Could textural analysis of MR images reduce overdiagnosis and overtreatment in prostate cancer?



Abstract ID: ICMP23_ABS_E2227

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International Conference on Medical Physics - 2023



ovations in Radiation Technology & Medical Physics ecember 2023 | Venue: DAE Convention Centre, Anushaktina

Poster ID:

DWI images

Maximum

p-value

0,012

0,041

0,006

0,000

0,002

0,002

0,001

0,000

0,029

0,000

0,003

0,004

0 001

Introduction

- Exclusion of clinically significant prostate cancer is essential to reduce harm from overtreatment.
- An accurate, non-invasive method is therefore required for identifying benign prostatic disease and for the active surveillance of localised prostate cancer (in case of clinically insignificant prostatic cancer).
- This work explores if textural analysis of acquired MR images can serve these goals.

Methodology

Image Acquisition





Data Integration & Analysis Feature Extraction

Results & Discussion

Surface Volume Ratio, 10Percentile & Imc1 derived from T2w and Range, Sphericity, Difference Variance & Skewness extracted from DW



- Study group: Following ethical approval, 89 subjects with increased prostate specific antigen (PSA) and prostatic hyperplasia (n=54) or neoplasia (n=35) were enrolled.
 - MR scanning: Subjects were scanned at 3.0T (Achieva TX Philips) using a multiparametric protocol according to the Prostate Imaging Reporting and Data System (PIRADSv2).
 - **Histopathology:** All subjects underwent transrectal ultrasound guided biopsy following MRI, whilst the Gleason score (GS) and the International Society of Urological Pathology (ISUP) grade group were recorded.
 - **Segmentation:** T2-weighted (T2w) and diffusion-weighted (DW) images were reviewed by an experienced radiologist and the dominant visible lesion was segmented (ITK-snap).

- images revealed a significant correlation (p<0,01) with GS.
- Regression analysis in 35 cancer cases with GS≥6 showed significant linear correlations (p<0,01) between GS and T2w-extracted Surface Volume Ratio & Imc1 and DW-extracted Interguartile Range.
- 5 T2w-extracted features could discriminate between low (1) and high (>1) ISUP scores, with corresponding ROC curves shown below:



- 14 DW-extracted features could discriminate between low and high
- Feature extraction: First-order, shape and gray-level cooccurrence matrix (GLCM) features (n=54) of segmented regions were extracted (Pyradiomics v3.1.0) for both T2w & DW images, following bias field correction, voxel rescaling to 1x1x1mm³, as well as image normalization and discretization.
- **Linear Regression:** Extracted features were correlated to Gleason score (SPSS).
- Intercomparison: The independent samples t-test (SPSS) was used to intercompare features between low (<6) & high (≥6) GS and low (1) & high (>1) ISUP grade categories.
- AU-ROC: The area under the receiver operating characteristic (AU-ROC) curve was used to quantify the discriminative power of statistically significant features for patient stratification (GS<6 vs GS \geq 6) and ISUP grade categorization (1 vs >1).

Conclusions

 Obtained results suggest that radiomic features could differentiate normal (GS<6) vs abnormal (≥6) prostatic tissue and low (1) vs high (>1) ISUP score in cancerous tissue, although powerful features seem to differ in the two scenarios.

ISUP scores (i.e., clinically insignificant vs significant cancer), with corresponding ROC curves presented below:



- T2w- and DW-extracted features with discriminative power are different and, thus, an approach similar to that of PI-RADS (DW domination for peripheral zone lesions and T2w domination for transition zone lesions) may have to be adopted.
- Textural analysis could potentially serve the goal of overdiagnosis and Limita overtreatment reduction in prostate ca, although different features may have to be employed for patient stratification, disease burden assessment, lesion monitoring and treatment outcome evaluation.

Acknowledgment

This work has been financed by the Horizon 2020 Framework Programme of the European Commission (project code 952179 – NKUA 16619)

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discriminators between	Skewness	0,025	MeanAbsoluteDeviation	0,002
normal (<6) and	Total Energy	0,009	Range	0,000
abnormal (≥6) GS.	Idm	0,03	RobustMeanAbsoluteDeviation	0,003
	Idmn	0,024	TotalEnergy	0,000
	Id	0,035	Uniformity	0,020
	Idn	0,013	Variance	0,003
	Inverse Variance	0,024	Autocorrelation	0,005
mitations Standardization of methodology and extending the study to a large sample size with longitudinal MRI scans are warranted.			JointAverage	0,006
			ClusterProminence	0,032
			ClusterTendency	0,004
			Correlation	0,004
			DifferenceEntropy	0,043
			DifferenceVariance	0,042
			JointEntropy	0,004
			Idmn	0,005
			Idn	0,002
			SumEntropy	0,003
			SumSquares	0,004

RootMeanSquared

0,039