**Supplementary**

**Table 1**  Correlation analysis results in the DIP group

|  |  |  |
| --- | --- | --- |
| **r value** | 　 | **p value** |
| 　 | MoCa | HAMD | HAMA | UPDRS I | UPDRS II | UPDRS III |  | 　 | MoCa | HAMD | HAMA | UPDRS I | UPDRS II | UPDRS III |
| L\_Whole\_hippocampus | 0.57 | -0.49 | -0.2 | -0.5 | -0.6 | -0.49 | 　 | L\_Whole\_hippocampus | 0.022 | 0.056 | 0.449 | 0.05 | 0.013 | 0.052 |
| L\_parasubiculum | -0.21 | -0.12 | -0.17 | -0.07 | 0.5 | 0.42 |  | L\_parasubiculum | 0.425 | 0.651 | 0.529 | 0.794 | 0.048 | 0.107 |
| L\_presubiculum | 0.39 | -0.25 | -0.31 | -0.52 | -0.56 | -0.51 |  | L\_presubiculum | 0.136 | 0.356 | 0.246 | 0.04 | 0.023 | 0.045 |
| L\_subiculum | 0.58 | -0.13 | -0.2 | -0.41 | -0.7 | -0.62 |  | L\_subiculum | 0.019 | 0.643 | 0.456 | 0.118 | 0.002 | 0.01 |
| L\_CA1 | 0.31 | -0.5 | -0.13 | -0.52 | -0.46 | -0.23 |  | L\_CA1 | 0.246 | 0.051 | 0.627 | 0.038 | 0.07 | 0.394 |
| L\_CA3 | 0.46 | -0.56 | -0.06 | -0.43 | -0.56 | -0.33 |  | L\_CA3 | 0.071 | 0.026 | 0.82 | 0.094 | 0.025 | 0.214 |
| L\_CA4 | 0.52 | -0.52 | -0.19 | -0.51 | -0.61 | -0.45 |  | L\_CA4 | 0.038 | 0.04 | 0.479 | 0.042 | 0.011 | 0.077 |
| L\_molecular\_layer\_HP | 0.54 | -0.37 | -0.11 | -0.48 | -0.7 | -0.54 |  | L\_molecular\_layer\_HP | 0.031 | 0.158 | 0.679 | 0.059 | 0.003 | 0.03 |
| L\_ GC-ML-DG | 0.53 | -0.55 | -0.23 | -0.48 | -0.56 | -0.5 |  | L\_ GC-ML-DG | 0.033 | 0.029 | 0.394 | 0.057 | 0.025 | 0.05 |
| L\_HATA | 0.09 | -0.63 | -0.32 | -0.59 | -0.5 | -0.34 |  | L\_HATA | 0.736 | 0.008 | 0.227 | 0.016 | 0.05 | 0.193 |
| L\_fimbria | 0.41 | 0 | -0.04 | 0.18 | -0.38 | -0.65 |  | L\_fimbria | 0.113 | 0.986 | 0.877 | 0.513 | 0.151 | 0.007 |
| L\_Hippocampal\_tail | 0.53 | -0.53 | -0.13 | -0.25 | -0.21 | -0.23 |  | L\_Hippocampal\_tail | 0.035 | 0.036 | 0.637 | 0.359 | 0.426 | 0.382 |
| L\_hippocampalfissure | 0.02 | 0.03 | -0.02 | -0.33 | -0.64 | -0.25 |  | L\_hippocampalfissure | 0.956 | 0.903 | 0.937 | 0.214 | 0.007 | 0.355 |
| R\_Whole\_hippocampus | 0.69 | -0.35 | -0.09 | -0.32 | -0.57 | -0.6 |  | R\_Whole\_hippocampus | 0.003 | 0.184 | 0.751 | 0.23 | 0.02 | 0.015 |
| R\_parasubiculum | -0.06 | 0.02 | 0.1 | -0.04 | 0.5 | 0.43 |  | R\_parasubiculum | 0.815 | 0.952 | 0.725 | 0.885 | 0.05 | 0.101 |
| R\_presubiculum | 0.45 | -0.21 | -0.14 | -0.41 | -0.37 | -0.48 |  | R\_presubiculum | 0.082 | 0.427 | 0.613 | 0.111 | 0.155 | 0.059 |
| R\_subiculum | 0.56 | -0.13 | -0.1 | -0.4 | -0.63 | -0.47 |  | R\_subiculum | 0.025 | 0.625 | 0.722 | 0.129 | 0.008 | 0.064 |
| R\_CA1 | 0.58 | -0.27 | 0 | -0.22 | -0.6 | -0.45 |  | R\_CA1 | 0.018 | 0.31 | 0.997 | 0.414 | 0.013 | 0.082 |
| R\_CA3 | 0.46 | -0.32 | 0.17 | -0.01 | -0.34 | -0.25 |  | R\_CA3 | 0.072 | 0.233 | 0.522 | 0.97 | 0.196 | 0.344 |
| R\_CA4 | 0.59 | -0.33 | 0.12 | -0.16 | -0.52 | -0.42 |  | R\_CA4 | 0.017 | 0.218 | 0.645 | 0.554 | 0.04 | 0.105 |
| R\_molecular\_layer\_HP | 0.26 | -0.42 | -0.53 | -0.23 | 0.06 | -0.29 |  | R\_molecular\_layer\_HP | 0.338 | 0.109 | 0.037 | 0.398 | 0.824 | 0.272 |
| R\_GC-ML-DG | 0.65 | -0.33 | 0.12 | -0.16 | -0.51 | -0.48 |  | R\_GC-ML-DG | 0.007 | 0.205 | 0.663 | 0.559 | 0.046 | 0.061 |
| R\_HATA | 0.29 | -0.35 | -0.18 | -0.16 | -0.14 | -0.33 |  | R\_HATA | 0.276 | 0.182 | 0.502 | 0.563 | 0.593 | 0.219 |
| R\_fimbria | 0.38 | 0.12 | 0.12 | 0.2 | -0.38 | -0.72 |  | R\_fimbria | 0.145 | 0.651 | 0.663 | 0.45 | 0.152 | 0.002 |
| R\_Hippocampal\_tail | 0.67 | -0.27 | 0.04 | -0.31 | -0.65 | -0.58 |  | R\_Hippocampal\_tail | 0.005 | 0.308 | 0.897 | 0.237 | 0.007 | 0.019 |
| R\_hippocampalfissure | -0.13 | 0.57 | 0.2 | 0.28 | -0.37 | -0.12 | 　 | R\_hippocampalfissure | 0.639 | 0.021 | 0.448 | 0.286 | 0.156 | 0.648 |

**Table 2** Correlation analysis results in the PD group

|  |  |  |
| --- | --- | --- |
| **r value** | 　 | **p value** |
| 　 | MoCa | HAMD | HAMA | UPDRS I | UPDRS II | UPDRS III |  | 　 | MoCa | HAMD | HAMA | UPDRS I | UPDRS II | UPDRS III |
| L\_Whole\_hippocampus | 0.72 | -0.42 | 0.23 | 0.38 | -0.17 | -0.27 | 　 | L\_Whole\_hippocampus | 0.001 | 0.092 | 0.377 | 0.134 | 0.518 | 0.286 |
| L\_parasubiculum | 0.27 | -0.04 | 0.27 | 0.3 | -0.1 | -0.24 |  | L\_parasubiculum | 0.288 | 0.875 | 0.288 | 0.25 | 0.69 | 0.356 |
| L\_presubiculum | 0.44 | -0.25 | 0.2 | 0.31 | -0.18 | -0.08 |  | L\_presubiculum | 0.075 | 0.328 | 0.436 | 0.223 | 0.494 | 0.771 |
| L\_subiculum | 0.73 | -0.24 | 0.32 | 0.47 | -0.02 | -0.07 |  | L\_subiculum | 0.001 | 0.345 | 0.207 | 0.058 | 0.933 | 0.776 |
| L\_CA1 | 0.71 | -0.32 | 0.24 | 0.43 | -0.12 | -0.29 |  | L\_CA1 | 0.001 | 0.211 | 0.357 | 0.084 | 0.639 | 0.262 |
| L\_CA3 | 0.51 | -0.18 | 0.16 | 0.28 | 0.03 | -0.24 |  | L\_CA3 | 0.037 | 0.487 | 0.527 | 0.285 | 0.921 | 0.346 |
| L\_CA4 | 0.7 | -0.39 | 0.2 | 0.39 | -0.14 | -0.34 |  | L\_CA4 | 0.002 | 0.123 | 0.436 | 0.12 | 0.583 | 0.184 |
| L\_molecular\_layer\_HP | 0.73 | -0.32 | 0.3 | 0.48 | -0.04 | -0.16 |  | L\_molecular\_layer\_HP | 0.001 | 0.207 | 0.239 | 0.049 | 0.87 | 0.54 |
| L\_GC-ML-DG | 0.68 | -0.39 | 0.16 | 0.37 | -0.18 | -0.36 |  | L\_GC-ML-DG | 0.003 | 0.118 | 0.542 | 0.148 | 0.489 | 0.159 |
| L\_HATA | 0.45 | -0.26 | 0.04 | 0.11 | -0.31 | -0.46 |  | L\_HATA | 0.07 | 0.316 | 0.88 | 0.672 | 0.233 | 0.064 |
| L\_fimbria | 0.42 | -0.53 | -0.39 | -0.31 | -0.25 | -0.33 |  | L\_fimbria | 0.089 | 0.028 | 0.117 | 0.226 | 0.327 | 0.201 |
| L\_Hippocampal\_tail | 0.32 | -0.56 | 0.05 | 0.01 | -0.28 | -0.26 |  | L\_Hippocampal\_tail | 0.213 | 0.02 | 0.862 | 0.956 | 0.282 | 0.305 |
| L\_hippocampalfissure | 0.68 | -0.32 | 0.29 | 0.28 | -0.1 | -0.5 |  | L\_hippocampalfissure | 0.003 | 0.213 | 0.251 | 0.27 | 0.694 | 0.039 |
| R\_Whole\_hippocampus | 0.68 | -0.31 | 0.4 | 0.44 | -0.13 | -0.24 |  | R\_Whole\_hippocampus | 0.003 | 0.233 | 0.11 | 0.078 | 0.619 | 0.347 |
| R\_parasubiculum | 0.33 | 0.01 | 0.4 | 0.25 | -0.22 | -0.25 |  | R\_parasubiculum | 0.203 | 0.971 | 0.111 | 0.337 | 0.397 | 0.325 |
| R\_presubiculum | 0.44 | 0.18 | 0.62 | 0.62 | 0 | -0.08 |  | R\_presubiculum | 0.075 | 0.48 | 0.007 | 0.008 | 0.991 | 0.747 |
| R\_subiculum | 0.71 | -0.06 | 0.5 | 0.6 | 0.2 | -0.01 |  | R\_subiculum | 0.001 | 0.809 | 0.043 | 0.011 | 0.445 | 0.96 |
| R\_CA1 | 0.74 | -0.3 | 0.43 | 0.43 | -0.13 | -0.36 |  | R\_CA1 | 0.001 | 0.25 | 0.088 | 0.083 | 0.618 | 0.152 |
| R\_CA3 | 0.37 | -0.58 | -0.15 | -0.05 | -0.25 | -0.17 |  | R\_CA3 | 0.143 | 0.015 | 0.571 | 0.835 | 0.33 | 0.509 |
| R\_CA4 | 0.63 | -0.4 | 0.24 | 0.36 | -0.18 | -0.22 |  | R\_CA4 | 0.006 | 0.112 | 0.344 | 0.157 | 0.488 | 0.388 |
| R\_molecular\_layer\_HP | 0.41 | -0.4 | 0.25 | 0.19 | -0.33 | -0.3 |  | R\_molecular\_layer\_HP | 0.105 | 0.113 | 0.332 | 0.472 | 0.193 | 0.247 |
| R\_GC-ML-DG | 0.64 | -0.41 | 0.2 | 0.38 | -0.17 | -0.2 |  | R\_GC-ML-DG | 0.005 | 0.098 | 0.446 | 0.134 | 0.524 | 0.431 |
| R\_HATA | 0.65 | -0.19 | 0.33 | 0.51 | -0.01 | -0.13 |  | R\_HATA | 0.004 | 0.474 | 0.192 | 0.038 | 0.956 | 0.616 |
| R\_fimbria | 0.13 | -0.37 | -0.4 | -0.32 | 0 | -0.03 |  | R\_fimbria | 0.61 | 0.138 | 0.107 | 0.213 | 0.987 | 0.919 |
| R\_Hippocampal\_tail | 0.68 | -0.28 | 0.43 | 0.48 | -0.01 | -0.17 |  | R\_Hippocampal\_tail | 0.003 | 0.28 | 0.087 | 0.053 | 0.978 | 0.505 |
| R\_hippocampalfissure | 0.56 | -0.04 | 0.45 | 0.35 | 0.03 | -0.36 | 　 | R\_hippocampalfissure | 0.021 | 0.872 | 0.071 | 0.166 | 0.915 | 0.156 |

**Table 3** Comparison of hippocampal volumes across different H-Y stages in DIP groups.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  |  |  |  |  |  |  |
| 　 | Grage 1(N=6) | 　 | Grade ≥ 2(N=13) | 　 | F value | p value |
|  | mean | sd |  | mean | sd |  |
| L\_Whole\_hippocampus | 3144.422  | 319.284  | 　 | 3315.779  | 375.370  | 　 | 0.899 | 0.359 |
| L\_parasubiculum | 60.637  | 10.445  |  | 71.173  | 16.936  |  | 0.759 | 0.398 |
| L\_presubiculum | 270.655  | 15.672  |  | 305.257  | 33.376  |  | 6.451 | 0.024 |
| L\_subiculum | 370.767  | 24.465  |  | 414.856  | 62.708  |  | 2.597 | 0.129 |
| L\_CA1 | 578.307  | 56.004  |  | 593.635  | 63.691  |  | 0.354 | 0.561 |
| L\_CA3 | 202.875  | 28.935  |  | 203.550  | 34.228  |  | 0.062 | 0.807 |
| L\_CA4 | 223.679  | 28.581  |  | 239.122  | 33.260  |  | 0.915 | 0.355 |
| L\_molecular\_layer\_HP | 495.812  | 51.462  |  | 523.816  | 64.725  |  | 1.145 | 0.303 |
| L\_GC-ML-DG | 260.740  | 33.832  |  | 272.542  | 40.604  |  | 0.469 | 0.505 |
| L\_HATA | 55.487  | 7.963  |  | 56.546  | 8.892  |  | 0.215 | 0.650 |
| L\_fimbria | 62.617  | 12.002  |  | 54.374  | 18.685  |  | 0.251 | 0.624 |
| L\_Hippocampal\_tail | 562.847  | 111.261  |  | 580.908  | 69.678  |  | 0.057 | 0.814 |
| L\_hippocampal-fissure | 136.145  | 30.858  |  | 151.124  | 23.655  |  | 2.599 | 0.129 |
| R\_Whole\_hippocampus | 3442.779  | 259.115  |  | 3409.467  | 375.455  |  | 0.010 | 0.920 |
| R\_parasubiculum | 59.945  | 17.470  |  | 60.856  | 14.805  |  | 0.472 | 0.503 |
| R\_presubiculum | 285.213  | 27.734  |  | 287.769  | 35.254  |  | 0.186 | 0.673 |
| R\_subiculum | 414.776  | 45.036  |  | 426.660  | 56.269  |  | 1.354 | 0.264 |
| R\_CA1 | 622.266  | 45.033  |  | 627.643  | 71.350  |  | 0.003 | 0.954 |
| R\_CA3 | 232.779  | 25.269  |  | 226.200  | 35.608  |  | 0.047 | 0.832 |
| R\_CA4 | 263.486  | 29.539  |  | 256.297  | 31.988  |  | 0.008 | 0.928 |
| R\_molecular\_layer\_HP | 594.283  | 58.416  |  | 585.511  | 76.290  |  | 0.120 | 0.734 |
| R\_GC-ML-DG | 301.572  | 35.132  |  | 287.919  | 39.234  |  | 0.152 | 0.703 |
| R\_HATA | 55.671  | 7.079  |  | 50.882  | 10.215  |  | 3.652 | 0.077 |
| R\_fimbria | 64.820  | 21.371  |  | 52.933  | 19.213  |  | 0.169 | 0.688 |
| R\_Hippocampal\_tail | 547.969  | 53.660  |  | 546.798  | 62.474  |  | 0.119 | 0.735 |
| R\_hippocampal-fissure | 161.174  | 31.638  | 　 | 183.131  | 24.632  | 　 | 4.387 | 0.055 |

**Table 4** Comparison of hippocampal volumes across different H-Y stages in PD groups.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  |  |  |  |  |  |  |
| 　 | Grage 1(N=12) | 　 | Grade ≥2(N=8) | 　 | F value | p value |
|  | mean | sd |  | mean | sd |  |
| L\_Whole\_hippocampus | 3490.018  | 242.296  | 　 | 3479.871  | 557.204  | 　 | 0.094 | 0.764 |
| L\_parasubiculum | 73.865  | 20.144  |  | 73.904  | 26.762  |  | 0.018 | 0.894 |
| L\_presubiculum | 330.326  | 44.919  |  | 315.462  | 71.255  |  | 0.003 | 0.954 |
| L\_subiculum | 442.802  | 47.649  |  | 451.123  | 100.550  |  | 0.508 | 0.487 |
| L\_CA1 | 623.095  | 33.882  |  | 638.441  | 120.387  |  | 0.150 | 0.704 |
| L\_CA3 | 206.630  | 12.609  |  | 212.025  | 26.390  |  | 0.281 | 0.604 |
| L\_CA4 | 250.574  | 15.651  |  | 251.985  | 35.993  |  | 0.148 | 0.706 |
| L\_molecular\_layer\_HP | 546.115  | 39.041  |  | 557.178  | 93.843  |  | 0.558 | 0.467 |
| L\_GC-ML-DG | 284.869  | 22.302  |  | 287.420  | 44.451  |  | 0.137 | 0.716 |
| L\_HATA | 58.962  | 9.039  |  | 54.051  | 10.776  |  | 1.209 | 0.289 |
| L\_fimbria | 68.528  | 21.446  |  | 63.776  | 17.016  |  | 0.262 | 0.616 |
| L\_Hippocampal\_tail | 604.252  | 67.012  |  | 574.506  | 80.841  |  | 0.143 | 0.711 |
| L\_hippocampal-fissure | 162.240  | 19.786  |  | 162.543  | 32.898  |  | 0.000 | 0.994 |
| R\_Whole\_hippocampus | 3633.348  | 246.371  |  | 3634.171  | 532.037  |  | 0.515 | 0.484 |
| R\_parasubiculum | 74.371  | 20.842  |  | 70.351  | 26.185  |  | 0.003 | 0.957 |
| R\_presubiculum | 313.859  | 27.164  |  | 315.550  | 68.819  |  | 0.796 | 0.386 |
| R\_subiculum | 447.412  | 45.670  |  | 457.432  | 90.071  |  | 1.498 | 0.240 |
| R\_CA1 | 654.969  | 54.861  |  | 659.820  | 87.867  |  | 0.602 | 0.450 |
| R\_CA3 | 237.055  | 17.195  |  | 228.304  | 27.871  |  | 0.642 | 0.436 |
| R\_CA4 | 267.931  | 17.711  |  | 270.778  | 42.211  |  | 0.293 | 0.596 |
| R\_molecular\_layer\_HP | 632.553  | 77.574  |  | 631.129  | 101.109  |  | 0.069 | 0.796 |
| R\_GC-ML-DG | 305.502  | 21.666  |  | 305.782  | 46.143  |  | 0.204 | 0.658 |
| R\_HATA | 59.632  | 6.406  |  | 57.539  | 11.283  |  | 1.294 | 0.273 |
| R\_fimbria | 68.713  | 18.477  |  | 60.664  | 16.736  |  | 0.009 | 0.924 |
| R\_Hippocampal\_tail | 571.351  | 47.274  |  | 576.821  | 77.244  |  | 1.140 | 0.303 |
| R\_hippocampal-fissure | 172.282  | 33.216  | 　 | 179.266  | 36.557  | 　 | 0.292 | 0.597 |