

Online Supplements

Transfer-Learning Deep Radiomics and Hand-Crafted Radiomics for Classifying Lymph Nodes from Contrast-Enhanced Computed Tomography in Lung Cancer

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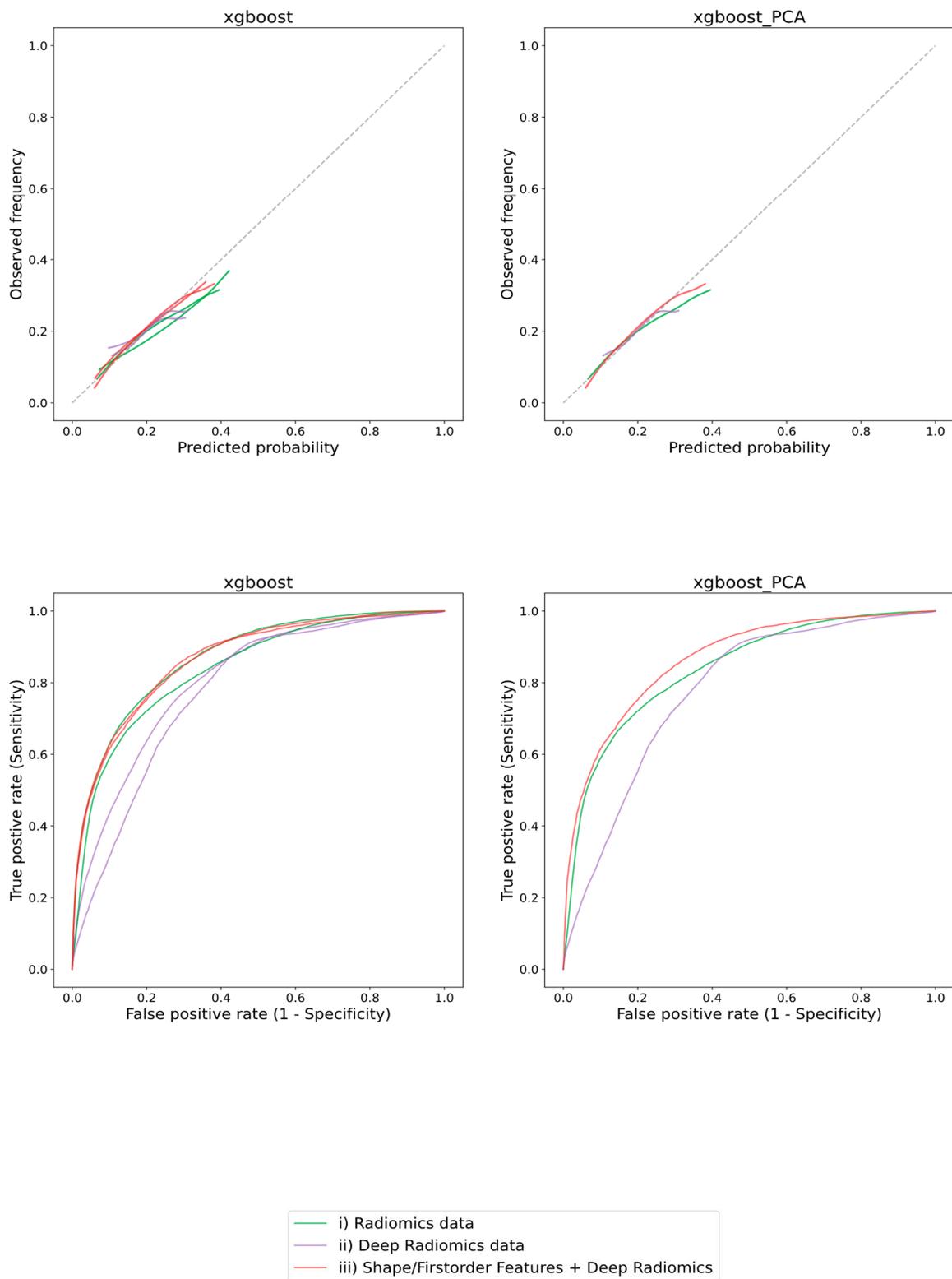


Figure S1. Calibration and discrimination for classification of PET-positivity for different radiomics approaches. Results of xgboost are shown. In the lowest-smoothed calibration plot (first row), the observed outcome frequency is plotted against the predicted outcome probability. The closer the curve is to the diagonal, the better the calibration. *The receiver-operating-characteristic (second row) plots the true positive rate against the false positive rate by varying thresholds (not shown). Discrimination is best for the curve that is closest to the left upper corner.* Legend: PCA = principal component analysis.

Table S1. Classification results using XGBoost for “traditional” hand-crafted, deep and hybrid features.

Metric Feature type Model	AUC Radiomics	Brier score (BS) Radiomics	R ² (scaled BS) Radiomics	Sensitivity Radiomics	Specificity Radiomics
XGBoost: i) Radiomics data	0.86 (0.838-0.875)	0.119 (0.116-0.121)	26.6 (25.1-28.3)	0.76 (0.711-0.799)	0.803 (0.782-0.823)
XGBoost: ii) Deep Radiomics data	0.807 (0.801-0.812)	0.129 (0.126-0.131)	20.5 (19.5-21.6)	0.784 (0.764-0.804)	0.705 (0.684-0.725)
XGBoost: iii) Shape/Firstorder Features + Deep Radiomics	0.868 (0.862-0.873)	0.105 (0.102-0.108)	35 (33.7-36.5)	0.821 (0.794-0.845)	0.763 (0.739-0.789)
XGBoost + PCA: i) Radiomics data	0.839 (0.832-0.845)	0.121 (0.118-0.125)	25 (23.2-26.7)	0.723 (0.703-0.747)	0.817 (0.795-0.836)
XGBoost + PCA: ii) Deep Radiomics data	0.776 (0.77-0.782)	0.138 (0.136-0.141)	14.6 (13.5-15.6)	0.853 (0.828-0.876)	0.609 (0.585-0.636)
XGBoost + PCA: iii) Shape/Firstorder Features + Deep Radiomics	0.864 (0.857-0.87)	0.107 (0.104-0.11)	34 (32.6-35.5)	0.798 (0.769-0.825)	0.775 (0.747-0.801)