Rescue of male fertility following faecal microbiota transplantation from alginate oligosaccharide-dosed mice

A very recent publication in Gut highlights that faecal microbiota transplantation (FMT) from alginate oligosaccharide (AOS)-dosed animals improves mouse sperm quality and spermatogenesis after busulfan treatment. The results suggest the potential of FMT for the improvement of infertility, since worldwide 10%–15% of couples are infertile and many of them have failed spermatogenesis. In addition, many investigations have found that gut microbiota may affect male or female reproduction. Although the improvement of male infertility is an
emerging novel area of interest and many investigations have attempted to ameliorate spermatogenesis by various methods, little progress has been achieved.\textsuperscript{5, 6} In the study done by Zhang et al.,\textsuperscript{1} FMT from AOS-dosed animals increased spermatozoa quality and the process of spermatogenesis; however, that gut microbiota from AOS-dosed animals can actually increase fertility rate is as yet unknown.

To confirm the beneficial advantages of FMT from AOS-dosed animals, we set out to explore the fertility rate (pregnancy rate and number of live pups/litter) following FMT from AOS-dosed animals to busulfan-treated mice (online supplementary file 1 and online supplemental figure 1). We found that B+A10-FMT (busulfan plus gut microbiota from AOS 10 mg/kg mice) significantly increased pregnancy rate (10-fold) and number of live pups/litter (twofold) compared with busulfan (B-sa; figure 1A,B). Notably, the number of live pups/litter was almost the same for B+A10-FMT and control (Con-sa; blank control) which suggested that A10-FMT had a strong potential for rescuing male fertility. However, B+Con-FMT (busulfan plus gut microbiota from control mice) did not significantly increase the pregnancy rate or number of live pups/litter compared with busulfan (figure 1A,B). At the same time, we compared the beneficial advantages of AOS 10 mg/kg (A10) and A10-FMT after busulfan treatment. A10 and A10-FMT produced a similar improvement on the pregnancy rate and number of live pups/litter (figure 1A,B). In our earlier studies,\textsuperscript{1, 5} we discovered that AOS 10 mg/kg improves the gut microbiota to, in turn, improve spermatogenesis and semen quality. Furthermore, A10-FMT similarly benefited gut microbiota through an increase in the ‘beneficial’ bacteria\textsuperscript{7} Bacteroidales,\textsuperscript{8} Bifidobacteriales,\textsuperscript{9} Campylobacteriales,\textsuperscript{10} Sphingomonadales and Campylobacterales which have beneficial effects such as protecting the intestinal barrier,\textsuperscript{8} production of anti-oxidant compounds\textsuperscript{9} and the possession of reduction enzymes.\textsuperscript{10} It is also interesting that the microbes from A10-FMT showed a good correlation with sperm concentration/motility, blood metabolome and testis metabolome.\textsuperscript{1} It is even more profoundly important that the microbiota from A10-dosed mice and A10-FMT-treated mice were well correlated.\textsuperscript{1, 5} Moreover, A10, A10-FMT and Con-FMT did not affect the fertility rate of control mice (without busulfan; figure 1C,D) which indicated that these treatments did not pose a disadvantage for male animal reproduction. In the current investigation, spermatogenesis was significantly improved by A10-FMT as shown by the germ cell marker VASA (figure 2).

There were almost no VASA-positive cells in the busulfan group (B-sa) and a very small number in B+Con-FMT group; however, a significant number of VASA-positive cells were found in the B+A10FMT and B+A10 groups (figure 2), which suggested that spermatogenesis was improved by A10-FMT, since busulfan mainly disrupted germ cells.\textsuperscript{2, 5, 6} At the same time, protein levels of the sperm cell marker PGK2 were determined by immunofluorescence staining.\textsuperscript{1, 2, 5} There were also no PGK2-positive cells in the B+A10FMT and B+A10 groups (figure 2). The data further revealed that A10-FMT rescued busulfan disrupted spermatogenesis, while Con-FMT did not. The data in this investigation confirmed that gut microbiota from AOS-dosed mice had the potential to improve spermatogenesis and then to increase male fertility rate.

**Figure 1** Mouse pregnancy rate and the number of live pups/litter. (A) Mouse pregnancy rate (number of pregnant mice/total mice in each group: Con-sa (experiment II), B-sa (experiment II), B+A10 (experiment III), B+A10 FMT (experiment II) and B+Con FMT (experiment II); n=60/group). a,b,c Means not sharing a common superscript are different (p<0.05). (B) The number of live pups/litter in each group: Con-sa (experiment II), B-sa (experiment II), B+A10 (experiment III), B+A10 FMT (experiment II) and B+Con FMT (experiment II); (n=60/group). a,b,c Means not sharing a common superscript are different (p<0.05). (C) Mouse pregnancy rate (number of pregnant mice/total mice in each group: Con (experiment II), A10 (experiment III), A10-FMT (experiment II) and Con-FMT (experiment II); (n=60/group). a,b,c Means not sharing a common superscript are different (p<0.05). (D) The number of live pups/litter. The average number of live pups/litter in each group: Con (experiment II), A10 (experiment III), A10-FMT (experiment II) and Con-FMT (experiment II); (n=60/group). a,b,c Means not sharing a common superscript are different (p<0.05). Note: (1) Con-sa (dosed with saline); (2) B-sa (busulfan (a single injection 20 mg/kg body weight of busulfan) plus saline); (3) B+A10 (busulfan plus AOS 10 mg/kg); (4) B+Con-FMT (busulfan plus gut microbiota from regular mice); (5) A10-FMT (busulfan plus gut microbiota from AOS 10 mg/kg dosed mice); (6) Con (dosed with saline); (7) A10 (dosed with AOS 10 mg/kg); (8) Con-FMT (dosed with gut microbiota from regular mice); (9) A10-FMT (dosed with gut microbiota from AOS 10 mg/kg dosed mice). See more detailed information in online supplemental file 1). AOS, alginate oligosaccharide; FMT, faecal microbiota transplantation.

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Figure 2  VASA and PGK2 staining of mouse testicular samples. (A) Germ cell marker VASA staining of mouse testicular samples. (B) Sperm cell marker PGK2 staining of mouse testicular samples. (C) Quantification data for VASA staining. (D) Quantification data for PGK2 staining. Note: (1) Con-sa (dosed with saline); (2) B-sa (busulfan (a single injection 20 mg/kg body weight of busulfan) plus saline); (3) B+ A10 (busulfan plus AOS 10 mg/kg); (4) B+ Con-FMT (busulfan plus gut microbiota from regular mice); (5) A10-FMT (busulfan plus gut microbiota from AOS 10 mg/kg dosed mice). For more details, see information in online supplemental file 1). The letters a, b, c, d and e indicate a significant difference among different treatments (p<0.05). AOS, alginate oligosaccharides; FMT, faecal microbiota transplantation.

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REFERENCES

1. Zhang P, Peng Y, Li L, et al. Improvement in sperm quality and spermatogenesis following faecal microbiota transplantation from algaline oligosaccharide dosed mice. Gut 2021;70:222–5.

2. Wang M, Liu X, Chang G, et al. Single-cell RNA sequencing analysis reveals sequential cell fate transition during human spermatogenesis. Cell Stem Cell 2018;23:599–614.

3. Ding N, Zhang X, Zhang XD, et al. Impairment of spermatogenesis and sperm motility by the high-fat diet-induced dysbiosis of gut microbes. Gut 2020;69:1608–19.

4. Qi X, Yun C, Sun L, et al. Gut microbiota-bile acid-interleukin-22 axis orchestrates polymeric ovary syndrome. Nat Med 2019;25:1225–33.

5. Zhao Y, Zhang P, Ge W, et al. Alginate oligosaccharides improve germ cell development and testicular microenvironment to rescue busulfan disrupted spermatogenesis. Theranostics 2020;10:3308–24.

6. Liu F-J, Dong W-Y, Zhao H, et al. Effect of molybdenum on reproductive function of male mice treated with busulfan. Theriogenology 2019;126:49–54.

7. Zhang C, Zhang W, Zhang J, et al. Gut microbiota dysbiosis in male patients with chronic traumatic complete spinal cord injury. J Transl Med 2018;16:353.

8. Siddaramappa S, Viswanathan V, Thiagarajan S, et al. Genomewide characterisation of the genetic diversity of carotenogenesis in bacteria of the order Sphingomonadales. Microb Genom 2018;4. doi:10.1099/mgen.0.000172. [Epub ahead of print: 05 Apr 2018].

9. Kaakoush NO, Sterzenbach T, Miller WG, et al. Identification of disulflde reductases in Campylobacteria: a bioinformatics investigation. Antonie Van Leeuwenhoek 2007;92:429–41.

10. Valdes AM, Walter J, Segal E, et al. Role of the gut microbiota in nutrition and health. BMJ 2018;361:k2179.

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