



In vivo Imaging of a Novel Strain of *Bacteroides fragilis* via Metabolic Labeling

Wenye Xu^{1†}, Peizhu Su^{1,2†}, Lijun Zheng^{3†}, Hongying Fan⁴, Ye Wang³, Yangyang Liu³, Yuqing Lin³ and Fachao Zhi^{1*}

¹ Guangdong Provincial Key Laboratory of Gastroenterology, Department of Gastroenterology, Institute of Gastroenterology of Guangdong Province, Nanfang Hospital, Southern Medical University, Guangzhou, China, ² Department of Gastroenterology, First People's Hospital of Foshan Affiliated to Sun Yat-sen University, Foshan, China, ³ Guangzhou Zhiyi Biotechnology Co., Ltd., Guangzhou, China, ⁴ Guangdong Provincial Key Laboratory of Tropical Disease Research, School of Public Health, Southern Medical University, Guangzhou, China

Non-toxigenic Bacteroides fragilis is regarded as a potential candidate for probiotic

OPEN ACCESS

Edited by:

Eugenia Bezirtzoglou, Democritus University of Thrace, Greece

Reviewed by:

Jozsef Soki, University of Szeged, Hungary Yiannis Kourkoutas, Democritus University of Thrace, Greece

*Correspondence:

Fachao Zhi zhifc41532@163.com [†]These authors have contributed equally to this work

Specialty section:

This article was submitted to Food Microbiology, a section of the journal Frontiers in Microbiology

Received: 22 June 2018 Accepted: 07 September 2018 Published: 01 October 2018

Citation:

Xu W, Su P, Zheng L, Fan H, Wang Y, Liu Y, Lin Y and Zhi F (2018) In vivo Imaging of a Novel Strain of Bacteroides fragilis via Metabolic Labeling. Front. Microbiol. 9:2298. doi: 10.3389/fmicb.2018.02298 owing to its various advantages. We previously isolated a new strain of B. fragilis (ZY-312) and verified its biosafety and capability of inhibiting the growth of pathogens in vivo. However, the colonization of ZY-312 in gastrointestinal (GI) tract remains to be determined. To track the colonization of ZY-312, mice were gavaged with ZY-312 labeled by means of metabolic oligosaccharide engineering and bioorthogonal click chemistry or given AF647-dibenzocyclooctyne (DIBO) directly. Then the fluorescence was detected in GI tract, spleen and kidneys. Results showed that ZY-312 could be labeled by metabolic oligosaccharide engineering, and the optimal incubation time with AF647-DIBO was 5 h in vitro. Following oral gavage with AF647-DIBO labeled ZY-312 or AF647-DIBO alone, mice were subjected to in vivo imaging and the fluorescence intensity was similar in both groups 3 h, 6 h, and 12 h post the gavage. The fluorescence of AF647-DIBO group disappeared 24 h post gavage which was probably due to the excretion via GI tract. While the fluorescence of AF647-DIBO labeled ZY-312 retained in the cecum for as long as 48 h. Immunofluorescence assay further confirmed that labeled ZY-312 transiently colonized not only in cecum but also in stomach, ileum and colon of mice 48 h postgavage and that no massive accumulation of ZY-312 was detected in other organs such as kidneys and spleen. In conclusion, ZY-312 could transiently colonize in GI tract, mainly in cecum, for at least 48 h, and it hardly disseminate to other organs, which shed new light on the future development of *B. fragilis* as a probiotic product.

Keywords: Bacteroides fragilis, ZY-312, probiotic, colonization, in vivo imaging

INTRODUCTION

The gut microbiota consists of around 2,000 different bacterial species (Boyce and Bertozzi, 2011), which are influenced by age, infection, and drugs, etc. Gut microbiota dysbiosis which is characterized by imbalance in the composition and function of these intestinal microbes is associated with a wide spectrum of diseases ranging from localized gastrointestinal disorders to neurologic, respiratory, metabolic, hepatic, and cardiovascular diseases (Lynch and Pedersen, 2016). Therefore, increased attention has been drawn on developing probiotics to help reestablish

micro-ecological balance. *In-vivo* colonization is an important characteristic of probiotics, which means strains can survive *in vivo*, repair damaged intestinal mucosa and defend the invasion of pathogenic bacteria.(Deriu et al., 2013) Moreover, the ability of probiotics to colonize in intestine is considered to be a critical factor in immune regulation.(Lebeer et al., 2010) Therefore, many *in-vitro* models have been established to examine the adhesion and colonization ability of bacteria in different tissues and cell lines including partial intestinal tissue resected from human, mucus isolated from feces, and ileostomy effluent and cells such as HT-29 and Caco-2 (Ouwehand et al., 1995; Wang et al., 2018).

However, the models mentioned above did not mimic the mucus layer, which is the first barrier inoculated strains meet with. Mucin-secreting HT-29-MTX cells might overcome the shortcomings to some extent (Martín et al., 2015). Nevertheless, the composition of mucin produced by HT-29-MTX cells is different from that of mucin derived from the human intestinal mucosa. Moreover, current methods aimed to detect colonization of probiotics such as Gram staining, sequencing and fluorescent labeling still exists several limitations. First of all, similar strains cannot be distinguished by Gram staining. Secondly, sequencing is a time-consuming method. Conventional fluorescent labeling is useful but also limited, for fluorescent proteins require aerobic conditions, but most gut commensals are anaerobes (Craggs, 2009).

To overcome those limitations, we used metabolic oligosaccharide engineering (MOE) and bioorthogonal click chemistry (BCC) (Boyce and Bertozzi, 2011) to label and track live ZY-312. In this method, a small functional group is incorporated in biomolecules of the target strain by a cell's endogenous biosynthetic function, and through BCC, this group reacts with a second chemical group, forming a stable covalent bond. (Sletten and Bertozzi, 2009) This method has been used to study glycoconjugates and polysaccharides in living systems, (Dehnert et al., 2011) including in prokaryotic organisms (Dumont et al., 2012; Kaewsapsak et al., 2013), *Staphylococcus aureus, Bacteroides ovatus*, etc.

In this study, we aimed to examine the colonization and dissemination of an emerging probiotic candidate ZY-312, a new strain of *Bacteroides. fragilis* through MOE and BCC method.

MATERIALS AND METHODS

Mice

Male and female 6 to 8-week-old C57BL/6 mice were purchased from Beijing Vital River Laboratory. Mice were housed in SPF conditions with food and water *ad libitum*. The animal experimental procedures were performed in accordance with the Guide for the Care and Usage of Laboratory Animals published by the US National Institutes of Health (NIH Publication No.85–23, revised 1996), and approved by Nanfang Hospital Animal Ethics Committee (protocol # NFYY-2014-123). Mice were randomly assigned to three groups: AF647–DIBO labeled ZY-312 group, AF647-DIBO group, and negative control group. Mice in AF647–DIBO labeled ZY-312 group and AF647-DIBO group were inoculated with labeled strains or AF647–DIBO respectively, for various time periods, including 3 h, 6 h, 12 h, 24 h, 48 h, 60 h, 72 h, and 96 h. And negative control group didn't receive any intervention.

Bacterial Strains and Culture Media

Bacteroides. fragilis strain ZY-312 was provided from Zhiyi Biological Technology Co., Ltd. [Guangzhou, Guangdong province, (Deng et al., 2016; Wang et al., 2017)]. The strain was cultured under anaerobic conditions (80% N2, 10% H2, 10% CO2) in basal peptone-yeast broth with azide-modified tetraacetylated-N-azidoacetylgalactosamine (GalNAz) at a final concentration of 100 µM at 37°C in an anaerobic incubator (Bugbox, Ruskinn) during the whole study. Basal peptoneyeast broth contains (per liter) 5 g yeast extract, 20 g proteose peptone, 5 g NaCl, 5 mg hemin, 0.5 mg vitamin K1 and 5 g K₂HPO₄. Hemin, vitamin K1 and K₂HPO₄ were added through a filter after the basal medium had been autoclaved. Ten-fold serial dilutions of the cultures were plated on trypticase soya agar containing 5% fresh sheep blood to obtain single colony following microscopic examination (TSA; Oxoid, Basingstoke, United Kingdom).

MOE Labeling of Bacteria

ZY-312 was cultured for 65 h at 37 °C under anaerobic conditions (80% N₂, 10% H₂, 10% CO₂) in an anaerobic glove box in basal peptone-yeast broth (OXOID, United Kingdom) with GalNAz (Thermofisher, United States) at a final concentration of 100 µM. Bacterium were spun down and washed three times in $1 \times$ phosphate-buffered saline (PBS) supplemented with 1% bovine serum albumin (BSA) when the OD₆₀₀ value ranged from 0.7 to 1.4. The bacteria pellet was resuspended in 10 ml PBS supplemented with 1% BSA (Sigma, Germany) and 20 mM AF647-DIBO (100 µl per vial) and each vial of the bacterium suspension was kept separately. To determine the optimal incubation time, bacteria suspensions were incubated for 3 h, 5 h, 7 h, and overnight in dark on a shake flask. After incubation, bacteria were pelleted and washed five times with PBS supplemented with 3% BSA. Finally, the bacteria were resuspended either in PBS for in vivo tracing or in 90% glycerol for fluorescent microscope observation.

In vivo Optical Imaging

Mice were divided into *B. fragilis* group and AF467-DIBO group and each group received gavage (300 μ l) at different time point (t = -96 h, -72 h, -60 h, -48 h, -24 h, -12 h, -6 h, and -3 h) prior to the *in vivo* observation (three mice for each group and time point). Mice were caged separately according to the group and time point to avoid the intake of feces cross the group. For the *in vivo* observation, all mice were anesthetized with isoflurane followed by the *in vivo* imaging using Xtreme imaging system (Bruker) All mice were sacrificed after the *in vivo* imaging and the intestine, stomach, kidney and spleen were collected with adherent connective tissues resected. The distributions of the fluorescence-labeled ZY-312 in those organs were also determined through Xtreme imaging system. Image analysis was performed using Molecular Imaging Software version 7.2. A region of interest (ROI) was drawn around fluorescent signals of one representative mouse using the ellipse option in MI software. This ROI was then saved as template and applied to the other images.

Frozen Sections

Tissue samples of different organs (three parallel samples for each organ) were collected in eppendorf tubes and washed with PBS to remove dirt and intestinal contents. Samples were dried on a clean filter and loaded on the pre-cooled embedding agent and covered with extra embedding agent. Cryosections were performed and by a freezing microtome and the slides were imaged on a microscope.

Immunofluorescence

Slides were placed at room temperature for ten minutes and washed with PBS for three times (5 min per time) and dried with a filter paper. Then they were blocked with goat serum for 1 h at room temperature and incubated with actin antibody (diluted with 5%BSA and Triton at 1:250) in a humidified chamber overnight at 4°C. Subsequently, the slides were washed with PBS and incubated with FITC-conjugated secondary antibodies (Abcam) for 1 h at room temperature. Finally, the samples were covered with mounting medium containing DAPI and examined under laser confocal fluorescence microscopy.

Imaging by Laser Confocal Fluorescence Microscopy

We respectively, chose red, blue, green filters of the confocal fluorescence microscopy to detect the fluorescence. FITC was detected through the green filter (excitation at 488 nm, emission at 525 nm), DAPI was through the blue filter (excitation at 340 nm, emission at 488 nm) and fluorescence of labeled ZY-312 was through red filter (excitation at 650 nm, emission at 665 nm).

Statistical Analysis

Statistical analysis was done with Prism using unpaired Student's *t*-test.

RESULTS

In vitro Labeling of ZY-312 With AF647-DBIO

To construct the MOE labeling of ZY-312, we incubated ZY-312 and AF647-DIBO with shaking and blocked from light for various time. Results showed that ZY-312 was amenable to AF647-DBIO labeling and the optimal incubation time could be 5 h (**Figure 1**).

Visualization of Labeled ZY-312 in vivo

We administered labeled ZY-312 to C57BL/6 mice by gavage and analyzed colonization of the bacteria in the GI tract. Generally,

in vivo imaging showed that the fluorescence of both AF647-DIBO group and labeled ZY-312 group gradually declined over time. For AF647-DIBO administration group, the fluorescence could be detected at 3 h but faded away at 24 h and later, which might be attributed to the possibility that the dye should be metabolized or excreted completely after 24 h. To our surprise, labeled ZY-312 was detectable from 3 h to 48 h post-gavage, even longer than that of AF647-DIBO itself, indicating that ZY-312 was capable of colonizing in the GI tract for at least 48 h (**Figures 2A,C**).

To further evaluate the spatial distribution of labeled ZY-312, fluorescence of specific organs was imaged. As for the GI tract, fluorescence intensity of both groups was declined with time and fluorescence distribution in both groups was also similar in the first 12 h. At 3 h, the fluorescent signal was most prominent in the stomach and in middle and lower small intestine, and by 6 h it reached the cecum. Finally at 12 h, the fluorescence of both groups had gradually decreased and focused on the cecum and the colon, probably after which AF647-DIBO was excreted later owing to the absence of fluorescence throughout the GI tract at 24 h. However, there still existed fluorescent signal in the cecum of labeled ZY-312 treated mice at 24 h and even later at 48 h. Besides, no signal was detected in organs including the kidneys and spleens (**Figure 2B**).

Collectively, it was demonstrated that ZY-312 was likely to colonize in the GI tract, mainly in cecum, for at least 48 h.

Imunofluorescence Assay of Labeled ZY-312 Colonizing in GI Tract

To further validate the colonization of ZY-312 in the GI tract, we observed the frozen sections of the excised stomach, ileum, cecum, and colon with immunofluorescence assay. Results showed that red fluorescence signal could be detected in the stomach, villi of ileum, cecum and colon only in AF647-DIBO labeled ZY-312 group but not in AF647-DIBO or negative control at 24 h and 48 h (**Figure 3**). Besides, no fluorescence signal was detected in the kidneys or spleen of mice gavaged with AF647-DIBO labeled ZY-312 or AF647-DIBO alone at 24 h and 48 h (data not shown).

DISCUSSION

Non-toxigenic *B. fragilis* is regarded as a probiotic candidate owing to its immunoregulatory property and other healthpromoting benefits (Hsiao et al., 2013). Its protective effect is confirmed in mouse experimental colitis model (Mazmanian et al., 2005; Round and Mazmanian, 2010) and autoimmune encephalomyelitis model (Ochoarepáraz et al., 2010) by increasing the proportion of Foxp3⁺ regulatory T lymphocytes. It's also capable to correct systemic immune defects in germfree mice by stimulating maturation of splenic CD4⁺ T cells (Mazmanian et al., 2005). Recent study reveals that the tumors model of antibiotic-treated or germ-free mice do not respond to cytotoxic T-lymphocyte-associated protein 4 (CTLA-4) blockade treatment. However, the defect was overcomed by gavage with



FIGURE 1 | Fluorescence intensity of the labeled ZY-312 co-incubated with AF647-DIBO at various time points. ZY-312 was grown 65 h in basal peptone-yeast broth with GalNAz and the culture was co-incubated with AF647-DIBO for 3 h, 5 h, 7 h, and 24 h, respectively. When co-incubated for 5 h, Fluorescence intensity was the strongest among all time points.



B. fragilis, immunization with *B. fragilis* polysaccharides, or adoptive transfer of *B. fragilis*-specific T cells, indicating that antitumor effect of CTLA-4 blockade is also associated with *B. fragilis* (Vétizou et al., 2015). ZY-312 is a new strain of

non-enterotoxigenic *B. fragilis*, isolated from the feces of a healthy breast-fed infant. Previously we have demonstrated that ZY-312 possesses several probiotic properties, including tolerance to air, simulating gastric fluid (pH 3.0), intestinal



fluid and bile (pH 6.8). It's also capable of adhere to colon cells without cytotoxicity in vitro. (Deng et al., 2016) ZY-312 possess similar characteristics to standard B. fragilis strains ATCC 25285 in morphology, growth kinetics, metabolic, and genetic profile and has been proved to be non-toxigenic in both normal and immune-deficient mice (Wang et al., 2017). Furthermore, ZY-312 has been shown to be able to inhibit the growth and adhesion of Vibrio parahaemolyticus (V. parahaemolyticus) in vitro and protect both RAW 264.7 and LoVo cells from damage caused by V. parahaemolyticus (Li et al., 2017). Recently, we discovered the modulatory effect of ZY-312 on gut microbiota in rats, which contributes to the amelioration of antibiotic-associated diarrhea (Zhang et al., 2018). In general, ZY-312 has the potential to become the first probiotic representative of the dominant Bacteroidetes phylum.

The ability to colonize to gut is a crucial characteristic of probiotic. Recently, Geva-Zatorsky's team revealed a new approach called MOE-BCC to label and track anaerobic commensals without affecting the viability, growth and carbohydrate metabolism of bacteria. Therefore, we adopted MOE-BCC to label and track ZY-312 in the host in real time. To track live ZY-312 *in vivo*, we incubated ZY-312 with DIBO derivatives, whose combination is copper-independent and is

non-toxigenic to cells and bacteria. Our in vivo imaging study showed that labeled ZY-312 mainly distributed in stomach and ileum by 3 h, in cecum by 6 h and in cecum as well as colon by 12 h, which are roughly consistent with the distribution of B. fragilis reported in previous study (Gevazatorsky et al., 2015). Of note, while the AF647-DIBO alone was excreted via GI tract at 24 h, AF647-DIBO labeled ZY-312 transiently colonized in the cecum and colon for as long as 48 h. In addition, immunofluorescence assay convinced us that labeled ZY-312 transiently colonized not only in the cecum but also in other regions of GI tract including the stomach, ileum and colon in mice 48 h post-gavage, regardless of the possibility that the fluorescence might be weakened somewhat due to metabolism in gut. Taken together, we believe that ZY-312 is able to transiently colonize in GI tract, particularly in cecum, for at least 48 h. To test whether ZY-312 could colonize in the GI track for even longer, a more stable and sensitive labeling system is required.

Our experiments also suggest that oral administration of ZY-312 is relatively safe and feasible. As is shown, labeled ZY-312 was not detected in organs including the spleen and kidneys at 24 h and 48 h post gavage, suggesting that ZY-312 could be only limited in the GI tract, which minimizes the chances of bacterial translocation, bacteremia and other adverse effects. As probiotics do not enter the blood circulation, it is hard to perform regular pharmacokinetical study to determine their dosage and administration frequency. However, our results demonstrated that ZY-312 still massively concentrated in cecum and colon 24 h after administration, where gut microbiota are most abundant, and most of them were excreted after 48 h. Therefore, we speculate that a proper frequency for oral administration of live ZY-312 should be once to twice a day to create a stable and long-term colonization of ZY-312 in the cecum and colon, where it plays its role in regulating intestinal flora like other probiotics do, and could also prevent bacterial over-accumulation at the same time.

Nevertheless, our experiments still have some other limitations. Once given, the progenies of bacteria cannot be labeled by themselves due to the lack of azide-modified sugars in GI tract. Therefore, we can only investigate the migration and colonization of initial labeled ZY-312 but not the progeny strains. However, given that *B. fragilis* only makes up a very small part of the human intestinal commensal bacteria (Wang et al., 2017), it is unlikely that descendants of ZY-312 would be proliferate massively in the intestine.

In summary, our study shows that ZY-312 could transiently colonize in the GI tract, mainly in cecum, for at least 48 h, and that oral administration of ZY-312 is a relatively safe and feasible way for them to colonize in the large intestine and avoid its massive accumulation in GI tract and other organs which might

REFERENCES

- Boyce, M., and Bertozzi, C. R. (2011). Bringing chemistry to life. *Nat. Methods* 8, 638–642. doi: 10.1038/nmeth.1657
- Craggs, T. D. (2009). Cheminform abstract: green fluorescent protein: structure, folding and chromophore maturation. *Chem. Soc. Rev.* 38, 2865–2875. doi: 10.1039/b903641p
- Dehnert, K. W., Beahm, B. J., Huynh, T. T., Baskin, J. M., Laughlin, S. T., Wang, W., et al. (2011). Metabolic labeling of fucosylated glycans in developing zebrafish. *ACS Chem. Biol.* 6, 547–552. doi: 10.1021/cb100284d
- Deng, H. M., Li, Z. C., Tan, Y. F., Guo, Z. B., Liu, Y., Wang, Y., et al. (2016). A novel strain of *Bacteroides fragilis* enhances phagocytosis and polarises M1 macrophages. *Sci. Rep.* 6:29401. doi: 10.1038/srep 29401
- Deriu, E., Liu, J. Z., Pezeshki, M., Edwards, R. A., Ochoa, R. J., Contreras, H., et al. (2013). Probiotic bacteria reduce *salmonella typhimurium* intestinal colonization by competing for Iron. *Cell Host Microbe* 14, 26–37. doi: 10.1016/ j.chom.2013.06.007
- Dumont, A., Malleron, A., Awwad, M., Dukan, S., and Vauzeilles, B. (2012). Clickmediated labeling of bacterial membranes through metabolic modification of the lipopolysaccharide inner core. *Angewandte Chem.* 124, 3197–3200. doi: 10.1002/ange.201108127
- Gevazatorsky, N., Alvarez, D., Hudak, J. E., Reading, N. C., Erturk-Hasdemir, D., Dasgupta, S., et al. (2015). *In vivo* imaging and tracking of host-microbiota interactions via metabolic labeling of gut anaerobic bacteria. *Nat. Med.* 21, 1091–1100. doi: 10.1038/nm.3929
- Hsiao, E. Y., Mcbride, S. W., Hsien, S., Sharon, G., Hyde, E. R., McCue, T., et al. (2013). The microbiota modulates gut physiology and behavioral abnormalities associated with autism. *Cell* 155, 1151–1463. doi: 10.1016/j.cell.2013. 11.024
- Kaewsapsak, P., Esonu, O., and Dube, D. H. (2013). Recruiting the host's immune system to target *helicobacter* pylori's surface glycans. *Chembiochem* 14, 721–726. doi: 10.1002/cbic.201300006

do harm to health. Our work sheds new light on the development of *B. fragilis* as an ideal probiotic product for future clinical trial.

AUTHOR CONTRIBUTIONS

WX conducted the experiments with mice and wrote the manuscript. PS did the experiments with bacteria and contributed to revising the manuscript. LZ helped with performing the experiments and contributed to revising the manuscript. HF, YW, YYL, and YQL designed the experiments and contributed to revising the manuscript. FZ provided overall directions and contributed to revising the manuscript.

FUNDING

This work was supported by National High Technology Research and Development Program 863 (No. 2015AA020702), National Natural Science Foundation of China (No.81770530), Science and Technology Program of Guangdong, China (No. 2016B090918064&2016A020217010) Science and Technology Program of Guangzhou, China (No. 201605122107130), and Medical Scientific Research Foundation of Guangdong Province, China (No. A2018291).

- Lebeer, S., Vanderleyden, J., and De keersmaecker, S. C. (2010). Host interactions of probiotic bacterial surface molecules: comparison with commensals and pathogens. *Nat. Rev. Microbiol.* 8, 171–184. doi: 10.1038/nrmicro 2297
- Li, Z. C., Deng, H. M., Zhou, Y. Z., Tan, Y. F., Wang, X., Han, Y., et al. (2017). Bioluminescence imaging to track *Bacteroides fragilis* inhibition of vibrio parahaemolyticus infection in MICE. *Front. Cell. Infect. Microbiol.* 7:170. doi: 10.3389/fcimb.2017.00170
- Lynch, S. V., and Pedersen, O. (2016). The human intestinal microbiome in health and disease. New Engl. J. Med. 375, 2369–2379. doi: 10.1056/NEJMra1600266
- Martín, R., Sánchez, B., Urdaci, M. C., Langella, P., Suárez, J. E., and Bermúdez-Humarán, L. G. (2015). Effect of Iron on the probiotic properties of the vaginal isolate *Lactobacillus jensenii* CECT 4306. *Microbiology* 161, 708–718. doi: 10. 1099/mic.0.000044
- Mazmanian, S. K., Liu, C. H., Tzianabos, A. O., and Kasper, D. L. (2005). An immunomodulatory molecule of symbiotic bacteria directs maturation of the host immune system. *Cell* 122, 107–118. doi: 10.1016/j.cell.2005.0 5.007
- Ochoarepáraz, J., Mielcarz, D. W., Ditrio, L. E., Burroughs, A. R., Begum-Haque, S., Dasgupta, S., et al. (2010). Central nervous system demyelinating disease protection by the human commensal *Bacteroides fragilis* depends on polysaccharide a expression. *J. Immunol.* 185, 4101–4108. doi: 10.4049/ jimmunol.1001443
- Ouwehand, A. C., Conway, P. L., and Salminen, S. J. (1995). Inhibition of S-fimbriamediated adhesion to human ileostomy glycoproteins by a protein isolated from bovine colostrum. *Infect. Immun.* 63, 4917–4920.
- Round, J. L., and Mazmanian, S. K. (2010). Inducible foxp3+ regulatory T-cell development by a commensal bacterium of the intestinal microbiota. *Proc. Natl. Acad. Sci. U.S.A.* 107, 12204–12209. doi: 10.1073/pnas.09091 22107
- Sletten, E. M., and Bertozzi, C. R. (2009). Bioorthogonal chemistry: fishing for selectivity in a sea of functionality. *Angew. Chem. Int. Ed. Engl.* 48, 6974–6998. doi: 10.1002/anie.200900942

- Vétizou, M., Pitt, J. M., Daillère, R., Lepage, P., Waldschmitt, N., Flament, C., et al. (2015). Anticancer immunotherapy by CTLA-4 blockade relies on the gut microbiota. *Science* 350, 1079–1084. doi: 10.1126/science.aad 1329
- Wang, J., Lu, R., Fu, X., Dan, Z., Zhang, Y. G., Chang, X., et al. (2018). Novel regulatory roles of Wnt1 in infection-associated colorectal Cancer. *Neoplasia* 20, 499–509. doi: 10.1016/j.neo.2018.03.001
- Wang, Y., Deng, H. M., Li, Z. C., Tan, Y. F., Han, Y., Wang, X., et al. (2017). Safety evaluation of a novel strain of *Bacteroides fragilis*. *Front. Microbiol.* 8:435. doi: 10.3389/fmicb.2017.00435
- Zhang, W. D., Zhu, B., Xu, J. H., Liu, Y. Y., Qiu, E., Li, Z., et al. (2018). Bacteroides fragilis protects against antibiotic-associated diarrhea in rats by modulating intestinal defenses. Front. Immunol. 9:1040. doi: 10.3389/fimmu.2018.01040

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

The reviewer YK and handling Editor declared their shared affiliation.

Copyright © 2018 Xu, Su, Zheng, Fan, Wang, Liu, Lin and Zhi. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.