

PCPro as a prognostic plasma lipidomic biomarker in TheraP (ANZUP 1603): a randomised trial of [177Lu]Lu-PSMA-617 (LuPSMA) vs cabazitaxel in metastatic castration resistant prostate cancer (mCRPC)

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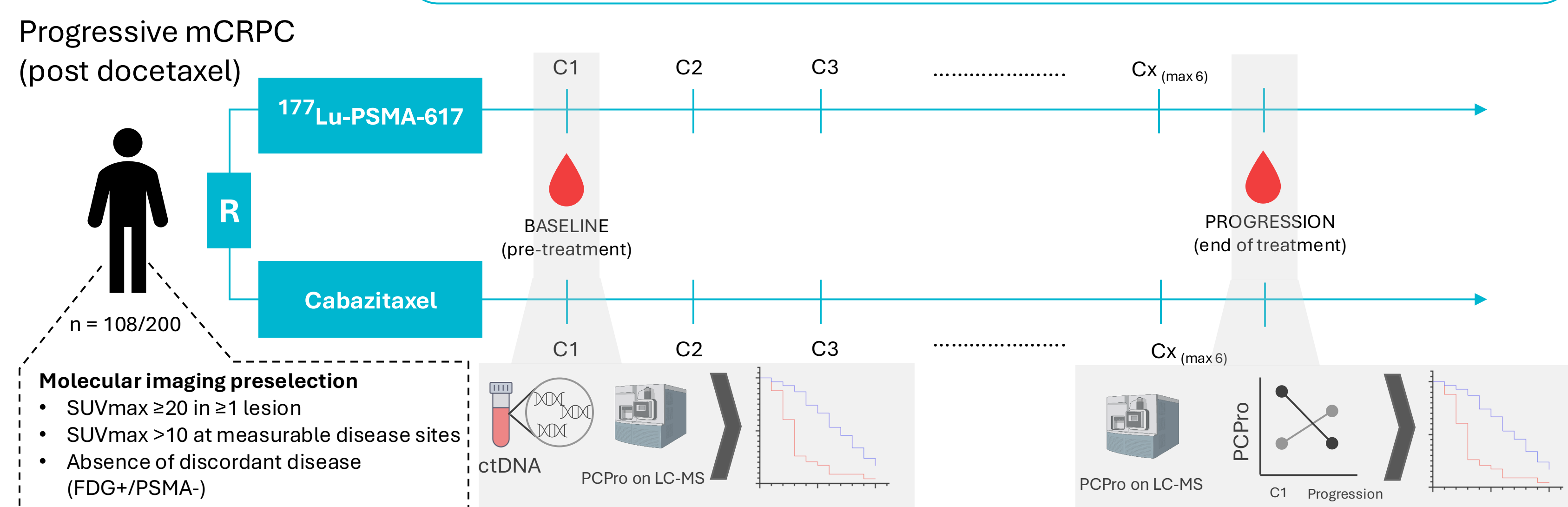
Background

- Lutetium-177[¹⁷⁷Lu]Lu-PSMA-617 is a recently established standard-of-care therapy in patients with metastatic castration-resistant prostate cancer (mCRPC).
- The TheraP trial showed that LuPSMA improves PSA response rate (RR), objective tumour RR and radiographic progression free survival (rPFS), compared with cabazitaxel in people with PSMA-positive, non-FDG-discordant mCRPC progressing after docetaxel.^{1,2}
- Elevated circulating sphingolipids, including ceramides, are associated with shorter PFS and OS in mCRPC treated with docetaxel or ARPIs.³
- PCPro is a validated, plasma lipid biomarker, developed in accordance with CLIA/NATA guidelines, comprising Ceramides: Cer(d18:1/18:0), Cer(d18:1/24:0), Cer(d18:1/24:1), total cholesterol and triglycerides.⁴
- PCPro positive patients with mCRPC have shorter rPFS and overall survival (OS) when treated with ARPIs and shorter OS when treated with docetaxel.
- Quantitative PET/CT and ctDNA biomarkers have demonstrated predictive and prognostic capability.^{5,6}
- This is the first report of the association of PCPro status with clinical outcomes in people treated with LuPSMA or cabazitaxel.

Study Design

Study Aims

- Describe the association between PCPro with clinical outcomes in participants treated with LuPSMA or cabazitaxel in the TheraP study
- Evaluate the association of PCPro with established molecular imaging prognostic thresholds and ctDNA% categories



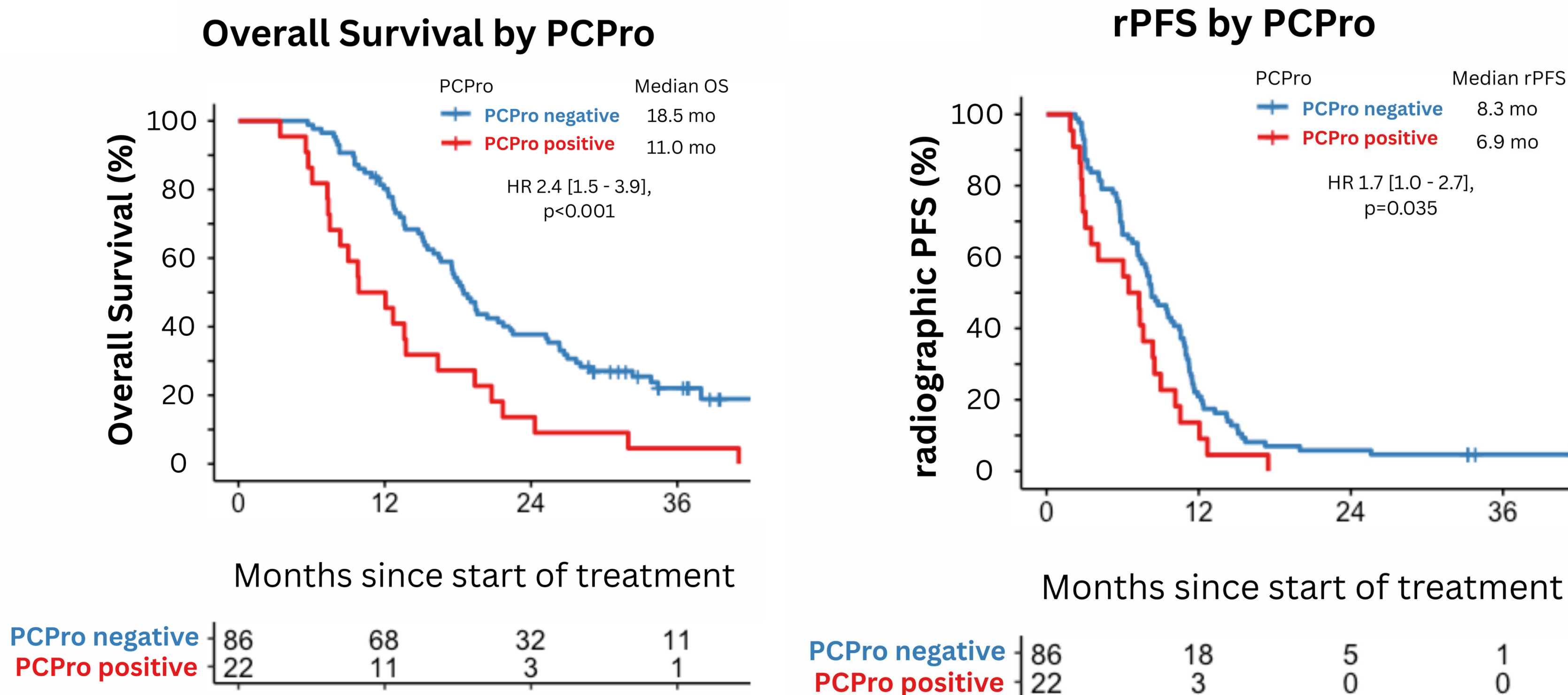
Patient Characteristics

	LuPSMA (biomarker cohort) (n=60)	Cabazitaxel (biomarker cohort) (n=49)	LuPSMA (whole cohort) (n=99)	Cabazitaxel (whole cohort) (n=101)
Age, median (IQR)	72 (67 – 76)	72 (68 – 76)	72 (67 – 77)	72 (67 – 77)
> 20 metastases*, n (%)	45 (75)	38 (78)	77 (78)	79 (78)
ECOG, N (%)				
0	30 (50)	29 (59)	42 (42)	44 (44)
1-2	30 (50)	20 (42)	57 (58)	56 (55)
Unknown	0	0	0	1 (1)
PSA (ng/mL), median (IQR)	73.6 (38.5-128.15)	95.3 (34.7-230)	93.5 (44-219)	110 (64-245)
Alkaline phosphatase (U/L), median (IQR)	112 (88-174)	133 (83-190)	111 (83-199)	130 (79-187)
Disease location, n (%)				
Lymph node only	6 (10)	5 (10)	7 (7)	9 (9)
Bone metastases	52 (87)	43 (88)	90 (91)	90 (89)
Visceral metastases	4 (7)	4 (8)	7 (7)	13 (13)
Previous ARPI, n (%)				
Abiraterone only	15 (25)	13 (27)	21 (21)	24 (24)
Enzalutamide only	12 (20)	4 (8)	49 (50)	58 (57)
Both	29 (48)	27 (55)	21 (21)	9 (9)
PSMA SUVmean, median (IQR)	9.0 (7.1-11.7)	8.5 (6.9-10.2)	8.5 (7.1-11.5)	8.5 (6.8-10.5)
FDG-PET MTV (mL), median (IQR)	101.5 (29.8-266.3)	66.6 (14.3-224)	100 (23-252.5)	78 (22-248)
PSMA SUVmean ≥10, n (%)	22 (37)	15 (31)	37 (37)	33 (33)
FDG MTV ≥200mL, n (%)	19 (32)	13 (27)	31 (31)	30 (30)
Median rPFS, months (95% CI)	8.43 (7.29 – 9.53)	7.89 (7.36 – 8.57)	8.51 (6.47 – 10.97)	7.95 (7.20 – 9.13)
HR rPFS (95% CI), p value	0.78 (0.59 – 1.03), p=0.075		0.68 (0.50 – 0.93), p=0.015	
Median OS, months (95% CI)	16.4 (13.7 – 21.2)	18.4 (15.5 – 24.3)	16.36 (13.73 – 19.58)	19.42 (15.51 – 23.06)
HR OS (95% CI), p value	0.98 (0.64 – 1.49), p=0.9		0.98 (0.71 – 1.36), p=0.9	

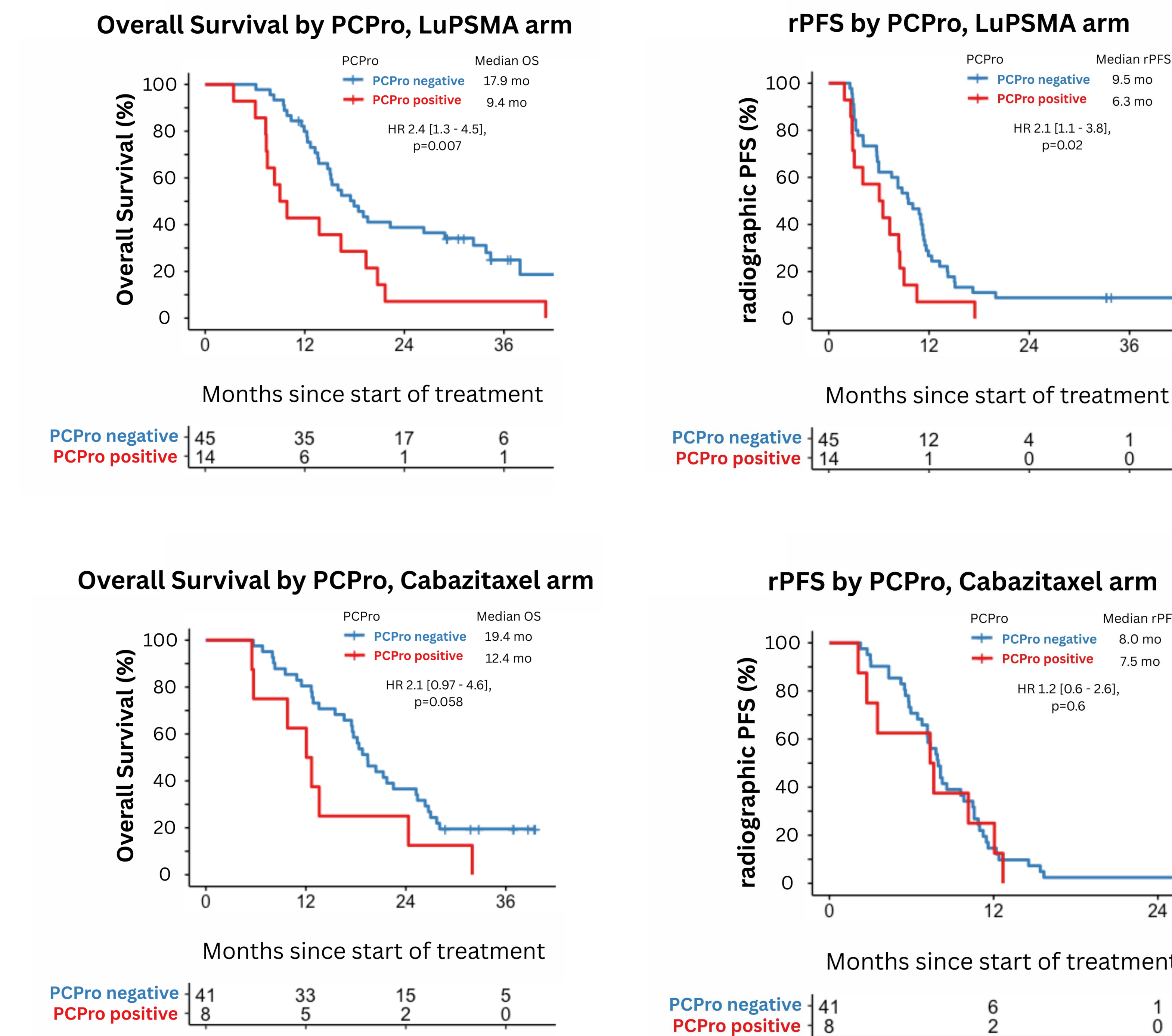
Table 1: Patient characteristics of biomarker and trial cohorts

* Disease burden as assessed by [¹⁸G]Ga-PSMA-11

PCPro is prognostic for OS and rPFS



PCPro is prognostic independent of treatment arm



Variable	Hazard Ratio OS [95% CI], p value	Interaction variable p value OS	Hazard Ratio rPFS [95% CI], p value	Interaction variable p value, rPFS
PCPro, positive	2.40 [1.47-3.92], p<0.001	0.8	1.67 [1.04 – 2.68], p=0.034	0.2
Treatment arm, LuPSMA	0.92 [0.60 – 1.41], p=0.8		0.78 [0.53 – 1.15], p=0.2	

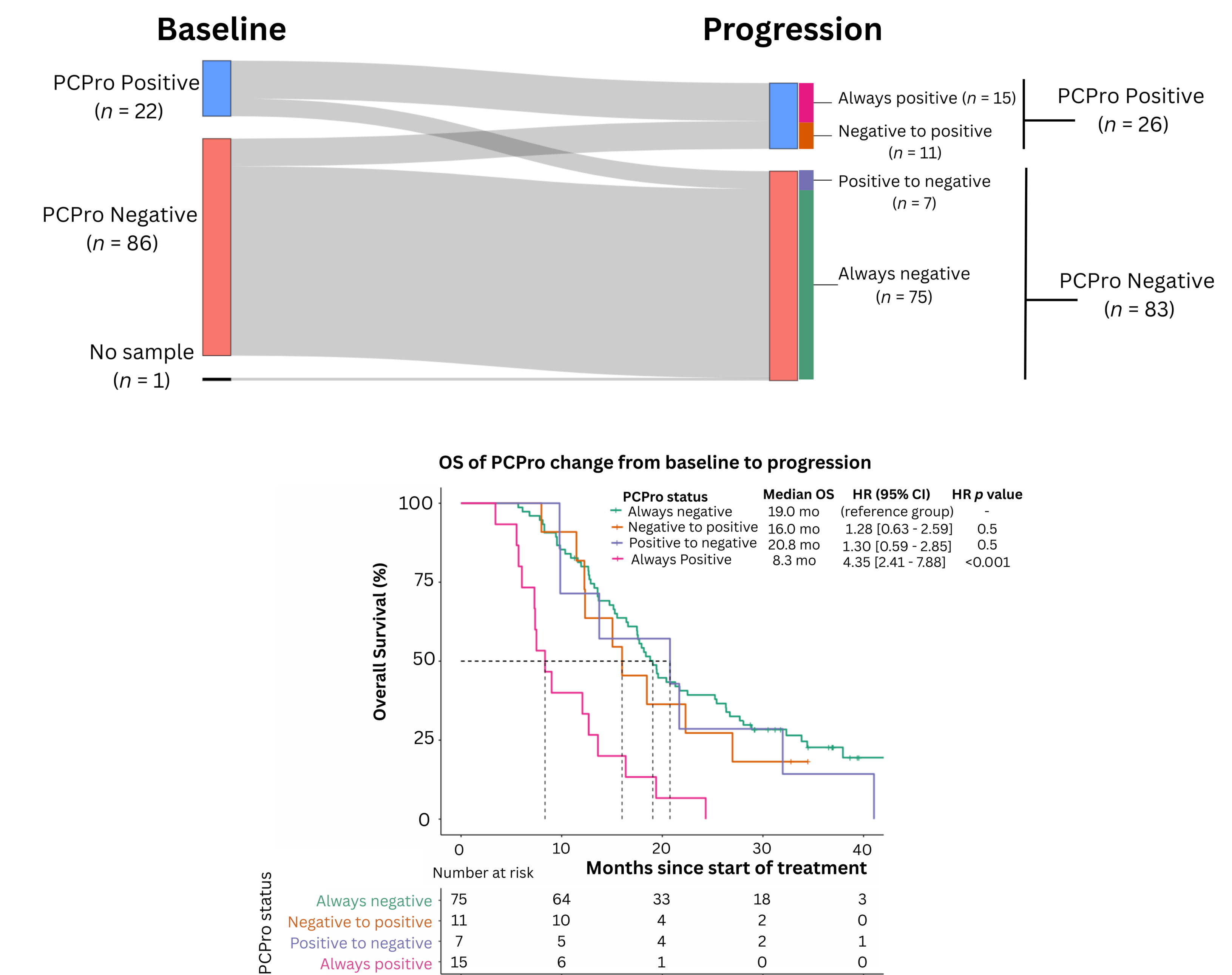
Table 2: Bivariable Cox regression of PCPro and Treatment arm

PCPro is prognostic independent of other prognostic biomarkers

Variable	Hazard Ratio OS [95% CI]	OS p value
PCPro, positive	1.78 [1.05 – 3.03]	0.032
PSMA PET SUV mean <10	1.39 [0.82 – 2.34]	0.2
FDG PET Mean Tumour Volume ≥ 200	2.29 [1.40 – 3.75]	0.001
ctDNA fraction	(Reference)	(Reference)
ctDNA 2-30%	4.07 [1.55 – 10.7]	0.004
ctDNA >30%	8.91 [3.17 – 25.1]	<0.001

Table 3: Multivariable analysis for PCPro with other prognostic variables for OS

Patients who remain PCPro positive at progression have the worst prognosis



Conclusions

- PCPro positive status was an independently significant prognostic factor for shorter OS and rPFS in participants treated in the TheraP trial with either LuPSMA or cabazitaxel
- Participants who remained PCPro positive at baseline and at progression had the worst prognosis
- These data support further research of PCPro as a prognostic biomarker in people with mCRPC being treated with LuPSMA or taxane chemotherapy.

References

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More Information

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Acknowledgements

Conflicts of interest

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