

Contents lists available at ScienceDirect

Biochimica et Biophysica Acta

journal homepage: www.elsevier.com/locate/bbadis

Green tea phenolics inhibit butyrate-induced differentiation of colon cancer cells by interacting with monocarboxylate transporter 1

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ARTICLE INFO

Article history:

Received 16 May 2013

Received in revised form 24 July 2013

Accepted 20 August 2013

Available online 28 August 2013

Keywords:

Epicatechin

Epigallocatechin gallate

Butyrate

Differentiation

Colon cancer

ABSTRACT

Diet has a significant impact on colorectal cancer and both dietary fiber and plant-derived compounds have been independently shown to be inversely related to colon cancer risk. Butyrate (NaB), one of the principal products of dietary fiber fermentation, induces differentiation of colon cancer cell lines by inhibiting histone deacetylases (HDACs). On the other hand, (–)-epicatechin (EC) and (–)-epigallocatechin gallate (EGCG), two abundant phenolic compounds of green tea, have been shown to exhibit antitumoral properties. In this study we used colon cancer cell lines to study the cellular and molecular events that take place during co-treatment with NaB, EC and EGCG. We found that (i) polyphenols EC and EGCG fail to induce differentiation of colon adenocarcinoma cell lines; (ii) polyphenols EC and EGCG reduce NaB-induced differentiation; (iii) the effect of the polyphenols is specific for NaB, since differentiation induced by other agents, such as trichostatin A (TSA), was unaltered upon EC and EGCG treatment, and (iv) is independent of the HDAC inhibitory activity of NaB. Also, (v) polyphenols partially reduce cellular NaB; and (vi) on a molecular level, reduction of cellular NaB uptake by polyphenols is achieved by impairing the capacity of NaB to relocate its own transporter (monocarboxylate transporter 1, MCT1) in the plasma membrane. Our findings suggest that beneficial effects of NaB on colorectal cancer may be reduced by green tea phenolic supplementation. This valuable information should be of assistance in choosing a rational design for more effective diet-driven therapeutic interventions in the prevention or treatment of colorectal cancer.

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1. Introduction

Colorectal cancer constitutes one of the most frequent malignancies worldwide and is one of the prevalent causes of cancer-related mortality in the western world [1]. Therefore, further development of therapeutic and preventive approaches to control this disease is clearly needed. A diet rich in fiber and plant-derived compounds present in tea, fruits and vegetables has been inversely associated with the risk of colorectal cancer [2,3]. Furthermore, the protective effect of the bioactive compounds present in these dietary constituents has been shown to be related to the activity of human intestinal microbiota. In this

regard, consumption of natural phenolics has been shown to increase microbial fermentation products such as butyrate (NaB – sodium butyrate) from dietary fiber [4–6], thereby providing a beneficial effect to the host. However, the mechanism by which NaB and phenolics interact at the cellular level has not been satisfactorily addressed.

NaB has been described as a potent antitumoral agent against colon cancer, and has been used in clinical trials for treating cancers [7]. NaB is a four-carbon fatty acid that represents a major oxidative fuel for colonic epithelial cells [8]. Previous studies have demonstrated that deficiency in the availability or utilization of NaB causes colitis and may be involved in ulcerative colitis and colon carcinogenesis [9,10]. Moreover, NaB induces apoptosis and cell cycle arrest in the G1/G0 phase, accompanied by terminal cell differentiation in several colon cancer cell lines [11–13]. The mechanism by which NaB induces differentiation primarily involves epigenetic regulation of gene expression through the inhibition of histone deacetylases (HDACs) [14], which remove acetyl groups from lysine residues of histones and decrease the affinity of protein transcription complexes for DNA.

Numerous studies have evaluated the antitumor activities of green tea phenolics in different experimental systems. The general consensus is that these tea components promote cancer cell growth inhibition and

Abbreviations: AP, alkaline phosphatase; NaB, butyrate; EC, (–)-epicatechin; EGCG, (–)-epigallocatechin gallate; HDAC, histone deacetylase; MCT1, monocarboxylate transporter 1; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; TSA, trichostatin A

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apoptosis, and reduce invasion, angiogenesis and metastasis [15]. A plethora of molecular mechanisms of tea phenolics has been suggested, including anti-oxidant and pro-oxidant effects, inhibition of mitogen-activated protein kinases, or modulation of growth factor receptor tyrosine kinases and the activity of transporters through alteration of lipid rafts by tea catechins (reviewed in Ref. [16]).

The effect of green tea catechins on differentiation has been addressed in several cell lines [17–19], including colon cancer cells [20], in which their effects appear to be cell line-specific. Moreover, a recent study has investigated the impact of green tea phenolics on NaB-induced colon cancer cells differentiation and concluded that changes in NaB uptake in Caco2 cells induced by phenolics do not correlate with changes upon cell differentiation [21]. In the present study we evaluated the effect of the major green tea phenolics (–)-epigallocatechin gallate (EGCG) and (–)-epicatechin (EC) on NaB-induced differentiation in human colon adenocarcinoma HT29 cells. We demonstrate that phenolics interfere with NaB induced differentiation, and we propose a mechanism for this inhibition based on the altered localization of a monocarboxylate transporter in plasma membrane lipid rafts.

2. Materials and methods

2.1. Chemicals and cell culture conditions

All chemicals were purchased from Sigma-Aldrich Co. (St. Louis, MO), unless otherwise specified. HT29 human colon adenocarcinoma cells (obtained from the American Type Culture Collection) were grown in Dulbecco's modified Eagle medium 25 mM D-glucose supplemented with 10% heat-inactivated fetal calf serum (PAA Laboratories, Pasching, Austria) and 0.1% antibiotics (100 U/mL penicillin and 100 µg/mL streptomycin) (Invitrogen, Paisley, UK). Caco2 cells were maintained in Dulbecco's modified Eagle medium 25 mM D-glucose, 20% fetal calf serum, 2 mM glutamine, and 1% antibiotics (100 U/mL penicillin and 100 µg/mL streptomycin). NCM460 (INCELL Corporation, San Antonio, Texas, USA) is an epithelial cell line derived from the healthy colon mucosa of a 68-year-old Hispanic male [22]. These cells were grown as a monolayer culture in M3Base medium (which contains growth supplements and antibiotics) supplemented with 10% heat-inactivated FCS and 2.5 mM of D-glucose. Cells were cultured at 37 °C in a humidified atmosphere with 5% CO₂.

2.2. Determination of cell viability

Assays were performed using a variation of the MTT assay described by Mosmann [23]. HT29 cells were seeded at 3×10^3 cells/well in 96-well flat-bottom plates. After 24 h of incubation at 37 °C, fresh media containing (–)-epicatechin (EC) and (–)-epigallocatechin gallate (EGCG) at different concentrations were added. After 72 h, the media was removed, and 50 µL of MTT (1 mg/mL in phosphate buffered saline) with 50 µL of fresh medium was added to each well and incubated for 1 h. The MTT reduced to blue formazan and the precipitate was dissolved in 100 µL of dimethyl sulfoxide. Absorbance values were measured on an ELISA plate reader (550 nm) (Tecan Sunrise MR20-301, Tecan, Salzburg, Austria).

2.3. Alkaline phosphatase (AP) activity assay

AP activity was measured using p-nitrophenyl phosphate as a substrate according to the published procedures [24]. HT29 cell cultures were started with 6×10^5 in 60-cm² petri dishes and incubated for 24 h at 37 °C. New medium containing phenolics, butyrate (NaB) and NaB + phenolics was added and incubated for 24, 48 and/or 72 h at 37 °C. The medium was changed every 24 h. After incubation, the cells were washed with phosphate buffered saline (PBS), detached from the flasks using 0.025% trypsin-EDTA (Invitrogen) and then resuspended in lysis buffer (1 mM dithiothreitol, 1 mM EDTA, 0.02% Triton X-100,

0.02% sodium deoxycholate, 0.2 mM phenylmethylsulfonyl fluoride, 1% sodium azide and 20 mM Tris-HCl, pH 7.5). Cells were homogenized using a laboratory sonicator (1/2 Liter Branson 200 Ultrasonic bath, 5 min, 40 kHz, 4 °C) and immediately ultracentrifuged at 105,000 ×g for 1 h at 4 °C. The supernatant was separated and used for the determination of AP activity using a Cobas Mira Plus chemistry analyzer (HORIBA ABX, Montpellier, France). The enzyme activity was estimated by measuring the absorbance at 405 nm due to formation of p-nitrophenol and was expressed as mU/mg of protein. Protein determination was performed in the same lysates using the BCA protein assay (Pierce Biotechnology, Rockford, IL).

2.4. Histone deacetylase (HDAC) assay

HT29 cells were incubated in 60-cm² petri dishes for 48–72 h at 37 °C (65–85% confluence). Next, cells were washed in PBS pH 7.4 followed by incubation in hypotonic buffer (20 mM HEPES pH 7.6, 20% glycerol, 10 mM NaCl, 1.5 mM MgCl₂, 0.2 mM EDTA, 0.1% Triton X-100) for 5 min. Then, cells were collected and nuclei pelleted at 1000 rpm in the microfuge for 10 min. Purified nuclei were resuspended in hypertonic buffer (20 mM HEPES pH 7.6, 20% glycerol, 450 mM NaCl, 1.5 mM MgCl₂, 0.2 mM EDTA, 0.1% Triton X-100) and gently shaken for 1 h at 4 °C. After centrifuging at 13,000 rpm in the microfuge for 5 min at 4 °C, the supernatant obtained was the nuclear extract. Then, nuclear extracts of non-treated HT29 cells were quantified using a standard BCA Protein Assay (Pierce Biotechnology, Rockford, IL) and an equivalent quantity of protein was subjected to treatment with NaB and NaB/phenolics for 30 min at 37 °C. HDAC activity was measured employing a Fluorometric Assay Kit (Biovision), following manufacturer's instructions. The procedure involves the use of the HDAC substrate, which consists of an acetylated lysine side chain, and incubation with a sample containing nuclear extract. Deacetylation sensitizes the substrate, and treatment with the lysine developer produces a fluorophore, which can be analyzed with a fluorometer (Ex/Em = 350 – 380/440 – 460 nm). A HeLa cell nuclear extract was used as a positive control. Percent inhibition of treated cells was compared with HT29 untreated controls.

2.5. [¹⁴C]-NaB uptake

HT29 cells were seeded at 2×10^4 cells/well in 24-well plates. After 24 h of incubation at 37 °C, fresh media containing NaB and NaB + phenolics was added and incubated for 48 h at 37 °C. The medium was changed after 24 h of incubation and left 24 h more. Next, cells were incubated at room temperature for 20 min in tracer-free buffer containing (in mM): 110 NaCl, 1 CaCl₂, 4 KCl, 0.44 K₂HPO₄, 1 MgSO₄, 5 glucose, 50 mannitol and 5 HEPES, pH 7.4. Cells were then washed and incubated with buffer containing (in mM): 259 mannitol, 20 HEPES, pH 6.5, 2 [¹⁴C]-NaB (1 µCi/mL) for 5 min. The uptake was stopped by washing the cells twice with ice-cold PBS. Finally, cells were solubilized with 0.5 N NaOH for at least 4 h. The protein concentration was measured by the method of BCA. Incorporated radioactivity was counted by a Tri-CARB 1600-TR liquid scintillation counter (Packard Instruments, Downers Grove, IL). The values were expressed as nmol/mg protein/5 min.

2.6. Cell lysates and Western blotting

6×10^5 HT29 cells were plated on 60-cm² petri dishes and incubated for 24 h. Then, fresh medium with NaB and NaB + phenolics was added and incubated for 48 h. After incubation, the cells were washed with ice-cold PBS and lysed in 20 mM Tris-HCl, pH 7.5, 150 mM NaCl, 1% Triton X-100, 1 mM EDTA, 1 mM EGTA, and 1× complete protease inhibitor cocktail. The lysate was sonicated and centrifuged at 5,000 ×g for 5 min at 4 °C, and protein concentration was determined by Bradford. The samples obtained above were subjected to 10% SDS-PAGE and

transferred to nitrocellulose membranes. MCT1 expression was detected utilizing human anti MCT1 antibody (Santa Cruz Biotechnology, sc-50324). Flotillin expression was analyzed using human anti flotillin antibody (BD Transduction Laboratories, 610820). β -Actin was used as a loading control (MP Biomedicals, Eschwege, Germany, 69100).

2.7. Rafts isolation and biochemical characterization

Lipid rafts were isolated by floatation on OptiPrep density gradient as previously described [25]. HT29 human colon adenocarcinoma cells were started in 60-cm² petri dishes with the same number of cells (6×10^5) and incubated for 24 h at 37 °C. Then, cells were exposed to or not exposed to NaB or NaB + phenolics for 48 h. After incubation, cells were resuspended and incubated for 30 min at 4 °C in TNE buffer containing (in mM) 25 Tris (pH 7.4), 150 NaCl, 5 EDTA, and 1% Triton X-100 supplemented with $1 \times$ complete protease inhibitor cocktail. The membranes were then adjusted to 40% final concentration of OptiPrep and layered at the bottom of density gradient with steps of final concentrations of 35, 30, 25, and 20% of OptiPrep in TNE buffer. TNE buffer was laid on the top of the gradient, which was then centrifuged at 48,000 rpm for 4 h at 4 °C. Fractions of 1 mL were collected from the top to the bottom of the gradient and then analyzed by Western blotting (see above). MCT1 and the described marker for lipid rafts, flotillin, were analyzed in each fraction [26].

2.8. Data presentation and statistical analysis

Data are given as the means \pm S.D. (standard deviation). For each assay, the parametric unpaired two-tailed independent sample *t*-test was used and differences were considered to be significant when $p < 0.05$ or $p < 0.001$.

3. Results

3.1. Viability of HT29 cells treated with (–)-epicatechin (EC) and (–)-epigallocatechin gallate (EGCG)

To determine a non-toxic but still active concentration of EC and EGCG, HT29 cell viability was determined in the presence of different phenolic concentrations (Fig. 1). From the obtained dose–viability curve we estimated the inhibitory concentration 20 (IC20), defined as the concentration of product that causes 20% inhibition of cell viability with respect to viability of control (non-treated) cells after 72 h. Although increasing concentrations of both EC and EGCG produced a dose-dependent decrease in cell viability, EGCG was much more efficient at doing so. The 72 h IC20 values obtained were 100 μ M for EC and 20 μ M for EGCG, which were the concentrations used throughout the study.

3.2. Phenolics reduce butyrate (NaB)-induced differentiation in colorectal adenocarcinoma cell lines

Firstly, we studied whether EC and EGCG affected NaB-induced differentiation. Cells were exposed to 2 mM NaB for 24, 48 and 72 h, alone or in the presence of 100 μ M EC or 20 μ M EGCG. NaB-induced differentiation measured as an increase in alkaline phosphatase (AP) activity, was reduced by both phenolics at 48 and 72 h of combined treatment (Fig. 2A). EC also reduced NaB-induced AP activity at 24 h (Fig. 2A). It should be noted that treatments with phenolics alone had no impact on differentiation (Fig. 2B). To rule out a direct effect of the phenolics on AP activity, we repeated the experiments measuring the activity of another differentiation marker (aminopeptidase N) at 48 h. Consistently, phenolics reduced NaB-induced differentiation (Supplemental Fig. 1A) and did not show an increase in differentiation when used alone (Supplemental Fig. 1B). This effect of phenolics was also extended to another epithelial colorectal adenocarcinoma cell line, Caco2,

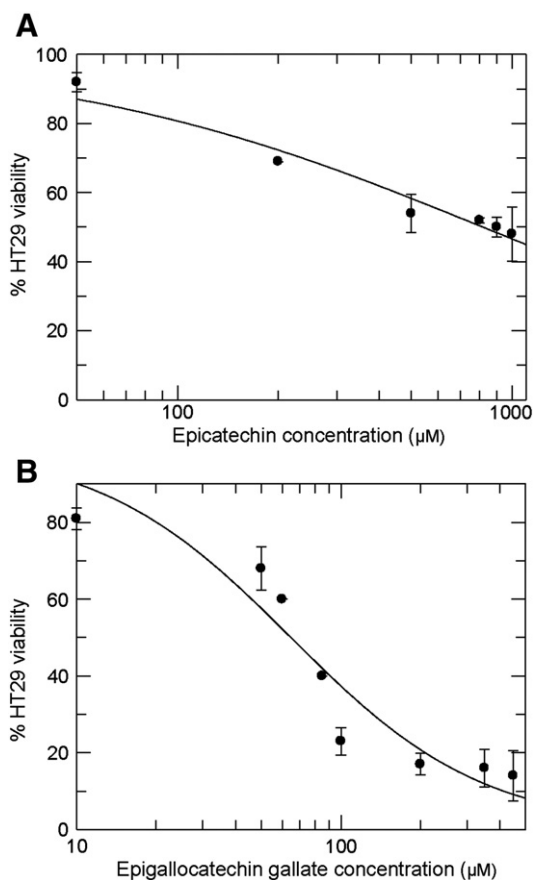


Fig. 1. Dose–effect curves of (–)-epicatechin and (–)-epigallocatechin gallate on cell viability. HT29 cell cultures were treated with increasing doses of (–)-epicatechin (EC) (A) or (–)-epigallocatechin gallate (EGCG) (B) as indicated on the x axis for 72 h. Cell viability was expressed as a percentage with respect to untreated control. IC20 (EC) = 100 μ M and IC20 (EGCG) = 20 μ M.

which showed the same differentiation profile with respect to NaB and phenolics treatment (Supplemental Fig. 2A and B).

To verify whether our observations were cancer cell line-specific, we determined NCM460 cell differentiation, measured as AP activity, after incubation with 2 mM NaB, alone or in the presence of 100 μ M EC or 20 μ M EGCG, for 24, 48 and 72 h. Supplemental Fig. 3 shows that cell differentiation was not significantly altered when NCM460 cells were treated with NaB and phenolics.

3.3. The effect of tea phenolics on differentiation is not related to histone deacetylase (HDAC) activity modulation

Given that the NaB-induced differentiation is related to its inhibition of HDAC activity [27], we decided to study the effects of EC and EGCG in HDAC-related differentiation. Firstly, we studied whether phenolics modified differentiation induced by trichostatin A (TSA), another well-described HDAC inhibitor. HT29 cells were exposed to 180 nM TSA for 48 h, alone or in presence of 100 μ M EC and 20 μ M EGCG. The addition of phenolics to TSA had no impact on TSA-induced differentiation, measured as AP activity (Fig. 3A).

To confirm these results, we wondered whether phenolic compounds could directly alter the HDAC activity of HT29. We determined the HDAC activity of nuclear extracts after incubation with NaB and phenolics alone or in combination. As expected, NaB was found to be a potent HDAC inhibitor, significantly decreasing HDAC activity by 55% and 67% at 500 μ M and 2 mM of NaB, respectively (Fig. 3B). Phenolics had no effect with respect to HDAC activity, both when incubated alone (Fig. 3B) and incubated simultaneously with NaB (Fig. 3C). These results

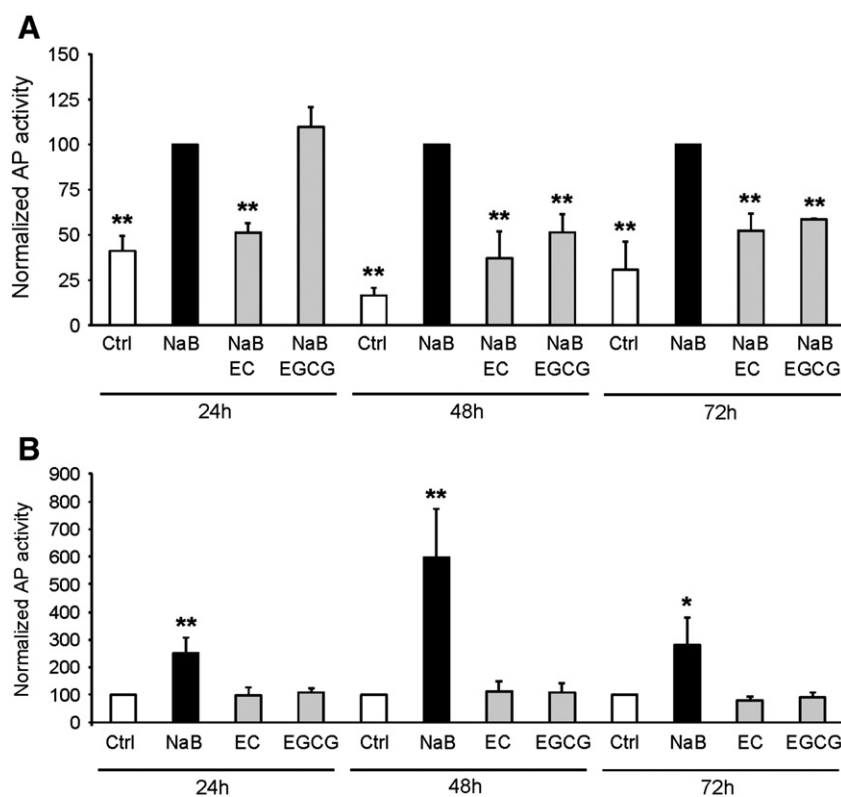


Fig. 2. Phenolics reduce butyrate-induced differentiation. (A) HT29 cells were treated with butyrate (NaB) 2 mM or with NaB and phenolics (EC 100 μ M and EGCG 20 μ M) for 24, 48 and 72 h and alkaline phosphatase (AP) activity was measured and normalized by protein level. The data are normalized and statistically tested with respect to NaB-treated cells. Mean \pm SD; $n \geq 3$; **Different from NaB, $p < 0.01$. (B) HT29 cells were treated with NaB 2 mM or with phenolics alone (EC 100 μ M and EGCG 20 μ M) for 24, 48 and 72 h and AP activity was measured. The data are normalized and statistically tested with respect to Ctrl cells. Values are mean \pm SD; $n \geq 3$; /*Different from Ctrl, $p < 0.05/p < 0.01$.

led us to conclude that phenolics do not affect NaB differentiation by directly modulating HDAC activity.

3.4. Phenolics impair NaB entry to the cell

To study the mechanism of interference between NaB and phenolics we studied the cellular entry of NaB into HT29 cells. HT29 cells were incubated with 2 mM NaB alone or in the presence of 100 μ M EC and 20 μ M EGCG for 48 h and acute [14 C]-NaB incorporation was measured. We observed that NaB treatment enhanced its own transport (Fig. 4). Moreover, we detected a significant decrease in [14 C]-NaB cellular entry after NaB and phenolics incubation relative to NaB-treated cells (Fig. 4).

3.5. The effect of tea phenolics on NaB-induced differentiation is not related to monocarboxylate transporter 1 (MCT1) expression, but with its membrane localization

Next, we evaluated the role of the intestinal transporter MCT1, which is known to be involved in NaB transport [28]. We examined MCT1 protein expression in response to NaB and phenolics. Western blot analysis of MCT1 showed no differences at protein level in any of the treatments (Fig. 5A).

Since phenolics have been shown to be lipid raft regulators [29], we next investigated whether EC and EGCG caused any alterations in the association of MCT1 with lipid rafts. Lipid raft microdomains were isolated by OptiPrep gradient ultracentrifugation and proteins in the gradient fractions were analyzed by SDS-PAGE and Western blot. Flotillin rich fractions account for the lipid rafts. As shown in Fig. 5B, MCT1 was predominantly expressed in these high-density non-raft fractions of control HT29 cells, even though some MCT1 expression was found in lipid rafts-corresponding fractions. NaB treatment for 48 h enhanced the presence

of MCT1 in low-density fractions representing lipid rafts. When phenolics were added to NaB, MCT1 was redistributed in all fractions, counteracting NaB-mediated enhanced localization of the transporter in the lipid rafts (Fig. 5B).

4. Discussion

In this study we analyzed the effect of phenolics (–)-epicatechin (EC) and (–)-epigallocatechin gallate (EGCG) on butyrate (NaB)-induced differentiation. We decided to use the IC₂₀ concentrations at 72 h (EC: 100 μ M, EGCG: 20 μ M) to avoid massive cell damage by phenolics. We used a NaB concentration of 2 mM, which has been demonstrated to induce differentiation in HT29 colon cancer cells [9]. Although treatment with EC and EGCG alone did not change cell differentiation, NaB-induced differentiation was reduced by both phenolics in colon cancer cells (Fig. 2, Supplemental Fig. 1, Supplemental Fig. 2). In contrast, previous studies using colon cancer Caco2 cells showed that EGCG was not able to revert the differentiation induced by NaB [21]. This discrepancy could be explained because a larger concentration of NaB (5 mM) and lower concentrations of EGCG (0.1–10 μ M) were used. Worthy of note, since NaB effects were not exerted on the non-tumoral NCM460 cells (Supplemental Fig. 3), the observed effects are cancer cell specific.

The mechanism of action of NaB in colon cancer mainly includes effects on differentiation via its inhibition of histone deacetylases (HDACs). EGCG has also been identified as an inhibitor of HDAC activity in prostate, skin and breast cancer cells [30]. However, studies in HT29 cells have found no significant change in the HDAC activity of cytoplasmic or nuclear fractions after sulforaphane and EGCG treatment [31]. Similarly, we also failed to detect any significant inhibition of HDAC activity in vitro by EC and EGCG (Fig. 3B), and differentiation was not impaired in response to treatment with trichostatin A (TSA) (Fig. 3A),

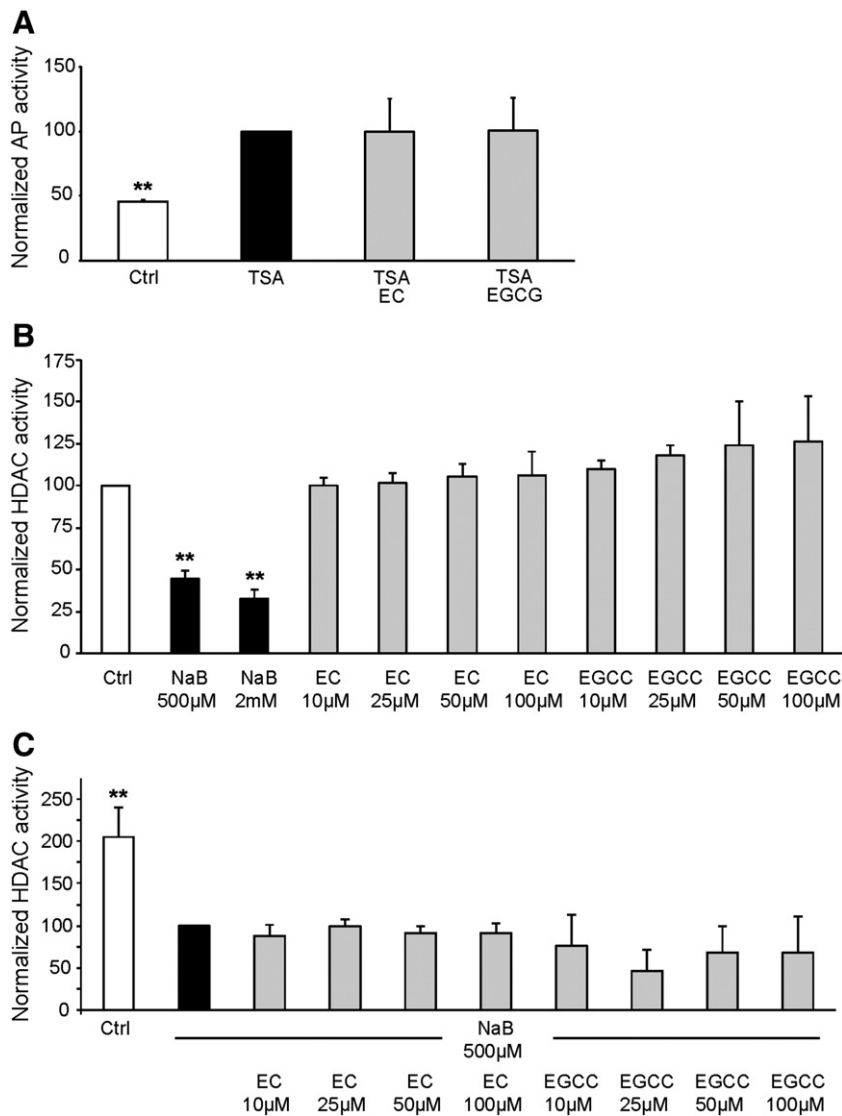


Fig. 3. Phenolics effect on differentiation is not histone deacetylase-related. (A) HT29 cells were treated with TSA at 180 nM or with TSA in the presence of phenolics (EC 100 µM and EGCC 20 µM) for 48 h and AP activity was measured. Data are normalized and statistically tested with respect to TSA-treated cells. Mean \pm SD; $n = 3$; **Different from TSA, $p < 0.01$. (B) Histone deacetylase (HDAC) activity determined in nuclear extracts from HT29 cells after treatment with NaB or NaB with phenolics for 30 min. The data are normalized and statistically tested with respect to Ctrl cells. Values are mean \pm SD; $n \geq 2$; **Different from Ctrl, $p < 0.01$. (C) HDAC activity determined in nuclear extracts from HT29 cells after treatment with NaB or phenolics for 30 min. The data are normalized and statistically tested with respect to NaB cells. Mean \pm SD; $n \geq 2$; **Different from NaB, $p < 0.01$.

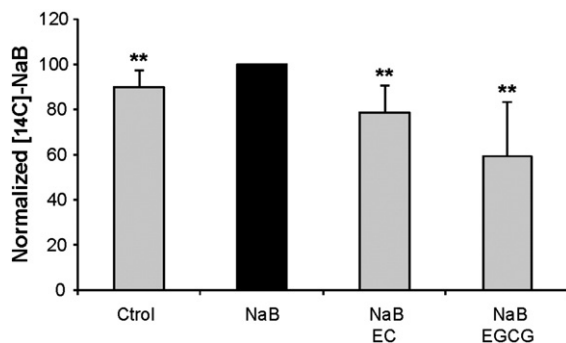


Fig. 4. Phenolics impair NaB influx into the cells. HT29 cells were treated with NaB 2 mM or with NaB and phenolics (EC 100 µM or EGCC 20 µM) for 48 h. Acute [¹⁴C]-NaB uptake was subsequently measured as described in Materials and methods. Results are normalized and statistically tested with respect to NaB cells. Values are mean \pm SD; $n \geq 2$; **Different from NaB, $p < 0.01$.

another well-known HDAC inhibitor also proposed as anti-tumor agent [32].

The effect of phenolics on NaB-induced differentiation could be due to an interaction between NaB and phenolics that prevents the entry and cellular action of NaB. Determination of [¹⁴C]-NaB uptake showed that although NaB treatment stimulated its own transport, phenolic treatment impaired NaB uptake (Fig. 4). It has been previously reported that phenolics differentially affect NaB uptake depending on the studied compound, the time of incubation and the NaB concentration used. Nevertheless, on the contrary to our observations, the differences in NaB uptake induced by phenolics did not correlate with changes in the anticarcinogenic activity including effects on cell differentiation [21].

The impairment of NaB uptake by phenolics led us to study NaB intestinal transporters. NaB enters into colonocytes by two major carrier-mediated mechanisms, the Na⁺-coupled monocarboxylate transporter 1 (SMCT1) and the aforementioned H⁺-coupled monocarboxylate

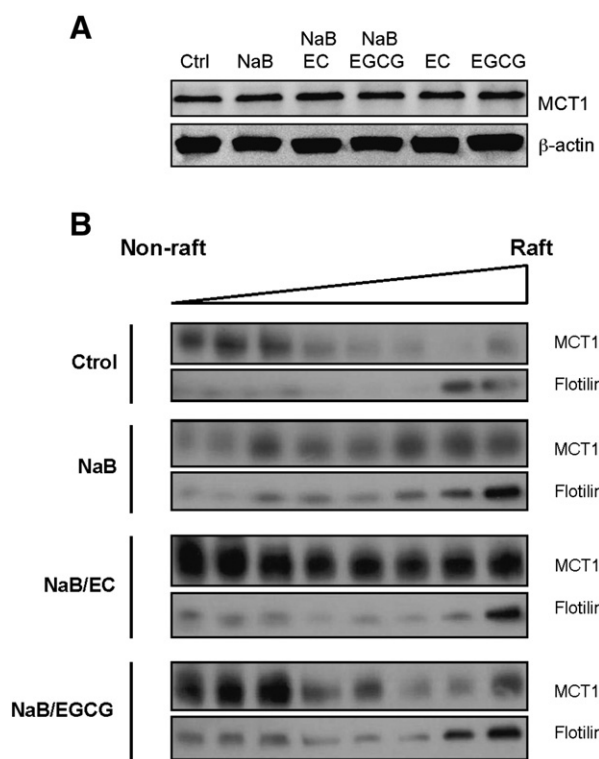


Fig. 5. Effect of NaB and phenolics on monocarboxylate transporter 1 (MCT1). (A) After 48 h of incubation with NaB, phenolics or both, HT29 lysates were probed against MCT1 in a Western blot, using β -actin as a loading control. A representative blot is shown. (B) EC and EGCG antagonize plasma membrane redistribution of MCT1 caused by NaB. HT29 cells were incubated with NaB or NaB/phenolics for 48 h and then lysed and laid at the bottom of an OptiPrep density gradient. After ultracentrifugation, fractions were collected from the top (lipid rafts low-density fractions) to the bottom of the gradient (high-density non-raft fractions). Proteins in the fractions were separated on 10% SDS-PAGE and blots were probed with anti-MCT1 or anti-flotillin antibodies.

transporter 1 (MCT1). Since their substrate, NaB, prevents and inhibits colon carcinogenesis, both transporters have been proposed to function as tumor suppressors [33]. Accordingly, MCT1 expression has been shown to be down-regulated during transition from normalcy to malignancy in colonic tissues. Regarding SMCT1 expression, it has been found to be silenced in colorectal carcinoma and colon cancer cell lines such as HT29 and Caco2 [34]. Moreover, given that some phenolics have been described to interplay with MCT1 [35–38], the next step was the evaluation of MCT1 expression. A substrate-induced MCT1 activity by NaB at higher concentrations has been previously demonstrated in AA/C1 human colonic epithelial cells [39] and colon cancer Caco2 cells [40]. However, no change in MCT1 protein levels was induced by NaB, phenolics, or both (Fig. 5A).

Since we did not detect an enhanced MCT1 protein, we investigated the possibility of another mechanism to explain the effects of phenolics on NaB-induced differentiation. This mechanism may imply alterations in transduction pathways or cellular membrane characteristics that affect the intrinsic activity of the transporter [21]. Recent reports have indicated that optimal function of many transporters is dependent on their association with lipid rafts [41]. Lipid rafts are defined as microdomains within the lipid bilayer of cellular membranes that assemble subsets of transmembrane or glycosylphosphatidylinositol-anchored proteins and lipids (cholesterol and sphingolipids) and experimentally resist extraction in cold detergent. Interestingly, lipid rafts have been related to some of the biological effects induced by tea phenolics [42]. EGCG has been shown to prevent activation of c-Met receptor [43] and epidermal growth factor receptor (EGFR) [44] via perturbations of the membrane lipid rafts. Similarly, our analysis of lipid raft-dependent MCT1 function in HT29 cells suggests that NaB activates

MCT1 functions at least in part by enhancing its distribution in lipid rafts, and that tea phenolics produce a redistribution of MCT1 in the non-lipid raft fractions (Fig. 5B). These observations led us to hypothesize that EGCG and EC might inhibit MCT1-mediated NaB transport by altering lipid raft organization. At the same time, this provides an explanation for the observed increase in NaB uptake after NaB treatment (Fig. 4), which could not be explained by changes in the quantity of MCT1 (Fig. 5A), but due to the modulation of lipid rafts by 2 mM NaB (Fig. 5B).

It has been shown that the chemical structure of phenolics is related to their biological activity [45]. In green tea catechins, the most bioactive catechin has been described to be EGCG, which contains a trihydroxyl structure in the D ring (gallate) as well as a pyrogallol B-ring, followed closely by ECG with a gallate group, and then to a lesser extent EGC and EC that possess a basic structure [46]. We have observed that although the global trend of EC and EGCG treatments is similar, small changes regarding the efficiency of both compounds can be distinguished. Apart from differences in the chemical structure, differences in the modulation of signal transduction pathways that affect the activity of the transporter or the process of differentiation may also account for the differences observed throughout the study.

As previously mentioned, several studies (4–6) have described an increase in cecal concentration of short chain fatty acids in response to phenolics. However, the mechanism by which phenolics induce this increase is not known. First of all, we hypothesize that modulation of intestinal microbiota by phenolics to increase production of fermentation products, such as NaB, might compensate the inhibition of cellular NaB uptake described in this study. On the other hand, we propose that the increase in intestinal NaB concentration found in previous studies may be, at least in part, a consequence of the inhibitory action of phenolics on NaB intestinal uptake.

5. Conclusions

The present study provides novel evidence that the tea phenolics EC and EGCG impair NaB uptake and the subsequent NaB-induced differentiation in HT29 cells. These novel findings suggest that although both NaB and green tea catechins have been reported to have a wide range of beneficial effects for human health, when used in tandem the functions of the two compounds interfere at a cellular level. This suggests that NaB and green tea catechins should be used separately. Further studies may be required to investigate the physiological significance of our findings, which provides a better understanding about the interactions of prebiotics and may aid in preparing a rational design for preventive and therapeutic interventions.

Supplementary data to this article can be found online at <http://dx.doi.org/10.1016/j.bbadis.2013.08.009>.

Conflict of interest

The authors declare that they have no conflict of interest.

Acknowledgements

The authors thank Ursula Valls Benavides, Marta Camps Camprubí, Mireia Pérez Verdaguer and Anna Oliveras Martínez for technical support in the experiments and all Pradeep K. Dudeja lab members, especially Arivarasu N. Anbazhagan, for help and support during the stay in their city and laboratory. Funding was provided by Spanish government (grants SAF2011-25726, AGL200912374-C03-03/ALI and personal financial support-FPU program); Spanish Networks RTICC RD06/0020/0046; AGAUR—Generalitat de Catalunya (grants 2009SGR1308 and Icrea Academia award 2010 to M.C.); NIDDK grants DK54016, DK81858 and DK92441; and European Commission (FP7) (ETHERPATHS KBBE-grant agreement 222639 and COSMOS KBBE-grant agreement 312941).

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