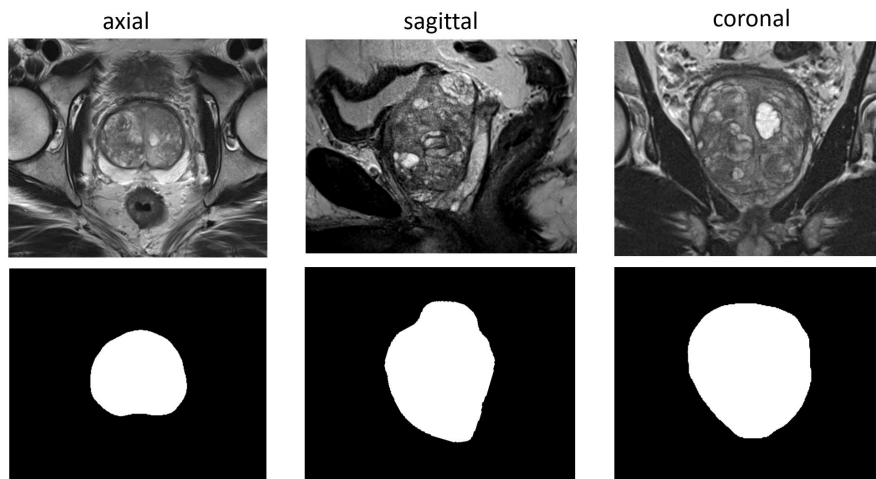


# High Resolution Prostate Segmentations for the ProstateX-Challenge

## Summary

We created 66 high resolution segmentations for randomly chosen T2-weighted volumes of the [SPI E-AAPM-NCI PROSTATEx Challenges](#). The high resolution segmentations were obtained by considering the three scan directions: for each scan direction (axial, sagittal, coronal), the gland was manually delineated by a medical student, followed by a review and corrections of an expert urologist. These three anisotropic segmentations were fused to one isotropic segmentation by means of shape-based interpolation in the following manner: (1) The signed distance transformation of the three segmentations is computed. (2) The anisotropic distance volumes are transformed into an isotropic high-resolution representation with linear interpolation. (3) By averaging the distances, smoothing and thresholding them at zero, we obtained the fused segmentation. The resulting segmentations were manually verified and corrected further by the expert urologist if necessary. Serving as ground truth for training CNNs, these segmentations have the potential to improve the segmentation accuracy of automated algorithms. By considering not only the axial scans but also sagittal and coronal scan directions, we aimed to have higher fidelity of the segmentations especially at the apex and base regions of the prostate.



The segmentations to standard DICOM representation were created with [dcmqi](#).

## Acknowledgements

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## Data Access

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## Detailed Description

### Detailed Description

Image Statistics	
Modalities	SEG
Number of Patients	66
Number of Studies	66
Number of Series	66
Number of Images	66
Image Size (GB)	0.119

**Note:** Please contact [help@cancerimagingarchive.net](mailto:help@cancerimagingarchive.net) with any questions regarding usage.

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**i Publication Citation**

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**i TCIA Citation**



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