# 开源数据库结肠炎和结肠癌的菌群差异 分析

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- 1 摘要
- 2 前言
- 3 材料和方法

# 3.1 材料

Other data obtained from published article (e.g., supplementary tables):

- Supplementary file from article refer to TransplantationSinha2022<sup>1</sup>.
- Supplementary file from article refer to TargetedSuppreFederi2022<sup>2</sup>.
- Supplementary file from article refer to DepressionAndYuan2021<sup>3</sup>.
- Supplementary file from article refer to AnIntegratedTRoelan2023<sup>4</sup>.
- Supplementary file from article refer to LocationAndCoSambru2023<sup>5</sup>.

# 3.2 方法

Mainly used method:

- R version 4.3.2 (2023-10-31); Other R packages (eg., dplyr and ggplot2) used for statistic analysis or data visualization.
- 4 分析结果
- 5 结论
- 6 附:分析流程
- 6.1 数据来源
- 6.1.1 结肠炎数据
- 6.1.1.1 TargetedSuppreFederi2022 结肠炎

'TargetedSuppreFederi2022 data'数据已全部提供。

#### (对应文件为 Figure+Table/TargetedSuppreFederi2022-data)

注: 文件夹 Figure+Table/TargetedSuppreFederi2022-data 共包含 4 个文件。

- 1. 1 Corrected France.csv
- $2.\ 2$ \_Corrected Israel.csv
- 3. 3\_Corrected US.csv
- 4. 4\_Corrected Germany.csv

#### 6.1.1.2 DepressionAndYuan2021 结肠炎

Table 1 (下方表格) 为表格 DepressionAndYuan2021 data 概览。

# (对应文件为 Figure+Table/DepressionAndYuan2021-data.xlsx)

注: 表格共有 91 行 4 列,以下预览的表格可能省略部分数据;表格含有 91 个唯一'Taxonomy'。

Table 1: DepressionAndYuan2021 data

Taxonomy	p.value	MRA.in.UC	MRA.in.HC
pGemmatimonadetes	0.00049553252446827	0.00171119907683608	0.000160190585722501
pActinobacteria	0.00834784591709094	0.0617033057472256	0.0389112516772091
pFirmicutes	0.00478686441960355	0.589610871565727	0.499806265231797
pBacteroidetes	$9.60337999115942 \mathrm{e}\hbox{-}05$	0.250752809840626	0.403687121772228
punidentified	$1.40986578996555\mathrm{e}\text{-}07$	0.000640301130981834	$2.05372545798078 \mathrm{e}\text{-}06$
cActinobacteria	0.00353055167754535	0.046725454341533	0.0237116295626934
cLongimicrobia	7.16632518063251e-07	0.00145496100157866	0
cBacteroidia	$6.38198131772293 \mathrm{e}\text{-}05$	0.24518920382048	0.403287329883075
cDeinococci	3.17450175630301e- $06$	0.00131123215732429	0
cBacilli	0.00330587139663399	0.0649165901033199	0.0154563377967633
cCytophagia	$2.10636789003367\mathrm{e}\text{-}05$	0.0016664310495747	$4.24436594649361\mathrm{e}\text{-}05$
cunidentified	$2.64154914758447\mathrm{e}\text{-}08$	0.0016652538833675	8.83101946931736e-05
cFlavobacteriia	$5.26714172360768\mathrm{e}\text{-}05$	0.00167703418517471	$5.95580382814426 \mathrm{e}\text{-}05$
oOceanospirillales	0.00495867783478051	0.000461817587592187	$2.05372545798077 \mathrm{e}\text{-}06$
oBifidobacteriales	0.00354779153342515	0.0443044462799416	0.0224465346805772

# 6.1.1.3 TransplantationSinha2022 结肠炎

Table 2 (下方表格) 为表格 TransplantationSinha2022 data 概览。

# (对应文件为 Figure+Table/TransplantationSinha2022-data.csv)

注:表格共有 141 行 12 列,以下预览的表格可能省略部分数据;表格含有 83 个唯一'W'。

Table 2: TransplantationSinha2022 data

W	detect2	detect3	detect4	detect5	Kingdom	Phylum	Class	Order	Family	
121	FALSE	TRUE	TRUE	TRUE	dBac	Bacter	Bacter	Bacter	Rikene	
89	FALSE	FALSE	FALSE	TRUE	dBac	Bacter	Bacter	Bacter	Rikene	
87	FALSE	FALSE	FALSE	TRUE	$\mathbf{d}\underline{\hspace{0.3cm}}\mathbf{Bac}$	Bacter	Bacter	Bacter	Marini	
94	FALSE	FALSE	FALSE	TRUE	dBac	Bacter	Bacter	Bacter	Barnes	

W	detect2	detect3	detect4	detect5	Kingdom	Phylum	Class	Order	Family	
125	FALSE	TRUE	TRUE	TRUE	dBac	Bacter	Bacter	Bacter	Barnes	
94	FALSE	FALSE	FALSE	TRUE	dBac	Bacter	Bacter	Bacter	Barnes	
95	FALSE	FALSE	FALSE	TRUE	dBac	Bacter	Bacter	Bacter	Barnes	
135	TRUE	TRUE	TRUE	TRUE	dBac	Bacter	Bacter	Bacter	Prevot	
138	TRUE	TRUE	TRUE	TRUE	dBac	Bacter	Bacter	Bacter	Prevot	
103	FALSE	FALSE	TRUE	TRUE	dBac	Firmic	Bacilli	Lactob	Entero	
134	TRUE	TRUE	TRUE	TRUE	$\mathrm{d}\_\_\mathrm{Bac}$	Firmic	Bacilli	Erysip	Erysip	
118	FALSE	TRUE	TRUE	TRUE	$\mathrm{d}\_\_\mathrm{Bac}$	Firmic	Bacilli	Erysip	Erysip	
90	FALSE	FALSE	FALSE	TRUE	dBac	Firmic	Bacilli	Erysip	Erysip	
87	FALSE	FALSE	FALSE	TRUE	dBac	Firmic	Bacilli	Erysip	Erysip	
95	FALSE	FALSE	FALSE	TRUE	dBac	Firmic	Bacilli	Erysip	Erysip	

# 6.1.2 结肠癌数据

# 6.1.2.1 AnIntegratedTRoelan2023 结肠癌

Table 3 (下方表格) 为表格 AnIntegratedTRoelan2023 data 概览。

(对应文件为 Figure+Table/AnIntegratedTRoelan2023-data.xlsx)

注:表格共有74行15列,以下预览的表格可能省略部分数据;表格含有1个唯一'sheet'。

Table 3: An<br/>IntegratedTRoelan<br/>2023 data  $\,$ 

sheet	Taxonomy	p_val	FDR	wilcox	mean_N	mean_T	median_N	median_T	Direction
Supple	D_0_B	2.8875	4.2447	1965	0.0121	0.0524	0.0006	0.0059	Enrich
Supple	D_0B	6.6405	4.8807	657	0.0018	0.0095	0	0	Enrich
Supple	D_0B	7.1087	3.4833	21023.5	0.0190	0.0126	0.0128	0.0067	Enrich
Supple	D_0B	2.8160	1.0348	18604	0.0132	0.0087	0.0078	0.0041	Enrich
Supple	D_0B	1.7251	5.0718	11985	0.0070	0.0043	0.0034	0.0019	Enrich
Supple	D_0B	5.0924	1.2476	7227.5	0.0126	0.0315	0.0031	0.0076	Enrich
Supple	D_0B	9.2218	1.9365	223.5	0.0002	0.0052	0	0	Enrich
Supple	D_0B	1.5535	2.8547	16631.5	0.0106	0.0076	0.0064	0.0041	Enrich
Supple	D_0B	2.1475	3.5075	1494	0.0022	0.0076	0	0	Enrich
Supple	D_0B	4.3690	6.2979	62	0.0001	0.0026	0	0	Enrich
Supple	D_0B	4.7127	6.2979	20341	0.0280	0.0232	0.0228	0.0170	Enrich
Supple	D_0B	7.9474	9.7356	237.5	0.0008	0.0058	0	0	Enrich
Supple	D_0B	1.6820	1.9019	4806.5	0.0040	0.0026	0	0	Enrich
Supple	D_0B	1.5690	1.5376	4009	0.0025	0.0039	0.0006	0.0013	Enrich
Supple	D 0 B	1.5362	1.5376	12107.5	0.0075	0.0059	0.0017	0.0012	Enrich

sheet	Taxonomy	p_val	FDR	wilcox	mean_N	mean_T	median_N	median_T	Direction

# 6.1.2.2 LocationAndCoSambru2023 结肠癌

Table 4 (下方表格) 为表格 LocationAndCoSambru2023 data 概览。

(对应文件为 Figure+Table/LocationAndCoSambru2023-data.csv)

注: 表格共有 44 行 2 列, 以下预览的表格可能省略部分数据; 表格含有 44 个唯一'tax\_id'。

Table 4: LocationAndCoSambru2023 data

tax id	taxon name
	— — — — — — — — — — — — — — — — — — —
40545	Sutterella_wadsworthensis
214856	Alistipes_finegoldii
328814	Alistipes_shahii
674529	Bacteroides_faecis
333367	$[{\it Clostridium}]\_{\it asparagiforme}$
437898	$Sutterella\_parvirubra$
74426	$Collinsella\_aerofaciens$
1531	$[{\it Clostridium}]\_{\it clostridioforme}$
239935	Akkermansia_muciniphila
901	Desulfovibrio_piger
1892897	$Shigella\_sp.\_FC569$
68259	$Streptomyces\_purpurogene is c$
1450439	${\bf Bacteroides\_sp.\_UW}$
585543	$Bacteroides\_sp.\_D20$
1581131	Actinomyces_spHMSC08A01

# 6.2 微生物注释

# Reference

- 1. Sinha, A. et al. Transplantation of bacteriophages from ulcerative colitis patients shifts the gut bacteriome and exacerbates the severity of dss colitis. Microbiome 10, (2022).
- 2. Federici, S. et al. Targeted suppression of human ibd-associated gut microbiota commensals by phage consortia for treatment of intestinal inflammation. Cell 185, 2879–2898.e24 (2022).
- 3. Yuan, X. et al. Depression and anxiety in patients with active ulcerative colitis: Crosstalk of gut microbiota, metabolomics and proteomics. Gut microbes 13, (2021).

- 4. Roelands, J. et al. An integrated tumor, immune and microbiome atlas of colon cancer. Nature medicine 29, 1273–1286 (2023).
- 5. Sambruni, G. et al. Location and condition based reconstruction of colon cancer microbiome from human rna sequencing data. Genome medicine 15, (2023).