

# Dementia classification based on brain age estimation

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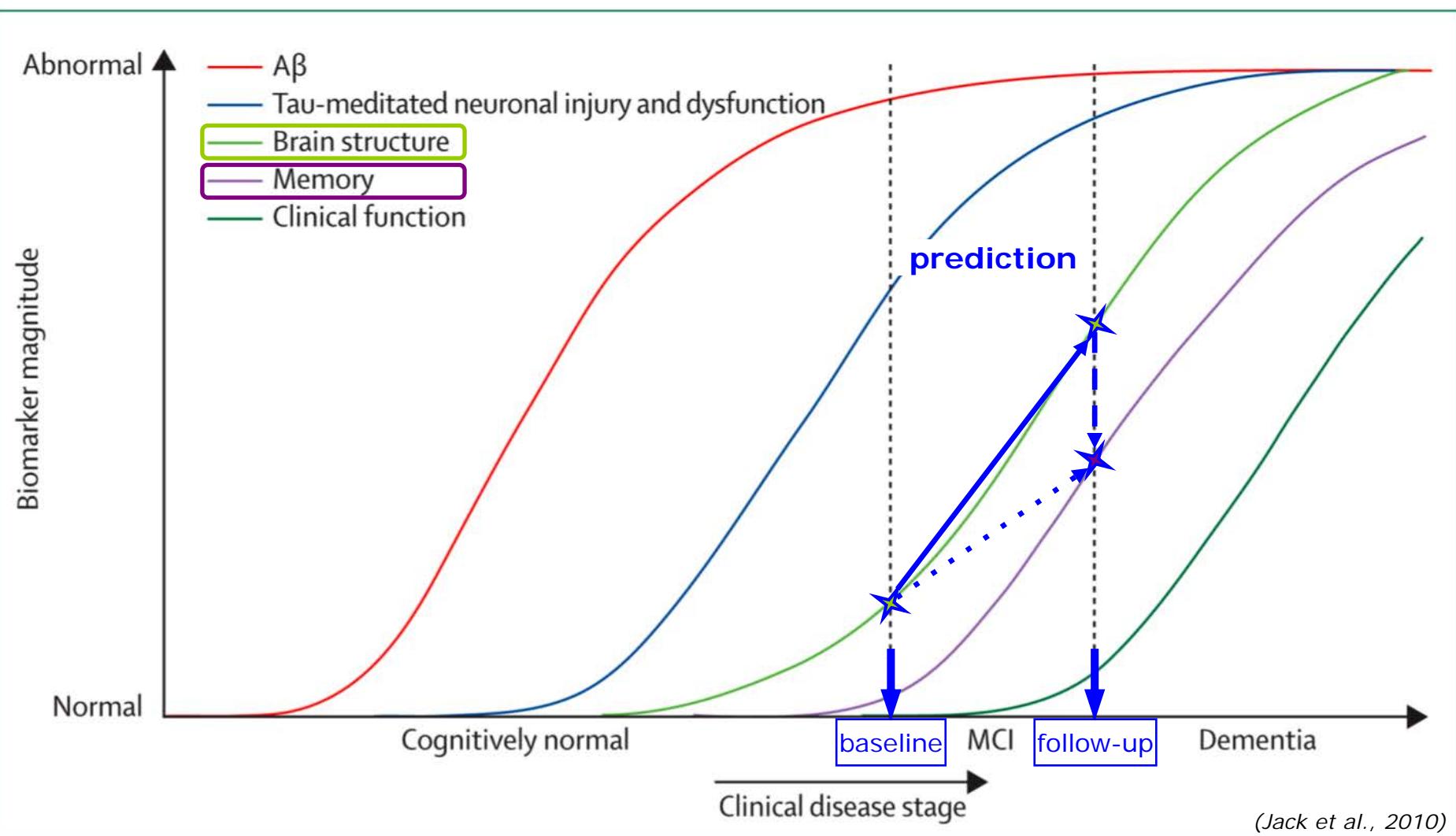
Structural Brain Mapping Group  
Depts. of Neurology & Psychiatry, Jena University Hospital, Germany



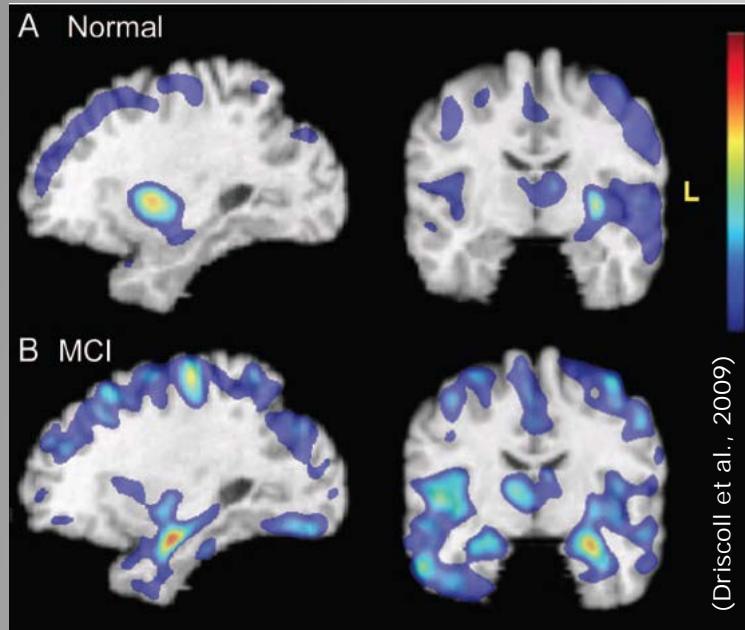
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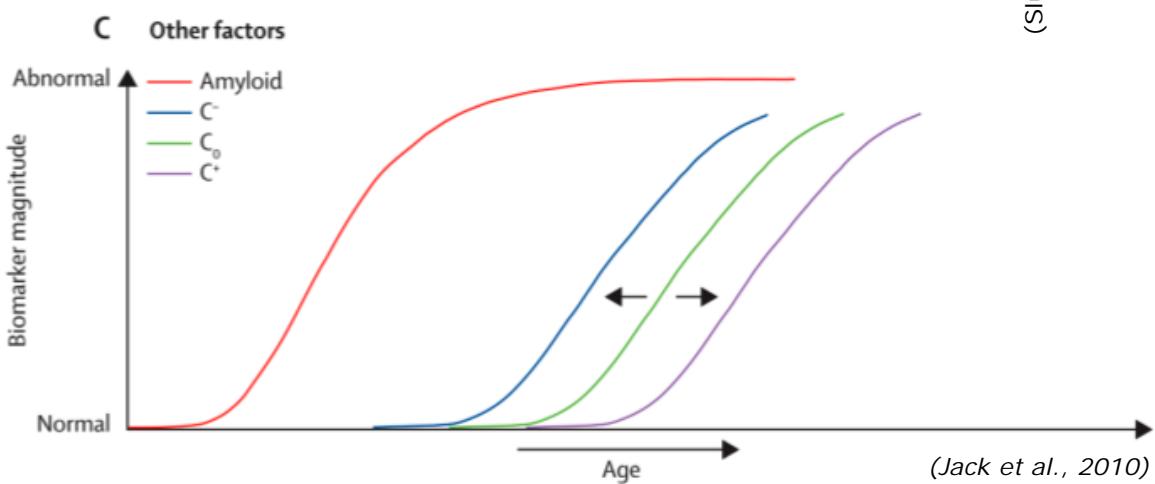
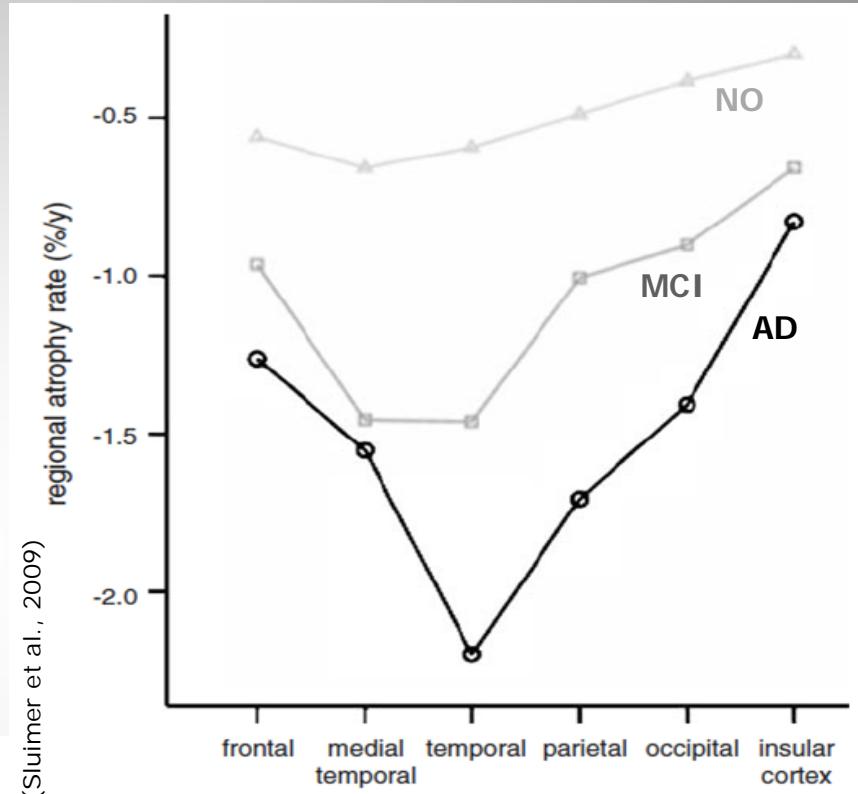
# Dynamic biomarkers of the Alzheimer's pathological cascade



# Advanced atrophy in MCI & Alzheimer's disease



→ complex & **characteristic** atrophy patterns over time



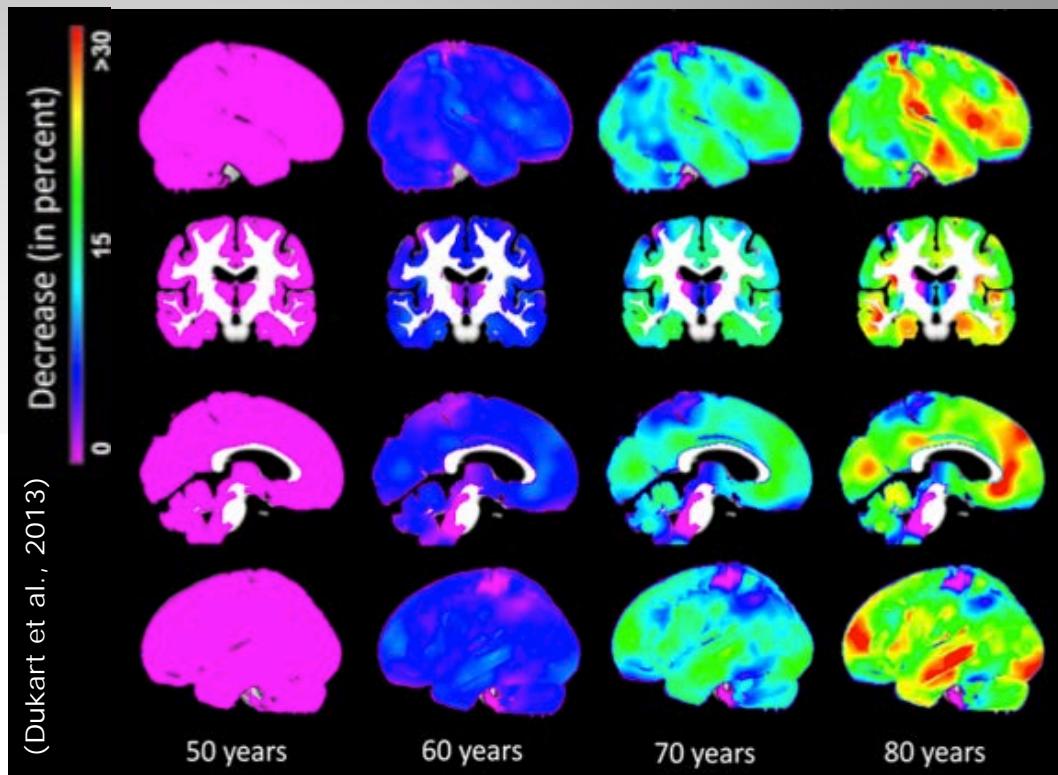
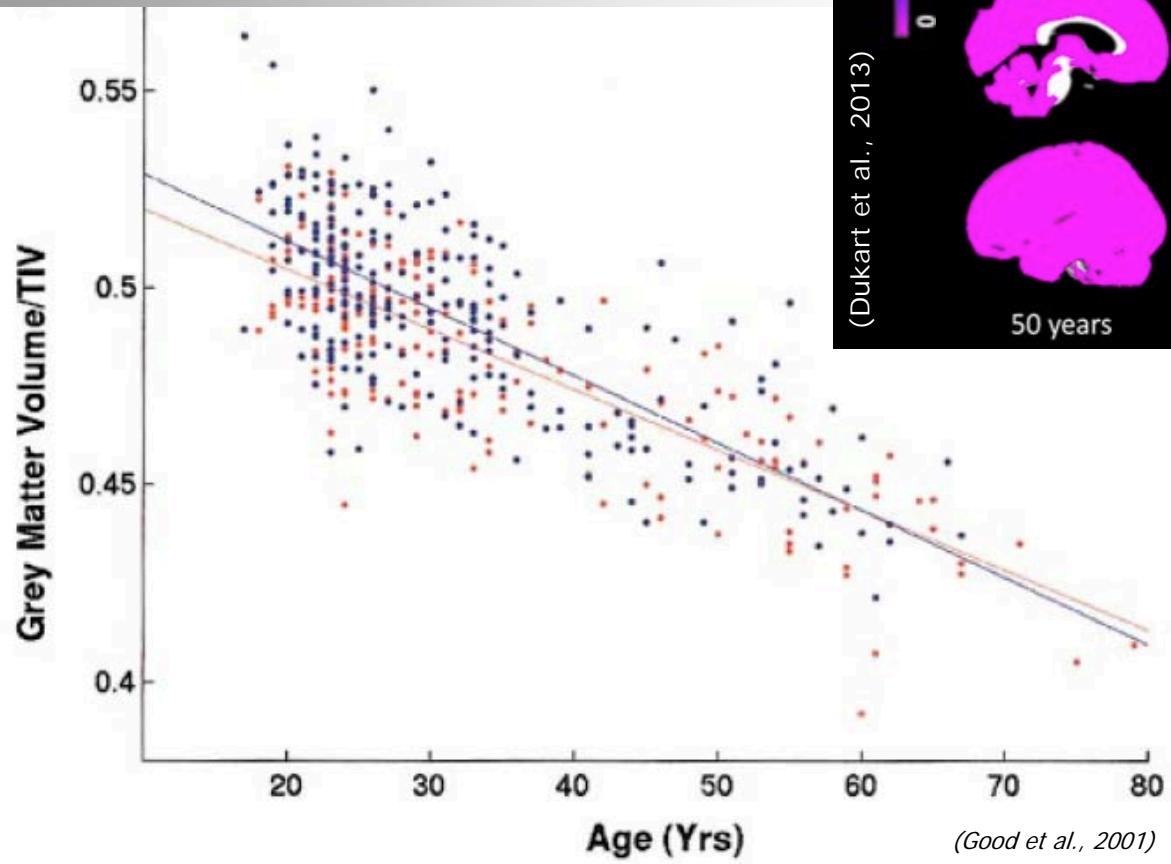
→ with **intraindividual** variability

→ with **interindividual** variability within the disease course

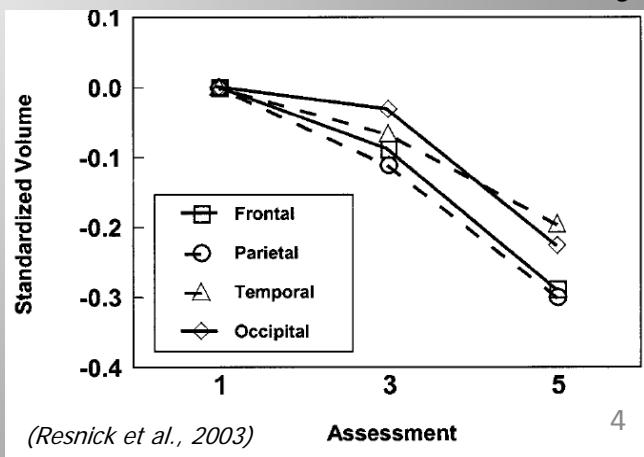
# Healthy aging → Characteristic atrophy patterns

→ complex & **characteristic** atrophy patterns over time

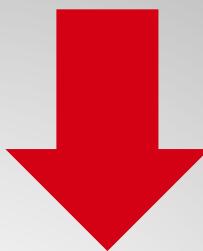
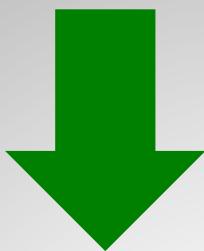
→ with **inter**individual variability



→ with **intraindividual** variability



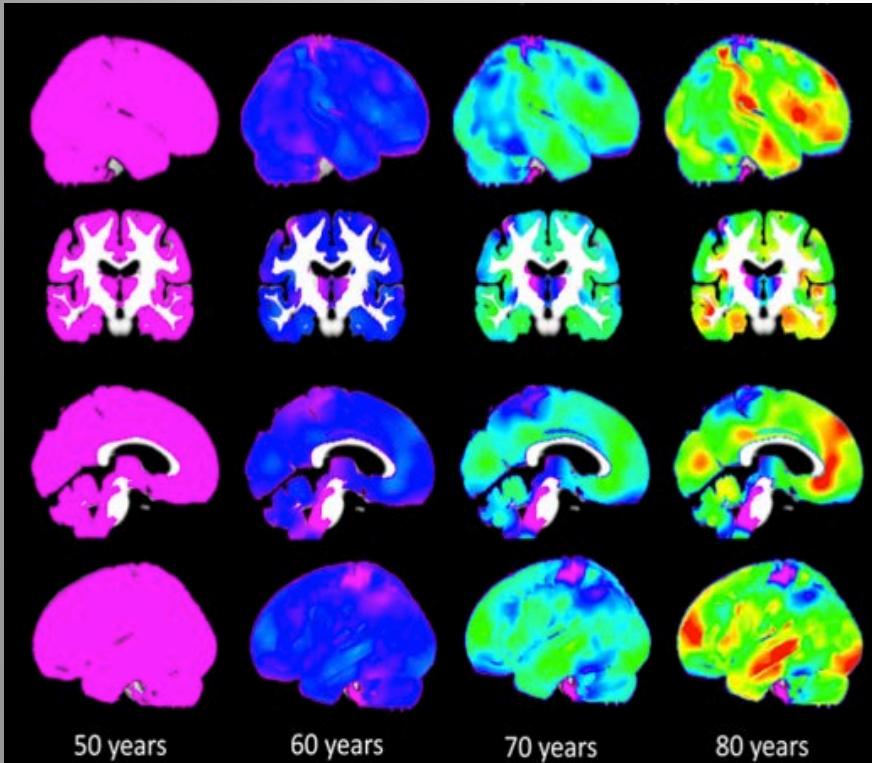
# Brain atrophy in Alzheimer's dementia



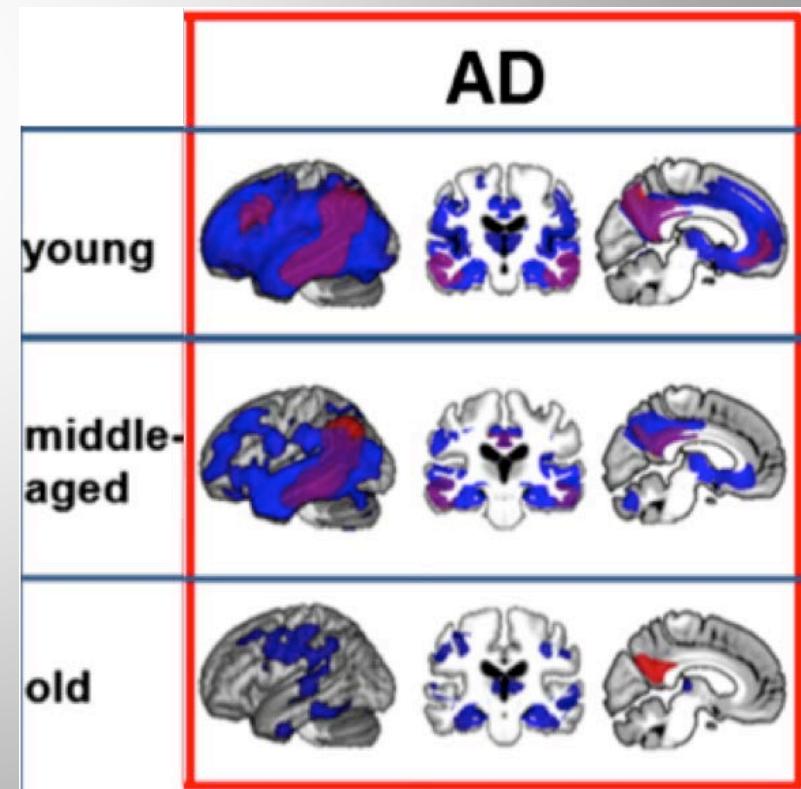
(Advanced) age-related atrophy

+

Disease-specific atrophy



(Dukart et al., 2013)



(Dukart et al., 2013)

# Prediction based on healthy brain aging

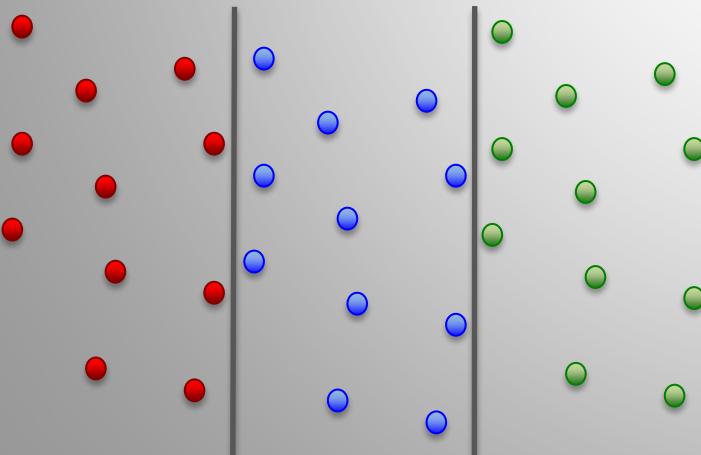
Baseline data (structural MR image)



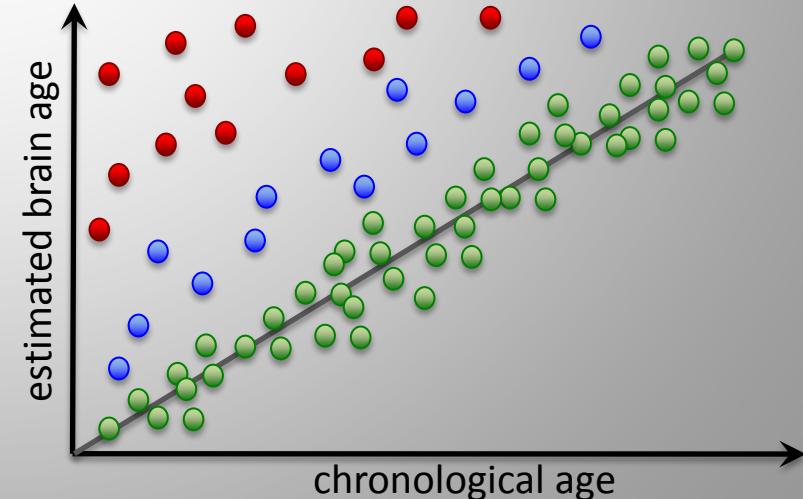
Individual prediction: **AD**, **MCI**, or **healthy**?



Typical approach: classification of disease-specific atrophy patterns



BrainAGE approach: deviation from normal aging-related atrophy patterns



# Modeling of structural brain aging

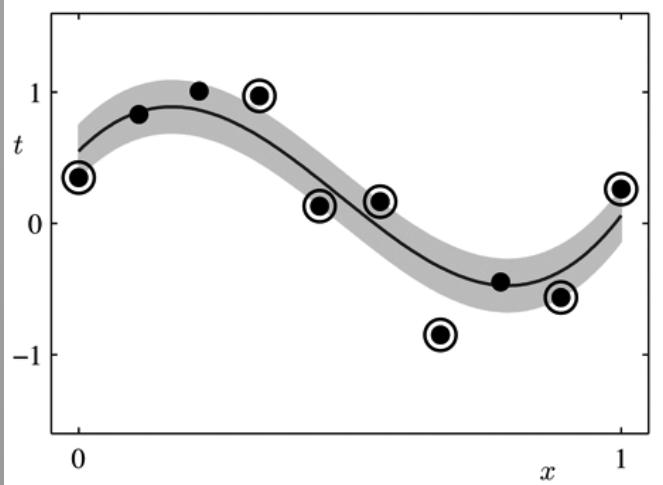
complex & characteristic atrophy patterns



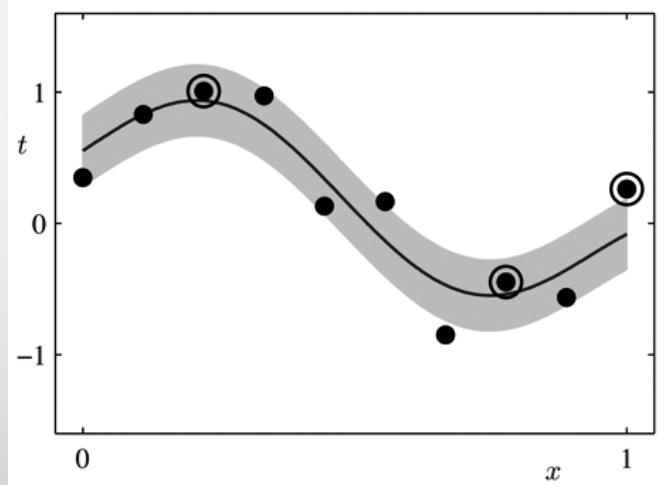
Pattern recognition / Machine-learning



Support Vector Regression

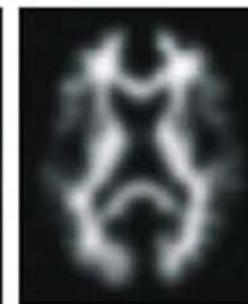
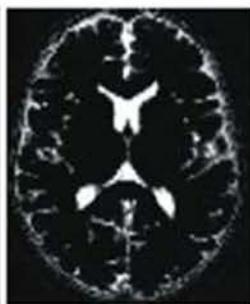
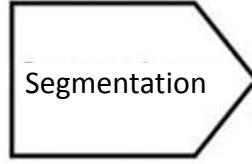


Relevance Vector Regression



- Bayesian approach
- crossvalidation NOT necessary
- sparse

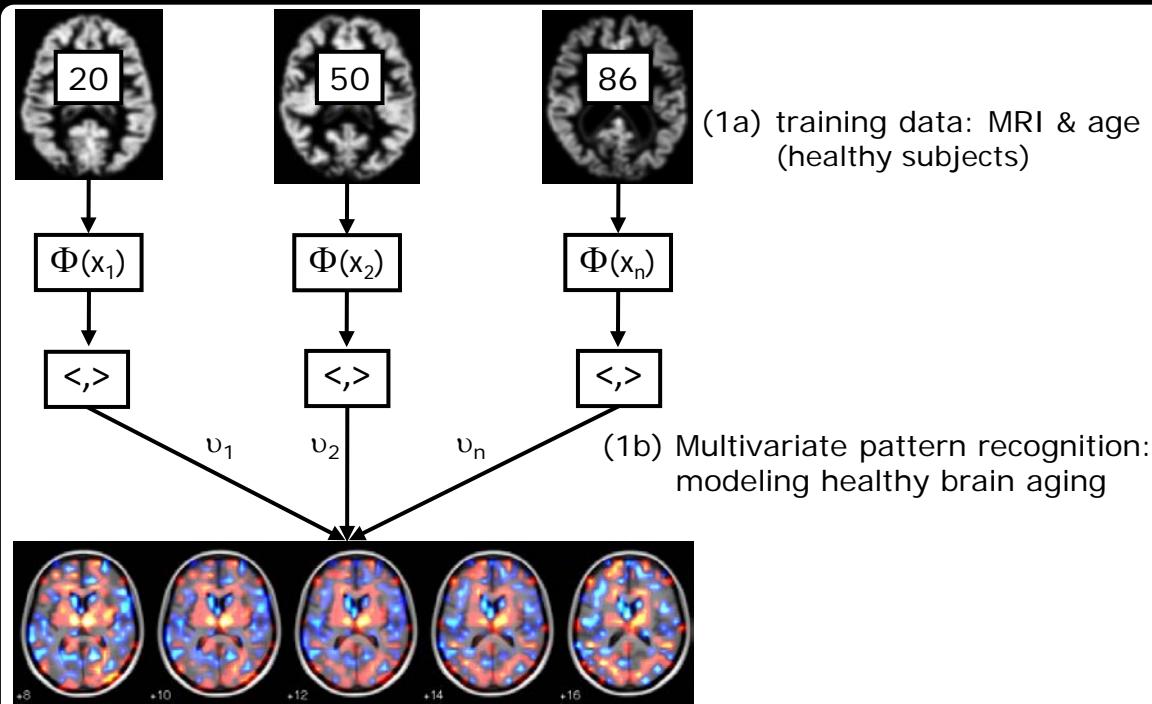
# Preprocessing of MR images

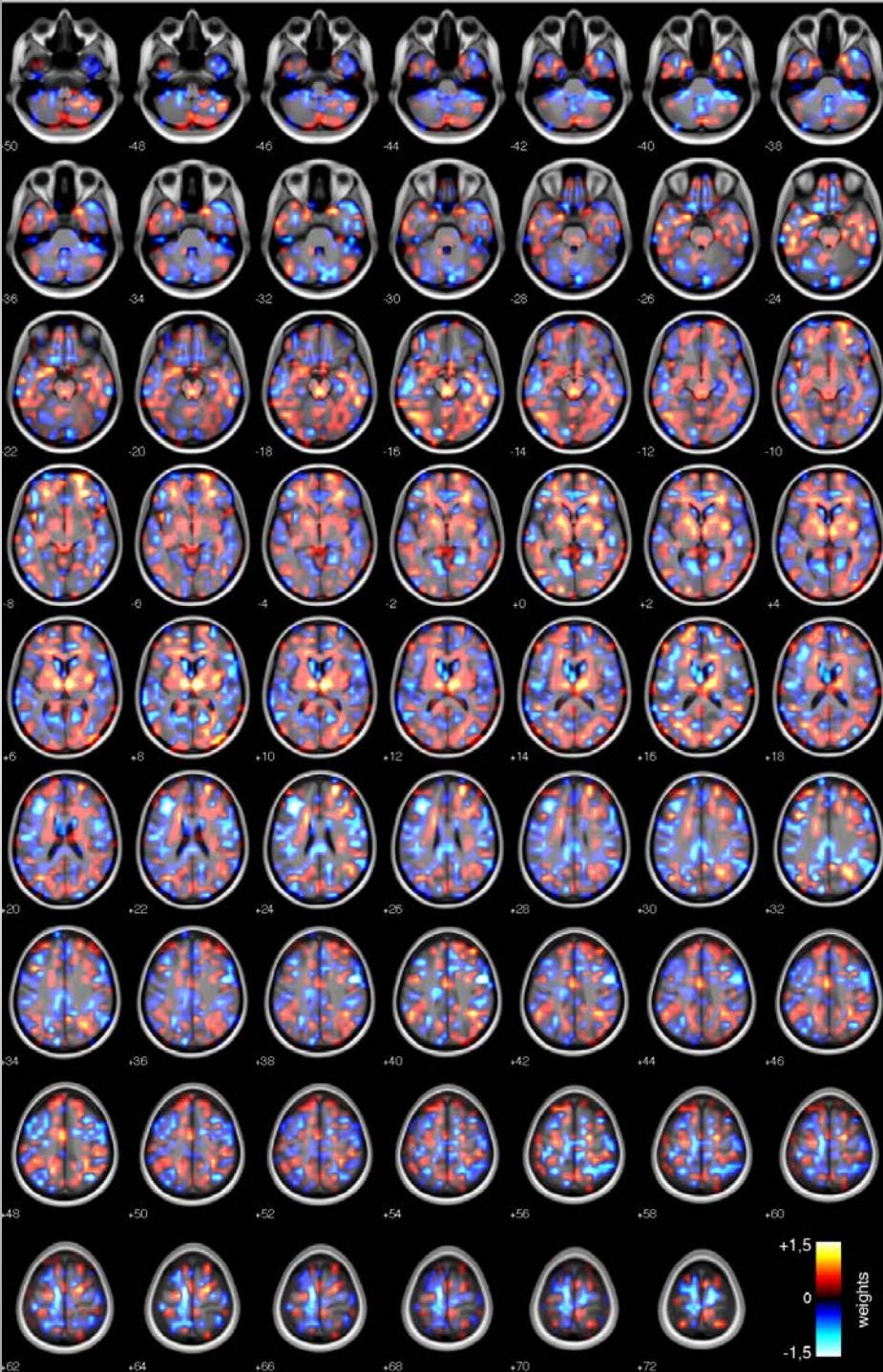


# Identification of individual atrophy patterns

## Modeling of normal brain aging

[with Relevance Vector Regression (RVR; Tipping, 2000)]

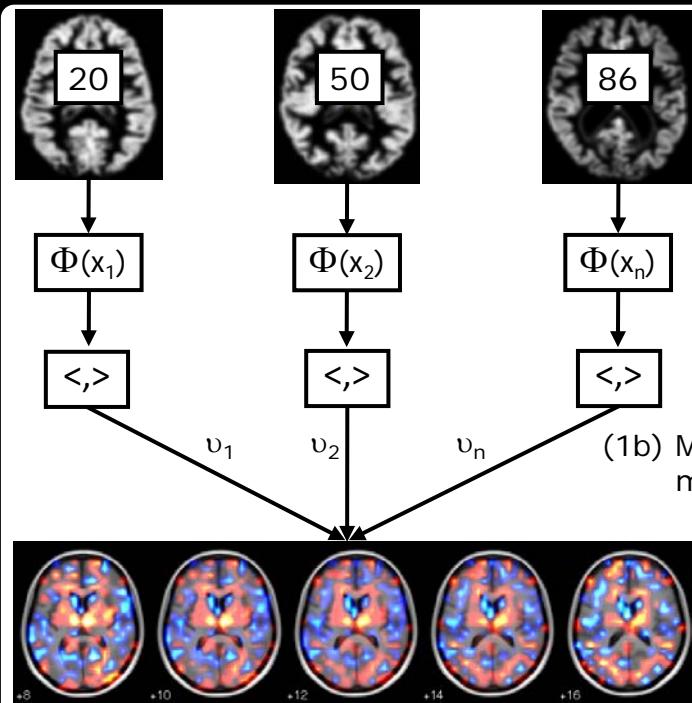




# Identification of individual atrophy patterns

## Modeling of normal brain aging

[with Relevance Vector Regression (RVR; Tipping, 2000)]

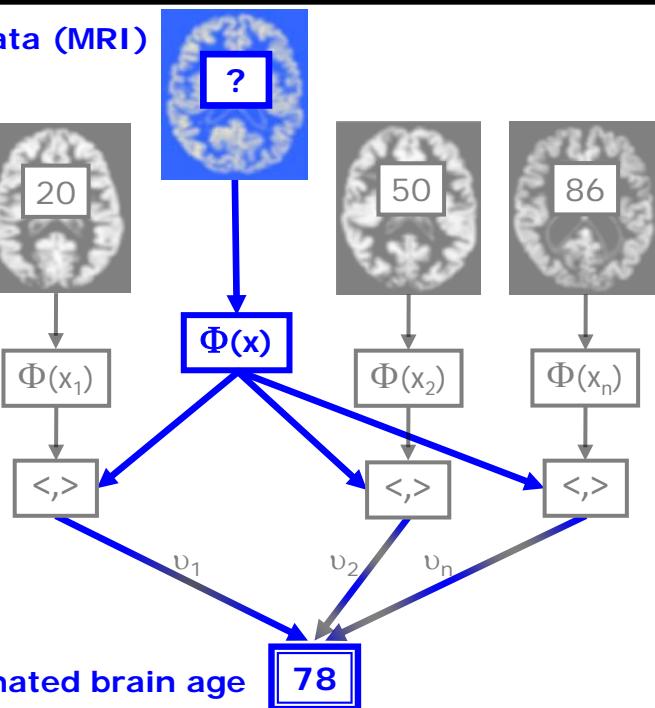


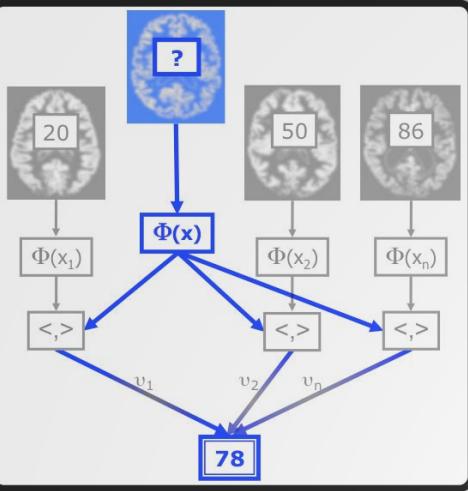
(2a) Input: test data (MRI)

(1a) training data: MRI & age (healthy subjects)

(1b) Multivariate pattern recognition: modeling healthy brain aging

(2b) Output: estimated brain age





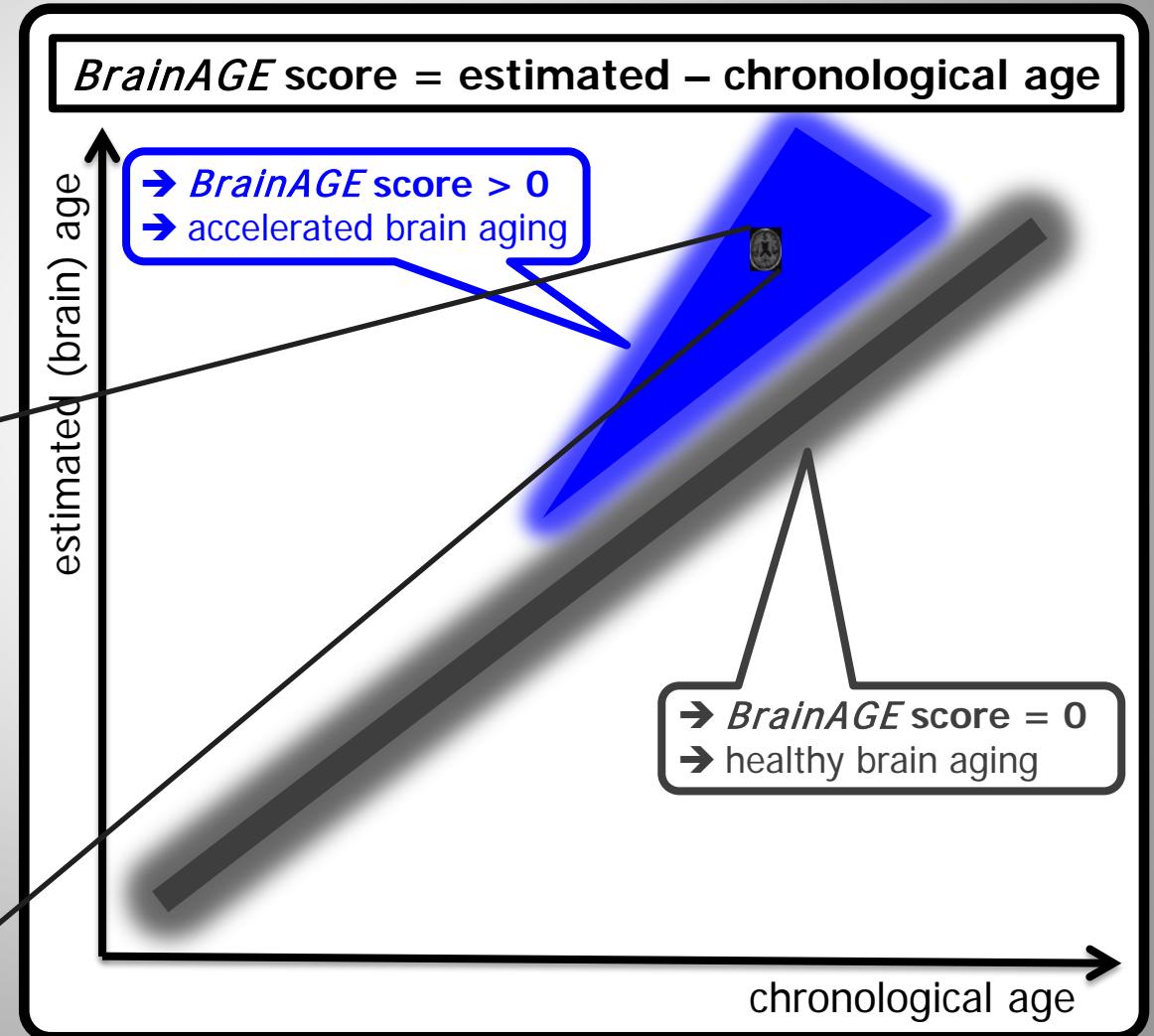
**Deviation from the individual age-related atrophy pattern?**

### Example

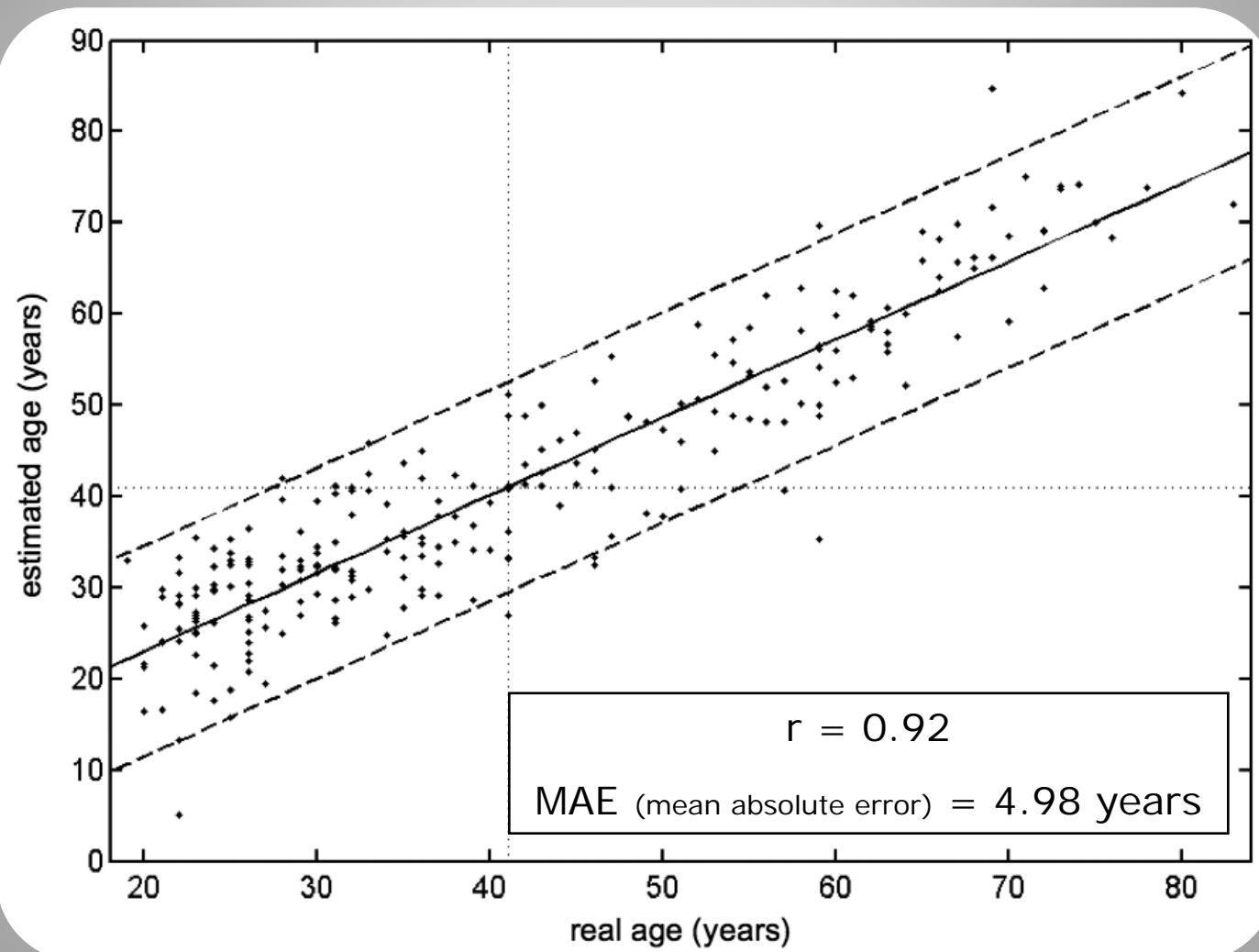
Chronological age: 68 years

Estimated age: 78 years

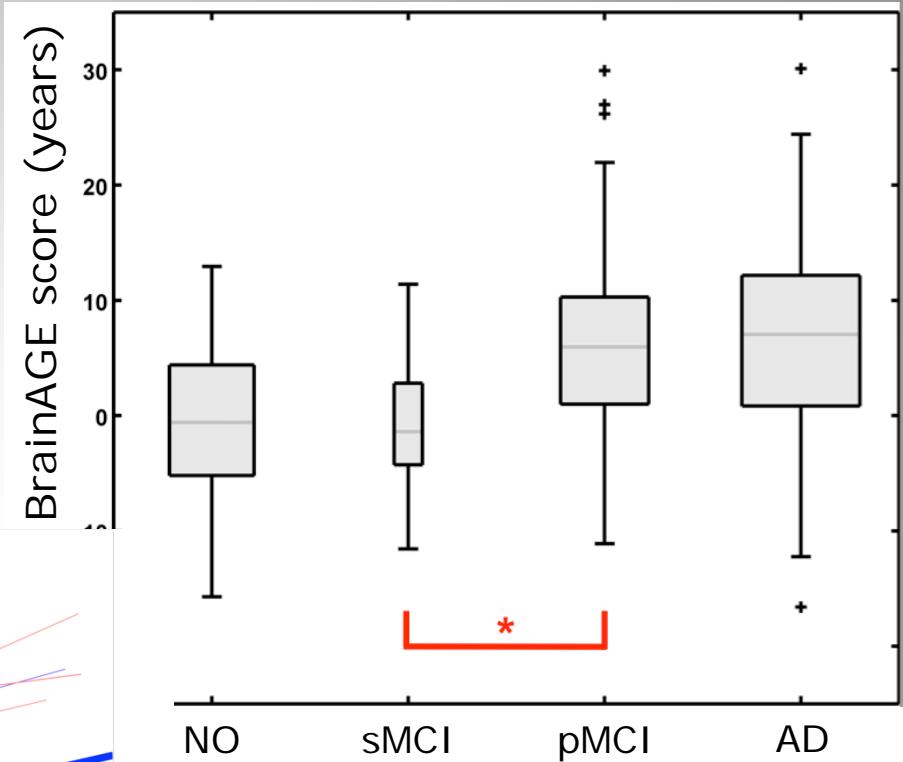
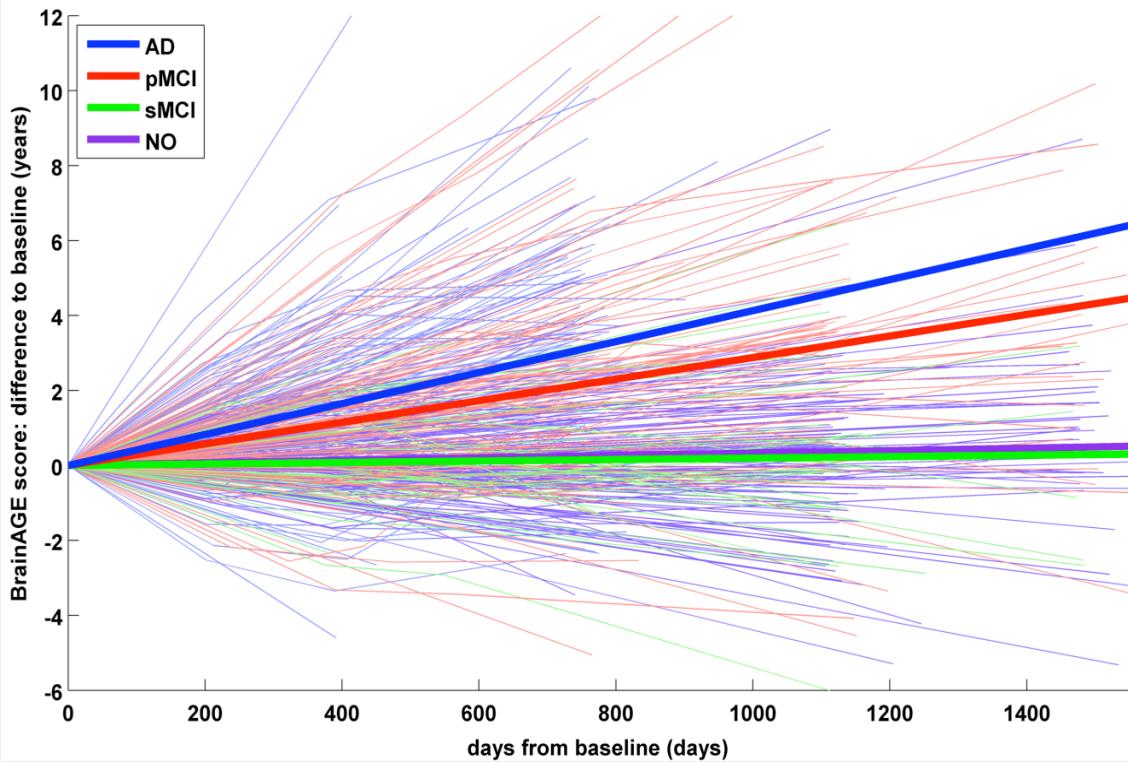
→ BrainAGE score: +10 years



## Reference curve for structural brain aging (Franke et al., 2010)



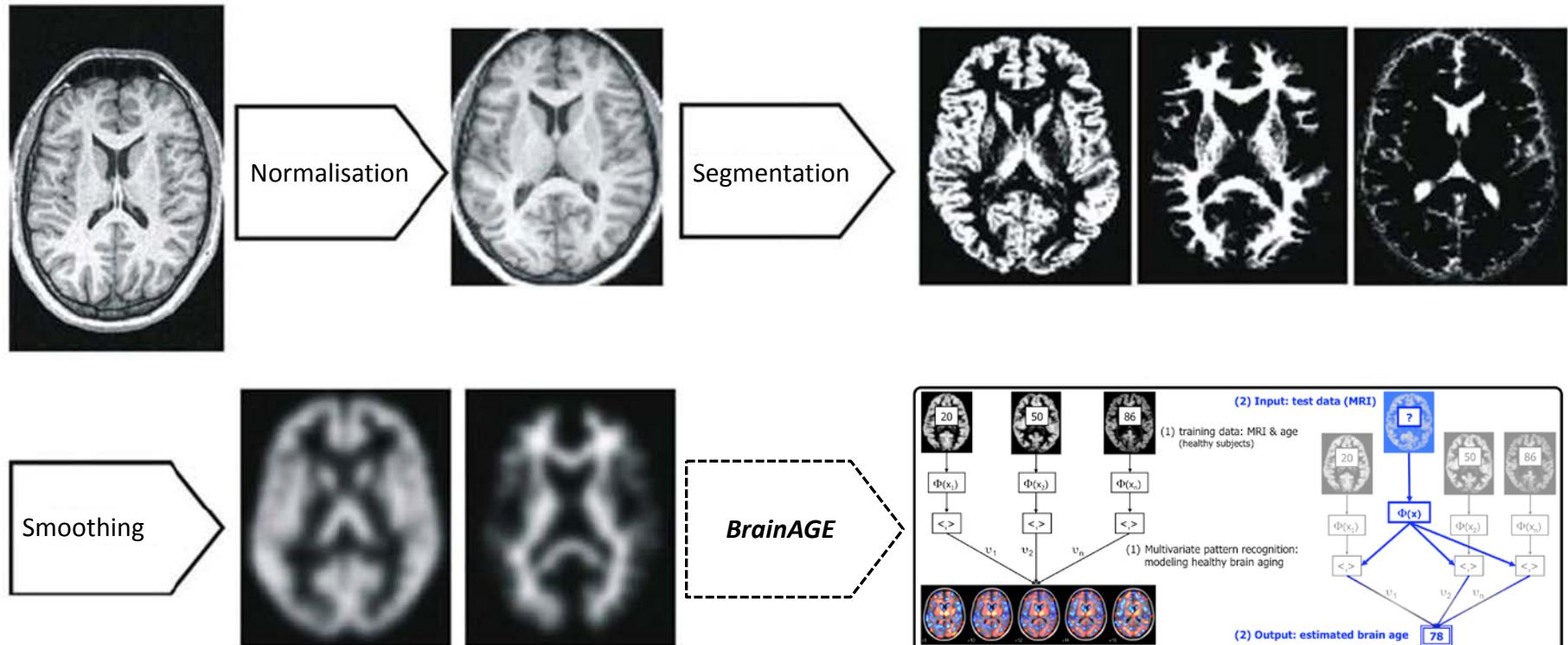
# Advanced brain aging in progressive MCI & AD (Franke & Gaser, 2012)



*BrainAGE score =  
estimated – chronological age*

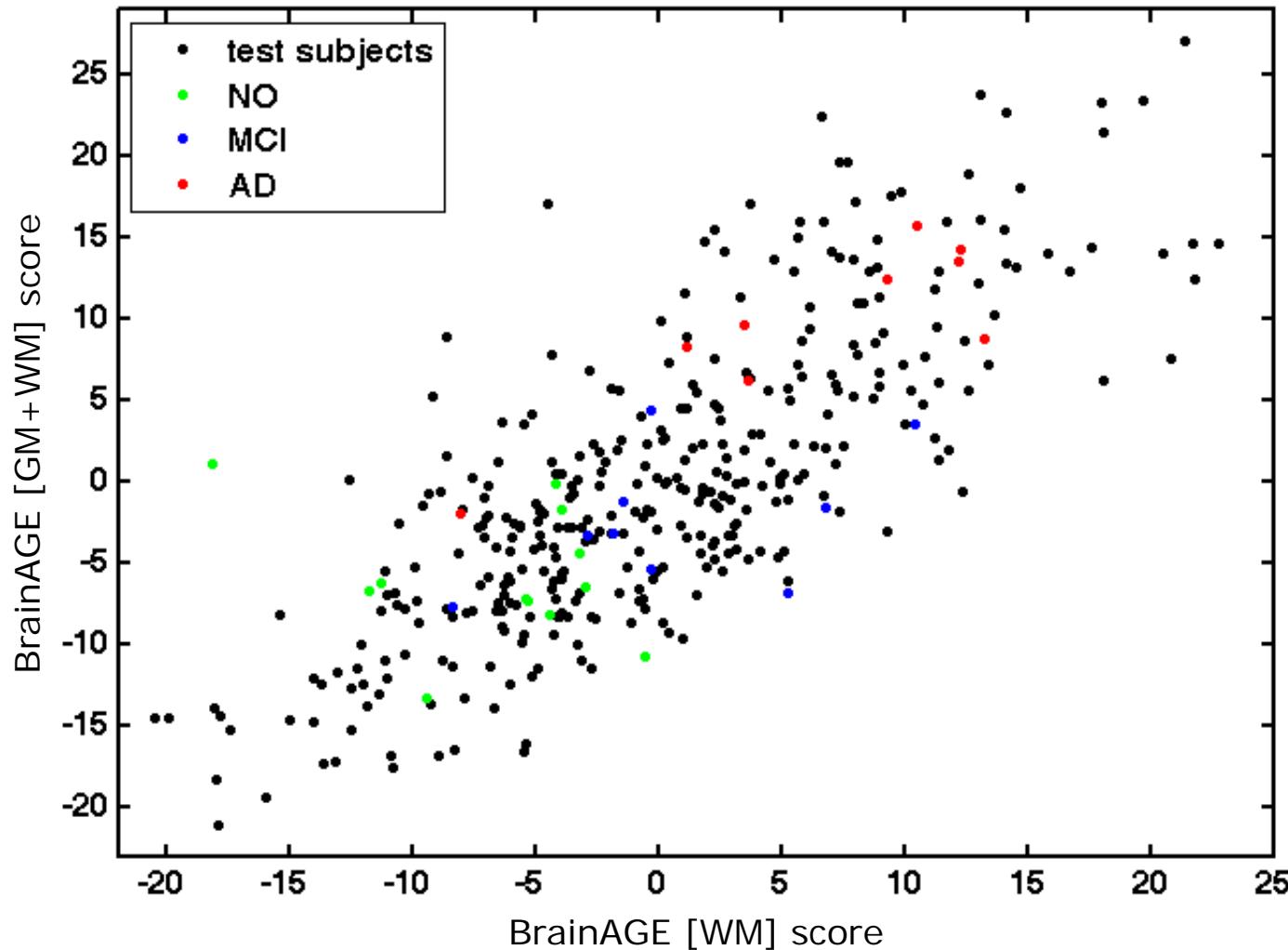
# Pipeline for *BrainAGE* approach in CAD Dementia

- MRI preprocessing with voxel-based morphometry [affine registration, smoothing with **(i)** 4mm & **(ii)** 8mm FWHM smoothing kernels]
- Modeling of structural brain aging with 561 healthy subjects [aged 20-86 years]
- *BrainAGE* estimation with **(i)** linear combination of GM + WM images & **(ii)** WM images



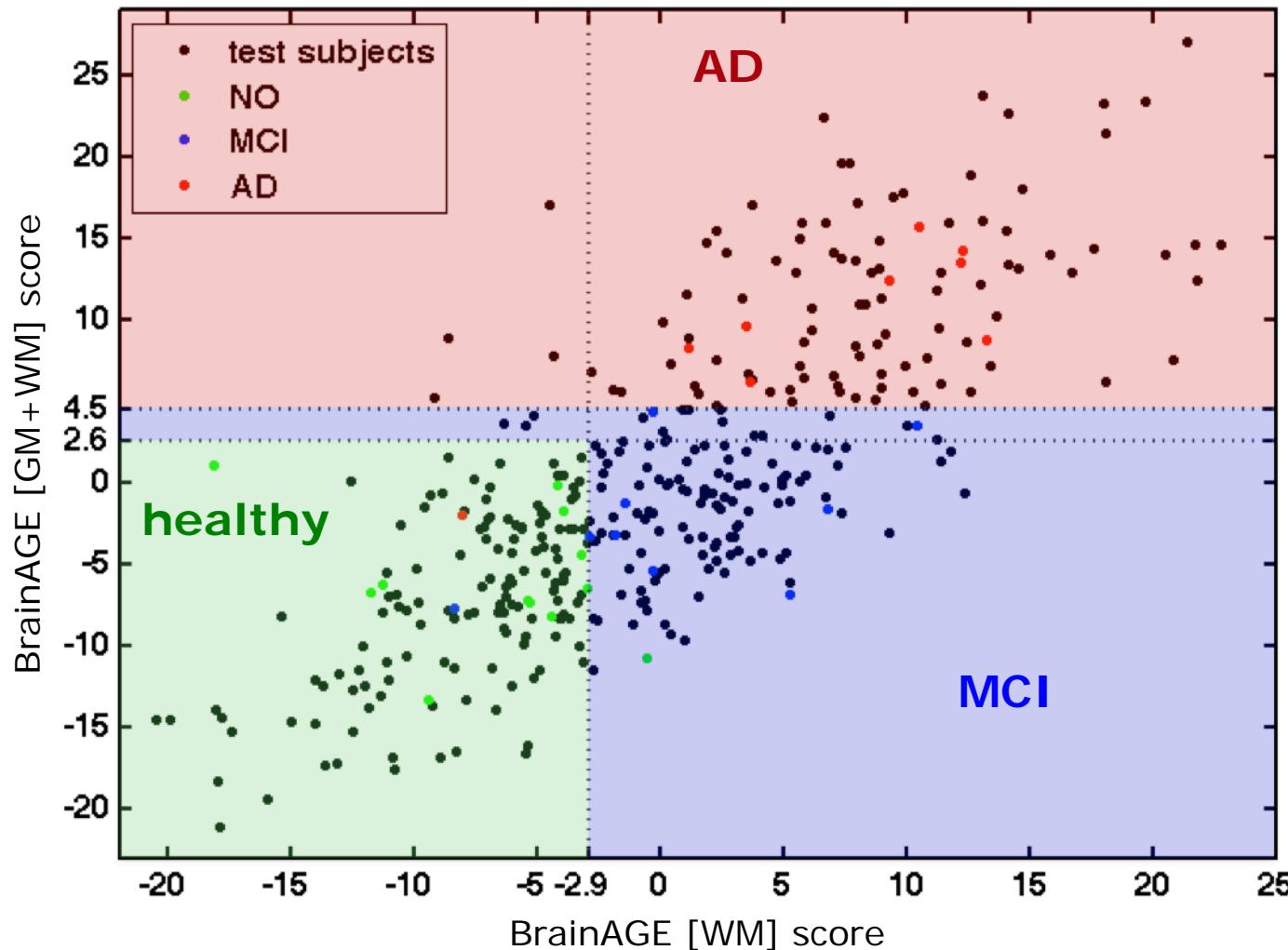
# *BrainAGE scores in CAD Dementia*

[*BrainAGE score = estimated – chronological age*]



# Thresholds for individual classification

[*BrainAGE* score = estimated – chronological age]



→ Classification accuracy in CAD Dementia training sample: **90%**

# Thank you!

This work was funded by the German Ministry for Education and Research (Bundesministerium für Bildung und Forschung – BMBF) [01EV0709 to C.G.; 01GW0740 to K.F.].

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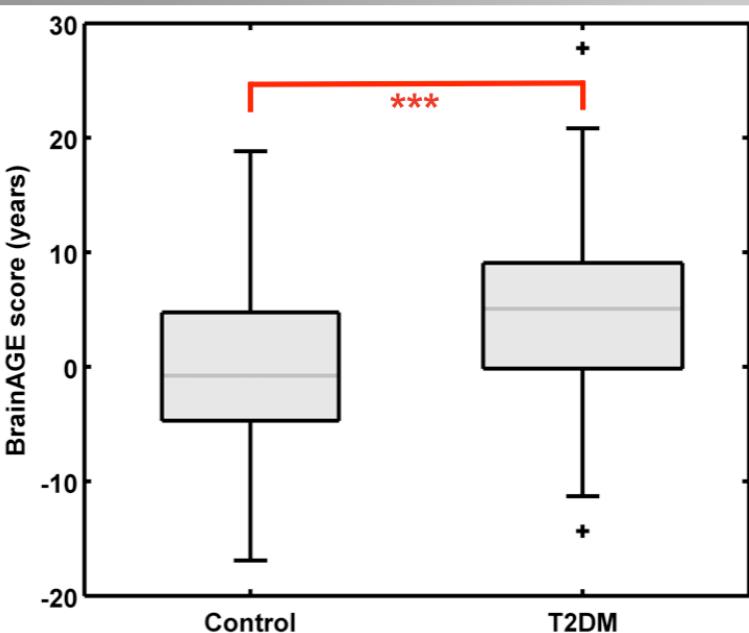
<http://dbm.neuro.uni-jena.de>





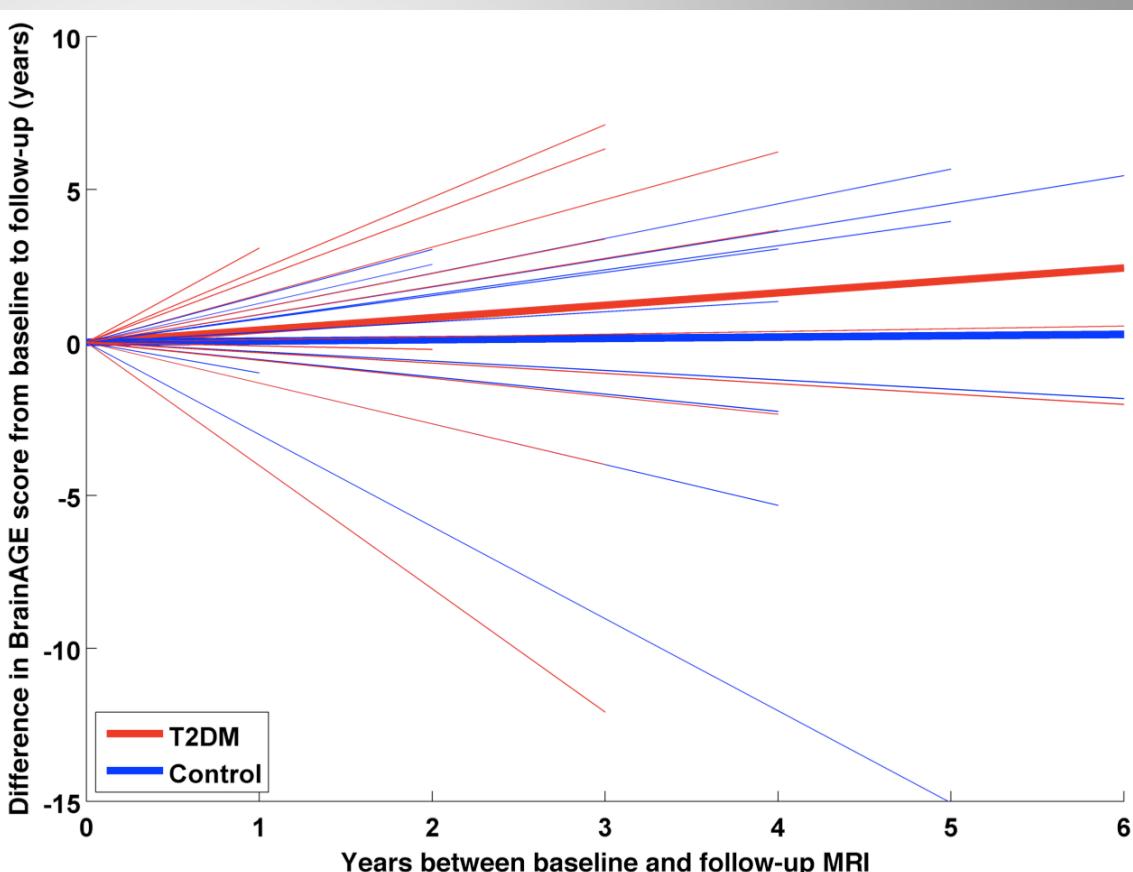
# Anwendung: Advanced BrainAGE in older adults with type 2 diabetes mellitus

(Franke et al., sub.)



Box plots of *BrainAGE* scores for controls and patients with type 2 diabetes mellitus (T2DM). The *BrainAGE* scores differed between groups, with mean ( $\pm$  SD) *BrainAGE* scores of  $0.0 \pm 6.7$  years in healthy controls and  $4.6 \pm 7.2$  years in T2DM subjects ( $P < 0.001$ ). Gray boxes contain the values between the 25th and 75th percentiles of the groups, including the median (dashed line). Lines extending above and below each box symbolize data within 1.5 times the interquartile range (outliers are displayed with a +). Width of the boxes indicates the group size.

Beschleunigte Gehirnalterung in pre-symptomatischen Probanden mit Typ 2 *diabetes mellitus*



Longitudinal changes in *BrainAGE* scores for control subjects (blue) and T2DM patients (red). Thin lines represent individual changes in *BrainAGE* over time; thick lines indicate estimated average changes for each group. The change in *BrainAGE* over time was dependent upon group, providing preliminary longitudinal evidence that T2DM accelerates brain aging.

# Alters-abhängige und Krankheits-spezifische Atrophie bei Alzheimer Demenz

AD-Schweregrad

