>18</sup>F-FDG PET scans provide semi-quantitative parameters used to characterize disease burden and treatment response across different patients as well as across different lesions within the same patient

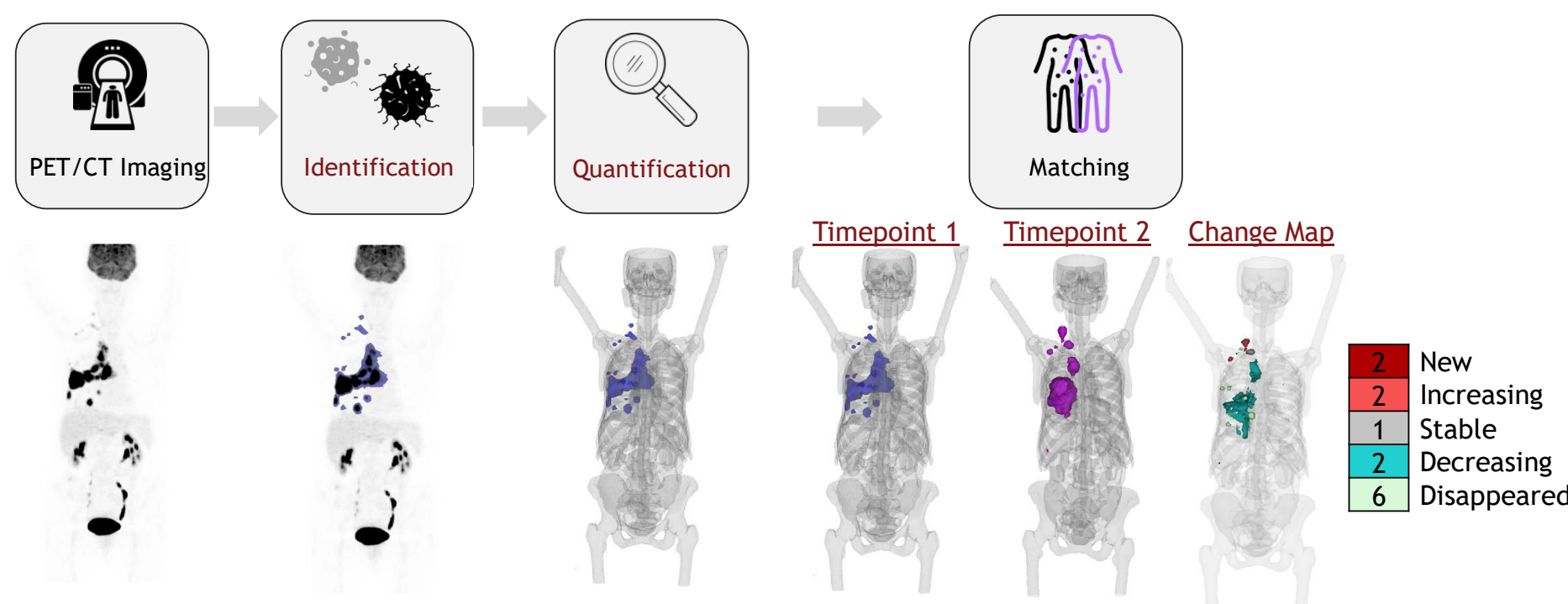
## Aim:

In this study, we aim to determine which <sup>18</sup>F-FDG PET parameter provides the best prognostic information for patients with metastatic GU malignancies

## Methods:

- 101 pts with metastatic GU malignancies from two separate, prospective trials with at least two <sup>18</sup>F-FDG PET scans were included in the study
- All regions of interest (ROI) were contoured for both the baseline and follow-up <sup>18</sup>F-FDG PET scans.
- ROI were quantified and matched between timepoints using TRAQinform IQ technology (AIQ Solutions, Madison, Wisconsin)

### How TRAQinform IQ™ Works



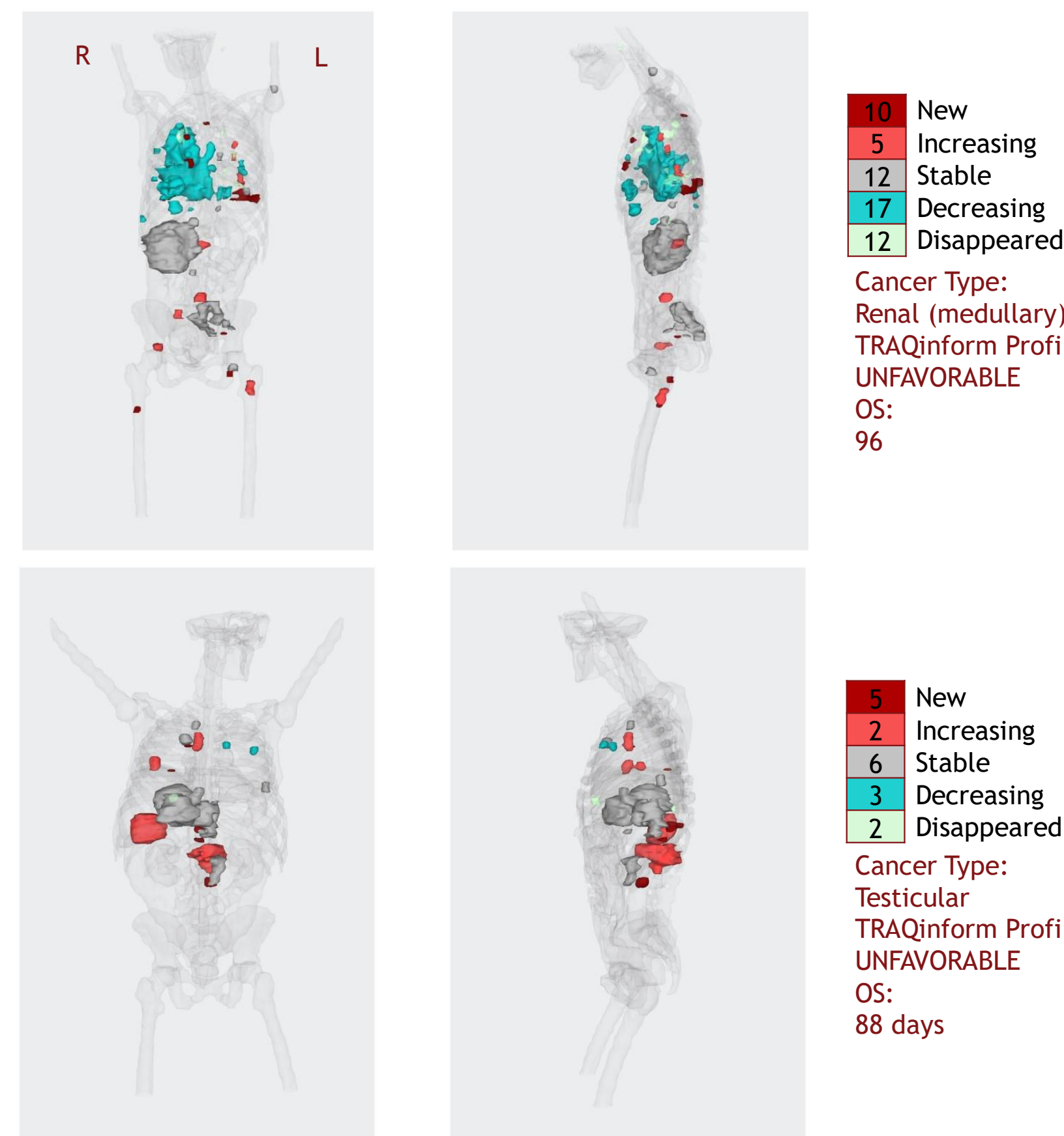
- Imaging features were extracted from each patient, including basic features (SUVmax, SUVmean, TLG, and changes in these features) and heterogeneity (inpatient heterogeneity of disease and response) features.
- Cox regression was used to determine univariate predictive power of each measure.
- The TRAQinform Profile was calculated to predict either progression-free (PFS) or overall survival (OS) using 5-fold cross-validation of a random survival forest.
- The performance of individual features and the TRAQinform Profile was evaluated using the c-index.

## Results:

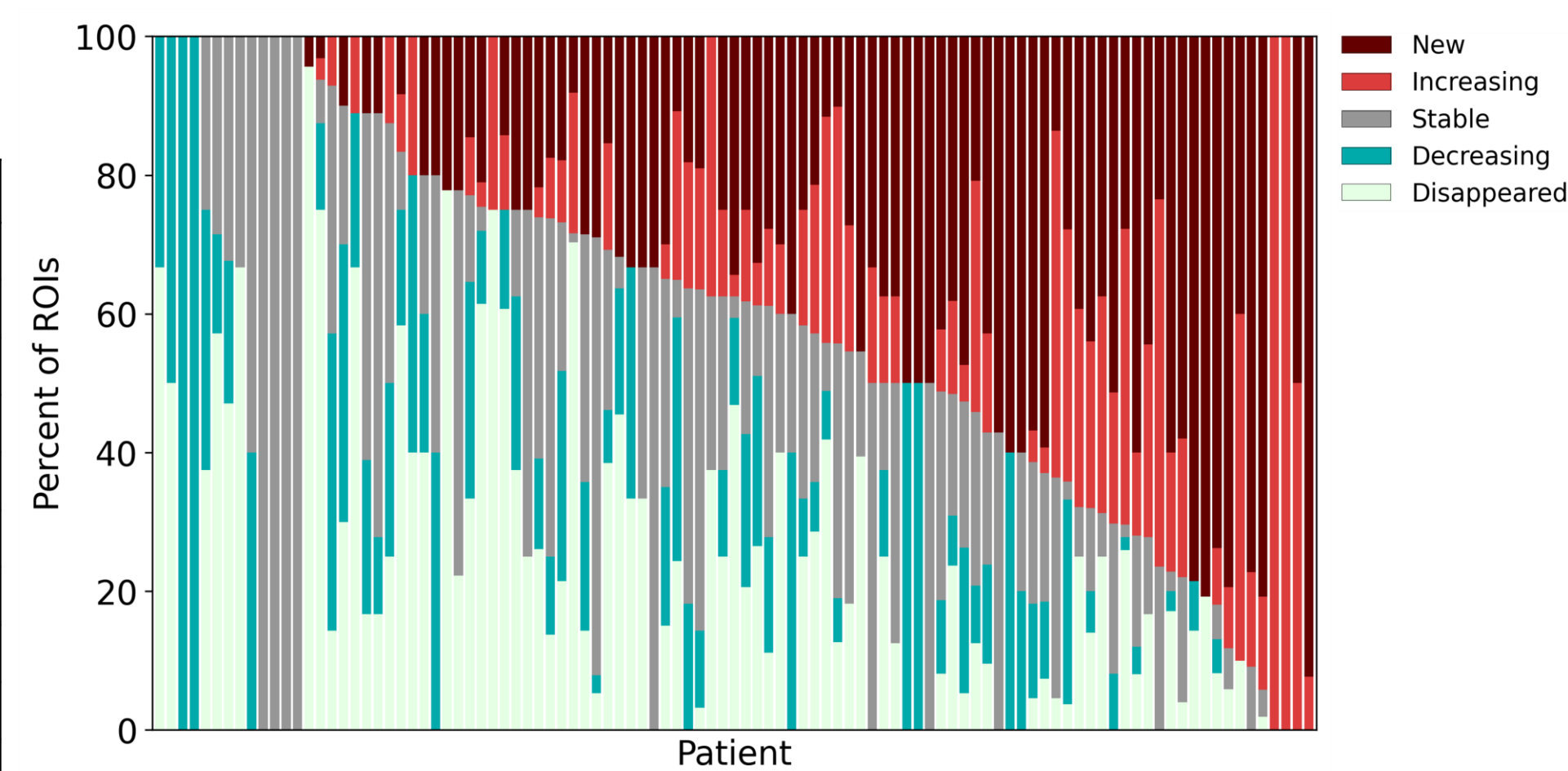
**Table 1.** Treatment and histology characteristics of patients analyzed in the study.

Characteristics	N (%)
<b>Treatment</b>	
Cabo only	54 (53.5)
Cabo, Nivo +/- Ipi	47 (46.5)
<b>Histology</b>	
Urothelial carcinoma	54 (53.5)
Bladder adenocarcinoma	8 (7.9)
Renal cell carcinoma (RCC)	6 (5.9)
Testicular	4 (4.0)
Renal medullary carcinoma	4 (4.0)
Bladder small cell carcinoma	4 (4.0)
Squamous cell carcinoma of the bladder	2 (2.0)
Penile	1 (1.0)
Sarcomatoid bladder	1 (1.0)
Sarcomatoid RCC	1 (1.0)
Prostate small cell carcinoma	1 (1.0)

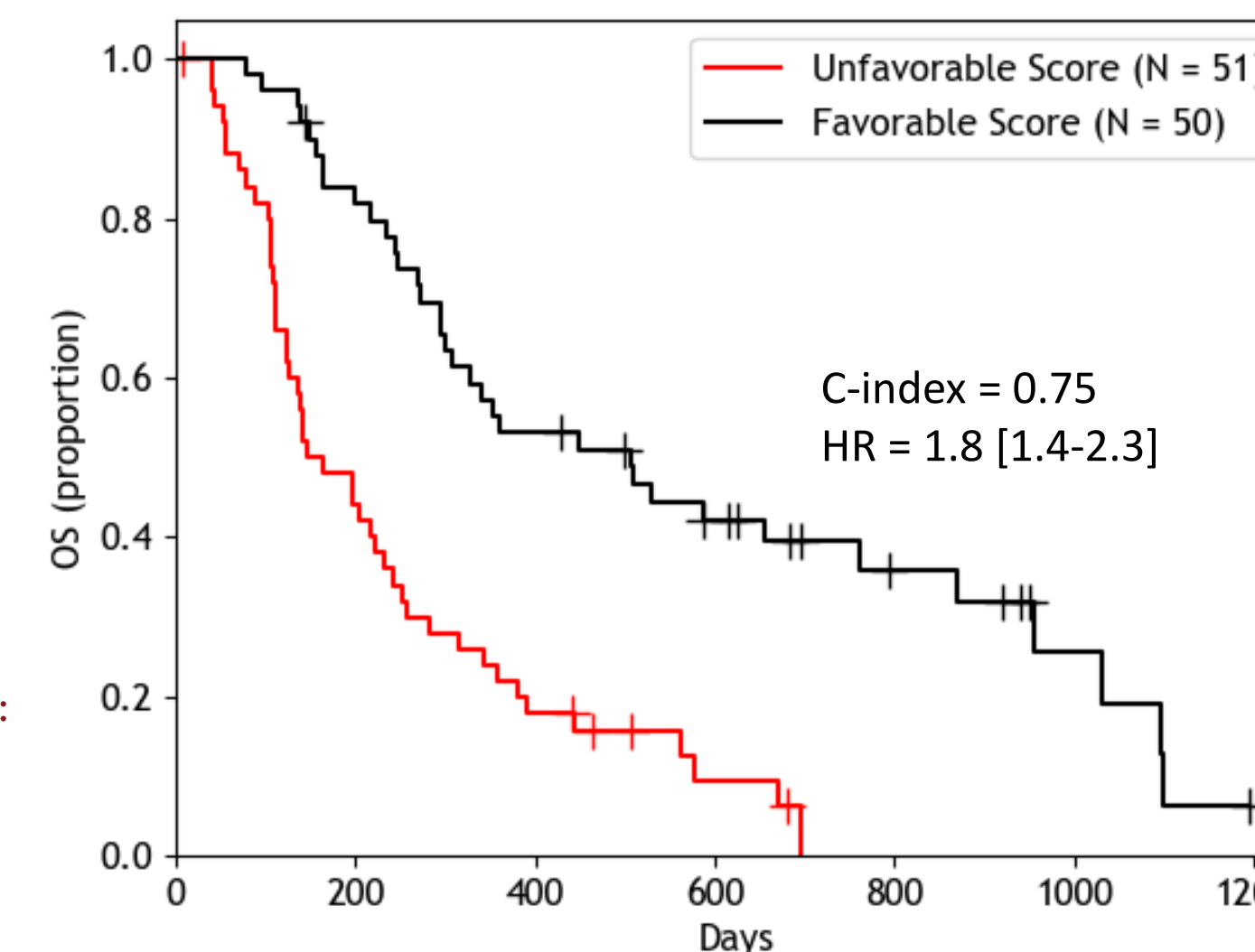
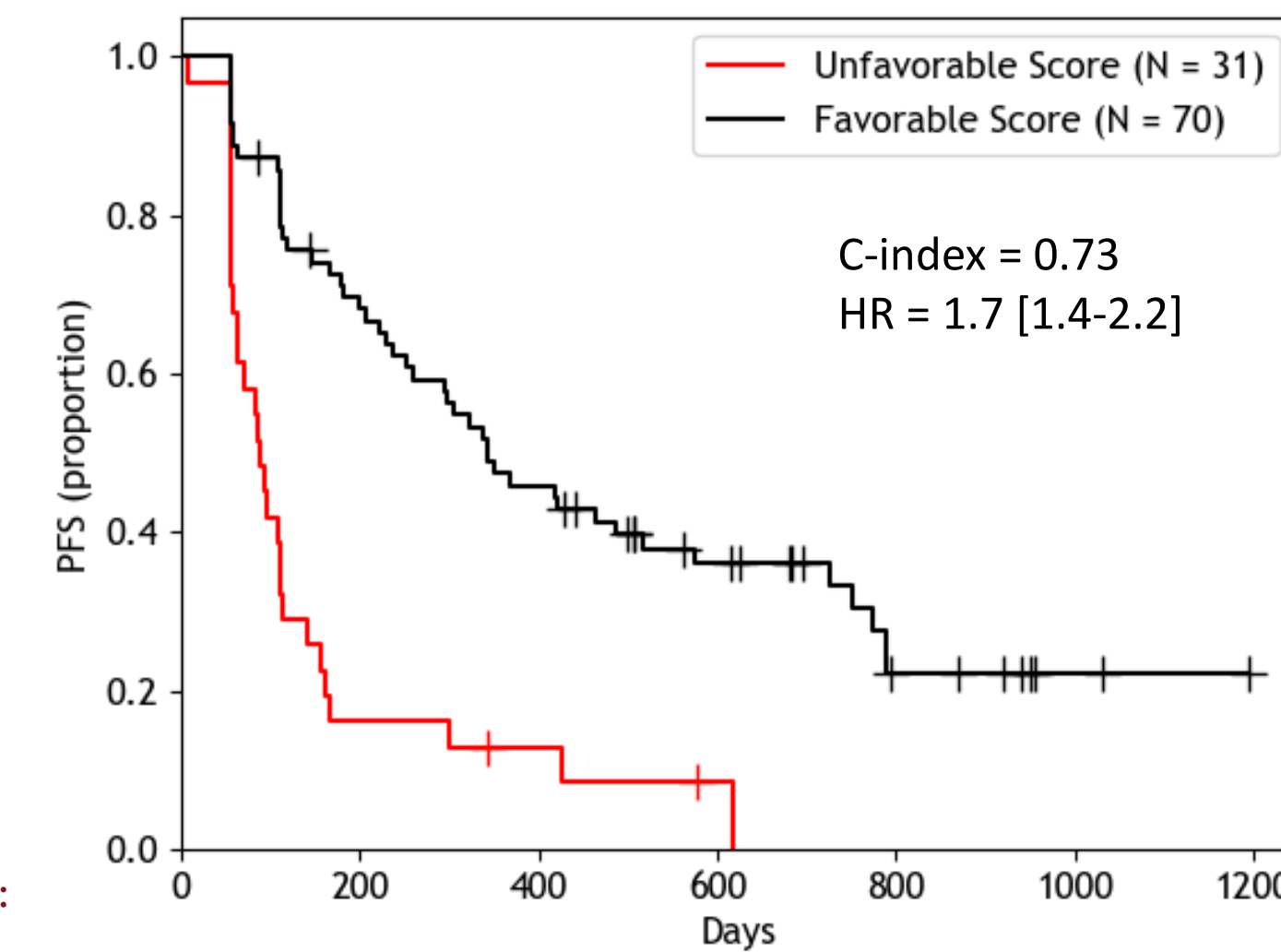
**Figure 1.** TRAQinform IQ generated map/images from two patients with metastatic GU enrolled in phase I or phase II study. Legends indicate the number of ROI that disappeared or are decreasing, stable, increasing or new



**Figure 2.** Heterogeneity plot of all mGU patients



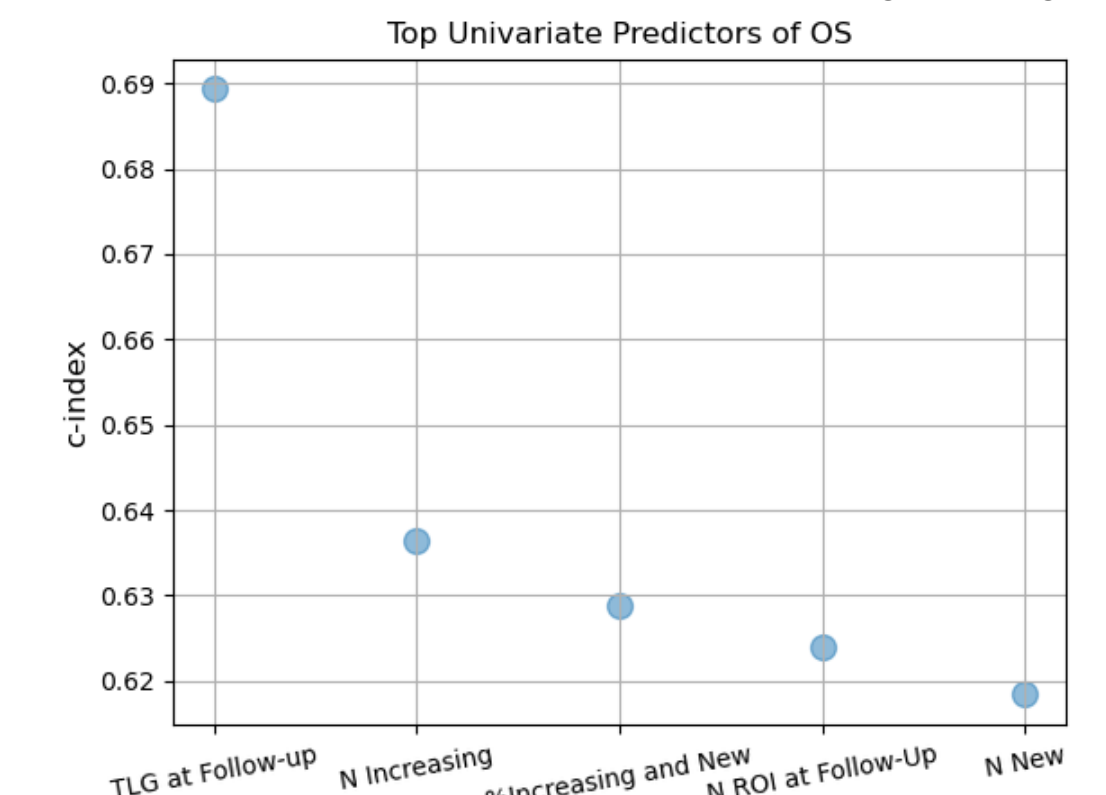
**Figure 4.** TRAQinform profile. Analysis performed with 5-fold cross validation



## Results cont.:

- 54 pts treated with Cabozantinib (Cabo) on phase II trial
- 47 pts treated with Cabo, Nivolumab (Nivo) +/- Ipilimumab (Ipi) on phase I trial
- The TLG at the follow-up image was the strongest predictor of both PFS (c-index = 0.62) and OS (0.69), followed by number of increasing ROI (0.61, 0.64), and percentage of all ROIs classified as new or increasing (0.57, 0.63).
- SUVmax (0.49, 0.47) and SUVmean (0.47, 0.56) at follow-up were found to be weaker predictors of PFS and OS.
- TRAQinform Profile was able to predict the responder's vs suboptimal responders to study treatment (c-index = 0.73, 0.75)

**Figure 3.** Univariate survival analysis plot of the top features for predicting OS. C-index was determined for each measure using Cox regression.



## Conclusion:

The TRAQinform Profile analysis of <sup>18</sup>F-FDG PET scans provided both prognostic and predictive information for patients with metastatic GU malignancies treated with either Cabo or CaboNivo +/- Ipi.

## Future Directions:

- Plan to further analyze data to determine correlation between site of disease response/progression and prognosis as well as examine the correlation between total body heterogeneity of response across all lesions and RECIST measurements