Machine learning and interpretability methods to investigate Alzheimer's disease

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Alzheimer's Disease Motivation Fachhochschule Dortmund University of Applied Sciences and Arts

Alzheimer's Disease (AD)

- Most frequent cause of dementia [1]
- Neurodegeneration starts decades before dementia symptoms occur
- At time of diagnosis, many neurons are irreversibly degenerated
- No cure, only reduction of symptoms [2]
- Early detection important but complex due to heterogenous disease profiles

Alzheimer's Diseas Motivation

Motivation

- Machine Learning (ML) to identify complex patterns in high-dimensional data
- Identifying complex patterns that improve the early prediction of AD
- Complex underlying systems require complex models
- Interpretable ML (IML) to explain decisions of black-box models and validated biological plausibility

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Research Approach

- 1. Train ML and Deep-Learning (DL) models to predict AD.
- 2. Check generalizability during internal and external validation.
- 3. Use interpretability methods to explain black-box models.
- 4. Compare explanations of ML- and DL- models to each other.
- 5. Validate biological plausability of explanations with a ground-truth Voxel-Based Morphometry (VBM) [3].

Research Approac Methods

Data

Table 1: Demographic data, and MRI field strength of the selected subjects, separated by diagnosis groups. For continuous features, mean and standard deviation are given.

Diagnosis	n	Age (years)	Females (%)	1.5 T (%)	3 T (%)
ADNI [4] (for training and internal validation)					
CN	512	74.20 ± 5.82	51.76	44.00	56.00
AD	335	74.95 ± 7.74	44.78	57.00	43.00
Σ	847	74.50 ± 6.66	49.00	49.00	51.00
AIBL [5] (for external validation)					
CN	446	72.53 ± 6.14	56.95	19.06	80.94
AD	71	73.26 ± 7.88	59.15	16.90	83.10
Σ	517	$\textbf{72.63} \pm \textbf{6.41}$	57.25	18.76	81.24
OASIS [6] (for external validation)					
CN	704	68.35 ± 9.27	58.66	12.36	87.64
AD	198	75.62 ± 7.92	48.48	10.61	89.39
Σ	902	69.94 ± 9.48	56.43	11.97	88.03

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Feature Extraction

- Classical ML models:
 - Volumes of brain-regions extracted from Magnetic-Resonance-Imaging (MRI) scans
 - Normalized by estimated Total Intracranial Volume (eTIV)
- Deep-Learning models:
 - Convolutional Neural Networks (CNNs) trained on skull-stripped 3D-MRI scans



Figure 1: T1-weighted MRI-scan segmented by FreeSurfer v6.0. Adapted from: [7].

Model Training

- ADNI dataset split in 80 % training and 20 % independent test set (stratified)
- Hyperparameter tuning: Grid-search including 5-fold-cross-validation (CV)
- Interpretable-by-design: Decision Trees (DTs), Logisitic Regression (LR)
- Black-Box: Support Vector Machines (SVMs) [8], Random Forest (RF) [9], eXtreme Gradient Boosting (XGBoost) [10], Light Gradient Boosting (LightGBM) [11]
- Deep Learning (CNNs): DenseNet [12], EfficientNet-B0 [13], Squeeze and Excitation (SE) [14]-ResNet [15], and -ResNeXt [16]
- Platt scaling [17] for model calibration

Research Approac Methods

Interpretability Methods

- Highly correlated features are consolidated into aspects [18]
- All models: SHapley Additive exPlanations (SHAP) [19], Local Interpretable Model-Agnostic Explanations (LIME) [20]
- Classical ML: Permutation-based feature importance
- Deep Learning: Gradient-weighted Class Activation Mapping (GradCAM) [21], GradCAM++ [22]
- Deep Learning explanations summarized for regions

Internal and External Validation Explain Model Decisions Comparison to Biologically Plausible Ground Truth Fachhochschule Dortmund University of Applied Sciences and Arts

Internal and External Validation



Figure 2: Plot showing performance of the trained ML and DL models.

- Performance of models that are interpretable-by-design show strong differences (performance of DT weak, performance of LR fair)
- Performance of DL models does not outperform classical ML models
- AIBL performances acceptable (generalizability for DL models worse than for classical ML)
- OASIS results acceptable but worse than remaining performances

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Explain Classical ML Model Decisions



Figure 3: SHapley Additive exPlanations (SHAP) [19] waterfall plot to explain individual decision of a subject with AD for LightGBM model.

- Explain the differences of the individual prediction (f(x) = 0.414) and the average model prediction (E(f(x)) = 0.885) using the model input features
- Feature expressions that increase the AD risk (aspect_27, aspect_24)
- Other feature expressions have a protective influence (aspect_17).

Internal and External Validation Explain Model Decisions Comparison to Biologically Plausible Ground Truth



Explain DL Model Decisions



Figure 4: Heatmap showing GradCAM++ results to explain individual decision of a subject with AD for the DenseNet model. Source: [23]

Internal and External Validation Explain Model Decisions Comparison to Biologically Plausible Ground Truth



Voxel-based-Morphometry Analysis



Figure 5: VBM analysis results visualize ground-truth relevant brain regions of a subject with AD. Source: [23]

Results and Conclusions

Comparison to Biologically Plausible Ground Truth

DenseNet

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Comparison to Biologically Plausible Ground Truth





SEResNet SEResNeXt VBM

Figure 6: Polar plot to compare classical ML model explanations to VBM ground truth.

Figure 7: Polar plot to compare Deep-Learning-Model explanations to VBM ground truth.

EfficientNet

Future Work

- Check why the localization of the Deep-Learning model explanations is rather unfocused (new information vs. overfitting / underfitting)
- Validation on clinically more relevant research questions (e.g., Mild cognitive impaired subjects, Amyloid-β-positivity, Tau-positivity)
- Use of multimodal input features

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Many thanks for your attention!

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