

·综述·

有氧运动改善脑卒中后抑郁与肠道菌群相关性的研究进展*

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脑卒中后抑郁(post-stroke depression, PSD),患者主要表现为情绪低落、对事物缺乏兴趣、焦虑及睡眠障碍,甚至出现厌世、自杀等一系列行为改变^[1]。研究报告称脑卒中后PSD患病率约为30%,其中严重抑郁症患病率为17.7%^[2-3],且患PSD患者死亡风险显著增高,严重影响脑卒中患者日常生活能力恢复,并给家庭及社会带来巨大经济负担^[4]。PSD发病机制尚不清楚,目前研究较多的有内源性生物机制学说和反应源机制学说^[1]。随着新一代测序技术及生物信息分析技术的快速发展,近年来肠道菌群在干预脑血管及神经疾患方面成为科研人员的关注热点。有研究认为,脑卒中后肠道菌群的紊乱与PSD的发生发展密切相关^[5],而有氧运动可以增加益生菌的定植,改善肠道菌群的构成及功能。因此探讨有氧运动、PSD与肠道菌群之间的内在机制联系,能够为以肠道菌群为治疗靶点,为PSD患者制定合理有效的运动处方提供理论证据。

1 肠道菌群与中枢神经系统疾病

人体肠道内存在数量庞大且功能复杂的微生物群落,其所编码的基因数量约为人类基因组的150倍,被视为人体“第二大脑”^[6]。随着有关肠道微生物对人类健康与疾病影响研究的深入,探讨肠道菌群与中枢神经系统功能之间的关系已成为研究热点。肠道菌群可通过脑-肠轴经神经、内分泌和免疫等多种途径与大脑进行双向信息交流。肠道菌群与脑卒中及其继发性中枢神经系统损伤、焦虑、抑郁、自闭症、阿尔茨海默病等一系列神经精神疾病的发生发展密切相关^[7],其中,脑卒中及抑郁与肠道菌群的关系是目前的研究热点。研究发现,脑卒中及抑郁患者存在明显的肠道菌群失调,而菌群失调及继发的肠道功能变化可导致患者预后不良。研究发现,可通过调节菌群构成及恢复肠道功能缓解卒中及抑郁病情程度^[8-9]。

1.1 脑卒中与肠道菌群

肠道菌群与高血压、糖尿病和高血脂等脑卒中高危诱因密切相关。肠道菌群代谢的短链脂肪酸(short chain fatty

acids, SCFAs)可调节血压,肠分泌物脑肠肽可维持血糖血脂稳态^[10-11]。研究发现,脑卒中高发人群肠道内肠杆菌科和韦荣氏菌科增多,产生SCFAs的毛螺菌科和疣微菌科含量降低^[12]。运动缺乏、应激事件诱发的肠菌群紊乱及肠道炎症可导致血脂异常,并参与高血压及糖尿病的发生发展^[13-14]。此外,肠道菌群可通过炎症、脂质代谢,三甲胺氧化物(trimethylamine-N-oxide, TMAO)产生等途径促进动脉粥样硬化形成与发展^[15]。研究发现,嗜酸小球菌可诱导耐受性树突状细胞减轻动脉粥样硬化(atherosclerosis, AS)^[16]。

肠道菌群与脑卒中的发病机制密切相关。肠道菌群通过代谢TMAO促使血栓形成外^[17],还可经白细胞介素17(interleukin 17, IL-17)的驱动发生血管炎并参与颅底动脉瘤的形成^[18-19]。肠道菌群释放的内毒素可通过激活Toll样受体(toll-like receptors, TLR)4导致脑血管畸形及增加AS破裂风险,促使脑卒中发生^[20-21]。卒中后炎症反应可破坏肠屏障及肠道菌群稳态,经肠屏障入血的内毒素及炎症因子可损害血脑屏障,加重脑组织氧化应激^[22-23]。激活的HPA轴及交感神经可加剧肠功能障碍,抑制肠道神经递质的合成及神经传导,导致脑卒中预后不良,增加卒中后精神疾病发生的风险。研究发现,乳酸杆菌可改善血管内皮功能,降低卒中发生风险,还可抑制氧化应激及神经细胞凋亡,减轻脑卒中损伤程度^[24-25]。

1.2 抑郁与肠道菌群

压力应激作为抑郁主要致病因素与肠道菌群密切相关,来自外源性及内源性应激可激活HPA轴,交感神经及肠神经系统,导致肠道运动及通透性改变,引发肠道炎症等肠功能障碍^[26]。研究发现,患有炎症性肠疾病的儿童及成年人出现焦虑及抑郁的风险增加^[27]。

肠道菌群与HPA轴、神经递质、炎症因子等抑郁发病机制密切相关。长期处于应激环境中可激活HPA轴损害肠屏障,促使细菌及内毒素进入体循环,诱发机体免疫并持续激活HPA轴加剧肠通透性,减少肠道5-羟色胺(5-hydroxytryptamine, 5-HT)、多巴胺(dopamine, DA)、 γ -氨基丁酸(γ -

DOI:10.3969/j.issn.1001-1242.2021.08.028

*基金项目:河南省科技攻关计划项目(192102310424);国家自然科学基金项目(81704135);河南省高等学校重点科研项目(18A360012)

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aminobutyric acid, GABA)等递质合成并抑制肠迷走神经活性,减少与中枢的信号传导^[28]。此外,肠道炎症引发的外周炎症可扩散入脑诱发神经毒性并阻碍相关神经递质及脑源性神经营养因子(brain-derived neurotrophic factor, BDNF)的合成,诱发抑郁情绪^[29-30]。而抑郁症患者持续存在的负面情绪可进一步触发对各种应激源的强烈反应及敏感性,形成恶性循环。研究发现,瑞士乳杆菌及双歧杆菌可恢复应激大鼠肠屏障,降低HPA轴及自主神经活性^[31]。抗抑郁药5-羟色胺再摄取抑制剂可抵抗革兰氏阳性菌,三环类抗抑郁药可阻止肠道病原菌的定植生长而维持肠道稳态^[32]。

1.3 脑卒中后抑郁与肠道菌群

目前研究较多的PSD发病机制主要涉及神经递质、炎症因子、HPA轴、BDNF等方面。脑卒中所致的肠功能障碍及肠道菌群失调可通过免疫、内分泌及神经系统等多种途径参与PSD的发病机制。

1.3.1 神经递质与肠道菌群:脑卒中后肠道菌群失调可影响神经递质合成及神经传导而诱发抑郁。脑卒中通过激活自主神经及HPA轴抑制肠蠕动、增加肠渗透性而破坏肠稳态^[33]。肠蠕动抑制可使肠球菌属、梭菌属及拟杆菌属等潜在致病菌过度繁殖,而使产生5-HT的植物乳杆菌、分泌GABA的双歧杆菌、合成去甲肾上腺素(norepinephrine, NE)的酵母菌等益生菌的减少。此外,肠道乳杆菌属及酪酸梭菌的减少可降低SCFAs生成而减少5-HT分泌。肠菌群紊乱可使外周5-HT水平失调并影响海马谷氨酸-谷氨酰胺-GABA循环而诱发异常情绪^[34-35]。研究发现,缺乏肠道菌的应激小鼠脑内多巴胺能转换率降低并伴情绪障碍^[36]。肠神经递质减少可减弱对肠内炎症的抑制,加重机体炎症对神经递质通路的破坏。此外,肠道菌群合成的神经递质可通过迷走神经与大脑沟通,而中枢神经系统损伤可使肠粘膜下和肌间神经元部分丢失,减少与迷走神经的突触连接,抑制与中枢的信号传输而诱发抑郁^[33]。研究发现,摄入鼠李糖乳杆菌可调节小鼠中枢GABA受体表达及情绪行为,而切断迷走神经作用则消失^[37]。

1.3.2 炎症细胞因子与肠道菌群:脑卒中后肠道免疫诱发的机体炎症可致脑部神经毒性而诱发抑郁。脑卒中可激活肠道髓样细胞触发受体-1促使肠通透性增高及菌群移位^[38]。革兰氏阴性菌、大肠杆菌及肠球菌等衍生的内毒素可渗入循环系统引发内毒素血症,诱导释放的肿瘤坏死因子- α (tumor necrosis factor- α , TNF- α)、 γ -干扰素(interferon- γ , IFN- γ)、白细胞介素1 β (interleukin-1 β , IL-1 β)、白细胞介素-6(interleukin-6),白介素18(interleukin 18, IL-18)等细胞因子可经体液或神经途径进入大脑,促使小胶质细胞转化为促炎的M1型,导致神经炎症促使抑郁发生^[39]。研究发现,罗伊氏乳杆菌和青春双歧杆菌可降低血液中脂多糖(lipopolysaccha-

ride, LPS), IL-6含量并抑制海马内小胶质细胞的活化而缓解焦虑及抑郁情绪^[40]。并且,渗入脑部的炎症因子可介导代谢犬尿氨酸为具有神经毒性的喹啉酸,其所诱导的谷氨酸兴奋性毒性是炎症引起抑郁的常见途径^[41]。此外,肠道菌群失调介导的神经炎症可阻碍5-HT、DA及NE的合成而诱发抑郁^[39]。研究发现,乳杆菌和长双歧杆菌可通过抑制肠道核因子 κ B(nuclear factor kappa-B, NF- κ B)活化和TNF- α 表达减轻抑郁^[42]。

1.3.3 HPA轴与肠道菌群:脑卒中后肠道菌群的变化可持续激活HPA轴,抑制海马神经再生而诱发抑郁。脑卒中可激活HPA轴释放GC,导致肠通透性增加及菌群失调。肠道菌群衍生物脂多糖、肽聚糖和肠道释放的IL-1 β 、IL-6和TNF- α 是HPA轴的有效激活剂,其与菌群紊乱形成恶性循环^[43]。肠有益菌的减少可减弱SCFAs对编码HPA轴中相关蛋白质基因表达的调控,从而增强对HPA轴的响应^[44]。大肠杆菌产生的酪蛋白水解酶B可刺激促肾上腺皮质激素(adreno-corticotrophic-hormone, ACTH)前体阿黑皮素原的释放,从而增强HPA轴活性^[45]。GC的持续增多,可致其受体抵抗或信号缺失,抑制海马神经元再生而触发抑郁情绪。将抑郁患者肠道菌群移植给无菌小鼠,发现GC受体下游Stat5a基因下调并伴异常行为^[46]。GC可降解色氨酸及酪氨酸,减少5-HT及NE合成而诱发抑郁。研究发现,香肠乳杆菌可维护肠屏障,降低脂多糖水平从而减弱HPA对应激的反应,减轻抑郁情绪^[47]。

1.3.4 BDNF与肠道菌群:脑卒中所致的肠道菌群紊乱可通过影响BDNF水平诱导抑郁产生。肠上皮细胞,肠神经胶质细胞和神经元中存在的BDNF及其酪氨酸激酶受体B(tyrosine kinase receptor B, TrkB)对调节肠道感觉和运动功能起着基本作用^[48]。脑卒中所致的菌群紊乱可改变肠道BDNF和TrkB表达,改变肠道运性及敏感性,所导致的胃肠疾病与情绪障碍的产生密切相关。脑卒中后肠道功能障碍诱发的神经炎症可导致脑内BDNF含量降低,抑制神经可塑性,诱导情绪障碍的发生。研究发现,摄入植物乳杆菌可通过调节海马TLR4/BDNF信号通路,降低海马TLR4表达,增加BDNF含量^[49]。约翰逊乳杆菌及干酪乳杆菌可改善肠道菌群组成及调节BDNF-TrkB信号通路改善抑郁行为^[50-51]。

2 有氧运动可有效改善PSD

研究发现,有氧运动可有效提高脑卒中患者日常生活能力、缓解脑卒中患者抑郁情绪并提高其生活质量^[52]。临床研究表明^[53-64]:干预PSD的有氧运动主要包括有氧耐力训练和肢体功能训练,如步行训练、气功训练、肢体主动训练等,结合患者病情和机能状况,运动负荷多为中等强度,频率每天15—60min,每周1—5次,疗程多在5—12周。运动对脑

卒中后抑郁情绪干预效果的 meta 分析研究结果显示:较高运动强度具有更好的改善效果^[65]。运动形式及运动负荷的选择需根据患者机体情况做出合理及时地调整,临床中进行有氧运动干预可选用最大摄氧量及心率对患者的运动强度及安全进行监测。

3 有氧运动通过肠道菌群改善 PSD 的作用机制

已有研究发现,有氧运动可有效改善脑卒中、抑郁及 PSD^[66-68],也可通过调节肠道菌群组成,维持肠道菌群稳态发挥对人体的有益作用。因此,探讨有氧运动、PSD、肠道菌群之间的联系对于寻找 PSD 治疗新靶点具有重要意义。

3.1 有氧运动可通过影响肠道菌群组成调节神经递质

有氧运动可增加脑内神经递质含量改善 PSD, Arida RM 等^[69]总结以往研究发现对大鼠进行跑步运动可调节脑内 5-HT、DA、GABA 及谷氨酸等神经递质水平缓解抑郁情绪,而对肠道菌群进行干预可调节神经递质水平^[70],有氧运动可促进肠道可调节神经递质水平的菌群及 SCFA 含量增加。Mailing LJ 等^[71]通过对比经常运动与久坐女性肠道菌发现,经常运动的女性肠道普氏栖粪杆菌和丁酸菌丰度增加,二者可增加具有影响神经递质释放功能的 SCFA 含量。动物研究发现,通过对大鼠进行 6 天自由轮转运动后大鼠肠道来自硬壁菌门的乳杆菌属和真杆菌属增加,放线菌门的双歧杆菌属增加,3 个属可促进 GABA 及 5-HT 合成,增加 SCFAs 含量而降低肠道 pH 值,抑制病原菌的生长及炎症反应。此外,乳酸杆菌,双歧杆菌水平与可调节食欲及饮食行为的血清瘦素含量成正相关^[72]。因此有氧运动可能通过调节肠道菌群的组成影响神经递质及行为的变化。

3.2 有氧运动可通过增加肠道抗炎有益菌发挥抗炎作用

有氧运动可通过抑制炎症反应改善 PSD, Li C 等^[68]发现对 PSD 大鼠进行 4 周的跑台训练可抑制磷酸酯酶与张力蛋白同源物升高介导的 TLR4、NF- κ B、NLRP3 信号上调而发挥对海马神经元的保护,改善 PSD。研究发现通过对大鼠进行自由轮转运动可减少肠道内 TNF- α 、促凋亡蛋白及 IL-17 含量并增加普氏栖粪杆菌含量。普氏栖粪杆菌产生的丁酸可调节 NF- κ B 活化、氧化应激及肠道免疫并可向周围环境分泌抗炎物质,抑制机体炎症发展^[73]。此外,有氧运动可抑制脑卒中后肠道菌群紊乱所诱发的炎症反应,Estaki M 等^[74]报道了运动受试者中产生 LPS 的革兰氏阴性菌减少,SCFA 增加,在恢复肠道屏障同时可抑制 LPS 易位并阻碍其与细胞外 TLR4 结合,减轻炎症反应。有氧运动可能通过抑制肠道炎症及继发的神经炎症改善 PSD。

3.3 有氧运动可增加肠道有益菌调节 HPA 轴

有氧运动可抑制 GC 水平上调,抑制 HPA 轴过度激活,缓解 PSD^[75]。肠道乳酸杆菌、双歧杆菌及鼠李糖乳杆菌等有

益菌可纠正 HPA 轴的过度激活, Mahdich MS 等^[76]通过临床研究证实有氧运动可增加人体肠道乳酸杆菌和双歧杆菌含量。Queipo-Ortuño MI 等^[72]研究也发现有氧运动可增加肠道双歧杆菌属及乳杆菌属含量。有氧运动并非都是有益的,强迫运动可被大鼠视为压力源,反而增强了 HPA 轴活性,降低了肠道内有益菌的含量,因此适度的有氧运动对于调节 HPA 轴活性具有较好的作用。

3.4 有氧运动可通过增加肠道有益菌上调 BDNF

有氧运动可增加 BDNF 与脑源性神经营养因子前体之比,上调 BDNF 水平,增强神经可塑性,改善 PSD^[77]。陈凤等^[78]通过动物实验证明肠道菌群的变化可影响 BDNF 的表达,研究发现,双歧杆菌和乳酸杆菌可上调 BDNF 水平,有助于神经元的增殖分化,从而发挥抗抑郁作用^[79]。有氧运动可增加上述益生菌的含量,上调 BDNF 水平,双歧杆菌及乳酸杆菌的增加可减轻肠道菌群紊乱,保护肠道屏障,维护与迷走神经的传导,调节大脑 BDNF 浓度^[80]。可知,有氧运动可通过增加有益菌调节 BDNF 浓度发挥抗抑郁作用。

4 小结

研究已证实肠道菌群与 PSD 紧密相关,但其深层机制尚未阐明。脑卒中可通过激活自主神经系统、释放 GC 增高肠通透性,升高粪肠球菌、变形杆菌及大肠杆菌等条件致病菌,降低合成神经递质的丁酸梭菌、乳酸菌及双歧杆菌等益生菌含量。肠道菌群失调诱发的肠道免疫通过肠脑轴引发神经毒性、破坏神经递质通路及神经营养因子含量、抑制 HPA 轴负反馈,分泌的 GC 不断诱发肠道菌群的失衡,导致肠道与大脑经肠脑轴相互影响的恶性循环,促使抑郁的发生发展。

有氧运动可增加肠道内双歧杆菌,乳酸菌等有益菌及 SCFAs 含量促进神经递质及 BDNF 含量增加。普氏栖粪杆菌及约翰逊乳杆菌的增多可抑制肠道炎症反应进而调节 HPA 轴活性,减少 GC 的释放,抑制机体炎症的发展而缓解卒中及 PSD 的严重程度。因此,适度的有氧运动可通过维护肠道菌群的平衡状态,发挥对 PSD 的有益作用。未来研究还需应用宏基因组学技术,通过有氧运动,揭示与 PSD 相关的菌群紊乱,并鉴别出有关的特定肠道细菌及功能,获得对 PSD 发生、发展的机制及诊断、治疗的新认识,以实现通过有氧运动调节 PSD 患者肠道菌群以改善患者健康的目的。

参考文献

- [1] Villa RF, Ferrari F, Moretti A. Post-stroke depression: mechanisms and pharmacological treatment[J]. Pharmacol Ther, 2018, 184: 131—144.
- [2] Schöttke H, Giabbiconi CM. Post-stroke depression and post-stroke anxiety: prevalence and predictors[J]. Int Psychogeriatr.

- atr,2015,27(11):1805—1812.
- [3] Mitchell AJ, Sheth B, Gill J, et al. Prevalence and predictors of post-stroke mood disorders: a meta-analysis and meta-regression of depression, anxiety and adjustment disorder[J]. *Gen Hosp Psychiatry*,2017,47:48—60.
 - [4] Oni OD, Olagunju AT, Olisah VO, et al. Post-stroke depression: prevalence, associated factors and impact on quality of life among outpatients in a Nigerian hospital[J]. *S Afr J Psychiatr*,2018,24:1058—1065.
 - [5] 范文涛, 闫咏梅, 别玉龙, 等. 脑卒中后抑郁症患者肠道菌群的多样性分析[J]. *南方医科大学学报*,2016,36(10):1305—1311.
 - [6] Qin J, Li R, Raes J, et al. A human gut microbial gene catalogue established by metagenomic sequencing[J]. *Nature*,2010,464(7285):59—65.
 - [7] Tyler Patterson T, Grandhi R. Gut microbiota and neurologic diseases and injuries[J]. *Adv Exp Med Biol*,2020,1238:73—91.
 - [8] Wallace CJK, Foster JA, Soares CN, et al. The effects of probiotics on symptoms of depression: protocol for a double-blind randomized placebo-controlled trial[J]. *Neuropsychobiology*,2020,79(1):108—116.
 - [9] Akhoundzadeh K, Vakili A, Shadnough M, et al. Effects of the oral ingestion of probiotics on brain damage in a transient model of focal cerebral ischemia in mice[J]. *Iran J Med Sci*,2018,43(1):32—40.
 - [10] Kang Y, Cai Y. Gut microbiota and hypertension: from pathogenesis to new therapeutic strategies[J]. *Clin Res Hepatol Gastroenterol*,2018,42(2):110—117.
 - [11] Canfora EE, Jocken JW, Blaak EE. Short-chain fatty acids in control of body weight and insulin sensitivity[J]. *Nat Rev Endocrinol*,2015,11(10):577—591.
 - [12] Zeng X, Gao X, Peng Y, et al. Higher risk of stroke is correlated with increased opportunistic pathogen load and reduced levels of butyrate-producing bacteria in the gut[J]. *Front Cell Infect Microbiol*,2019,9:4.
 - [13] Dunphy-Doherty F, O'Mahony SM, Peterson VL, et al. Post-weaning social isolation of rats leads to long-term disruption of the gut microbiota-immune-brain axis[J]. *Brain Behav Immun*,2018,68:261—273.
 - [14] Kriaa A, Bourgin M, Potiron A, et al. Microbial impact on cholesterol and bile acid metabolism: current status and future prospects[J]. *J Lipid Res*,2019,60(2):323—332.
 - [15] Jonsson AL, Bäckhed F. Role of gut microbiota in atherosclerosis[J]. *Nat Rev Cardiol*,2017,14(2):79—87.
 - [16] Mizoguchi T, Kasahara K, Yamashita T, et al. Oral administration of the lactic acid bacterium *pediococcus acidilactici* attenuates atherosclerosis in mice by inducing tolerogenic dendritic cells[J]. *Heart Vessels*,2017,32(6):768—776.
 - [17] Zhu W, Gregory JC, Org E, et al. Gut microbial metabolite TMAO enhances platelet hyperreactivity and thrombosis risk[J]. *Cell*,2016,165(1):111—124.
 - [18] Shikata F, Shimada K, Sato H, et al. Potential influences of gut microbiota on the formation of intracranial aneurysm[J]. *Hypertension*,2019,73(2):491—496.
 - [19] Karbach SH, Schönfelder T, Brandão I, et al. Gut microbiota promote angiotensin II-induced arterial hypertension and vascular dysfunction[J]. *J Am Heart Assoc*,2016,5(9):e003698.
 - [20] Ridler C. Cerebrovascular malformations: microbiota promotes cerebral cavernous malformations[J]. *Nat Rev Neurol*,2017,13(7):386.
 - [21] Jonsson AL, Bäckhed F. Role of gut microbiota in atherosclerosis[J]. *Nat Rev Cardiol*,2017,14(2):79—87.
 - [22] Houlden A, Goldrick M, Brough D, et al. Brain injury induces specific changes in the caecal microbiota of mice via altered autonomic activity and mucoprotein production[J]. *Brain Behav Immun*,2016,57:10—20.
 - [23] Singh V, Sadler R, Heindl S, et al. The gut microbiome primes a cerebroprotective immune response after stroke[J]. *J Cereb Blood Flow metab*,2018,38(8):1293—1298.
 - [24] Malik M, Suboc TM, Tyagi S, et al. *Lactobacillus plantarum* 299v supplementation improves vascular endothelial function and reduces inflammatory biomarkers in men with stable coronary artery disease[J]. *Circ Res*,2018,123(9):1091—1102.
 - [25] Chen R, Xu Y, Wu P, et al. Transplantation of fecal microbiota rich in short chain fatty acids and butyric acid treat cerebral ischemic stroke by regulating gut microbiota[J]. *Pharmacol Res*,2019,148:104403.
 - [26] Gao X, Cao Q, Cheng Y, et al. Chronic stress promotes colitis by disturbing the gut microbiota and triggering immune system response[J]. *Proc Natl Acad Sci USA*,2018,115(13):E2960—E2969.
 - [27] Mikocka-Walus A, Knowles SR, Keefer L, et al. Controversies revisited: a systematic review of the comorbidity of depression and anxiety with inflammatory bowel diseases[J]. *Inflamm Bowel Dis*,2016,22(3):752—762.
 - [28] Yong SJ, Tong T, Chew J, et al. Antidepressive mechanisms of probiotics and their therapeutic potential[J]. *Front Neurosci*,2020,13:1361.
 - [29] Bryleva EY, Brundin L. Suicidality and activation of the kynurenine pathway of tryptophan metabolism[J]. *Curr Top Behav Neurosci*,2017,31:269—284.

- [30] Peirce JM, Alviña K. The role of inflammation and the gut microbiome in depression and anxiety[J]. *J Neurosci Res*, 2019, 97(10):1223—1241.
- [31] Ait-Belgnaoui A, Colom A, Braniste V, et al. Probiotic gut effect prevents the chronic psychological stress-induced brain activity abnormality in mice[J]. *Neurogastroenterol Motil*, 2014, 26(4):510—520.
- [32] Koopman M, El Aidy S, MIDTrauma consortium. Depressed gut? the microbiota-diet-inflammation triad in depression[J]. *Curr Opin Psychiatry*, 2017, 30(5):369—377.
- [33] Li XJ, You XY, Wang CY, et al. Bidirectional brain-gut-microbiota axis in increased intestinal permeability induced by central nervous system injury[J]. *CNS Neurosci Ther*, 2020, 26(8):783—790.
- [34] Yano JM, Yu K, Donaldson GP, et al. Indigenous bacteria from the gut microbiota regulate host serotonin biosynthesis[J]. *Cell*, 2015, 161(2):264—276.
- [35] Zheng P, Zeng B, Liu M, et al. The gut microbiome from patients with schizophrenia modulates the glutamate-glutamine-GABA cycle and schizophrenia-relevant behaviors in mice[J]. *Sci Adv*, 2019, 5(2):eaau8317.
- [36] Crumeyrolle-Arias M, Jaglin M, Bruneau A, et al. Absence of the gut microbiota enhances anxiety-like behavior and neuroendocrine response to acute stress in rats[J]. *Psychoneuroendocrinology*, 2014, 42:207—217.
- [37] Bravo JA, Forsythe P, Chew MV, et al. Ingestion of lactobacillus strain regulates emotional behavior and central GABA receptor expression in a mouse via the vagus nerve[J]. *Proc Natl Acad Sci USA*, 2011, 108(38):16050—16055.
- [38] Liu Q, Johnson EM, Lam RK, et al. Peripheral TREM1 responses to brain and intestinal immunogens amplify stroke severity[J]. *Nat Immunol*, 2019, 20(8):1023—1034.
- [39] Peirce JM, Alviña K. The role of inflammation and the gut microbiome in depression and anxiety[J]. *J Neurosci Res*, 2019, 97(10):1223—1241.
- [40] Jang HM, Lee KE, Kim DH. The preventive and curative effects of lactobacillus reuteri NK33 and bifidobacterium adolescentis NK98 on immobilization stress-induced anxiety/depression and colitis in mice[J]. *Nutrients*, 2019, 11(4):819.
- [41] Raison CL, Dantzer R, Kelley KW, et al. CSF concentrations of brain tryptophan and kynurenines during immune stimulation with IFN- α : relationship to CNS immune responses and depression[J]. *Mol Psychiatry*, 2010, 15(4):393—403.
- [42] Han SK, Kim DH. Lactobacillus mucosae and bifidobacterium longum synergistically alleviate immobilization stress-induced anxiety/ depression in mice by suppressing gut dysbiosis[J]. *J Microbiol Biotechnol*, 2019, 29(9):1369—1374.
- [43] de Punder K, Pruimboom L. Stress induces endotoxemia and low-grade inflammation by increasing barrier permeability[J]. *Front Immunol*, 2015, 6:223.
- [44] van de Wouw M, Boehme M, Lyte JM, et al. Short-chain fatty acids: microbial metabolites that alleviate stress-induced brain-gut axis alterations[J]. *J Physiol*, 2018, 596(20):4923—4944.
- [45] Breton J, Tennoune N, Lucas N, et al. Gut commensal E. coli proteins activate host satiety pathways following nutrient-induced bacterial growth[J]. *Cell metab*, 2016, 23(2):324—334.
- [46] Luo Y, Zeng B, Zeng L, et al. Gut microbiota regulates mouse behaviors through glucocorticoid receptor pathway genes in the hippocampus[J]. *Transl Psychiatry*, 2018, 8(1):187.
- [47] Ait-Belgnaoui A, Durand H, Cartier C, et al. Prevention of gut leakiness by a probiotic treatment leads to attenuated HPA response to an acute psychological stress in rats[J]. *Psychoneuroendocrinology*, 2012, 37(11):1885—1895.
- [48] Grider JR, Piland BE, Gulick MA, et al. Brain-derived neurotrophic factor augments peristalsis by augmenting 5-HT and calcitonin gene-related peptide release[J]. *Gastroenterology*, 2006, 130(3):771—780.
- [49] Mohammed SK, Magdy YM, El-Waseef DA, et al. Modulation of hippocampal TLR4/BDNF signal pathway using probiotics is a step closer towards treating cognitive impairment in NASH model[J]. *Physiol Behav*, 2020, 214:112762.
- [50] Lee HJ, Lim SM, Kim DH. Lactobacillus johnsonii CjLJ103 attenuates scopolamine-induced memory impairment in mice by increasing BDNF expression and inhibiting NF- κ B activation[J]. *J Microbiol Biotechnol*, 2018, 28(9):1443—1446.
- [51] Tian P, Zou R, Song L, et al. Ingestion of bifidobacterium longum subspecies infantis strain CCFM687 regulated emotional behavior and the central BDNF pathway in chronic stress-induced depressive mice through reshaping the gut microbiota[J]. *Food Funct*, 2019, 10(11):7588—7598.
- [52] Zou L, Yeung A, Zeng N, et al. Effects of mind-body exercises for mood and functional capabilities in patients with stroke: an analytical review of randomized controlled trials[J]. *Int J Environ Res Public Health*, 2018, 15(4):721.
- [53] 杜仁仁, 孙文江, 蔡珍珍, 等. 传统功法结合康复治疗对脑卒中恢复期患者轻度抑郁、焦虑及日常生活活动能力的疗效及机制分析[J]. *辽宁中医药大学学报*, 2020, 22(12):152—160.
- [54] 张保国, 开绍江, 张园园, 等. 中等强度运动疗法联合抗抑郁药物治疗卒中后抑郁的疗效观察[J]. *实用临床医药杂志*, 2019, 23(9):44—47.
- [55] Chen CH, Hung KS, Chung YC, et al. Mind-body inter-

- active qigong improves physical and mental aspects of quality of life in inpatients with stroke: a randomized control study[J]. *Eur J Cardiovasc Nurs*, 2019, 18(8):658—666.
- [56] 毛林波, 吴嘉琪. 运动疗法在脑卒中后抑郁治疗体会(附30例)[J]. *健康之路*, 2018, 17(9):397.
- [57] 李晓荟, 刘平, 熊斌, 等. 改良24式太极拳对脑卒中后抑郁的临床研究[J]. *中华保健医学杂志*, 2018, 20(5):434—435.
- [58] Morais VAC, Tourino MFDS, Almeida ACS, et al. A single session of moderate intensity walking increases brain-derived neurotrophic factor (BDNF) in the chronic post-stroke patients[J]. *Top Stroke Rehabil*, 2018, 25(1):1—5.
- [59] 赵彬, 唐强, 王艳, 等. 太极拳对卒中后抑郁患者运动功能及抑郁状态的影响[J]. *中国康复理论与实践*, 2017, 23(3):334—337.
- [60] McDonnell MN, Mackintosh SF, Hillier SL, et al. Regular group exercise is associated with improved mood but not quality of life following stroke[J]. *Peer J*, 2014, 2:e331.
- [61] 李玉玲, 胡杏娟, 温馨. 坐式太极运动在脑卒中后抑郁患者康复训练中的应用[J]. *齐鲁护理杂志*, 2013, 19(23):4—6.
- [62] Stuart M, Benvenuti F, Macko R, et al. Community-based adaptive physical activity program for chronic stroke: feasibility, safety, and efficacy of the Empoli model[J]. *Neurorehabil Neural Repair*, 2009, 23(7):726—734.
- [63] Smith PS, Thompson M. Treadmill training post stroke: are there any secondary benefits? a pilot study[J]. *Clin Rehabil*, 2008, 22(10—11):997—1002.
- [64] 陈丽萍, 杨萍. 运动疗法对脑卒中后抑郁症的影响[J]. *中西医结合心脑血管病杂志*, 2006, 4(3):270—271.
- [65] Eng JJ, Reime B. Exercise for depressive symptoms in stroke patients: a systematic review and meta-analysis[J]. *Clin Rehabil*, 2014, 28(8):731—739.
- [66] Linder SM, Davidson S, Rosenfeldt A, et al. Forced and voluntary aerobic cycling interventions improve walking capacity in individuals with chronic stroke[J]. *Arch Phys Med Rehabil*, 2021, 102(1):1—8.
- [67] Luo J, Tang C, Chen X, et al. Impacts of aerobic exercise on depression-like behaviors in chronic unpredictable mild stress mice and related factors in the AMPK/PGC-1 α pathway[J]. *Int J Environ Res Public Health*, 2020, 17(6):2042.
- [68] Li C, Xu X, Wang Z, et al. Exercise ameliorates post-stroke depression by inhibiting PTEN elevation-mediated up-regulation of TLR4/NF- κ B/NLRP3 signaling in mice[J]. *Brain Res*, 2020, 1736:146777.
- [69] Arida RM, Cavalheiro EA, Scorza FA. From depressive symptoms to depression in people with epilepsy: contribution of physical exercise to improve this picture[J]. *Epilepsy Res*, 2012, 99(1—2):1—13.
- [70] Strandwitz P. Neurotransmitter modulation by the gut microbiota[J]. *Brain Res*, 2018, 1693(Pt B):128—133.
- [71] Mailing LJ, Allen JM, Buford TW, et al. Exercise and the gut microbiome: a review of the evidence, potential mechanisms, and implications for human health[J]. *Exerc Sport Sci Rev*, 2019, 47(2):75—85.
- [72] Queipo-Ortuño MI, Seoane LM, Murri M, et al. Gut microbiota composition in male rat models under different nutritional status and physical activity and its association with serum leptin and ghrelin levels[J]. *PLoS One*, 2013, 8(5):e65465.
- [73] Campbell SC, Wisniewski PJ, Noji M, et al. The effect of diet and exercise on intestinal integrity and microbial diversity in mice[J]. *PLoS One*, 2016, 11(3):e0150502.
- [74] Estaki M, Pither J, Baumeister P, et al. Cardiorespiratory fitness as a predictor of intestinal microbial diversity and distinct metagenomic functions[J]. *Microbiome*, 2016, 4(1):42.
- [75] 张一清, 朱晓军, 王彤. 运动训练对大鼠缺血性卒中后抑郁的发生及下丘脑-垂体-肾上腺皮质轴的影响[J]. *中国康复医学杂志*, 2014, 29(4):306—310.
- [76] Mahdich MS, Maryam J, Bitar B, et al. A pilot study on the relationship between lactobacillus, bifidobacterium counts and inflammatory factors following exercise training[J]. *Arch Physiol Biochem*, 2021, 16:1—10.
- [77] Luo L, Li C, Du X, et al. Effect of aerobic exercise on BDNF/proBDNF expression in the ischemic hippocampus and depression recovery of rats after stroke[J]. *Behav Brain Res*, 2019, 362:323—331.
- [78] 陈凤, 王龙娟, 李贞贞, 等. 大鼠肠道菌群变化影响其大脑海马源性神经营养因子的表达[J]. *中国微生态学杂志*, 2015, 27(1):10—13.
- [79] Wang H, Lee IS, Braun C, et al. Effect of probiotics on central nervous system functions in animals and humans: a systematic review[J]. *J Neurogastroenterol Motil*, 2016, 22(4):589—605.
- [80] O'Leary OF, Ogbonnaya ES, Felice D, et al. The vagus nerve modulates BDNF expression and neurogenesis in the hippocampus[J]. *Eur Neuropsychopharmacol*, 2018, 28(2):307—316.