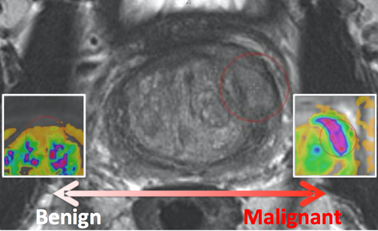
PROSTATEx Challenge 2017

# SPIE-AAPM-NCI Prostate MR Classification Challenge

# Test Set Release



URL: [https://spie.org/conferences-and-exhibitions/medical-imaging/prostatex-grand-challenge](%20https://spie.org/conferences-and-exhibitions/medical-imaging/prostatex-grand-challenge)

This document provides information about the annotated prostate MR images provided with the ProstateX challenge and available on TCIA under that same name.

This document should be accompanied by two files: ProstateX-Findings-Test.csv and ProstateX-Images-Test.csv.

H.J.Huisman, December 2016 

# Images (see the file ProstateX-Images-Test.csv)

Prostate MR in accordance with ACR PIRADS2.0 comprises at least 3 types of images or parameters that should be jointly analyzed for the assessment of prostate cancer. The prostate MR imaging was performed at the Radboud University Medical Centre (Radboudumc) in the Prostate MR Reference Center under supervision of prof. dr. Barentsz. The Radboudumc is located in Nijmegen, The Netherlands. The dataset was collected and curated for research in computer-aided diagnosis of prostate MR under supervision of Dr. Huisman, Radboudumc as documented in:

G. Litjens, O. Debats, J. Barentsz, N. Karssemeijer and H. Huisman. "Computer-aided detection of prostate cancer in MRI", *IEEE Transactions on Medical Imaging* 2014;33:1083-1092.

If you use this data for research, then please refer to the above publication.

The images come in two encodings. The acquired MR is provided in DICOM encoding. Additionally Ktrans images are provided. They come in mhd format. Ktrans is a key pharmacokinetic parameter computed from the available dynamic contrast-enhanced T1-weighted series. Each patient has one study with several DICOM images and one Ktrans image. The Ktrans image is encoded in two files ProstateX-[ProxID]-Ktrans.[mhd/zraw], where ProxID is the ProstateX patient identifier. The DICOM images comprise several series each comprising several instances. The DICOM files are documented in the ProstateX-Images-Test.csv file. The columns in this file encode the following:

* ProxID – ProstateX patient identifier
* Name – Series description
* fid – Finding ID
* Pos – Scanner coordinate position of the finding
* WorldMatrix – Matrix describing image orientation and scaling
* ijk – Image column (i),row (j), and slice (k) coordinates of the finding. Using the VTK/ITK/Python array convention, (0,0,0) represents the first column and first row of the first slice.
* TopLevel
  + 0 - Series forms one image
  + 1 – A set of series forms a 4D image (e.g. Dynamic MR)
  + NA – Series form one image, but is part of a Level 1 4D image
* SpacingBetweenSlices – Scalar spacing between slices
* VoxelSpacing – Vector with x,y,z spacing scalars
* Dim – Vector with 4D dimensions of the image
* DCMSerDescr – Original DICOM series description
* DCMSerNum – DICOM series number

For example, to get the ADC image of Patient ProstateX-0204 do the following. After you imported the DICOM files into your environment, go to patient ProstateX-0204 and find the series with ADC in it. In this case it is ‘ep2d\_diff\_tra\_DYNDIST\_ADC’. It has SeriesNumber 7. The DICOM images in that series form the ADC image for this challenge. Image slice k at coordinate i,j contains a finding fid. See findings for more details.

The most reliable way to match the images with the information in the spreadsheet is to match the image series description DICOM tag (0008,103E) with the column DCMSerDescr in the spreadsheet.

# Findings (see the file ProstateX-Findings-Test.csv)

The findings are documented in the ProstateX-Findings-Test.csv file. Documentation for the columns in this file is as follows:

* ProxID – ProstateX patient identifier
* fid - Finding ID
* pos - Scanner coordinate position of the finding
* zone – Anatomic zone containing the lesion within the prostate
  + AS – Anterior fibromuscular stroma
  + PZ – Peripheral zone
  + SV – Seminal vesicle
  + TZ – Transition zone

# Additional Information

To organize the key information that has been posted to the Challenge website Forum, the organizers have created a “Frequently Asked Questions” thread (see <http://spiechallenges.cloudapp.net/forums/6/16/>) in which the essential elements of other threads will be re-posted.

As you prepare to download the test cases, please note the following very important policy: Once you as a participant submit your test set classification output to the challenge organizers, you will be considered fully enrolled in the Challenge, and your performance results (without links to your identity) will become part of any presentations, publications, or subsequent analyses derived from the Challenge at the discretion of the organizers. You will not be able to withdraw from the Challenge once you submit your test set classification results.

The test set cases are to be manipulated, processed, and analyzed without human intervention, although manual or human-supervised delineation of the prostate gland border or the gross lesion margin will be allowed, if necessary. Manual delineation of other structures or anatomic regions is not permitted.