Effects of dihydroartemisinin, a metabolite of artemisinin, on colon cancer chemoprevention and adaptive immune regulation

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Abstract

Background Artemisinin (ART) is an anti-malaria natural compound with a moderate anticancer action. As a metabolite of ART, dihydroartemisinin (DHA) may have stronger anti-colorectal cancer (CRC) bioactivities. However, the effects of DHA and ART on CRC chemoprevention, including adaptive immune regulation, have not been systematically evaluated and compared.

Methods Coupled with a newly-established HPLC analytical method, enteric microbiome biotransformation was conducted to identify if the DHA is a gut microbial metabolite of ART. The anti-CRC potential of these compounds was compared using two different human CRC cell lines for cell cycle arrest, apoptotic induction, and anti-inflammation activities. Naive CD4+ T cells were also obtained for testing the compounds on the differentiation of Treg, Th1 and Th17.

Results Using compound extraction and analytical methods, we observed for the first time that ART completely converted into its metabolites by gut microbiome within 24 h, but no DHA was detected. Although ART did not obviously influence cancer cell growth in the concentration tested, DHA very significantly inhibited the cancer cell growth at relatively low concentrations. DHA included G2/M cell cycle arrest via upregulation of cyclin A and apoptosis. Both ART and DHA downregulated the pro-inflammatory cytokine expression. The DHA significantly promoted Treg cell proliferation, while both ART and DHA inhibited Th1 and Th17 cell differentiation.

Conclusions As a metabolite of ART, DHA possessed stronger anti-CRC activities. The DHA significantly inhibited cell growth via cell cycle arrest, apoptosis induction and anti-inflammation actions. The adaptive immune regulation is a related mechanism of actions for the observed effects.

Keywords Dihydroartemisinin · Artemisinin · Cell cycle and apoptosis · Colorectal cancer · T helper 17 cells · Anti-inflammation and adaptive immunity
Introduction

Artemisia annua L., or artemisia, is an herb traditionally used in Chinese medicine to treat fever, inflammation, and malaria infections [1]. One of its active compounds, artemisinin, a sesquiterpene trioxane lactone agent, was identified by Chinese scientists in the 1970s. This novel compound has been used in combination with other drugs to treat drug-resistant malaria and has saved millions of lives. The 2015 Nobel Prize in Physiology or Medicine was awarded to Professor Youyou Tu for her key contributions to the discovery of artemisinin (ART) [2, 3]. Afterwards, Professor Tu discovered dihydroartemisinin (DHA), which greatly improved ART’s water solubility and treatment efficacy against malaria [4].

Any given botanical likely consists of many chemically known or unknown single constituents, and some of the constituents could be pharmacologically categorized as bioactive compounds. Meanwhile, medicinal herbs and their derivatives, including ART, are almost always taken orally. After intake, their parent compounds could be bio-transformed by the enteric microbiome, and the metabolites could possess even higher bioactivities compared with the parent ones [5, 6]. Studies have shown that after oral ART administration, DHA can be detected in plasma, suggesting that the latter is a metabolite of the former [7, 8]. However, after ART oral ingestion, whether the parent compound ART is metabolized by the enteric microbiome into DHA has not been examined.

When botanical-origin novel compounds introduced for certain medical treatment in humans, researchers often start to look for their additional therapeutic capabilities, especially from the herb’s original and empirical indications, such as using artemisia against inflammation. Consequently, both ART and DHA have been used to evaluate their effects on inflammatory bowel disease and variable activities have been reported [9, 10].

Previous data indicated that inflammatory bowel disease (IBD), including ulcerative colitis and Crohn’s disease, is a risk factor for colon cancer initiation and development [11, 12]. Epidemiological, experimental, and clinical studies provide evidence that anti-inflammatory phytochemicals possess unique modes of action for the body’s homeostasis to prevent the disease and its progression [13]. Interestingly, salicylate, the active metabolite of aspirin, and the antidiabetic drug metformin are both originally derived from botanicals, which activate the energy sensing kinase AMPK and show anti-inflammation and anticancer effects, while ART and its analogs also activate the AMPK pathway [14, 15].

Artemisia has been reported to possess both anti-inflammatory and anti-cancer actions with a moderate efficacy against colorectal cancer (CRC) [16]. With its established safety record in millions of malarial patients, ART is also being investigated in diseases like infections, inflammation, and cancers [17]. Due to its relatively low anticancer potential, the perspective of cancer chemoprevention of ART is restricted. However, the effects of DHA, a metabolite of ART, on CRC chemoprevention have not been systematically evaluated especially in comparison with the ART. In this study, using a newly-established HPLC analytical method, enteric microbiome biotransformation will be conducted to identify if the DHA is a gut microbial metabolite of ART. Then, the anti-CRC potential of the two compounds will be compared using two different human colon cancer cell lines for cell cycle arrest and induction of apoptosis. Furthermore, their mechanisms of action in relation to the anti-inflammation and adaptive immune regulation will be examined.

Materials and methods

Reagents

Artemisinin and dihydroartemisinin are obtained from Sigma-Aldrich (St. Louis, MO). The chemicals and solvents used were all for high performance liquid chromatography or HPLC grade. Plasticware was purchased from Falcon Labware (Franklin Lakes, NJ). Test media, insulin trypsin, glutamine, and buffered saline were purchased from Mediatech (Herndon, VA). Antibiotics, such as streptomycin and penicillin, were purchased from Sigma-Aldrich (St. Louis, MO). Fetal bovine serum (FBS) was purchased from Life Technologies Corporation (Grand Island, NY). For the cell proliferation study, the MTS kit was purchased from Promega (Madison, WI). Cell cycle staining buffer was purchased from Biosciences Pharmingen (San Diego, CA), and the cell apoptosis detection kit was purchased from Biosciences (Rockville, MD).

Biotransformation and sample processing

The human fecal specimens were collected from five healthy adult male American subjects (no smoking and alcohol, no current medication, no use of antibiotics) and stored in a −85 °C freezer. The fecal specimens were mixed, and the mixture was suspended well in phosphate buffer to obtain fecal slurry. The slurry was filtered through muslin to remove particulate material. The resulting fecal slurry was used as intestinal microbiome fraction. 1 μl of the fecal slurry was mixed with 14 ml anaerobic medium containing 3.75 mg of ART. They were anaerobically incubated at 37 °C for 12 and 24 h. Then, the reaction mixture was extracted with ethyl ether. The ethyl ether layer was collected.
and dried in a water bath (37 °C). Then, the residue was dissolved in methanol. The methanol solution was centrifuged at 13,000×g for 10 min before HPLC analysis.

**HPLC analysis**

Our laboratory uses a Waters instrument 2965 (Milford, MA) for HPLC analysis. This system (Model 996) was used for constituent peak identification and integration, supported by Waters Empower software. The separation was carried out on a Prodigy ODS (2) column (5 µm, 250 × 3.2 mm I.D.) (Phenomenex, Torrance, CA) with a guard column (Phenomenex ProdigyTM, 3.0 × 4.0 mm I.D.). For HPLC analysis, a 20-µl sample was injected into the column and eluted at room temperature with a constant flow rate of 1.0 ml/min. Acetonitrile (solvent A) and water (solvent B) were used as the mobile phase. Gradient elution started with 40% solvent A and 60% solvent B, changed to 60% A for 10 min and held for 2 min, then changed to 90% A for 3 min and held for 3 min. Finally, it was changed to 40% A for 4 min and held for 3 min to rebalance the separated column. The detection wavelength was set to 210 nm.

**Cancer cell lines and cell culture**

HCT-116 and HT-29 human colorectal cancer cells and rat small intestine epithelial cell line IEC-6 cells were purchased from American Type Culture Collection (Manassas, VA). HCT-116 and HT-29 cells were grown in McCoy’s 5A medium, and IEC-6 cells were grown in DMEM media, with 10% FBS and 50 IU streptomycin-penicillin in the humidified air at 37 °C with 5% CO₂.

**Cell proliferation analysis**

The solvent used to dissolve ART and DHA were DMSO. The 96-well plates were used to seed cells and test response to drug treatment. Drug treatment durations were 24, 48, or 72 h. MTS assay was employed to determine cell proliferation. As our previous publications, absorbance was recorded at 490 nm [18, 19]. DMSO (0.5%) was used to dissolve the test drugs. Controls were exposed to culture medium containing 0.5% DMSO without drugs. The 0.5% DMSO did not affect the proliferation of both cell lines. Data was expressed as percent of control (DMSO vehicle) and set at 100% [5, 20].

**Cell cycle and cyclin A assays**

HCT-116 cells were seeded on 24-well plates. On day 2, the medium was changed, during which different ART and DHA concentrations were used to treat the cells. After 48 h, the cells were harvested and fixed with 80% ethanol. After 0.25% Triton X-100 treatment, the cells were resuspended in PBS containing PI, RNase, and cyclin A-FITC. FACScan flow cytometer (Becton Dickinson, Mountain View, CA) and FlowJo 10.7.1 software (Tree Star, Ashland, OR) were used to assay cell cycle and cyclin A [21]. At least 10,000 cells were recorded for each assay.

**Apoptosis analysis**

HCT-116 cells were seeded on 24-well plates. On day 2, the medium was changed, during which different ART and DHA concentrations were used to treat the cells. After 48 h, both floating and adherent cells were harvested. The cells were stained with PI and annexin V. A FACScan flow cytometer were used to assay apoptotic cells. At least 20,000 cells were recorded for each assay.

**IL-8 secretion assay**

Using 24-well plates, HT-29 cells were cultured for 48 h. Fresh medium containing lipopolysaccharide (LPS, 100 ng/ml) was added, which served as a control. Different concentrations of ART/DHA and LPS added in the medium were set as the experimental group. After incubating for 6 h, the secreted IL-8 in the medium was determined by enzyme-linked immunosorbent assay.

**Naive CD4 cell isolation and differentiation of T-helper cells**

Single cell suspensions were prepared by using 4–6 week old naive C57BL/6 (B6) mouse spleens. Pooled splenocytes (after erythrocyte lysis) were depleted of CD11b + , CD8α + , and CD19 + cells using biotinylated primary antibodies (BioLegend, San Diego, CA, USA) and streptavidin-coated secondary magnetic particles ( Stem Cell Technologies, Vancouver, BC, Canada). T cell differentiation was based on our previous publications [22, 23]. Test compounds were initially added into the culture medium. After 3 days, the differentiated cells were harvested, then in the presence of Brefeldin A, the cells were restimulated for 4 h with PMA (50 ng/ml) and ionomycin (750 ng/ml) (Sigma-Aldrich).

**Statistical analysis**

Data were expressed as mean ± SE. In general, a one-way ANOVA was used to define the data significant level. In some instances, Student’s t test was employed for two-group comparison. In all cases, P < 0.05 was considered as statistically significant.
Fig. 1 Chemical structures of artemisinin (ART) and dihydroartemisinin (DHA), and gut microbiome transformation of ART. A The in vivo metabolism from ART to DHA. B Purity determination of ART and DHA. HPLC chromatograms record at 210 nm, and UV spectra from 196 to 400 nm are presented. C Gut microbiome biotransformation of ART. Chromatograms for gut microbiome control and ART incubated at 37 °C for 0, 12 and 24 h are presented. ART is absolutely converted to its metabolites in 24 h, including a compound at Rt=7.8 min, while DHA is not detected. The UV spectra of the ART metabolite (7.8 min) and two gut microbiome metabolites (9.9 and 11.2 min) are presented.
Results

HPLC analysis of purities of ART and DHA

The chemical structures of ART and DHA are shown in Fig. 1A. An analytical method was developed that determined these ART and DHA simultaneously using a reverse phase column with gradient elution. To date, almost all reports on the HPLC analysis of ART and DHA employed reverse phase column with an isocratic mobile phase of acetonitrile:water (60:40) [24–26]. Using our reverse phase column with the isocratic mobile elution, the retention time for ART was 4.904 min, while for DHA were 4.202 and 4.869 min. Although the two peaks of DHAα and DHAβ can be separated well, in this condition, ART cannot be separated with DHAβ. Thus, the reported method has significant defects in simultaneous determination of ART and DHA.

To increase the resolution of separation, in some reports, buffers or acids are used in these elution systems. However, the buffer cannot be stored in the column for a long time, and a complex, time-consuming process has to be conducted each time the assay is complete. To avoid buffer, acetonitrile and water, two simple, easily prepared eluents that are safe for the column were used. In this study, a gradient elution program was developed without buffer to separate ART and DHA. As shown in Fig. 1B, using our method, the retention time (Rt) for ART is 10.566 min, while Rt for DHAα and DHAβ are 6.671 and 8.905 min. Those three peaks are baseline separated. The purity of ART and DHA are 98.7% and 98.5%, respectively. The contents of DHAα and DHAβ in DHA are 70.2% and 28.2%, respectively, and the ratio for α and β form of DHA is 2.49:1.

Gut microbiome biotransformation of ART

Based on previous reports, DHA is an in vivo metabolite of ART [7, 8]. Like many other herbal medicines, A. annua is taken orally. After oral ingestion, the main constituent artemisinin is exposed to gut microflora in the gastrointestinal tract. Whether DHA is converted by the gut microbiome or by enzymes in vivo from ART is still unknown. Since there is no enteric microbial converts report on ART, gut microbiome biotransformation tests were performed on ART.

As shown in Fig. 1C, for the control groups at 0, 12 and 24 h, since there is no peak at the retention times of ART and DHA, the background of the gut microbiome did not influence the determination of ART and DHA. When ART was incubated with gut microbiome for 12 h, compared to the chromatogram at 0 h, the peak height of ART was reduced, suggesting ART was transformed by the gut microbiome to its metabolites, and two new peaks were detected. After 24 h incubation, ART was not detected, suggesting that ART was absolutely transformed to its metabolites in 24 h, while three new peaks were detected. UV spectra of those three peaks suggested that the compound at Rt = 7.823 min is likely a metabolite of ART. Because UV spectra of other two peaks are much different to ART or DHA, those two peaks should be produced by the gut microbiome, which was supported by the chromatogram from the control group, in which a compound at Rt = 11.116 min was detected. Our data firstly proved that DHA is not a gut microbial metabolite of ART.

Antiproliferative effects of ART and DHA

To investigate the anticancer potential of ART and DHA, the antiproliferative effects were evaluated using a two-step strategy. In the first step, one time point (48 h) was investigated with a relatively wide range of the compound concentration. As shown in Fig. 2A, for the cell line HCT-116, DHA showed significant antiproliferative effect in concentrations over 10 μM, with an IC50 of 28.2 μM. Morphological observation was performed after crystal violet staining, showing that ART did not influence HCT-116 cell growth, while DHA obviously inhibited cell growth at 15 and 25 μM (Fig. 2B). Similar effects were observed in the HT-29 cell line, but the active concentration of DHA was 30 μM, which is higher than that on HCT-116 cells. A rat small intestine epithelial cell line, IEC-6, was used to evaluate if they were affected by the drugs. At tested concentrations (1–30 μM), both ART and DHA did not inhibit cell growth, indicating that ART and DHA did not affect normal intestinal cells.

Subsequently, antiproliferative potential was investigated by measuring time- and dose-dependent effects. Since DHA showed strong inhibition activities on HCT-116 cells, this cell line was employed for further studies. In the second step, cell proliferations were observed at 24, 48, and 72 h, and better selected concentration ranges were employed. The treatment concentrations for both compounds were 5–25 μM. As shown in Fig. 2C, after treatment with DHA, by doses over 10 μM, dose- and time-dependent antiproliferative effects were observed. However, ART did not inhibit cancer cell growth (Fig. 2C).

ART and DHA on cancer cell cycle

Based on MTS results, DHA showed significant anti-proliferation effects on HCT-116 cells with the doses over 10 μM. Thus, concentrations used in cell cycle tests were of 10–25 μM for 48 h. Figure 3A shows that 10–25 μM of DHA treatment changed cell cycle profiles.
At 10 μM, DHA obviously reduced S-phase proportion and increased G2/M-phase proportion. Treatment with 15–25 μM showed more significant changes. After treatment with 20 μM of DHA for 48 h, in comparison to the control (48.5% of G1, 29.5% of M, and 17.9% of G2/M), cells in G1-phase were 32.7% (P < 0.01), M-phase 25.0% (P < 0.01), and in G2/M phase 40.8% (P < 0.01). DHA treatment significantly decreased G1- and S-phase cells and increased G2/M-phase cells. However, cell cycle effects of ART in HCT-116 cells were not found (Fig. 3B).
Effects of ART and DHA on expression of cyclin A

Cell cycle progression is regulated by cyclins, particularly cyclin A for the S-phase and passage through the G2/M-phase. The observation of G2/M phase cancer cell cycle arrest through GBC led to further investigation of cell cycle regulation through evaluating the expression of cyclin A. Using HCT-116 cells, for the control group, the cell proportion for cyclin A positive cells was 14.7%. After 48 h treatment with 10, 15, 20, and 25 μM of DHA, the cyclin A positive proportions increased to 33.3%, 38.3%, 44.1%, and 51.0%, respectively (all \( P < 0.01 \)) (Fig. 3B). On the other hand, with ART treatment at same concentrations, the expression of cyclin A was not changed. Therefore, DHA treatment induced an obvious increase in cyclin A expression on HCT-116 cells.

Apoptotic induction of ART and DHA on colorectal cancer cells

To further investigate mechanisms of DHA’s cell growth inhibition, an apoptotic assay was conducted within the staining of PI and annexin V. For PI and annexin V staining, negative for both are viable cells, while positive for both are late apoptotic or necrotic cells. PI-positive and annexin V-negative are cells that underwent necrosis, while annexin V-positive and PI-negative are early apoptotic cells. As shown in Fig. 4A, the early and late apoptotic cells for the control were 5.3% and 3.4%, respectively. Treatments with DHA at 20–30 μM increased apoptotic cells significantly. For example, 30 μg/ml of DHA treatment for 48 h resulted in an increasing of early and late apoptotic cells to 14.5% and 26.3%, respectively (both \( P < 0.01 \)). However, ART treatment did not induce cancer cell apoptosis (Fig. 4A). Compared to control, after ART and DHA treatment, we did not observe obvious differences of cell cycle and apoptosis in IEC-6 normal intestinal cells.

ART and DHA on pro-inflammatory cytokine expression

Previous research has reported that the pro-inflammatory cytokine IL-8 recruited and activated neutrophils, which contribute to the gut mucosal damage. Extract from the plant *A. annua* showed anti-inflammatory activities in different acute and chronic inflammatory models. To test the anti-proliferative potentials of *A. annua*’s major constituent and its metabolite, in this study, the anti-inflammatory effects of ART and DHA on LPS-induced IL-8 secretion in HT-29 cells were investigated. As shown in Fig. 4B, both ART and DHA significantly inhibited IL-8 secretion at the concentrations of 4–10 uM (both \( P < 0.01 \)). Our data suggests that both ART and DHA have similar anti-inflammatory potential.

Effects of ART on Treg, Th1 and Th17 cell differentiation

In our previous anti-inflammatory cytokine expression investigation, both ART and DHA showed obvious inhibitory effects on IL-8 secretion. To compare the effects with DHA, the effects of ART on T cell differentiation were assayed. As shown in Fig. 5, ART did not influence Treg cell differentiation. However, ART influenced Th1 and Th17 cell differentiation. For the Th1 cells, ART inhibited the cell differentiation at concentrations of 0.5–2 μM. For Th17 cells, inhibitory effects of ART were observed at the concentrations of 1 and 2 μM. Therefore, we observed dose-dependent effects of ART on Th1 and Th17 cell differentiation.

DHA increased Treg differentiation and decreased Th17 differentiation

To investigate DHA on adaptive immunity, GBC on T cell differentiation was evaluated. Treg cells act as suppressor cells with unique immune regulation. The normality of Treg cells plays crucial role in the maintenance of the body’s immune tolerance. It has been shown that suppression of Th17-cell differentiation from naïve CD4+ T cells has beneficial effects on inflammatory disease management. To test DHA on the differentiation of Treg, Th1, and Th17, CD4+ T cells were maintained under Treg, Th1, and Th17-polarizing conditions. At the treatment concentrations of 0.5 μM, DHA did not influence Treg, Th1, and Th17 cell differentiation obviously. For Th1 cells, only 2 μM of DHA showed some effects. For the Treg cells, interestingly, 1 and 2 μM of DHA significantly increased Treg cell differentiation (both \( P < 0.01 \)). However, for the Th17 cell differentiation, approximately 30.4% of CD4+ T cells were IL-17+ in the control group. One μM of DHA treatment inhibited Th17 differentiation to 22.1% \( (P < 0.01) \). Furthermore, 2 μM of DHA treatment inhibited Th17 differentiation down to 9.5% \( (P < 0.01) \) (Fig. 5C). Concentration-dependent positive regulation in Treg and negative regulation in Th17 differentiation by DHA treatment were observed.
**Discussion**

Colorectal cancer (CRC) remains a leading cause of morbidity and mortality worldwide [27]. In spite of the continual advances in developing treatment strategies for CRC patients, dose-limiting toxicity and other severe side effects remain as limitations for chemotherapies. The consequences of the drugs can end in refusal to continue with chemotherapy. Cancer patients often try to treat its side effects through complementary and alternative medicinal means [28, 29]. Furthermore, botanical extracts enhance chemopreventive effects of chemotherapeutic agents [23]. As a tremendous amount of chemical diversity is found in millions of species of plants, botanicals, their extracts, and single compounds have been a valuable source of new therapeutic candidate compounds for CRC management.

*Artemisia annua* L. is an aromatic annual herbaceous plant. In oriental countries, this herbal medicine traditionally used for fever, inflammation, and malaria-related symptoms [1, 30]. Phytochemical studies have identified various constituents from this herb including terpenoids, steroids, coumarins, phenolics, flavonoids, purines, etc. [31]. ART is a sesquiterpene lactone isolated from the aerial parts of *A. annua*. In addition to antimalaria action, since ART is a hydrophobic compound, it is thus capable of permeating through the cellular membrane to elicit anti-cancer functions. Although ART’s anticancer activities have been reported, its effects were relatively unimpressive [16, 17].

Dihydroartemisinin (DHA) is a metabolite of artemisinin [7, 8]. To date, many artemisinin derivatives, including its metabolite DHA, have been evaluated for variable medical conditions, including inflammation and malignancies. Compared to ART, DHA has a stronger anti-malaria activity and has the advantage of better water solubility. Together with its established safety record in anti-malarial treatment, DHA possesses promising drug profiles in the crossover usage as a potential anticancer agent. However, previous studies focus more on direct cancer cell inhibition mechanisms. The impact of inflammation in cancer development, especially the link of adaptive immune response in cancer chemoprevention, were neglected. Furthermore, published reports focused on single compound pharmacology, so no bioactive comparison between ART and DHA has been performed.

To qualitative and quantitative determination of ART and its metabolites, an HPLC method was developed to determine the purity of ART and DHA. Literature reported methods for the determination of ART and DHA based on WHO criteria, in which a isocratic mobile phase is used, with the composition of a 60:40 ratio of acetonitrile:water [24] and with buffer to increase the separate resolution [25, 26]. Based on our observations, ART and DHAβ could not be separated with the isocratic mobile phase. In this study, a gradient elution program without buffer was developed, with which ART and DHA are baseline separated, and they can be determined accurately. The purities of both compounds were over 98%, suggesting the high purity of the compound used in this study.

Although DHA is a metabolite of ART, whether the metabolism is mainly after its gastrointestinal absorption or mediated via the biotransformation of gut microbiome is not clear. In this study, gut microbial biotransformation of ART was conducted. Due to the thermostability issues of ART, the temperature during sample preparation needs to be controlled strictly. Thus, 37 °C was selected to evaporate solvent. Based on previous reports, ethyl acetate was used to extract ART and DHA from the biological fluid [8]. However, because of the high boiling point of ethyl acetate, after extraction, overnight waiting was needed to remove ethyl acetate. Since ART and DHA easily dissolved in ethyl ether, ethyl ether was selected to extract ART and its metabolites. After extraction, ethyl ether could be evaporated on 37 °C water bath in only 20 min. Therefore, our method saved time, significantly increasing the efficacy of sample processing.

Up to date, there is no enteric microbiota biotransformation report on ART. Our gut microbiome transformation results showed that ART was completely converted to its metabolites in 24 h, but DHA is not an ART’s gut microbial metabolite. Although DHA was not detected in ART’s reaction solution, for the first time, our results suggested that ART can be biotransformed by gut microbiome in 24 h, and one metabolite was detected. The whole picture of identification of gut microbial metabolites in ART is an important task for the future studies.
Fig. 4  Apoptotic and anti-inflammatory analyses of HCT-116 cells treated with ART and DHA. A HCT-116 cells were treated with 15, 20, 25 and 30 μM of ART and DHA for 48 h. Apoptosis was quantified using annexin V-FITC/propidium iodide (PI) staining followed by flow cytometric analysis. Upper panel: Representative scatter plots of PI (y-axis) versus annexin V (x-axis). Lower panel: Percentage of viable, early apoptotic and late apoptotic cells. B Effects of ART and DHA on inflammatory cytokine IL-8 secretion in HT-29 cells. The basal level of IL-8 secretion from HT-29 cells was very low (<20 pg/ml). LPS-induced IL-8 secretion was inhibited by both compounds. Data are presented as the mean ± standard error of triplicate experiments. *P < 0.05, **P < 0.01 vs. control.
Using MTS assay, antiproliferative effects of ART and DHA were evaluated. DHA showed stronger anti-CRC cell activities than ART. The two cell lines used in this study varied in tumor suppressor gene p53 expression. HCT-116 is a p53 wild type, while HT-29 cells contain a p53 mutation. The p53 antigen is overproduced in the cell line HT-29, and there is a G → A mutation in codon 273 of the p53 gene resulting in an Arg → His substitution [32]. Cancer cells with p53 mutations are resistant to many chemotherapeutic agents, including cisplatin, alkylating agents (temozolomide), anthracyclines (doxorubicin), antimetabolites (gemcitabine), antiestrogens (tamoxifen) and EGFR-inhibitors (cetuximab) [33]. It is well known that p53 suppresses tumor formation and renders protection against DNA damage. The way in which p53 influences drug resistance depends on several different parameters including the mode of action of the drug, genetic alterations during carcinogenesis, and the type of cancer [33, 34]. We observed that DHA had greater antiproliferative abilities in the p53 wild type cell line (HCT-116) than that in the p53 mutation cell line (HT-29). Further mechanistic studies should be performed to explore the details of p53 mutation mediated DHA’s antiproliferation resistance. Nevertheless, data obtained from this study suggested that p53 plays a role in DHA’s anticancer activities.

The analysis on cell cycle distribution demonstrated that DHA target a cell cycle checkpoint for G2/M arrest. Cyclin A plays an important role in controlling the G2/M transition of the cell cycle by binding to CDK2 [35, 36]. Our results showed that DHA up-regulated cyclin A, indicating that DHA could facilitate the cell’s progression through the S phase into the G2/M phase and lead to the accumulation of cell distribution in the G2/M phase. Apoptosis is caspase-dependent cell death, which is considered an important pathway in the inhibition of cancer cells by many anticancer agents [37]. Caspases are a family of cysteine proteases that play essential roles in cell apoptosis [38]. DHA induced cancer cell apoptosis, suggesting that antiproliferation of DHA on human colorectal cancer cells is mediated in part by the induction of apoptosis (Fig. 6).

Since IBD and chronic inflammation is recognized as a risk factor for CRC development [39], an effective prevention method of malignant colon tumor formation and progression is targeting inflammatory pathways [40]. The adaptive immune response has been considered to play a critical role in the pathogenesis of IBD. T lymphocytes represent the key cell population of the adaptive immunity arm [41]. The original T cells amplify and are differentiated into different subsets such as Th1, Th17, and Treg cells after stimulation of antigens. Th1 cells eliminate pathogens inside cells, Th17 cells remove extracellular bacteria and fungi, and Treg cells promote tissue repairs. However, disorders of T cell responses and imbalance of T cell subsets induce excessive releases of cytokines and chemokines, which lead to inflammation [41]. To explore ART and DHA on adaptive immune responses, the effects of those two compounds on the differentiation of Th1, Th17, and Treg cells have been investigated.

Based on our results, although ART did not change Treg cell differentiation, it significantly inhibited Th1 and Th17 cell differentiation. On the other hand, DHA treatment moderately down-regulated Th1 cell differentiation. Interestingly, DHA significantly increased Treg differentiation and decreased Th17 differentiation in a dose-dependent manner. Compared to ART, DHA showed more potent inhibitory effects on Th17 than ART did. Our data suggested that DHA promoted Treg differentiation and suppressed differentiation of Th17 cells, which are two important Th subset connects to gut inflammation.

Activated Th17 cells mainly function through secreting cytokines, such as IL-17, IL-21, and IL-22. IL-17 induces immune cells transferring into peripheral tissues and binding with IL-17 receptors, then activates NF-κB and promotes the release of a variety of pro-inflammatory factors. Treg cells inhibit other Th cells such as Th1 and Th17 via direct contacts with cells and releases of cytokines such as IL-10 and TGF-β to maintain immune tolerances. Th17 cells and Treg cells are in dynamic equilibrium under normal circumstances. The balance is broken with the over-increase of Th17 cells, over-enhancement of immunogenicity, and decrease or abnormal function of Treg cells, causing intestinal mucosal damages. Based on our data, DHA increased Treg differentiation and inhibited Th17 differentiation (Fig. 6). The ratio of Th17/Treg in IBD patients’ blood is significantly higher, suggesting that transformation imbalance of Th17/Treg plays an important role on pathogenesis of IBD. Therefore, regulation of balance between Th17/Treg has become a new approach for IBD management [42]. Our data suggested that DHA may have the potential to regulate the balance of Th17/Treg in order to inhibit gut inflammation and to prevent colon cancer initiation and development.
Conclusion

In this study, a HPLC method for ART and DHA identification was developed, and accompanying this method with ethyl ether extraction significantly increased the efficacy of sample processing. Gut microbiome biotransformation of ART was performed, and, for the first time, ART was observed to be completely converted to its metabolites within 24 h, but no DHA was detected as a gut microbial metabolite of ART. Using human colorectal cancer cells, ART did not influence cancer cell growth in the tested concentrations, but DHA significantly inhibited cancer cell growth. The DHA induced G2/M cell cycle arrest via upregulation of cyclin A and apoptosis. As a metabolite of ART, DHA also downregulated the expressions of pro-inflammatory cytokine IL-8. Furthermore, DHA promoted Treg cell proliferation, while both ART and DHA inhibited Th1 and Th17 cell differentiation. Our results suggest that DHA is an active anticancer metabolite of ART. While more DHA research should be conducted on cancer chemoprevention, derivatives including novel DHA metal complexes should be considered via chemical synthesis, since these complexes have been proven to be effective in cancer therapeutics.

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Authors contributions Project administration: CZW, CSY. Participated in research design: CFZ, QHZ, LC, ZL, LH. Conducted experiment: CZW, CW, YL, DHW. Wrote an original draft preparation: CZW, ML, CSY. Performed data analysis: CW, YL. Edited and reviewed: CZW, CHL, TLJ, CSY.

Data availability The analyzed data sets generated during the study are available from the corresponding author upon reasonable request.

Declarations

Conflict of interest The authors declare that there are no conflict of interest.

Ethical approval All procedures for experimental protocols of the present study involving animals and cells were performed in accordance with the ethical standards of the institution of practice at which the studies were conducted. The need for ethics approval for human fecal-related study was waived by the local medical ethics committee and is deemed unnecessary according to local guidelines. It did not pertain to any (clinical/biological) investigation of any human tissue at any level. Verbal and written consent was obtained from all participants. Ethics approval was not required by the local medical ethics committee, as it was deemed unnecessary according to the committee’s requirements.

Consent to participate All authors have agreed to participate in the manuscript.

Consent for publication All authors have agreed to publish the manuscript.

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