ORIGINAL RESEARCH

A purified membrane protein from Akkermansia munciniphila or the pasteurised bacterium blunts colitis associated tumourigenesis by modulation of CD8+ T cells in mice

Lijuan Wang,1 Lei Tang,1 Yiming Feng,1 Suying Zhao,2 Mei Han,2 Chuan Zhang,3 Gehui Yuan,1 Jun Zhu,1 Shuyuan Cao,1 Qian Wu,1 Lei Li,1 Zhan Zhang 1

ABSTRACT

Objective Gut microbiota have been linked to inflammatory bowel disease (IBD) and colorectal cancer (CRC). Akkermansia munciniphila (A. munciniphila) is a gram-negative anaerobic bacterium that is selectively decreased in the faecal microbiota of patients with IBD, but its causative role and molecular mechanism in blunting colitis-associated colorectal cancer (CAC) remain inconclusive. This study investigates how A. munciniphila engages the immune response in CAC.

Design Mice were given dextran sulfate sodium to induce colitis, followed by azoxymethane to establish CAC with or without pasteurised A. munciniphila or a specific outer membrane protein (Amuc_1100) treatment. Faeces from mice and patients with IBD or CRC were collected for 16S RNA sequencing. The effects of A. munciniphila or Amuc_1100 on the immune response in acute colitis and CAC were investigated.

Results A. munciniphila was significantly reduced in patients with IBD and mice with colitis or CRC. A. munciniphila or Amuc_1100 could improve colitis, with a reduction in infiltrating macrophages and CD8+ cytotoxic T lymphocytes (CTLs) in the colon. Their treatment also decreased CD16/32+ macrophages in the spleen and mesenteric lymph nodes (MLN) of colitis mice. Amuc_1100 elevated PD-1+ CTLs in the spleen. Moreover, A. munciniphila and Amuc_1100 blunted tumourigenesis by expanding CTLs in the colon and MLN. Remarkably, they activated CTLs in the MLN, as indicated by TNF-α induction and PD-1 downregulation. Amuc_1100 could stimulate and activate CTLs from splenocytes in CT26 cell conditioned medium.

Conclusions These data indicate that pasteurised A. munciniphila or Amuc_1100 can blunt colitis and CAC through the modulation of CTLs.

INTRODUCTION

Colorectal cancer (CRC) is the fourth leading cause of cancer-associated death worldwide. The initiation and progression of CRC involves complex interactions between genetics, diet, lifestyle and environmental factors. Inflammatory bowel disease (IBD), including Crohn’s disease and ulcerative colitis (UC), is a known risk factor for the development of CRC, namely, CAC. For every 10 units that in the inflammatory burden index increases in patients with IBD, the risk of CRC is increased by 2.1%. Changes in the composition of the gut microbiota have been associated with an increased risk of IBD and CRC. Fusobacterium nucleatum can increase CRC cell proliferation and tumour development and also promote CRC chemoresistance by modulating autophagy. Peptostreptococcus anaerobius could promote colorectal carcinogenesis and modulate tumour immunity. However, in many studies, the focus has been on the putative harmful effects of the microbiota, while their beneficial effects are somewhat neglected.
Figure 1 Decreased A. muciniphila was associated with colitis and CRC in human and mice. (A) Heatmap of faecal microbiota from patients with UC, Ad, CRC and healthy controls. (B) Comparison of the faecal microbiota among UC, Ad, CRC and controls at the phylum level. (C) FISH detection of A. muciniphila in the colon of human patients using a FITC-conjugated A. muciniphila specific probe (green). Scale bars, 50 µm. (D) qPCR validation of the relative abundance of A. muciniphila in human patients. (E) ROC curves of A. muciniphila in predicting UC. (F) Heatmap of faecal microbiota from mice with CAC at different time points. (G) Comparison of faecal microbiota in CAC mice at the phylum level. (H) FISH detection of A. muciniphila in the mice with colitis and CAC (at week 23). (I) The relative abundance of A. muciniphila in the mice with colitis and CAC (at week 23). (J) Comparison of faecal microbiota in CAC mice at the phylum level. The results were expressed as the mean±SEM and were analysed by ordinary one-way ANOVA with Tukey’s multiple comparisons. ***P<0.001. Ad, adenoma; ANOVA, analysis of variance; CAC, colitis-associated colorectal cancer; CRC, colorectal cancer; UC, ulcerative colitis.

Akkermansia muciniphila is a gram-negative anaerobic bacterium, which is the first and only representative member of the Verrucomicrobia phylum found in the human intestinal tract. It has been considered as a promising probiotics. The abundance of A. muciniphila is significantly reduced in metabolic disorders, such as diabetes and obesity. Supplementation with pasteurised A. muciniphila improved colonic infiltration of macrophages and the proportion of CD16/32⁺ macrophages in the spleen and mesenteric lymph nodes (MLN) of colitis mice. A. muciniphila also significantly decreased colonic infiltrating cytotoxic T lymphocytes (CTLs), as well as the proportion of CTLs in the spleen, while it increased the PD-1⁺ CTL subset. For CAC, pasteurised A. muciniphila/Amuc_1100 treatment increased CTLs in colon tumours and the MLN. Their treatment increased TNF-α⁺ CTLs and decreased the proportion of PD-1⁺ CTLs in the MLN. A. muciniphila increased the proportion of CTLs in primary splenocytes cultured in CT26 cell conditioned medium in vitro. Furthermore, these CTLs induced higher apoptotic rates than cells without A. muciniphila treatment. Collectively, these results highlight the protective effects of A. muciniphila in CAC, which occur through modulation of the proportion and function of CTLs. This study provides a novel evidence that A. muciniphila can serve as a potential therapeutic target in patients with CRC.

MATERIALS AND METHODS
Culture and pasteurisation of A. muciniphila
A. muciniphila MuCT (ATCC BAA-835) (online supplementary figure 1A) was cultured in brain heart infusion broth containing 10 mg/L resazurin (an oxidation-reduction indicator) under strict anaerobic conditions. A representative culture stock was used to determine the CFU/mL under anaerobic conditions by plate counting using mucin media containing 1% agarose. This culture was diluted with anaerobic PBS containing 2.5% glycerol to a final concentration of 1.5×10⁸ CFU/100 µL. Then, A. muciniphila was inactivated by pasteurisation for 30 min at 70°C. After pasteurisation, no viable A. muciniphila could be recovered in culture.

Expression and purification of Amuc_1100 in vitro
Amuc_1100 protein was produced according to a previous study. The PCR product of a His-tagged Amuc_1100 was cloned into the pET-26b (+) vector (GenScript, Nanjing, China) to construct the expression plasmid (online supplementary figure 1B and C), which was validated by visualising Mlu I and Xho I digested fragments on a agarose gel (online supplementary figure 1D). The plasmid pET-26b-Amuc_1100 was transformed into E. coli BL21 (DE3), which was cultured in LB broth with kanamycin (50 µg/mL) at 37°C and 220 rpm shaking. Isopropyl β-D-1-thiogalactopyranoside (1 mM) was added to the LB broth to induce Amuc_1100 expression during the logarithmic growth phase. The bacteria were harvested when the OD600 value was over one and then lysed by sonication, the centrifugal supernatant was used to purify the Amuc_1100 protein with Ni-16NTA His-Bind (Novagen, Merck Millipore, Massachusetts, USA). The result of Coomassie brilliant blue staining (online supplementary...
Figure 2  Supplementation with pasteurised *A. muciniphila* or Amuc_1100 improved colitis induced by DSS. (A) Weight changes are expressed as the mean change from the starting weight. (B) (DAI), (C) colon length and (D) spleen weight were analysed. (E) Representative histological images of colon tissues by H&E staining. Scale bars, 200 µm. (F) IHC staining and quantitation of cleaved-caspase 3 in the proximal colon. Scale bars, 50 µm. (G) The mRNA expression of proinflammatory cytokines in colon tissue, including TNF-α, IFN-γ, IL-1β, IL-6, IL-18 and IL-33. Data are presented as the means±SEM and were analysed by ordinary one-way ANOVA with Tukey’s multiple comparisons. *P<0.05, **p<0.01, ***p<0.001, compared with respective control; ##p<0.01, ###p<0.001, compared with DSS group. ANOVA, analysis of variance; DAI, disease activity index; DSS, dextran sulfate sodium; IHC, immunohistochemistry.

Patient

Human stool sample from active patients with UC, adenoma (Ad) and CRC as well as healthy controls were obtained from Jiangsu Province Hospital of Chinese Medicine (Nanjing, China) with patient informed consent under an approved institutional review board protocol. The characteristics of the patients are listed in online supplementary table 1. All of the patients with CRC were first diagnosed and then histologically confirmed. Exclusion criteria were predefined as follows: diabetes, hypertension, coronary heart disease and a history of CRC.

Animal experiments

Male C57BL/6J mice aged to 6–8 weeks were housed in groups of five mice per cage in a 12/12 hour day/night cycle with free access to food and water under conditions of controlled humidity (50%±5%) and temperature (22±2°C).

Acute colitis was induced by 2% dextran sulfate sodium (DSS) (MP Biomedicals, molecular weight 35–50 kDa) in the drinking water ad libitum for eight consecutive days (online supplementary figure 2A). To explore the effects of *A. muciniphila* on colitis, the mice were treated with oral administration of pasteurised *A. muciniphila* (1.5×10⁸ CFU) or the protein Amuc_1100 (3 µg) from 2 weeks before DSS treatment to sacrifice. Body weight and disease activity index (DAI) were observed every day according our previous study.²² The average weight loss, diarrhoea and bleeding comprised the DAI (online supplementary table 2).

The mouse CAC model was established by one intraperitoneal injection (i.p.) of 10 mg/kg azoxymethane (AOM) followed by three cycles of 2% DSS treatment (1 week per cycle) (online supplementary figure 3A). To test the impact of *A. muciniphila* on CAC, the mice were treated with pasteurised *A. muciniphila* (1.5×10⁷ CFU) or the Amuc_1100 (3 µg) from 2 weeks before AOM injection until sacrifice. Body weight and DAI were observed every day. Stool samples were collected at the 7th, 10th, 23rd weeks. The mice were euthanised at the 8th, 12th and 23rd weeks.

16S rRNA sequencing

Microbial DNA from stool samples was extracted, then 16S rRNA was amplified at the V3 hypervariable region. Sequencing was conducted by an Illumina MiSeq (PE300) and the data were analysed by Majorbio Bio-pharm Biotechnology (www.i-sanger.com). Operational taxonomic units (OTUs) were used for alpha diversity, richness and rarefaction curve analyses with a 0.97 threshold. In addition, partial least squares discriminant analysis (PLS-DA) was performed by Bray-Curtis based on OTU level.

*A. muciniphila* quantification

The abundance of *A. muciniphila* in stool samples was quantified by quantitative PCR (qPCR) in accordance with our previous study.¹⁹ Localisation of *A. muciniphila* at the surface of the intestinal mucosa in colitis and CRC was performed by fluorescent in situ hybridisation (FISH) as described previously.¹ The fluorescein isothiocyanate (FITC)-labelled *A. muciniphila* probe (FITC-CCTTGGCGTTGGCTTCAGAT) was synthesised.
Figure 3  Supplementation with pasteurised A. muciniphila or Amuc_1100 suppressed the activation and infiltration of CD8⁺ CTLs and macrophages in colitis mice. Representative FCM analysis and quantification of CTLs (A) and PD1⁺ CTLs (B) in the spleen. Representative FCM analysis and quantification of CD16/32⁺ macrophages (CD11b⁺F4/80⁺) (C) and PD1⁺ CD16/32⁺ macrophages (D) in the spleen. Scale bars, 50 µm. Representative FCM analysis and quantification of CD16/32⁺ macrophages (E) and PD1⁺ CD16/32⁺ macrophage (F) in the MLN. IHC staining of CD8a (G) and F4/80 (H) in the proximal colon. Data are presented as the means±SEM and were analysed by ordinary one-way ANOVA Tukey’s multiple comparisons. *P<0.05, **p<0.01, ***p<0.001, compared with respective control; #p<0.05, ##p<0.01, ###p<0.001, compared with DSS group. ANOVA, analysis of variance; CTL, cytotoxic T lymphocytes; DSS, dextran sulfate sodium; FCM, flow cytometry; IHC, immunohistochemistry; MLN, mesenteric lymph nodes.

Quantitative real-time polymerase chain reaction (qPCR)
Total RNA was extracted from colon tissues or CT26 cells and then the cDNA was synthesised with HiScript II Q RT SuperMix for qPCR (Vazyme Biotech). qPCR was performed using the QuantStudio 5 System (Thermo Fischer Scientific). The qPCR primers were designed and synthesised by GENERAY Biotechnology (Shanghai, China), and the primer sequences are listed in online supplementary table 3. GAPDH was used as the reference gene.

Histology and immunohistochemistry (IHC)
For histology assessment, colon tissue sections were fixed, dehydrated, embedded and sectioned (5 µm) for H&E staining. For IHC assays, the paraffin sections were deparaffinised, endogenous enzymes were inactivated and antigens were thermally repaired. The sections were then blocked and stained with CD8a, F4/80, cleaved-caspase 3, γH2AX or Ki67 antibodies, followed by corresponding secondary antibody and a Streptavidin Biotin Complex kit (Boster BioEngineering, Wuhan, China). The detailed information of antibodies for IHC analysis is listed in online supplementary table 4. Stained slides were scanned by Pannoramic SCAN (3DHISTECH Kft, Budapest, Hungary). IHC was quantified by Image-Pro-Plus software and the mean density was determined based on the rate of integral optical density sum and area.

Flow cytometry (FCM)
For the analysis of CTLs, splenocytes and MLNs cells were stained with fluorescently labelled antibodies (online supplementary table 4) including Fixable Viability Dye eFlour 780, PerCP-eFluor 710 anti-mouse CD3, FITC rat anti-mouse CD8a, PE rat anti-mouse TNF, BD Horizon BV421 hamster anti-Mouse CD279. To remove non-specific binding of fluorescently labelled antibodies to Fc receptors, the Fc receptors were blocked with purified anti-mouse CD16/32. For intracellular staining of TNF-α, the cells were treated with the Leucocyte Activation Cocktail for 12 hours. The cells were then fixed, permeabilised and washed using the Foxp3/Transcription Factor Staining Buffer Set Kit (#85-00-5523-00, eBioscience).
For the analysis of the macrophages, splenocytes and MLNs cells were stained with fluorescein labelled antibodies, including FITC rat anti-mouse CD11b, PE rat anti-mouse F4/80, PerCP-Cy5.5 anti-mouse CD16/32, anti-mouse CD279 (PD-1) and FcR block. Gates and quadrants were based on fluorescence minus one (FMO) control. The samples were detected with BD FACSVerse (BD Biosciences). The data were analysed by FlowJo_V10.

Coculture of primary mouse CTLs with CT26 cells
CT 26 cells were purchased from the Cell Bank of the Chinese Academy of Sciences (Shanghai, China) and cultured in RPMI-1640 medium containing 10% fetal bovine serum (FBS) at 37°C in a 5% CO2 humidified atmosphere. Splenocytes were isolated and cultured in fresh RPMI-1640 medium containing 20% FBS and CT26 conditioned medium (v/v, 1/1) with or without 10 µg/mL Amuc_1100 for 24 hours. Then, the cultured splenocytes were used to sort CTLs by FCM. These suspended CTLs and adherent CT26 cells were cocultured at 1:1 for 48 hours. The apoptotic rate of CT26 cells was examined with the FITC Annexin V Apoptosis Detection Kit I (#556547, BD Biosciences) according to the manufacturer’s instructions.

Statistical analysis
The receiver operating characteristic curve (ROC) analysis was performed by SPSS 22.0 software (Chicago, Illinois, USA). A two-tailed Student’s t-test was performed to compare the cell apoptotic rate and related gene expression in CT26 cells. All human and animal data were analysed by ordinary one-way analysis of variances with Tukey’s multiple comparisons. All plots are shown as the means±SEM. P<0.05 was considered as statistically significant.

RESULTS
The abundance of A. muciniphila was decreased in human and mice with colitis or CRC
16S rRNA sequencing indicated that the community composition could be clearly separated by PLS-DA and rarefaction curves between UC and healthy controls (online supplementary figure 4A-B). The diversity (Shannon index) and richness (Chao, ACE and Sobs indices) of the gut microbiota were significantly reduced in patients with UC in comparison with controls (online supplementary figure 4C-F). The sequencing data revealed a decreased abundance of A. muciniphila in patients with UC, Ad and CRC (figure 1A–B), which were verified by qPCR (figure 1C) and FISH analysis (figure 1D). ROC analysis revealed the predicted value of A. muciniphila in UC (AUC=0.674, p=0.0007) (figure 1E). In agreement with the human data, the diversity and richness of the microbiota in colitis mice were significantly decreased (online supplementary figure 2C-F). Unexpectedly, supplementation with Amuc_1100 increased the abundance of A. muciniphila in colitis mice (online supplementary figure 2G-I). There was a complete separation between the gut microbiota from CAC mice at different time points (online supplementary figure 5A-B). The diversity and richness of the gut microbiota in CAC mice were significantly reduced in the early stage (online supplementary figure 5C-F). During the process of CAC, A. muciniphila decreased in a time-dependent manner (figure 1F–G). FISH and qPCR data further confirmed that A. muciniphila abundance was decreased in colitis and CAC mice (figure 1H–I).

Pasteurised A. muciniphila or Amuc_1100 improved colitis in mice
Oral supplementation of pasteurised A. muciniphila or Amuc_1100 failed to recover the loss of body weight in colitis mice (figure 2A). However, Amuc_1100 significantly alleviated the DAI (figure 2B). Both pasteurised A. muciniphila and Amuc_1100 significantly relieved colon shortening (figure 2C) and splenomegaly (figure 2D). They also attenuated histological injuries in the proximal colon, such as the loss of histological structure, disruption of the epithelial barrier and pronounced inflammatory cell infiltration (figure 2E). IHC analysis demonstrated that pasteurised A. muciniphila or Amuc_1100 decreased the number of cleaved-caspase three positive cells (figure 2F). qPCR analyses of colon from colitis mice after their treatment revealed significant downregulation of TNF-α, IFN-γ, IL-1β, IL-6, IL-18 and IL-33 (figure 2G). Collectively, these results demonstrated that treatment with pasteurised A. muciniphila or Amuc_1100 could ameliorate DSS-induced acute colitis.

Reduced infiltration of CTLs and macrophages in the spleen and MLN by supplementation of Amuc_1100 or pasteurised A. muciniphila
To investigate the effects of A. muciniphila or Amuc_1100 on immune response, the proportion of CTLs and macrophages as well as their expressions of PD-1 and TNF-α were detected by FCM (online supplementary figure 6). The proportion of CTLs was significantly increased in the spleens of colitis mice (figure 3A). Oral supplementation of Amuc_1100 significantly decreased the proportion of CTLs (figure 3A) and rescued the reduced percentage of PD-1+ CTLs in the spleen (figure 3B). However, A. muciniphila and Amuc_1100 could not restore the reduction in CD4+ T cells (online supplementary figure 7A). They inhibited the elevation in proinflammatory CD16/32+ macrophages (CD11b+ F4/80+) in the spleen (figure 3C).

The proportion of CTLs (online supplementary figure 7B) and PD-1+ CTLs (online supplementary figure 7C) did not change in the MLN with or without pasteurised A. muciniphila or Amuc_1100 treatment. However, their treatment suppressed the increase in CD16/32+ macrophages (figure 3E), but they did not impact the expression of PD-1 on CD16/32+ macrophages in the spleen (figure 3D) or MLN (figure 3F). Furthermore, they alleviated the colonic infiltration of CTLs (figure 3G) and macrophages (figure 3H).

Supplementation with pasteurised A. muciniphila or Amuc_1100 improved clinical symptoms of CAC
Neither pasteurised A. muciniphila nor Amuc_1100 administration could offset the reduction in body weight observed in CAC mice (figure 4A). However, they significantly decreased the DAI, especially at week 23 (figure 4B). Pasteurised A. muciniphila or Amuc_1100 did not change the colon length (online supplementary figure 3B), while their treatment could mitigate splenomegaly in CAC mice (figure 4C). In addition, their treatment delayed the tumour formation (figure 4D-G) and online supplementary figure 3C) and decreased the tumour numbers and area by the 12th week (figure 4E-F). Their treatment also decreased the expression of γH2AX (figure 4H) and cleaved caspase 3 (figure 4I) in colon epithelial cells, which indicated that they could attenuate double-stranded DNA breaks and subsequent apoptosis in CAC mice. Furthermore, the expression of Ki67 was reduced on pasteurised A. muciniphila and Amuc_1100 treatment (figure 4J), suggesting that they may also inhibit cell proliferation. Collectively, these data demonstrated that oral
administration of pasteurised *A. muciniphila* or Amuc_1100 could prevent AOM/DSS-induced tumourigenesis by attenuating DNA damage, cell apoptosis and abnormal proliferation.

**Pasteurised *A. muciniphila* or Amuc_1100 modulated immune response in CAC mice**

Studies have shown that CTLs undergo aggressive proliferation, cloning and dynamic differentiation in the human tumour microenvironment. The proportion of CTLs in the MLN increased significantly only after the 23rd week (figure 5B and online supplementary figure 8B), while the proportion of PD-1<sup>+</sup> CTLs was significantly increased by the 12th week (figure 5A,C). Pasteurised *A. muciniphila* and Amuc_1100 significantly enhanced the increase in CTLs in the MLN of CAC mice (figure 5A–B), while they reduced the percentage of PD-1<sup>+</sup> CTLs starting in the 8th week (figure 5C). The TNF-α<sup>+</sup> CTLs cell subset percentage was significantly higher after their treatment for 8 and 12 weeks (figure 5A,D and online supplementary figure 8A). In the spleen, pasteurised *A. muciniphila* and Amuc_1100 significantly reduced the proportion of PD-1<sup>+</sup> CTLs by the 8th week (online supplementary figure 9). Furthermore, their treatment suppressed the colonic infiltration of CTLs in CAC mice (figure 5E). These data suggested that pasteurised *A. muciniphila* and Amuc_1100 increased the proportion of CTLs and enhanced their cytotoxic effect in CAC mice.

Pasteurised *A. muciniphila* and Amuc_1100 suppressed the increase percentage of macrophages in the spleen of CAC mice since by the 8th week (figure 6A,B), while they had no effect on the proportion of macrophages (online supplementary figure 10A) or CD16/32<sup>+</sup> macrophage in the MLN (online supplementary figure 10B). However, their treatment significantly inhibited the expression of PD-1 on CD16/32<sup>+</sup> macrophage in the MLN and spleen (figure 4B and online supplementary figure 10C) and alleviated the colonic infiltration of macrophages (figure 6C).

**Amuc_1100 induced activation of CTLs cultured in CT26 cell conditioned medium**

To explore the underlying antitumour mechanism of Amuc_1100, CTLs were isolated from the spleen of normal mice and cocultured with CT26 cells in vitro (figure 7A). Amuc_1100 significantly increased the proportion of CTLs in primary splenocytes cultured in CT26 cell conditioned medium (figure 7B). The purified CTLs pretreated with Amuc_1100 significantly increased the apoptosis of CT26 cells in the conditioned medium (figure 7C). qPCR of cocultured CT26 cells revealed a downregulation of Bcl-2 (figure 7D) and upregulation of caspase 3 (figure 7E). Intratumoural expression of cleaved-caspase 3 (figure 7F) was increased in CAC mice treated with pasteurised *A. muciniphila* or Amuc_1100.

**DISCUSSION**

The maintenance of intestinal homeostasis relies on coordinated interactions between the microbiota, the intestinal epithelium...
Gut microbiota

Figure 5 Pasteurised *A. muciniphila* or Amuc_1100 enhanced CTLs activation in CAC mice. FCM analysis of CTLs, PD-1⁺ CTLs and TNF-α⁺ CTLs in the MLN in the 12th week (A). Quantification of CTLs (B), PD-1⁺ CTLs (C) and TNF-α⁺ CTLs (D) in the 8th, 12th and 23rd week, respectively. (E) IHC staining of CD8a in the colon. Scale bars, 50 µm. Data are presented as the means±SEM and were analysed by ordinary one-way ANOVA with Tukey’s multiple comparisons. *P<0.05, **p<0.01, ***p<0.001, compared with control; #p<0.05, ##p<0.01, ###p<0.001, compared with AOM+DSS group.

ANOVA, analysis of variance; AOM, azoxymethane; CAC, colitis-associated colorectal cancer; CTL, cytotoxic T lymphocyte; DSS, dextran sulfate sodium; FCM, flow cytometry; IHC, immunohistochemistry.

The *A. muciniphila* type strain ATCC BAA-835 and a murine strain (designated 139) exerted anti-inflammatory effects on chronic colitis. Previous studies also revealed the protective effects of *A. muciniphila*-derived extracellular vesicles in DSS-induced colitis. In this study, oral administration of pasteurised *A. muciniphila* or Amuc_1100 significantly improved acute colitis, including DAI, splenomegaly, colon shortening and histological injury. *A. muciniphila* is known to be important in improving the host metabolic functions in obese and diabetic people. Thus, it seemed reasonable that *A. muciniphila* and Amuc_1100 failed to restore the body weight loss observed in colitis or CAC mice. Their treatment also inhibited colonic expression of proinflammatory cytokines, such as IFN-γ, TNF-α and IL-1β. However, the BAA-835 strain previously failed to promote short-term intestinal inflammation in gnotobiotic IL-10-deficient mice. Oral supplementation of pasteurised *A. muciniphila* or Amuc_1100 could not shape the microbial community in colitis mice, but they did enhance the abundance of *A. muciniphila*. It seems that the protective effects are dependent on the microbiota. CTLs are emerging as major mediators of human IBD, which is also prognostically linked to CTL profiling. The main contribution to colitis development was mediated by cytokine producing, proliferative, CTLs in the intraepithelial compartment of mice. Amuc_1100 significantly decreased the proportion of CTLs in the spleen and the colonic infiltration of CTLs. Moreover, Amuc_1100 inhibited the activation of CTLs through the upregulation of PD-1. Taken together, Amuc_1100 could improve DSS-induced colitis through the modulation of the population and activation of CTLs.

Beyond the effect on colitis, this study uncovered that Amuc_1100 provoked an antitumour immune response in CAC. Oral supplementation of pasteurised *A. muciniphila* or Amuc_1100 delayed tumour formation and reduced tumour...
number and size. As a known genotoxic stress, AOM can trigger DNA damage and subsequent cell apoptosis and abnormal proliferation.33 This study revealed that pasteurised A. muciniphila and Amuc_1100 inhibited the expression of γH2AX and Ki67 in colon tissue, while they triggered the apoptosis of tumour cells in vitro and in vivo. CTLs are considered the principal cellular mediators of antitumour immunity and a prognosis marker for CRC.34–36 Pasteurised A. muciniphila and Amuc_1100 significantly increased CTL percentage in the MLN and colon. Interestingly, they also upregulated TNF-α in CTLs, which aggravated tumour cells apoptosis.37 38 The different microenvironment of colitis and CAC may partially explain this discrepancy. A. muciniphila and Amuc_1100 might promote the recruitment of CTLs through increasing the secretion of specific chemokines in tumour cells, where they were activated due to MHCI upregulation.39 However, the underlying mechanisms need further investigation.

Immune checkpoint inhibitors targeting the PD-1/programmed death-ligand 1 (PD-D1-L1) axis are widely used.40 41 The composition of the commensal microbiota in patients is associated with the therapeutic efficacy of anti-PD-1 monoclonal antibody.42 Oral gavages with A. muciniphila increased the efficacy of a PD-1 blockade with respect to tumour growth.18 The expression of PD-1 was upregulated on mucosal CD4, CD8αβ and TCRαβ cells of the colon and the ileum, which correlated with CAC progression.43 Although the majority of CRCs with high CTLs infiltration have favourable outcomes, those with concurrently high PD-L1 gene expression have poor clinical outcomes.44 In the present study, the proportion of PD-1+ CTLs was significantly reduced in CAC mice treated with pasteurised A. muciniphila or Amuc_1100. T cells expressing lower levels of PD-1 produced higher levels of IFN-γ and TNF-α in comparison to those with high PD-1 expression, which might in turn trigger cell apoptosis and induce PD-L1 expression in the tumour microenvironment, thereby shaping T-cell PD-1 status.45

Macrophages are the gatekeepers of intestinal immune homeostasis, M1 macrophages elicit rapid proinflammatory responses to infection and tissue damage.46 47 In the present study, colitis was blunted by A. muciniphila or Amuc_1100 through a reduction in the proportion of macrophage, especially CD16/32+ macrophages (M1). Previous studies revealed that CTLs could promote the recruitment and activation of macrophages during inflammation.18 Thus, A. muciniphila or Amuc_1100 may inhibit the recruitment and activation of macrophages by the reduction of CTLs. Conversely, they could also inhibit the activation of macrophages by increased CTLs. Clinical and experimental evidence has shown that tumour-associated macrophages (TAMs) promote cancer initiation and progression. The proportion of macrophage in the spleen as well as the colonic infiltration of macrophages were significantly increased in CAC mice. A. muciniphila or Amuc_1100 treatment decreased these macrophage populations without disrupting their expression of CD16/32 or PD-1, indicating that their antitumour capabilities were independent of macrophage activation. A recent study reported that TAMs could promote CTL tumour infiltration through the production of CXCL9.49 Thus, A. muciniphila or Amuc_1100 may also inhibit macrophage functions, such as chemokines secretion, and enhance the tumour infiltration of CTLs. However, the interactions of macrophages and CTLs on A. muciniphila or Amuc_1100 treatment remain unclear.

This work reveals a key role for A. muciniphila as a critical regulator of intestinal homeostasis. Pasteurised A. muciniphila or Amuc_1100 could reshape the immune landscape to blunt
Amuc_1100 induced CTL activation in CT26 cell conditioned medium. (A) Schematic representation of CTL isolation and coculture with CT26 cells. (B) The fraction of CTLs in the splenocytes cultured with CT26 cell conditioned medium with or without Amuc_1100 for 72 hours. (C) The apoptotic rate of CT26 cells cocultured with sorted CTLs for 48 hours (n=3). The mRNA expression of Bcl-2 (D) and caspase 3 (E) and in CT26 cells was detected by qPCR. (F) Intratumoural cleaved-caspase 3 in CAC mice were analysed by IHC. Scale bars, 20 μm. Data are presented as the means±SEM and were analysed with t test. *P<0.05, **p<0.01, ***p<0.001, compared with control. CAC, colitis-associated colorectal cancer; CTL, cytotoxic T lymphocyte; IHC, immunohistochemistry.

Acknowledgements The authors would like to thank the participants that have made this research possible. We also thank the Majorbio Bio-pharm Biotechnology Co., Ltd for 16S sequencing and data analysis platform (www.i-sanger.com).

Contributors LW, LT and YF carried out most experiments and analysed the data. SZ, MH and CZ provided clinical sample and performed data analyses. GY and JZ helped with animal experiments and analysis. SC, QW, ZZ and LL designed the experiments and analysed data. ZZ and LL directed the research and wrote the manuscript with input from coauthors.

Funding This project was supported by the National Natural Science Foundations of China (81973096 and 81502801) and Postgraduate Research & Practice Innovation Program of Jiangsu Province (KYX18_1513).

Competing interests None declared.

Patient and public involvement Patients and/or the public were not involved in the design, or conduct, or reporting, or dissemination plans of this research.

Patient consent for publication Not required.

Ethics approval All animal procedures were approved by the Animal Ethical and Welfare Committee of Nanjing Medical University (IACUC-1812031).

Provenance and peer review Not commissioned; externally peer reviewed.

Data availability statement Data are available in a public, open access repository. All data relevant to the study are included in the article or uploaded as supplementary information. The 16S sequencing data from human and mice faeces had been submitted to NCBI (Sequence Read Archive, SRA). This BioProject accession number is PRJNA596333 (https://www.ncbi.nlm.nih.gov/bioproject/PRJNA596333).

Open access This is an open access article distributed in accordance with the Creative Commons Attribution Non Commercial (CC BY-NC 4.0) license, which permits others to distribute, remix, adapt, build upon this work non-commercially, and license their derivative works on different terms, provided the original work is properly cited, appropriate credit is given, any changes made indicated, and the use is non-commercial. See: http://creativecommons.org/licenses/by-nc/4.0/.

ORCID iD Zhan Zhang http://orcid.org/0000-0002-9685-5753
REFERENCES


Supplementary material Gut

Supplementary material Gut


PLS-DA on OTU level

Supplementary material Gut

DOI: 10.1136/gutjnl-2019-320105

Supplementary material

A

F4/80+CD11b+ cells in Lymphocytes (%)

B

CD16/32+ cells in CD11b+ F4/80+ cells (%)

C

PD-1+ cells in CD16/32+ CD11b+ F4/80+ cells (%)

Control
AOM+DSS
A. muciniphila+DSS
A. muciniphila+DSS
A. muciniphila+DSS
A. muciniphila+DSS
A. muciniphila+DSS

8 w
12 w
23 w

8 w
12 w
23 w

8 w
12 w
23 w

8 w
12 w
23 w

8 w
12 w
23 w

A purified membrane protein from *Akkermansia muciniphila* or the pasteurized bacterium blunts colitis associated tumourigenesis by modulation of CD8$^+$ T cells in mice

Lijuan Wang, Lei Tang, Yiming Feng, Suying Zhao, Mei Han, Chuan Zhang, Gehui Yuan, Jun Zhu, Shuyuan Cao, Qian Wu, Lei Li, Zhan Zhang

Supplementary data
MATERIALS AND METHODS

Western blotting

Amuc_1100 was a protein of 32 kDa, which was one of the most abundant outer membrane proteins encoded by a specific Type IV pili gene cluster in A. muciniphila. To test this protein, we added a His tag to the c-terminal of Amuc_1100. The purified Amuc_1100 with the His tag could be detected with a His-tag antibody. An equal amount of protein was subjected to 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis and transferred onto polyvinylidene difluoride membranes. The membranes were stained with a primary antibody toward the His tag (A100186-100, GenScript) and then incubated with secondary antibody. Protein bands were visualized by a gel imaging system (Bio-Rad).
**Figure legends**

Supplementary Figure 1. Expression and purification of Amuc_1100 in vitro. (A) Representative Gram-staining of *A. muciniphila*. (B) The coding sequence of Amuc_1100 amplified by PCR. (C) His-tagged Amuc_1100 was cloned into the pET-26b(+) vector using restriction endonuclease sites (NdeⅠ/XhoⅠ). (D) The recombinant plasmids were transformed into BL21 *E. coli*. Plasmid digested by MluⅠ and XhoⅠ were analysed by 1% agarose gel electrophoresis. (E) The purified Amuc_1100 protein was stained by Coomassie brilliant blue. His-tagged Amuc_1100 was eluted with different concentrations of imidazole. (F) The purified Amuc_1100 was detected by Western blotting.

Supplementary Figure 2. Effects of pasteurized *A. muciniphila* or Amuc_1100 on the faecal microbiota in colitis mice. (A) Schematic diagram of the experimental study design. (B) PLS-DA plot of the faecal microbiota colitis of mice treated with *A. muciniphila* or Amuc_1100. The Shannon (C), Chao (D), Ace (E) and Sobs (F) indices on the OUT level were analysed. (G) Relative abundance of bacterial species. (H) Ternary plots of the relative abundances of species in colitis mice. (I) Pie chart of the faecal microbiota composition on the species level. Data are presented as the mean± SEM and were analyzed ordinary one-way ANOVA with Tukey’s multiple comparisons. *P< 0.05, **P< 0.01, ***P< 0.001, compared with respective control.

Supplementary Figure 3. Supplementation with pasteurized *A. muciniphila* or
Amuc_1100 suppressed carcinogenesis in mice. (A) Schematic diagram of colitis associated colon cancer (CAC) model induced by AOM/DSS in mice. (B) The colon length at the 8th, 12th and 23rd week. (C) Tumour incidence.

Supplementary Figure 4. The microbial composition in control, ulcerative colitis (UC), adenoma (Ad) and CRC patients. (A) Partial least squares discriminant analysis (PLS-DA) plot of the faecal microbiota. (B) Rarefaction curves were used to evaluate the microbial richness of human faeces. (C) Shannon, (D) Chao, (E) ACE and (F) Sobs indices were analysed on OUT level in human. Sobs, Chao and Ace indices reflect community richness, and the Shannon index represents community diversity. Data are presented as the means± SEM and analyzed by ordinary one-way ANOVA with Tukey’s multiple comparisons. ***P < 0.001, compared with control; *P < 0.05, **P < 0.01, ***P < 0.001, compared with UC.

Supplementary Figure 5. The microbial composition of mice with CAC. (A) PLS-DA plot of the faecal microbiota. (B) Rarefaction curves of the microbial richness in mice. (C) Shannon, (D) Sobs, (E) Chao and (F) ACE indices on the OUT level were analyzed. Sobs, Chao and Ace indices reflect community richness, and the Shannon index represents community diversity. Data are presented as the means± SEM and were analysed by ordinary one-way ANOVA Tukey’s multiple comparisons. **P < 0.01, ***P < 0.001, compared with 0 w.
Supplementary Figure 6. FCM gating strategies. (A) Gating strategy for the analysis of the percentage of CD8+ T cells in CD3+ T cells and PD-1+ or TNF-α+ cells in CD8+ T cells. (B) Gating strategy for the analysis of the percentage of CD11b+F4/80+ cells in lymphocytes, CD16/32+ cells in CD11b+F4/80+ cells and PD-1+ cells in CD16/32+ CD11b+F4/80+ cells.

Supplementary Figure 7. Effects of A. muciniphila or Amuc_1100 on T cells in colitis mice. (A) The percentage of CD4+ T cells in CD3+ T cells in the spleen. (B) The fraction of CD8+ T cells in CD3+ T cells and (C) the expression of PD-1 in CD8+ T cells in the mesenteric lymph node (MLN). Data are presented as the means± SEM and were analysed by ordinary one-way ANOVA with Tukey’s multiple comparisons. **P< 0.01, ***P< 0.001, compared with respective control.

Supplementary Figure 8. A. muciniphila or Amuc_1100 enhanced CTLs activation in CAC mice. FCM analysis of CTLs, PD-1+ CTLs and TNF-α CTLs in the MLN in the 8th (A) and 23rd week (B).

Supplementary Figure 9. Effects of pasteurized A. muciniphila or Amuc_1100 on CD8+ T cells in the spleens of CAC mice. (A) The percentage of CD8+ T cells at 8th and 23rd week. (B) The fraction of TNF-α+ cells in CD8+T cells at 8th and 23rd week. (C) The fraction of PD-1+ cells in CD8+T cells at 8th and 23rd week. Data are presented as the means± SEM and were analysed with ordinary one-way ANOVA
with Tukey’s multiple comparisons. *$P<0.05$, **$P<0.01$, ***$P<0.001$, compared with control; *$P<0.05$, **$P<0.01$, compared with AOM+DSS group.

Supplementary Figure 10. Effects of pasteurized *A. muciniphila* or Amuc_1100 on macrophages in the MLN of CAC mice. (A) The percentage of macrophage (CD11b*F4/80*) cells in the 8th, 12th and 23rd week. (B) The expression of CD16/32 in macrophages. (C) The expression of PD-1 in macrophages. Data are presented as the means± SEM and were analyzed with ordinary one-way ANOVA with Tukey’s multiple comparisons. *$P<0.05$, **$P<0.01$, ***$P<0.001$, compared with control; *$P<0.05$, ###$P<0.001$, compared with AOM+DSS group.
**Supplementary Table 1. Patients Characteristic**

<table>
<thead>
<tr>
<th></th>
<th>Control (n=72)</th>
<th>Ulcerative colitis (n=58)</th>
<th>Adenomatous polyps (n=18)</th>
<th>Colorectal cancer (n=22)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gender, male/female</td>
<td>36/36</td>
<td>31/27</td>
<td>13/5</td>
<td>17/5</td>
</tr>
<tr>
<td>Age (year)</td>
<td>44.75±17.7</td>
<td>42.8±14.9</td>
<td>62.6±9.1</td>
<td>64.3±10.1</td>
</tr>
<tr>
<td>Smoking</td>
<td>13.9%</td>
<td>5.2%</td>
<td>22.2%</td>
<td>40.9%</td>
</tr>
<tr>
<td>Glucose (mmol/L) a</td>
<td>4.95±0.72</td>
<td>4.32±0.65</td>
<td>4.95±0.72</td>
<td>5.37±1.33</td>
</tr>
<tr>
<td>TG (mmol/L) a</td>
<td>1.58±1.21</td>
<td>1.22±0.69</td>
<td>1.95±1.03</td>
<td>1.38±0.21</td>
</tr>
</tbody>
</table>

* a, Mean±SEM
Supplementary Table 2. Scoring system for disease activity index

<table>
<thead>
<tr>
<th>Score</th>
<th>Body weight loss</th>
<th>Stool consistency</th>
<th>Bleeding</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>none</td>
<td>none</td>
<td>none</td>
</tr>
<tr>
<td>1</td>
<td>0-5%</td>
<td>---</td>
<td>trace</td>
</tr>
<tr>
<td>2</td>
<td>5-10%</td>
<td>loose stool</td>
<td>mild hemoccult</td>
</tr>
<tr>
<td>3</td>
<td>10-20%</td>
<td>---</td>
<td>obvious hemoccult</td>
</tr>
<tr>
<td>4</td>
<td>&gt;20%</td>
<td>diarrhea</td>
<td>gross bleeding</td>
</tr>
</tbody>
</table>
### Supplementary Table 3. Specific primers sequences

<table>
<thead>
<tr>
<th>Gene Name</th>
<th>Forward Sequence</th>
<th>Reverse Sequence</th>
</tr>
</thead>
<tbody>
<tr>
<td>A. <em>muciniphila</em></td>
<td>AGAGGTCTCAAGCGTTGTTCGGAA</td>
<td>TTTGCTCCCCTGGCCTTGTC</td>
</tr>
<tr>
<td>Amuc_1100</td>
<td>GGGTACCATATGATCGTCAATTCCAAACGC</td>
<td>CTTGCTCGAGATCTTCAGACGGTTCCTG</td>
</tr>
<tr>
<td>EUB</td>
<td>AGAGTTTGATCCTGGCTC</td>
<td>TGCTGCCTCCCTAGGGAGT</td>
</tr>
<tr>
<td>16S rRNA</td>
<td>GTGCCAGCMGCCGCGGTAA</td>
<td>GGACTACHVGGGTWTCTTAAT</td>
</tr>
<tr>
<td>Mus-TNF-α</td>
<td>CCCTCACACTCAAAAACAC</td>
<td>ACAAGGTACAACCCATCGGC</td>
</tr>
<tr>
<td>Mus-IL-6</td>
<td>GAGGATACACCCTCACCAGACC</td>
<td>AAGTGCATCATCGTTGTTCTACA</td>
</tr>
<tr>
<td>Mus-IL-18</td>
<td>GTTTACAAGCATCAGGCACAGC</td>
<td>GGTTTGAGGCGGCTTTCTTCTT</td>
</tr>
<tr>
<td>Mus-IFN-γ</td>
<td>GCCCAAGCATTCAATGAGC</td>
<td>ATCTCTTCCCACCCGGAAT</td>
</tr>
<tr>
<td>Mus-IL-1β</td>
<td>GCCACCTTTTAGACAGTGAGG</td>
<td>ATGTGCTGCTGCGAGATTTTC</td>
</tr>
<tr>
<td>Mus-IL-33</td>
<td>AAGTACAGCATCAGGCACAGC</td>
<td>TTATTTTGAAGGGGAGACC</td>
</tr>
<tr>
<td>Mus-Caspase-3</td>
<td>TCTGACTGGAAGCCGAAA</td>
<td>AGGGACTGGATGAACCACGAC</td>
</tr>
<tr>
<td>Mus-Bcl-2</td>
<td>ATGATAACCCGGAGATCGT</td>
<td>GTTCAGGTACTCAGTCACC</td>
</tr>
</tbody>
</table>
Mus-GAPDH AGGTAGTGAGTTGAGGTCA

TGTAGACCATTGAGTGAGGTCA

<table>
<thead>
<tr>
<th>Antibodies</th>
<th>Use</th>
<th>Source</th>
<th>Identifier</th>
</tr>
</thead>
<tbody>
<tr>
<td>CD3 PerCP-eFluor 710</td>
<td>Flow Cytometry</td>
<td>eBioscience</td>
<td>46-0032-80</td>
</tr>
<tr>
<td>CD16/32-PerCP-Cy5.5</td>
<td>Flow Cytometry</td>
<td>eBioscience</td>
<td>45-0161-80</td>
</tr>
<tr>
<td>FcR Block</td>
<td>Flow Cytometry</td>
<td>eBioscience</td>
<td>14-0161-86</td>
</tr>
<tr>
<td>Fixable Viability Dye eFlour™ 780</td>
<td>Flow Cytometry</td>
<td>eBioscience</td>
<td>65-0865</td>
</tr>
<tr>
<td>CD279 (PD-1)-BV421</td>
<td>Flow Cytometry</td>
<td>BD Biosciences</td>
<td>562584</td>
</tr>
<tr>
<td>CD4-FITC</td>
<td>Flow Cytometry</td>
<td>BD Biosciences</td>
<td>553046</td>
</tr>
<tr>
<td>CD8a-FITC</td>
<td>Flow Cytometry</td>
<td>BD Biosciences</td>
<td>553030</td>
</tr>
<tr>
<td>CD11b-FITC</td>
<td>Flow Cytometry</td>
<td>BD Biosciences</td>
<td>557396</td>
</tr>
<tr>
<td>F4/80-PE</td>
<td>Flow Cytometry</td>
<td>BD Biosciences</td>
<td>565410</td>
</tr>
<tr>
<td>TNF-PE</td>
<td>Flow Cytometry</td>
<td>BD Biosciences</td>
<td>561063</td>
</tr>
<tr>
<td>Annexin V/PI</td>
<td>Flow Cytometry</td>
<td>BD Biosciences</td>
<td>556547</td>
</tr>
<tr>
<td>CD8 (IHC)</td>
<td>IHC</td>
<td>Abcam</td>
<td>ab108343</td>
</tr>
<tr>
<td>F4/80 (IHC)</td>
<td>IHC</td>
<td>Santa Cruz Biotechnology</td>
<td>sc-377009</td>
</tr>
<tr>
<td>Antibody</td>
<td>Staining Method</td>
<td>Vendor</td>
<td>Catalog Number</td>
</tr>
<tr>
<td>-------------------------</td>
<td>-----------------</td>
<td>-------------------------------</td>
<td>----------------</td>
</tr>
<tr>
<td>Cleaved-caspase 3</td>
<td>IHC</td>
<td>Cell Signaling Technology</td>
<td>9664</td>
</tr>
<tr>
<td>γH2AX</td>
<td>IHC</td>
<td>Cell Signaling Technology</td>
<td>7631</td>
</tr>
<tr>
<td>Ki67</td>
<td>IHC</td>
<td>Cell Signaling Technology</td>
<td>9449</td>
</tr>
<tr>
<td>His-tag</td>
<td>WB</td>
<td>GenScript</td>
<td>A100186-100</td>
</tr>
</tbody>
</table>