Exploring the Generalizability of the Prognostic Value of Machine Learning Models Trained on Mid-Treatment PSMA PET/CT to End-of-Treatment PSMA PET/CT in mCRPC Patients Treated with ¹⁷⁷Lu-PSMA Radioligand Therapy Using the TRAQinform Profile: A **Retrospective, Single-Center Analysis**



Vishnu Murthy¹, Ojaswita Lokre², Timothy Perk², Pan Thin¹, Kathleen Nguyen¹, Lucia Chen³, Andrei Gafita¹, Johannes Czernin¹, Jeremie Calais¹ ¹ Ahmanson Translational Theranostics Division, David Geffen School of Medicine at UCLA, Los Angeles, CA, USA ²AIQ Solutions, Madison, WI, USA, ³Department of Medicine Statistics Core, David Geffen School of Medicine at UCLA, Los Angeles, CA, USA

OBJECTIVE

To explore the prognostic value of the TRAQinform Profile trained on mid-treatment prostate-specific membrane antigen (PSMA) PET/CT (PSMA-PET) patients when applied to end-of-treatment PSMA-PET (ePET) in patients with metastatic castration-resistant prostate cancer (mCRPC) treated with ¹⁷⁷Lu-PSMA Radioligand Therapy (PSMA-RLT)

METHODS

i) Patients

- mCRPC patients who underwent PSMA-RLT with available baseline PSMA-PET (bPET) and ePET within 6 mo of the last PSMA-RLT cycle were included in this retrospective analysis
- Overall survival (OS) from ePET was collected

ii) Analysis of PSMA PET Images

- TRAQinform IQ technology (AIQ Solutions-Madison, WI) was used to conduct lesion region of interest (ROI)-based analyses at bPET and ePET, and changes in SUV_{max} , SUV_{mean} , volume (cm³), and SUV_{total} (SUV_{mean} x volume) across all lesion ROI from bPET to ePET were extracted
- ROI were matched across time points and categorized into new, increasing, stable, decreasing, and disappeared based on changes in SUV_{total} (Figure 1)

iii) Statistical Analysis

- This study applied TRAQinform Profile (a random forest model), trained on 185 external PSMA-PET, to generate TRAQinform Profile scores for early identification of optimal vs. suboptimal responders to PSMA-RLT
- Associations between TRAQinform Profile scores and OS were evaluated by Kaplan-Meier analysis
- Associations between % new, % increasing, % stable, % decreasing, and % disappeared lesion and OS were evaluated using univariate Cox regression models







Patients

- Twenty mCRPC patients were included
- Twelve of 20 patients (60%) had died at the last follow-up, the median follow-up time from ePET for survivors was 31.2 mo (IQR, 6.8–40.7 mo), and the median survival time from ePET was 13.3 mo (IQR, 10.2– NR)

RESULTS

Image Analysis

 The median number of lesion ROI identified on bPET and ePET were 13.5 (IQR: 5.8-59.5) and 33 (IQR: 2.8-86.3) respectively • The median % changes in SUV_{max}, SUV_{mean}, volume, and SUV_{total} from bPET to ePET were -33.9%, -20.7%, 4.3%, and -13.9% respectively

Statistical Analysis

- An increase in percent new and percent new or increasing lesions was associated with a higher risk of death (HR = 1.03; p = 0.004, HR = 1.03; p = 0.01 respectively), while an increase in percent disappeared or decreasing lesions was associated with a lower risk of death (HR = 0.97; p = 0.024)
- The median TRAQinform Profile score was 0.27, and patients with a TRAQinform Profile score above the median had shorter OS compared with patients with a TRAQinform Profile score below the median (median OS, 10.9 mo [95% CI, 5.0-NR] vs. median OS, 22.0 mo [95% CI, 14.5-NR]; P = 0.049) (Figure 2). Figure 3 shows sample cases from our cohort.

CONCLUSION

- In this retrospective study of 20 mCRPC patients treated with PSMA-RLT, the TRAQinform Profile score was prognostic for OS Percent new, percent new or increasing, and percent
- disappeared or decreasing lesions were significantly associated with OS
- Validation in larger, prospective multicentric clinical trials is warranted