

Supplementary Files



Pre-Treatment T2-WI Based Radiomics Features for Prediction of Locally Advanced Rectal Cancer Non-Response to Neoadjuvant Chemoradiotherapy: A Preliminary Study

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PyRadiomics Configuration

setting:

Image discretization:# The gray values were discretized using a fixed-bin width of 5 binWidth: 5

Normalization: # Images were normalized before feature extraction to 0 mean and 100 standard deviation normalize: true normalizeScale: 100

Resampling:
Images were interpolated to isotropic voxels with 2 mm sides using a B-Spline interpolator interpolator: 'sitkBSpline'
resampledPixelSpacing: [2, 2, 2]

Resegmentation

Segmentations were resegmented by excluding voxels which differed > 3 sigma from the

resegmentRange: [-3,3] resegmentMode: sigma

first order specific settings:

When normalizing, gray values below the mean will be negative. Shifting by 300 (3 StdDevs * 100) ensures that the majority of voxels is positive (only outliers >3 SD lower than the mean will be negative).

voxelArrayShift: 300

Misc:# default label value.label: 1

imageType:

mean

Selection of filters
Original: {}

LoG:

sigma: [3.0, 5.0] Wavelet: {}

featureClass:

Selection of the extracted radiomics features

shape:

firstorder:

glcm:

- 'Autocorrelation'
- 'JointAverage'
- 'ClusterProminence'
- 'ClusterShade'
- 'ClusterTendency'
- 'Contrast'
- 'Correlation'
- 'DifferenceAverage'
- 'DifferenceEntropy'
- 'DifferenceVariance'
- 'JointEnergy'
- 'JointEntropy'
- 'Imc1'
- 'Imc2'
- 'Idm'
- 'Idmn'
- 'Id'
- 'Idn'
- 'InverseVariance'
- 'MaximumProbability'
- 'SumEntropy'
- 'SumSquares'
- glrlm:
- glszm:

gldm:



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