Supplementary Material

Differential progression of regional hippocampal atrophy in aging and Parkinson’s disease

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# Supplementary Methods 1

Libraries and functions used for the statistical analysis using R:

|  |  |
| --- | --- |
| *Demographic and clinical variables.**Descriptives* | **psych** library: describe.by() function for means, medians and SD.**stats** library: IQR() function for interquartile range.  |
| *Demographical and clinical variables.**Test stats* | **stats** library: chisq.test() function for chi-squared test; kruskal.test() function pairwise.wilcox.test() for multiple comparisons with Bonferroni P-adjusted. |
| *Multiple linear regression model* | **stats** library: lm() function and anova() to test for differences between segments and total volumes models.**MASS** library: stepAIC() function to fit the best model. |

# Supplementary Methods 2

Multiple linear regression models resulting from stepwise Akaike information criteria:

|  |  |  |
| --- | --- | --- |
|  | **time 1 hippocampal volumes ratios** | **time 2 minus time 1 hippocampal volumes ratios** |
| *Model 1 hippocampal segments* |
| **RAVLT total learning**  | lh CA3 + rh CA4 + bl CA1 + rh subiculum + rh presubiculum + bl HATA + rh hippocampal tail | education + bl CA1 + bl CA3 + rh CA4 + bl subiculum + bl presubiculum + bl parasubiculum + bl HATA + bl fimbria + bl hippocampal tail + bl fissure + bl molecular layer + bl GC-ML-DG |
| **RAVLT recall** | age + education + lh CA3 + rh CA1 + rh CA4 + bl parasubiculum + rh subiculum + rh fimbria + rh hippocampal tail + rh fissure + lh molecular layer | education + lh CA1 + lh CA3 + bl CA4 + lh subiculum + bl presubiculum + lh parasubiculum + lh HATA + bl fimbria + bl hippocampal tail + lh fissure + lh molecular layer + lh GC-ML-DG |
| **RAVLT recognition** | bl CA1 + rh CA4 + bl subiculum + rh HATA + bl hippocampal tail + bl molecular layer | age + education + bl CA1 + bl CA3 + bl CA4 + bl subiculum + bl presubiculum + bl parasubiculum + bl HATA + bl fimbria + bl hippocampal tail + bl fissure + bl molecular layer |
| *Model 2 whole hippocampi* |
| **RAVLT total learning**  | rh whole hippocampus | age + lh whole hippocampus |
| **RAVLT recall** | bl whole hippocampus | age + bl whole hippocampus |
| **RAVLT recognition** | No prediction model | lh whole hippocampus |

# Supplementary Table 1 Means and SD of hippocampal segments

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | **PD-NC****n = 28****mean (SD)** | **PD-MCI****n = 16****mean (SD)** | **PD all sample****n = 44****mean (SD)** | **Controls****n = 21****mean (SD)** |
| *Left hippocampal tail* | **time 1**  | 15.0 (1.5) | 13.8 (1.4) | 14.6 (1.6) | 14.6 (1.6) |
| **time 2**  | 14.8 (1.6) | 13.9 (1.5) | 14.5 (1.6) | 14.6 (1.9) |
| *Right hippocampal tail* | **time 1**  | 15.3 (1.3) | 15.0 (1.6) | 15.2 (1.4) | 15.5 (1.7) |
| **time 2**  | 15.1 (1.1) | 15.0 (1.7) | 15.1 (1.3) | 15.3 (1.8) |
| *Left CA1* | **time 1**  | 17.9 (0.9) | 18.7 (0.9) | 18.2 (1.0) | 18.3 (1.0) |
| **time 2**  | 18.3 (1.0) | 18.8 (0.9) | 18.5 (1.0) | 18.3 (1.0) |
| *Right CA1* | **time 1**  | 18.2 (1.1) | 18.6 (0.7) | 18.4 (1.0) | 18.1 (0.9) |
| **time 2**  | 18.5 (1.0) | 18.7 (1.0) | 18.6 (0.9) | 18.2 (1.1) |
| *Left CA3* | **time 1**  | 6.3 (0.6) | 6.4 (0.6) | 6.3 (0.6) | 6.7 (0.7) |
| **time 2**  | 6.4 (0.6) | 6.4 (0.7) | 6.4 (0.6) | 6.7 (0.7) |
| *Right CA3* | **time 1**  | 6.7 (0.7) | 6.8 (0.5) | 6.7 (0.6) | 6.8 (0.7) |
| **time 2**  | 6.8 (0.6) | 6.7 (0.5) | 6.8 (0.6) | 6.9 (0.8) |
| *Left CA4* | **time 1**  | 8.4 (0.5) | 8.1 (0.5) | 8.3 (0.5) | 8.4 (0.5) |
| **time 2**  | 8.4 (0.5) | 8.2 (0.6) | 8.3 (0.6) | 8.4 (0.5) |
| *Right CA4* | **time 1**  | 8.6 (0.5) | 8.3 (0.5) | 8.5 (0.5) | 8.6 (0.5) |
| **time 2**  | 8.7 (0.5) | 8.4 (0.4) | 8.6 (0.5) | 8.6 (0.5) |
| *Left subiculum* | **time 1**  | 11.9 (0.9) | 11.9 (0.7) | 11.9 (0.8) | 11.6 (0.7) |
| **time 2**  | 11.8 (0.9) | 12.0 (0.8) | 11.9 (0.9) | 11.6 (0.8) |
| *Right subiculum* | **time 1**  | 11.7 (0.8) | 11.7 (0.7) | 11.7 (0.8) | 11.5 (0.7) |
| **time 2**  | 11.7 (0.9) | 11.8 (0.8) | 11.7 (0.8) | 11.5 (0.7) |
| *Left presubiculum* | **time 1**  | 8.5 (1.1) | 8.9 (0.9) | 8.6 (1.0) | 8.4 (0.9) |
| **time 2**  | 8.4 (1.1) | 8.8 (0.8) | 8.5 (1.1) | 8.4 (0.8) |
| *Right presubiculum* | **time 1**  | 7.8 (1.0) | 7.9 (0.7) | 7.8 (0.9) | 7.8 (0.7) |
| **time 2**  | 7.7 (0.9) | 7.8 (0.8) | 7.7 (0.9) | 7.7 (0.8) |
| *Left parasubiculum* | **time 1**  | 1.7 (0.4) | 2.0 (0.5) | 1.8 (0.4) | 1.7 (0.5) |
| **time 2**  | 1.7 (0.4) | 2.1 (0.6) | 1.9 (0.5) | 1.7 (0.5) |
| *Right parasubiculum* | **time 1**  | 1.6 (0.3) | 1.6 (0.4) | 1.6 (0.3) | 1.6 (0.5) |
| **time 2**  | 1.6 (0.3) | 1.7 (0.4) | 1.6 (0.4) | 1.6 (0.5) |
| *Left fimbria* | **time 1**  | 2.3 (0.6) | 2.3 (0.5) | 2.3 (0.6) | 2.1 (0.4) |
| **time 2**  | 2.1 (0.6) | 2.2 (0.6) | 2.2 (0.6) | 2.1 (0.3) |
| *Right fimbria* | **time 1**  | 1.8 (0.5) | 2.0 (0.5) | 1.8 (0.5) | 1.8 (0.5) |
| **time 2**  | 1.7 (0.5) | 1.8 (0.6) | 1.7 (0.5) | 1.8 (0.5) |
| *Left fissure* | **time 1**  | 4.5 (0.9) | 4.9 (1.0) | 4.6 (0.9) | 4.8 (0.9) |
| **time 2**  | 4.7 (1.0) | 5.3 (1.2) | 4.9 (1.1) | 5.1 (0.8) |
| *Right fissure* | **time 1**  | 4.3 (0.9) | 5.1 (1.2) | 4.6 (1.1) | 4.8 (0.9) |
| **time 2**  | 4.9 (1.0) | 5.5 (1.1) | 5.1 (1.1) | 5.1 (0.9) |
| *Left HATA* | **time 1**  | 1.6 (0.2) | 1.6 (0.2) | 1.6 (0.2) | 1.7 (0.2) |
| **time 2**  | 1.6 (0.2) | 1.6 (0.3) | 1.6 (0.3) | 1.7 (0.2) |
| *Right HATA* | **time 1**  | 1.6 (0.2) | 1.6 (0.2) | 1.6 (1.2) | 1.6 (0.2) |
| **time 2**  | 1.6 (0.2) | 1.6 (0.3) | 1.6 (0.2) | 1.6 (0.3) |
| *Left molecular layer* | **time 1**  | 17.2 (0.4) | 17.3 (0.5) | 17.2 (0.5) | 17.3 (0.4) |
| **time 2**  | 17.2 (0.4) | 17.1 (0.5) | 17.2 (0.5) | 17.2 (0.4) |
| *Right molecular layer* | **time 1**  | 17.3 (0.5) | 17.4 (0.6) | 17.3 (0.6) | 17.3 (0.5) |
| **time 2**  | 17.3 (0.5) | 17.3 (0.5) | 17.3 (0.5) | 17.2 (0.5) |
| *Left GC-ML-DG* | **time 1**  | 9.2 (0.5) | 9.0 (0.4) | 9.1 (0.5) | 9.3 (0.5) |
| **time 2**  | 9.2 (0.5) | 9.0 (0.5) | 9.1 (0.5) | 9.3 (0.5) |
| *Right GC-ML-DG* | **time 1**  | 9.5 (0.5) | 9.2 (0.5) | 9.3 (0.5) | 9.4 (0.5) |
| **time 2**  | 9.5 (0.5) | 9.2 (0.5) | 9.4 (0.5) | 9.4 (0.6) |
| *Left whole hippocampus* | **time 1**  | 0.3 (0.0) | 0.2 (0.0) | 0.2 (0.0) | 0.2 (0.0) |
| **time 2**  | 0.2 (0.0) | 0.2 (0.0) | 0.2 (0.0) | 0.2 (0.0) |
| *Right whole hippocampus* | **time 1**  | 0.3 (0.0) | 0.2 (0.0) | 0.3 (0.0) | 0.3 (0.0) |
| **time 2**  | 0.3 (0.0) | 0.2 (0.0) | 0.2 (0.0) | 0.2 (0.0) |

CA, Cornu Ammonis; GC-ML-DG, Granule cells in the molecular layer of the dentate gyrus; HATA, Hippocampal Amygdala Transition Area; PD-MCI, Parkinson’s disease mild cognitive impairment; PD-NC, Parkinson’s disease normal cognition; SD, standard deviation.

Values are volumes in mm3. Hippocampal segments are ratios to whole hippocampal volumes ((lh or rh segment / lh or rh whole hippocampus) \* 100). Whole hippocampus volumes are ratios to estimated total intracranial volume ((lh or rh whole hippocampus / eTIV) \* 100).

# Supplementary Table 2 Multiple regression results of time 1 hippocampal volumes ratios as predictors of memory performance change over time

|  |  |
| --- | --- |
| *Model 1 hippocampal segments* | *Model 2 whole hippocampi* |
|  | **variables** | **t-stat (P-value)** | **variables** | **t-stat (P-value)** |
| RAVLT total learning | Non-significant model | right hippocampus | **2.841 (0.007)** |
| RAVLT recall | age | **-2.310 (0.028)** | left hippocampus | **-2.610 (0.013)** |
| education | -1.707 (0.098) | right hippocampus | **3.346 (0.002)** |
| left CA3 | **-3.040 (0.005)** |  |
| right CA1 | -1.428 (0.163) |
| right CA4 | **-3.561 (0.001)** |
| left parasubiculum | **-5.825 (<0.001)** |
| right subiculum | **-3.147 (0.004)** |
| right parasubiculum | -0.640 (0.527) |
| right fimbria | **-4.831 (<0.001)** |
| right hippocampal tail | **-5.798 (0.001)** |
| right fissure | **3.195 (0.003)** |
| left molecular layer | **-3.491 (0.001)** |
| RAVLT recognition | Non-significant model | No prediction model |

CA, Cornu Ammonis; RAVLT, Rey’s Auditory Verbal Learning test.

Regions that reached significance P<0.05 are marked in bold.

# Supplementary Table 3 Multiple regression results of time 2 minus time 1 hippocampal volumes ratios as explanatory variables of memory changes

|  |  |
| --- | --- |
| *Model 1 hippocampal segments* | *Model 2 whole hippocampi* |
|  | **variables** | **t-stat (P-value)** | **variables** | **t-stat (P-value)** |
| **RAVLT total learning** | Non-significant model |  | age | -1.600 (0.117) |
| left hippocampus | 1.688 (0.099) |
| **RAVLT recall** | education | 1.299 (0.205) | age | -1.376 (0.176) |
| left CA1 | 1.211 (0.237) | left hippocampus | **2.151 (0.038)** |
| left CA3 | 1.211 (0.237) | right hippocampus | -1.556 (0.128) |
| left CA4 | 1.211 (0.237) |  |
| right CA4 | 1.696 (0.102) |
| left subiculum | 1.211 (0.237) |
| left presubiculum | 1.211 (0.237) |
| left parasubiculum | 1.211 (0.237) |
| right presubiculum | 1.123 (0.272) |
| left HATA | 1.211 (0.237) |
| left fimbria | 1.211 (0.237) |
| right fimbria | **2.650 (0.014)** |
| left hippocampal tail | 1.211 (0.237) |
| right hippocampal tail | **2.986 (0.006)** |
| left fissure | **-2.888 (0.008)** |
| left molecular layer | 1.211 (0.237) |
| left GC-ML-DG | 1.211 (0.237) |
| **RAVLT recognition** | Non-significant model | left hippocampus | **2.567 (0.014)** |

CA, Cornu Ammonis; GC-ML-DG, Granule cells in the molecular layer of the dentate gyrus; HATA, Hippocampal Amygdala Transition Area; RAVLT, Rey’s Auditory Verbal Learning test.

Regions that reached significance P<0.05 are marked in bold.