

SUPPLEMENTARY MATERIALS

RQS Scoring System

The radiomics quality score (RQS) score consisted of 16 components. The reviewers performed RQS evaluation according to six domains as previously reported (1, 2). Prior to the evaluation, a research meeting was held to educate the reviewers on the RQS system.

Two reviewers (with 6 and 9 years of experience in radiology, respectively) independently scored the articles for each of the six domains using RQS (Supplementary Material). If disagreement occurred between the two reviewers, a final decision was made through a consensus. Domain 1 represented image protocol quality and stability of image and segmentation (0 to 5 points). It consisted of image protocol quality (+2: well-documented protocol [+1], usage of public image protocol [+1]), multiple segmentations (+1), phantom study on all scanners (+1), and imaging at multiple time points (+1). Domain 2 represented feature selection and validation (-8 to 8 points). It consisted of feature reduction or adjustment for multiple testing (-3 or +3 points) and validation (-5 to +5 points). Domain 3 represented biologic/clinical validation and utility (0 to 6 points). It consisted of multivariable analysis with non-radiomics feature (+1), detection or discussion of biologic correlates (+1), comparison to 'gold standard' (+2), and report of potential clinical utility (+2). Domain 4 represented model performance index (0 to 5 points). It consisted of cut-off analyses (+1), discrimination statistics (+2: report of discrimination statics [+1] application of resampling method [+1]) and calibration statistics (+2: calibration statics [+1], application of resampling method [+1]). Domain 5 represented high level of evidence (0 to 8 points). It consisted of prospective validation (+7) or cost-effectiveness analysis (+1). Domain 6 represented open science and data (0 to 4 points). It consisted of open source scans, segmentations, code, and data (+4).

Consensus Reached for RQS Scoring

The additional topics were discussed by the two reviewers and a consensus was reached for evaluation. RQS was scored according to the consensus for the following topics:

1) Image protocol quality (domain 1): As Alzheimer's Disease Neuroimaging Initiative (ADNI) is a widely used multicenter open source database (3), several radiomics studies which used the ADNI database did not document the image protocol in the manuscript. However, an additional point for well-documented protocol was still earned in case the ADNI database was used with a reference for the database depository (adni.loni.usc.edu or www.adni-info.org) as the ADNI imaging protocol is widely known.

2) Multiple segmentation (domain 1): In quantitative neurodegenerative studies, brain region segmentation is usually performed automatically rather than manually or semi-automatically by single or multiple readers. Thus, if well-known and validated automatic segmentation software for brain imaging such as FreeSurfer, FMRIB Software Library, Statistical Parametric Mapping, or volBrain were used to perform segmentation of brain regions (4-6), an additional point was earned as it was considered to have better segmentation reproducibility.

3) Validation (domain 2): If cross-validation or nested cross-validation was performed only within the training set, it was considered to be missing validation and scored -5 points, as previously described (1). When the study was validated by randomly splitting a multicenter open source data such as the ADNI database into training and test sets, a scoring for a validation based on a dataset from the another institute (+4) was applied rather than applying the full scoring for a validation based on three or more datasets from distinct institutes (+5). This scoring was applied because although MRIs from ADNI were obtained across different sites and different MRI vendors, the ADNI study uses a strictly controlled MRI protocol, resulting in less heterogeneity of radiomics features that is acquired from a true external validation (7). Also, because the data was randomly split without knowledge of the acquired primary site, there is a possibility that data from the identical site may have been included in both the training and test sets.

4) Comparison with the gold standard (domain 3): A notable difference between oncology and AD studies is that the histologic confirmation of the disease diagnosis is feasible ex vivo for oncology, but not for AD (8). Thus, for diagnostic studies the pathologic gold standard according to the amyloid, tau, and neurodegeneration (ATN) guideline for AD diagnosis recommended by the National Institute on Aging and Alzheimer's Association Research Framework was applied

(9); cerebrospinal fluid amyloid beta (A β) or amyloid PET are specific to aggregated A β , and tau PET reflects aggregated tau, whereas hippocampal atrophy reflects neurodegeneration. For prognosis studies, well-known predictors of cognitive conversion such as apolipoprotein E allele of the apolipoprotein gene, clinical severity, brain atrophy, cerebrospinal fluid biomarkers were applied (10-14).

5) Potential clinical utility (domain 3): The statement of the Food and Drug Administration and National Institutes of Health Working Group (15) suggested 'clinical utility' is thought to be accomplished when a biomarker guide to net improvement of health outcomes or give information valuable for prevention, diagnosis, management or treatment of a disease (15-17). If only the potential utility was discussed without proper analysis, no additional point was scored.

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