## **1** Supporting information

2 Figure S1 Effects of APH, APH-CA and CA on brain neurotransmitters of scopolamine

3 induced amnesia mice without changes.

4 Figure S2 The correlation analysis between gut microbiota and fecal metabolites on
5 Species level. The right side of the figure is the metabolite name, and the bottom is the
6 gut microbiota. Each grid in the figure represents the correlation between two attributes
7 (metabolites and gut microbiota), and different colors represent the correlation
8 coefficient between attributes.

9 Figure S3 The correlation analysis between gut microbiota and brain neurotransmitters
10 on Genus level. The x-axis and y-axis are neurotransmitters and microbiota on Genus
11 level, respectively. The correlation R value and P value are obtained through
12 calculation. The R value is shown in different colors in the figure. If the p value is less
13 than 0.05, it is marked with \* and the legend on the right is the color range of different
14 R values.

15 Figure S4 The correlation analysis between fecal metabolites and brain
16 neurotransmitters.

17 Figure S5 The distribution proportion of dominant species in each sample and the18 distribution proportion of each dominant species in different groups.

19





**S**1



S2

A



S2

В





С



- 31 Figure S2 D



Marvinbryantia Bacteroides Mucispirillum norank (\_\_Ruminococcaceae Roseburia Lachnospiraceae\_UCG-006 Ruminococcaceae\_UCG-013 norank (\_\_Desulfovbrinonaceae unclassified\_f\_Ruminococcaceae besulfovbrinonaceae unclassified\_f\_Ruminococcaceae norank (\_\_Desulfovbrinonaceae unclassified\_f\_Ruminococcaceae Aaerotruncus [Eubacterium]\_Nlanophilum\_group A2 Blautia Ruminiclostridium\_ Ruminiclostridium\_ Ruminiclostridium\_ Socilibacter Lachnospiraceae\_NK4A136\_group Ruminiclostridium\_ unclassified\_f\_Lachnospiraceae norank (\_\_Pepteoccaceae Alstipes Alstipes Odoribacter Maribaculum norank (\_\_Prospielotichaceae Ruminococcaceae\_UCG-014 Morabacteri Maribaculum norank (\_\_Prospielotichaceae Ruminococcaceae\_UCG-010 Aerococcus Psychrobacter Suphylococcus Corpobacterium\_1 Prevotellaceae\_UCG-010 Ruminococcus\_1 Parabacteroides unclassified\_o\_Bacteroidales [Eubacterium\_ventriosum\_group Parasutterella norank (\_\_Muribeculaceae Helicobacter Iactobacilus Candidatus\_Saccharimonas Enterorhabdus Akarmansia Biddobacterium\_Facealibaculum



33

34 Figure S3

35



Figure S4 A



Figure S4 B



Figure S4 C



Figure S4 D



Figure S5

| Sample\Estimators | sobs          | shannon   | ace           | chao          | shannoneven     |
|-------------------|---------------|-----------|---------------|---------------|-----------------|
| control           | 538.83 ±18.28 | 4.41±0.22 | 620.93±22.48  | 625.24±21.27  | 0.7±0.03        |
| model             | 510.33±51.55  | 4.26±0.28 | 592.91±38.84  | 593.42±39.55  | 0.68±0.04       |
| APH               | 446.15±196.02 | 3.67±1.67 | 512.47±234.72 | 509.98±234.03 | 0.58±0.27       |
| АРН-СА            | 540.00±29.28  | 4.40±0.18 | 619.49±45.78  | 635.11±51.55  | 0.70±0.03       |
| CA                | 464.50±55.07  | 3.97±0.48 | 556.05±40.33  | 557.70±55.60  | $0.65 \pm 0.07$ |

Table S1  $\alpha$  diversity index table