Baseline PSMA PET/CT parameters predict overall survival and

treatment response in metastatic castration-resistant prostate

cancer patients

ELECTRONIC SUPPLEMENTARY MATERIAL

Table S1

For patient-level analysis				
Parameter	Explanation (IBSI code [20])			
Morphological				
PSMA-TV	Total tumour volume in mL (RNU0)			
DmaxVox	The distance in cm between the two lesions that are the furthest apart, using the outermost voxels (no IBSI code)			
Intensity-based*				
SUV_{mean}	The mean grey level intensity within the total tumour volume (Q4LE)			
SUV _{max}	The maximum grey level intensity within the total tumour volume (84IY)			
TL-PSMA	The product of SUV _{mean} and total tumour volume in mL (no IBSI code)			
For losion-loyal analy	voic			
For lesion-level analy Parameter	Explanation (IBSI code [20])			
Parameter				
Parameter Morphological	Explanation (IBSI code [20])			
Parameter Morphological PSMA-TV	Explanation (IBSI code [20]) Lesion volume in mL (RNU0) A measure of the roundness of the shape of the tumour region relative to a sphere; higher values correspond to higher			
Parameter Morphological PSMA-TV Sphericity	Explanation (IBSI code [20]) Lesion volume in mL (RNU0) A measure of the roundness of the shape of the tumour region relative to a sphere; higher values correspond to higher			
Parameter Morphological PSMA-TV Sphericity Intensity-based*	Lesion volume in mL (RNU0) A measure of the roundness of the shape of the tumour region relative to a sphere; higher values correspond to higher roundness (QCFX)			
Parameter Morphological PSMA-TV Sphericity Intensity-based* SUV _{mean}	Lesion volume in mL (RNU0) A measure of the roundness of the shape of the tumour region relative to a sphere; higher values correspond to higher roundness (QCFX) The mean grey level intensity within the VOI (Q4LE)			
Parameter Morphological PSMA-TV Sphericity Intensity-based* SUV _{mean} SUV _{max}	Lesion volume in mL (RNU0) A measure of the roundness of the shape of the tumour region relative to a sphere; higher values correspond to higher roundness (QCFX) The mean grey level intensity within the VOI (Q4LE) The maximum grey level intensity within the VOI (84IY) A measure of peakedness of the grey level distribution; higher values mean that the mass of the distribution is concentrated towards the tail(s), lower values mean that the mass of the distribution is concentrated towards a spike near the mean value			

GLCM**	
Energy	Or Angular Second Moment; a measure of homogeneous patterns in the VOI, a greater energy implies that there is a higher frequency of neighbouring intensity value pairs (8ZQL)
Contrast	A measure of the local intensity variation; larger values correlate with a higher differences in grey level intensity values among neighbouring voxels (ACUI)
Correlation	The linear dependency of grey level values to their respective voxels in the GLCM; higher values mean higher correlation (NI2N)
Entropy	A measure of randomness in neighbourhood grey level intensity values (TU9B)
Homogeneity	Or Inverse Difference; a measure of homogeneity of grey level intensity values, higher values mean more uniform grey levels (IB1Z)

^{*}The intensity of a pixel or voxel is also called a grey level.

Table S2
Multivariate Cox regression analyses for overall survival with PSMA PET/CT and clinical parameters.

^{*} HR: hazard ratio, displayed per unit increase or, in case of log₂ transformation, per doubling. ** p-values are displayed when < 0.05.

		os	
Variable name (unit)	HR*	95% C.I.	p-value**
Log ₂ (TL-PSMA (SUV×mL))	1.402	1.161-1.693	< 0.001
Log ₂ (baseline PSA (ng/mL))	0.826	0.693-0.985	0.034
Baseline Hb (mmol/L)	0.647	0.464-0.903	0.010
Line of treatment (second-versus first-line)	3.406	1.679-6.909	< 0.001
DmaxVox (per 10 cm)	1.314	1.065-1.619	0.011
Log₂(baseline PSA (ng/mL))	0.864	0.724-1.030	
Baseline Hb (mmol/L)	0.695	0.496-0.975	0.035
Line of treatment (second- versus first-line)	3.180	1.521-6.645	0.002
Log ₂ (PSA density (ng/mL ²))	0.760	0.652-0.885	<0.001
Baseline Hb (mmol/L)	0.640	0.465-0.883	0.006
Line of treatment (second- versus first-line)	3.941	1.984-7.828	< 0.001

^{**} The GLCM is calculated from 13 different directions in 3D and the GLCM features are average values over these 13 directions.

PSMA-TV: PSMA-based tumour volume, SUV: standardised uptake value, VOI: volume of interest, TL-PSMA: PSMA-based total lesion uptake. GLCM: grey level co-occurrence matrix.

Table S3
Lesion characteristics (n = 241). Categorical data are presented as number (percentage), continuous data as median (interquartile range).

Characteristic		Value	
Received trea	tment		
ARTA		113	(47%)
Chemotherapy		128	(53%)
Line of treatme	ent		
1 st		122	(51%)
2 nd		119	(49%)
Lesion locatio	n		
Prostate		32	(13%)
Lympl	n node	64	(27%)
	N1	21	
	M1a	43	
Bone		139	(58%)
	Locoregional	55	
	Axial, non-locoregional	59	
	Appendicular	25	
Visce	ral	6	(3%)
SUV _{mean}		9.4	(7.3 – 13.6)
SUV _{max}		30.0	(16.3 – 47.5)
PSMA-TV (mL)		7.7	(3.4 – 21.9)
TL-PSMA (SUV×mL)		76.4	(30.4 – 239.4)

Table S4
Univariate logistic regression analyses for imaging-based lesion progression after 3-4
months of treatment, all corrected for line of treatment. * OR: odds ratio, displayed per unit
increase or, in case of log₂ transformation, per doubling. ** p-values are displayed when <
0.05. Ref = reference group.

	Lesion pr	Lesion progression (n = 241 lesions)		
Variable name (unit)	OR*	95% CI	p-value**	
Location			0.013	
Prostate	0.169	0.049-0.589	0.005	
Lymph node	0.553	0.282-1.086		
Bone	ref	ref		
Visceral	0.232	0.026-2.076		
Morphological				
Log ₂ (PSMA-TV (mL))	0.977	0.835-1.142		
Log ₂ (Sphericity)	1.162	0.472-2.859		
Intensity-based				
$Log_2(SUV_{mean)}$	1.973	1.265 -3.079	0.003	
$Log_2(SUV_{max})$	1.400	1.049-1.868	0.022	
Log ₂ (SUV-kurtosis)	0.808	0.614-1.064		
Log ₂ (SUV-IQR)	1.352	1.071 -1.707	0.011	
Log ₂ (TL-PSMA (SUV×mL))	1.056	0.915-1.219		
GLCM				
Log₂(Energy)	0.837	0.713-0.982	0.029	
Log₂(Contrast)	1.168	1.035-1.318	0.012	
Correlation	0.812	0.150-4.403		
Entropy	1.181	1.004 -1.388	0.044	
Homogeneity	0.065	0.005-0.802	0.033	