Regulation of Cysteinyl Leukotriene Receptor 2 Expression—A Potential Anti-Tumor Mechanism

Cecilia Magnusson¹, Astrid M. Bengtsson¹, Minghui Liu¹, Jian Liu¹, Yvonne Ceder³, Roy Ehrnström², Anita Sjölander¹*

¹Cell and Experimental Pathology, Department of Laboratory Medicine, Lund University, Skånes University Hospital, Malmö, Sweden, ²Pathology, Department of Laboratory Medicine, Lund University, Skånes University Hospital, Malmö, Sweden, ³Clinical Chemistry, Department of Laboratory Medicine, Lund University, Skånes University Hospital, Malmö, Sweden

Abstract

Background: The cysteinyl leukotrienes receptors (CysLTRs) are implicated in many different pathological conditions, such as inflammation and cancer. We have previously shown that colon cancer patients with high CysLT1R and low CysLT2R expression demonstrate poor prognosis. Therefore, we wanted to investigate ways for the transcriptional regulation of CysLT2R, which still remains to be poorly understood.

Methodology/Principal Findings: We investigated the potential role of the anti-tumorigenic interferon α (IFN-α) and the mitogenic epidermal growth factor (EGF) on CysLT2R regulation using non-transformed intestinal epithelial cell lines and colon cancer cells to elucidate the effects on the CysLT2R expression and regulation. This was done using Western blot, qPCR, luciferase reporter assay and a colon cancer patient array. We found a binding site for the transcription factor IRF-7 in the putative promoter region of CysLT2R. This site was involved in the IFN-α induced activity of the CysLT2R luciferase reporter assay. In addition, IFN-α induced the activity of the differentiation marker alkaline phosphatase along with the expression of mucin-2, which protects the epithelial layer from damage. Interestingly, EGF suppressed both the expression and promoter activity of the CysLT2R. E-boxes present in the CysLT2R putative promoter region were involved in the suppressing effect. CysLT2R signaling was able to suppress cell migration that was induced by EGF signaling.

Conclusions/Significance: The patient array showed that aggressive tumors generally expressed less IFN-α receptor and more EGFR. Interestingly, there was a negative correlation between CysLT2R and EGFR expression. Our data strengthens the idea that there is a protective role against tumor progression for CysLT2R and that it highlights new possibilities to regulate the CysLT2R.

Introduction

Prolonged inflammation is known to increase the risk of developing cancer [1,2]. It has been estimated that 15% of all cancers can be accredited to infectious agents [3]. Tumor microenvironment has often been associated with infiltrating leukocytes in the tumor tissue and the surrounding stroma [4]. Moreover, patients with ulcerative colitis display a 30 to 50% increased risk of developing colon cancer [5]. Elevated levels of inflammatory mediators, such as the cysteinyl leukotrienes (cysLTs), have been found in these patients [6]. CysLTs is the collective name for LTC₄, LTD₄ and LTE₄. They are lipid mediators derived from arachidonic acid through the 5-lipooxygenase (5-LO) pathway [7]. In addition to inflammatory bowel disease (IBD) [7], they have been implicated in the pathogenesis of several chronic inflammatory diseases, such as asthma, pulmonary fibrosis, and atherosclerosis [8,9]. The known biological effects of cysLTs are mediated through two different G-protein coupled receptors (GPCRs): CysLT1R [10], which is the high affinity receptor for LTD₄ and CysLT2R [11], which is the high affinity receptor for LTC₄. Recently, an orphan GPCR denoted as GPR17 was found to be an additional receptor for cysLTs. GPR17 can also be activated by uracil nucleotides [12]. There is a 38% sequence similarity between CysLT1R and CysLT2R and they are both able to induce calcium release upon activation [13,14]. However, they do not appear to be functionally interchangeable and they might even antagonize each other’s actions. For example, CysLT2R has been found to inhibit CysLT1R-induced proliferation in mast cells [15]. Colon tumors generally have an increased expression of CysLT1R [16] and a decreased expression of CysLT2R [17]. It has also been observed that colon cancer patients with high expression levels of CysLT1R have poor prognosis. Similar results are also seen in patients with breast cancer [18]. CysLT2R, on the other hand, has been implicated in...
the differentiation of Caco-2 cells [17] and is involved in vascular permeability [19]. This indicates that CysLT2-R has a more protective role in cancer development while CysLT1-R favors tumor progression.

Interferons (IFNs) are part of our defense system against viruses, bacteria, parasites, and malignant cells [20]. Both type I (for example IFN-α and IFN-β), and type II (IFN-γ) IFNs are produced during the early innate immune response. However, while IFN-γ is secreted mainly by natural killer cells and T-cells, the type I IFNs are produced by most cell types following virus infection or toll-like receptor activation [21]. IFN activation results in the increased activity of the transcription factor interferon regulatory factor 7 (IRF-7), which is involved in a positive feedback loop to augment a larger induction of IFN-α and IFN-β genes [22]. Even though both types of IFNs have been implicated to have anti-carcinogenic effects, the type I IFNs exhibit stronger anti-proliferative and anti-angiogenic effects and have been proposed to have apoptotic effects on cancer cells [23,24]. In a mouse model, the type I IFNs have also been implicated to have anti-inflammatory functions in colitis [25]. In addition, they have demonstrated some effectiveness in the treatment of IBD patients [26]. We located a predicted IRF-7 binding site in the putative promoter region of the CysLT2-R, which suggests that the CysLT2-R promoter might be regulated by the IFN-α. This is interesting given that the anti-proliferative and anti-tumorigenic effects attributed to the IFN-α are consistent with the observed characteristics of the CysLT2-R function [17]. IFN-γ has previously been reported to increase the expression of CysLT2-R mRNA and protein in eosinophils [27]. In addition it enhances the CysLT2-R induced inflammatory response of primary endothelial cells [28].

Epidermal growth factor (EGF) signaling via the EGFR is a known inducer of cell proliferation and tumor cell invasion [29,30]. Examining the putative promoter region for the CysLT2-R, we located four conserved E-box elements (consensus sequence CANNTG). EGF signaling has been seen to induce transcriptional repression through E-boxes [31].

In this study, we found novel regulation factors for the CysLT2-R. The anti-tumorigenic cytokine IFN-α is able to increase the transcription of the CysLT2-R while EGF, a known inducer of cell proliferation and migration, suppresses the CysLT2-R expression. This is consistent with the proposed anti-tumorigenic role for CysLT2-R in colon cancer.

Materials and Methods

Chemicals

The rabbit polyclonal anti-human CysLT2-R antibody (diluted 1:1000) was purchased from Innovagen (Lund, Sweden). The ligand LTC4 and Montelukast were from Cayman Chemicals Co. (Ann Arbor, MI). The mouse anti-actin (diluted 1:2000) was from Dako (Glostrup, Denmark). Methyl-3H-thymidine, the enhanced chemiluminescence (ECL) reagents and the Western blot detection reagents as well as the hyperfilm were from Amersham International (Buckinghamshire, UK). TaqMan primers and master mix for real-time PCR were purchased from Applied Biosystems (Cambridge, UK). The RNasey Plus Mini kit was from Qiagen GmbH (Hilden, Germany). Unless otherwise stated, all other reagents were of analytical grade and purchased from Sigma Chemicals Co. (St. Louis, MO) or from ICN (Temecula, CA).

Cell Culture

The non-transformed human intestinal epithelial cell line (Int 407) [32], which exhibits typical epithelial morphology and growth, was cultured as a monolayer for approximately five days in Eagle’s basal medium supplemented with 15% newborn calf serum. The two colon cancer cell lines, Caco-2 (DSMZ No: ACC 169) and SW 480 (DSMZ No: ACC 313), were respectively grown in Dulbecco's modified Eagle medium and RPMI 1640 supplemented with 10% and 20% fetal bovine serum respectively. All media was supplemented with 2 mM L-glutamine, 55 IU/mL penicillin and 55 μg/mL streptomycin. The cell lines were cultured at 37°C in a humidified atmosphere with 5% CO₂. The cells were regularly tested to ensure the absence of mycoplasma contamination.

Cell Lysates

Cells were washed, scraped with ice-cold 1×PBS, and centrifuged for 5 min at 1000 g. The cell pellets were lysed with lysis buffer [33] for 30 min on ice, homogenized 10 times with a 20G needle, and centrifuged for 5 min at 500 g.

Gel Electrophoresis and Immunoblotting

To ensure equal loading, all samples were evaluated and compensated for protein concentration using the Coomassie blue protein assay. Proteins were denatured by boiling in sample buffer for 5 min [34]. The samples were subjected to electrophoresis on 8% or 10% polyacrylamide gels in the presence of 10% SDS. The immunoblotting and developing were performed as described in Nielsen et al. [34].

Real-Time PCR Analysis

Cells were washed in PBS and immediately frozen at −80°C. Thereafter, they were scrapped in lysis buffer provided in the kit and homogenized 10 times with a 20G needle. RNA was purified on RNasey MiniElute Spin Columns and dissolved in RNase free H₂O. cDNA synthesis was performed using RevertAid H Minus M-MuLV reverse transcriptase (Fermentas Life Sciences). The mRNA expression levels of CysLT2-R, CysLT1-R, mucin-2 and the endogenous housekeeping gene, hypoxanthine phosphoribosyltransferase 1 (HPRT-1), were quantified using real-time PCR analysis (TaqMan Chemistry). cDNA was mixed with 0.9 nM TaqMan primers and master mix and amplified at 60°C in an Mx3005P thermocycler (Stratagene). The following primers were used: CysLT2-R (Hs00252658_s1), CysLT1-R (Hs00929113_m1), mucin-2 (Hs00159374_m1), and HPRT-1 (Hs00999999_m1). The samples were analyzed and normalized against HPRT-1 using the MxPro software (Stratagene).

Plasmid Constructs

Human CysLT2-R promoter fragments were subcloned into a pGL3-enhancer vector (Promega) containing a luciferase reporter gene. To produce construct I, one thousand base pairs upstream of the transcription start site of the CysLT2-R gene were amplified from genomic DNA extracted from the human Int 407 cell line by nested PCR. The following primers were used in the process:
forward 5'-ACATGCAGCAGCATATAATGT-3' and reverse 5'-GGGACATATTTTTCTTCCAT-3', followed. This was followed by a PCR that use the following primers; forward 5'-TTATGACGGGTCTTTTTTCCGTTTGCT-3' and reverse 5'-TTCCTATTGGCTGGGCTTAAGAGAAC-3'. All PCR products were gel purified using the QIAquick gel extraction kit (Qiagen). They were digested with restriction enzymes and ligated into the pGL3-enhancer reporter vector using the Quick ligation kit (New England Biolabs). The plasmids were transformed into SoloPack Gold super competent cells (Stratagene). To produce the clone with the deleted IRF-7 binding site (construct II), the site directed mutagenesis with the following primers were used: forward 5'-ATGGGCTATTTCTTCTTTTCAATGGTTATTTGAAAC-3' and reverse 5'-ACATTCGGTCACTTCTTTCAATTTTCCCCAATGTTAAGAAC-3'. The Quick change XL kit (Stratagene) was used according to standard protocol. Three deletion constructs were produced by digesting construct I with restriction enzymes. Construct III was digested with SacI and SpeI to produce a construct from -1 to -412. Construct IV was digested with HindIII and SpeI to remove the downstream part of the promoter, thereby creating a construct between sites -1012 and -413 of the CysLT2R promoter. Blunt end digestion was performed using T4 DNA polymerase (Promega) and the vector was ligated back together. The promoter construct V was digested with SacI and Smal to give a construct from -1 to -187. All plasmids were extracted using Endofree plasmid maxi kit (Qiagen) and were sequenced accordingly.

Transient Transfections and Measurement of Luciferase Activity

Transient transfections of Int 407 cells were carried out using Lipofectamine LTX and Plus Reagent (Invitrogen) in serum free medium according to the manufacturer’s instructions. The final DNA concentration used for transfections was 1 μg/ml for all plasmids except for the luciferase transfection control vector Renilla (0.2 μg/ml). The transfections were carried out for 4–6 h in 37°C, after which the medium was changed to 15% serum containing the medium. Forty-eight hours after transfection, the cells were washed with IFN-α (500–2000 U/ml) or EGF (100 ng/ml) in serum free medium for 24 h. The cells were lysed with passive lysis buffer (Promega) and stored in -80°C. Thawed lysates were centrifuged for 5 min at 1000 g and analyzed using the Dual Luciferase Reporter Assay System from Promega, following the manufacturer’s instructions.

Electromobility Shift Assay (EMSA)

Nuclear extracts were prepared from Int 407 cell treated with IFN-α (1000 U/ml) or EGF (100 ng/ml) for 24 h using the Nuclear Extraction Kit (Chemicon International). The process was performed according to the manufacturers’ instructions. In additional, 10 mM 1-Naphthyl phosphate monosodium salt monohydrate, a broad phosphatase inhibitor, (Sigma Aldrich) was added into both cytoplasmic and nuclear lysis buffer before use. Biotin 3′-end labeled single-stranded oligonucleotide corresponding to the IRF-7 binding region of CysLT2R putative promoter (sense: 5′-AAT CAG GAA ATT TAA ATT TAT TAT-3′ and antisense: 5′-ATA ATA AAT TTA AAT TTA AAT TTC GCT GTG-3′) and E-box binding site (sense: 5′-TTC TTT CCG CAT ATT TTA AAT TTGA AAA AAG TGG-3′ and antisense: 5′-CAG ATT TCT CAA ATG CTG AAA GAA-3′) were purchased from TAG Copenhagen A/S. DNA probes were then annealed in buffer containing 10 mM Tris, 1 mM EDTA, and 50 mM NaCl (pH 8.0) by incubating the oligonucleotides at 95°C for 5 min then gradually reducing the heat until the DNA reached room temperature. Binding reactions for EMSA were prepared using the LightShift Chemiluminescent EMSA kit (Pierce Thermo Scientific). The binding buffer for IRF-7 contained 10 mM Tris-HCl (pH 7.5), 1 mM EDTA, 50 mM NaCl, 2 mM dithiothreitol, 5% glycerol, 0.5% NP-40, and 10 μg/ml BSA. To reduce nonspecific binding, 62.5 μg of poly(dI-dC) was added per ml. Each reaction contains 20 μg nuclear extract. After 20 min of incubation with the probe at room temperature, extracts were loaded on a 5% polyacrylamide gel. After 1 h at 100 V, the gel was transferred onto a positive charged Nylon Membrane, (Thermo scientific) and detected according to kit instruction. The binding buffer for E-box contained 20 mM HEPES, pH 7.6, 150 mM KCl, 1 mM MgCl2, 10% glycerol, 0.2 mM ZnSO4, 0.3 mg/ml BSA and 1 μg/20 μl of poly(dIdC). The binding reaction was performed on ice for 30 min before being loaded onto the gel. To demonstrate the specificity of protein-DNA complex formation, a 1,000-fold molar excess of unlabeled oligonucleotides or IRF-7 (G-8) antibody (Santa Cruz) or SNAI 1 (E-18) antibody (Santa Cruz) was added to the binding reaction mixtures before the probe.

Alkaline Phosphatase Activity

The assay was performed as previously described [35]. Briefly, alkaline phosphatase activity was measured using disodium p-nitrophenyl phosphate as substrate. Caco-2 cells were seeded in petri dishes and incubated for 24 h at 37°C. Five hundred to 2000 U/ml of IFN-α and/or 40 nM of LTC4 was added to the cells and subsequently incubated for 72 h at 37°C. New LTC4 was added every 24 h. Sodium butyrate (2 mM) was used as a positive control (data not shown). The alkaline phosphatase activity was estimated by measuring the level of p-nitrophenol at 405 nm.

Thymidine Incorporation Assay

Caco-2 cells were grown in flat-bottomed 96 well plates for five days. The medium was changed to serum-free 2 h prior to stimulation. Cells were stimulated with LTC4 (40 nM), IFN-α (1000 U/ml) or EGF (100 ng/ml) for 48 h. DNA synthesis was assessed by adding 0.5 μCi of [3H] thymidine during the final 18 h of stimulation. The wells were washed once with PBS and incubated with 0.05% trypsin-EDTA solution/well for 5 min at 37°C. Cells were harvested and collected on a filter paper in a Perkin Elmer harvester. The filter paper was dried and [3H] thymidine incorporation was measured in a 1450 Microbeta Trilux liquid scintillation counter (Wallac).

Cell Migration Assay

Cell migration was analyzed in a modified Boyden chamber, which consisted of two chambers separated by a polycarbonate PVDF membrane (pore size of 8 μm) that was covered with a collagen I gel (3 mg/ml). Int 407 cells were grown for 18 h in the presence or absence of LTC4 (40 nM) and/or EGF (100 ng/ml) in medium containing 1.5% serum. Fifteen percent serum was used as a positive control. Cells were added to the upper well and allowed to migrate into the chamber for 18 h at 37°C. The migrated cells were fixed in 4% paraformaldehyde and the membranes were stained with a 1% crystal violet/10% in a methanol solution. The membranes were washed in PBS and the remaining dye was removed with 10% SDS. The absorbance was measured at 590 nm.

Patient Samples

Archival formalin fixed and paraffin embedded colon cancer and control colon specimens of colorectal cancer patients
collected in 1990 were obtained from the biobanks of Malmö University Hospital. Tissues from 78 patients with varying grades and stages of the disease were included. Grading of the tumors was performed using Dukes’ classification [36]. The matched control samples of normal colon tissue in this investigation were taken from the borders of the surgical specimens.

Ethics Statement
This study was performed after ethical permission from the Regional Ethical Review Board, Lund University # 367/2005. Archival tissue specimens from 78 colon cancer patients that were operated between 1990 and 1991 were used in the present study. Since the samples were old and taken from different parts of the region, it was not possible to obtain written consent from each patient. Detailed information describing the study and tissue microarray (TMA) construction was published in 2006 in a daily newspaper and patients were offered to contact us by mail or by phone if they had any objections. None of the 78 patients objected. This procedure was performed following strict guidelines from the Regional Ethical Review Board in Lund who approved the procedure.

Tissue Arrays and Immunohistochemistry
For histological assessment, the archival paraffin-embedded colorectal cancer and normal mucosa specimens were used and prepared as previously described [16]. The array was stained with anti-IFNα/βR1, EGFR and CysLT2R antibodies. Secondary peroxidase antibodies used were from Dako. After immunostaining, all slides were counterstained with Mayer’s hematoxylin.

Statistical Analyses
Statistical significance was determined as P<0.05 by two-tailed Student’s t test or one-way ANOVA. Pearson’s correlation test was used when comparing different sets of immunohistochemistry staining. SPSS software 16.0 (SPSS, Inc.) was used for the patient material analysis.

Results
An IRF-7 Binding Site and Four E-boxes Have Been Found to be Present in the CysLT2R Promoter Region
To increase the understanding of the role of cysteinyl leukotrienes in inflammation-induced colorectal carcinogenesis, we examined possible ways by which the expression of CysLT2R can be regulated, with one of the receptors generating the effects of leukotrienes. The promoter region of CysLT2R contains several interesting binding sites for known transcription factors and transcriptional repressors. Previous reports from our laboratory have shown that CysLT2R is down regulated in colon cancer [17,37] and that it is involved in colon cancer cell differentiation [17] as well as in decreased cell migration in breast cancer cells [18]. Therefore, it was of particular interest that a putative binding element for IRF-7 and four E-box elements were present in the promoter region of the CysLT2R. Five constructs of the CysLT2R promoter region were created and ligated into a pGL3-enhancer reporter vector to evaluate the importance of IRF-7 and the E-boxes for activation/transcription of CysLT2R (Fig. 1A). IFN-α can be used to activate the transcription factor IRF-7. However, the presence of the IFN-α receptor IFN-αR1 is needed to do this. The presence of the IFN-αR1 is confirmed in the intestinal cell lines intended for this study (Fig. 1B). Likewise, the expression of the EGFR (Fig. 1C) and the CysLT2R (Fig. 1D) was verified.

CysLT2R Promoter Activity is Activated by the IFN-α via an IRF-7 Binding Site and It is Repressed by the EGF Through the E-boxes Present in the Putative Promoter Region
We have seen that CysLT2R expressions were altered in patient’s tumor material, which might be due to a changed “microenvironment around the tumor”. Therefore, we decided to search for a Th1 regulatory cytokine motif in the promoter region of the CysLT2R. The presence of a putative IRF-7 binding site in the CysLT2R promoter region indicated that the Th1 pro-inflammatory cytokine IFN-α ought to induce CysLT2R promoter activity and transcription of the CysLT2R encoding gene. A dilution series with IFN-α showed that 1000 U/ml IFN-α generated an optimal activation of CysLT2R (Fig. 2A). To elucidate whether IFN-α induces the CysLT2R through IRF-7 binding to its putative binding site in the promoter region, the promoter activity for deletion mutant constructs I-V were tested (Fig. 2B). IFN-α induced a significant two-fold activation of the CysLT2R in the non-transformed cell line Int 407 (construct I). This activation was partly reduced in the ΔIRF7 mutant (construct II). IFN-α was unable to induce the CysLT2R promoter activity in constructs IV and V, which both lacked full-length IRF-7 response elements. However, a significant reduction in activation compared to wild-type construct I was also observed for construct III, which contains an intact IRF-7 response element but lacks the part upstream of bp –412. Taken together, this indicates that the IFN-α-induced activation of the CysLT2R requires the IRF-7 binding site for complete activation, but also needs additional unidentified binding elements located upstream of the IRF-7 site for full activation.

Further examination of the CysLT2R promoter sequence revealed the presence of four conserved E-box elements. EGF signaling can induce repressor elements that bind to mentioned sequences. Four of the CysLT2R luciferase reporter constructs were used to investigate the effects of EGF on the CysLT2R. We found that EGF suppressed CysLT2R luciferase reporter activity (Fig. 2C). The importance of the E-boxes was investigated by testing the ability of the EGF to suppress CysLT2R promoter activity in three deletion constructs, (construct III, with only E-box 4 present, construct IV containing three E-boxes, and construct V without any E-boxes). The EGF-induced reduction of CysLT2R promoter activity significantly declined with a decreasing number of E-boxes present in the putative promoter region. There was a statistically significant difference between the EGF-induced reductions of construct IV and construct V. However, the reduction is probably mediated both through an E-box-dependent as well as an E-box-independent pathway in the intestinal cells.

Analysis of IRF-7 and E-box DNA binding activities on the CysLT2R putative promoter region in Int 407 cells were performed to compliment the luciferase experiments (Fig. 2D). The EMSA results showed that IRF-7 binding complexes were detected in both untreated and IFN-α treated cells. We found an increased association upon the stimulation of the complexes. The addition of cold probe and anti-IRF-7 antibody competed with IRF-7 in binding the IRF-7 binding site on the promoter region. The addition of cold probe could completely inhibit probe-protein complex formation. Likewise, Snail-containing complexes were detected in both untreated and EGF treated cells. We found increased association upon stimulation of the complex. The addition of cold probe could completely inhibit the formation of probe-protein complex. Moreover, additional anti-Snail antibody reduced the probe-protein complex formation.

Next, we wanted to see that the regulatory effects for IFN-α and EGF observed in connection with CysLT2R luciferase reporter
EGFR and Snail activation is involved in EGF repressing CysLT2R

Stimulation with EGF leads to the activation of the EGFR (Fig. 3A) and to the activation of Snail (Fig. 3B), a transcriptional repressor known to bind to E-box elements. It is likely that the EGF-induced suppression of the CysLT2R is generated by Snail that binds to the E-box elements present in the promoter region of the CysLT2R. Interestingly, an induction of CysLT2R mRNA is observed after EGF treatment of intestinal cells at the same time as CysLT2R mRNA is suppressed (Fig. 3C), indicating, as earlier suggested, opposing effects of the CysLTRs [37].

IFN-α Can Induce Differentiation of Caco-2 Cells

Sodium butyrate is a known inducer of differentiation in Caco-2 cells [38], which is frequently used as a model system for differentiation in cell culture [39]. The activation of different luminal brush border proteins, such as alkaline phosphatase, can be used as a measurement for epithelial cell differentiation. Previously, we have shown that the CysLT2R ligand LTC4 is able to induce differentiation in Caco-2 cells [17]. IFN-α induced a small but significant increase in the activation of alkaline phosphatase, indicating increased differentiation of Caco-2 cells. IFN-α had, however, no additive effect to the alkaline phosphatase activity induced by LTC4 (Fig. 4A). To confirm the effect of alkaline phosphatase activity, an additional differentiation assay was used. Mucin-2 is a marker for intestinal cell differentiation [40]. Both LTC4 and IFN-α could significantly induce mucin-2 mRNA in Caco-2 cells (Fig. 4B). The CysLT2R antagonist AP-100984 could significantly block both LTC4 induction of mucin-2, which was not seen with the specific CysLT1R antagonist Montelukast. This strengthens the hypothesis that CysLT2R might have a protective role in colon cancer and in normal intestinal mucosa. We have previously shown that LTC4 does not induce proliferation in Caco-2 cells [17]. Here, we show that IFN-α does not induce a proliferative response either. EGF, on the other hand, is a known mitogen inducer [41] and was used here as a positive control (Fig. 4C). This corresponds well to the previous reported roles of the CysLTRs.

We next investigated the effect of CysLT2R signaling on Int 407 cell migration. We found that LTC4-induced activation of CysLT2R reduced Int 407 cell migration (Fig. 4D). Furthermore, we also found that EGF-induced Int 407 cell migration could be suppressed by CysLT2R activation (Fig. 4D).

EGF Inversely Correlates with CysLT2R in Colorectal Cancer

Next, we wanted to clarify if the levels of IFNα/βR1 and EGFR correlated with the CysLT2R pattern in a colon cancer patient array. The TMA contains material from 78 colon cancer patients and has previously been thoroughly described [16]. Representative stainings from the array for IFNα/βR1 and EGFR are shown in Fig. 5A. We have previously shown that CysLT2R is down-regulated in colon tumor tissue in patients from this array and that more aggressive tumors expressed less of the CysLT2R [17]. We observed that patients with more aggressive tumors generally had less expression of IFNα/βR1 while patients with less aggressive tumors generally exhibited a higher expression of IFNα/βR1. A more distinct trend was visualized when the patients were divided into subgroups with high (+++, ++++) or low (+, +++) IFNα/βR1 staining. Duke’s A patients had a higher percentage of increased IFNα/βR1 staining than Duke’s B and Duke’s C patients (Fig. 5B). The expression of the EGFR was generally up-regulated in colon cancer patients in this study, which is coherent with earlier published studies [42] (Fig. 5C). Representative pictures from a

activity also affected CysLT2R mRNA and protein expression. Consistent with the results from the gene activation assay, the IFN-α significantly up-regulated both the CysLT2R mRNA (peaking at 12 h) and the protein (maximum increase at 24 h) (Fig. 2E). The EGF induced a significant suppression of both the CysLT2R mRNA and the protein expression (Fig. 2F).
Duke’s C colon cancer patient are shown in Fig. 5D. They exhibit high expression of CysLT2R and IFN-α/βR1 in control tissue and decreased expression in the tumor. Conversely, EGFR expression was enhanced in the tumor material. A statistical significant negative correlation could be observed in the expression levels between CysLT2R and EGFR in this tissue array. Although IFN-α/βR1 and CysLT2R had similar expression patterns in relation to cancer staging, no significant correlation between the expression levels of these two receptors could be found (Table 1).

**Discussion**

Several different ways of regulating CysLT2R have previously been reported. For example, LTD4 can up-regulate the mRNA expression of CysLT2R in monocytes [43] and epithelial cells [44] and the cytokines IL-4, IL-8 and IFN-γ [28,45,46] can up-regulate the mRNA expression of CysLT2R in several different cell types. Since a binding site for IRF-7 was present in the putative promoter region of the CysLT2R, we hypothesized that the IFN-α would be able to induce CysLT2R expression via transcription factor IRF-7. The ability of the IFN-α to induce anti-proliferative and anti-tumor progression responses [24,47] made it an interesting candidate to study in relation to CysLT2R. IFNs have been reported to mediate their anti-tumor effects by altering immune responses, such as the suppression of cytokine IL-1 [48] and inducing TRAIL [49], and by inducing anti-angiogenic responses by inhibiting VEGF [50]. We show that the IFN-α induced a two-fold activation of the wild-type CysLT2R promoter and increased expression in the CysLT2R mRNA and protein. How the IFN-α mediates its effects, however, is not completely elucidated. The IRF-7 binding site is important and required for the optimal induction of the CysLT2R promoter by the IFN-α, but it is not the only crucial activation site. Additional transcription factors are probably engaged as well. We discovered that the promoter region of the CysLT2R contains four conserved E-
box elements. Interestingly, the mitogenic EGF suppressed both CysLT2R mRNA and protein expression as well as the promoter activity, strengthening the hypothesis that the CysLT2R is a potential tumor suppressor. To validate the importance of the E-boxes, CysLT2R promoter deletion constructs were used. EGF stimulation significantly decreased the CysLT2R promoter activity in all promoter constructs, but to a significantly reduced capacity in the deletion constructs. These results indicate that the E-box elements are important but are not the only mechanisms that allow the EGF to suppress the transcription of the CysLT2R. We cannot rule out that other transcription factors can also be activated by the EGF to reduce the CysLT2R luciferase reporter activity of construct V that lack the E-boxes. Mann et al. report that EGF, through the EGFR, can repress the proposed tumor suppressor prostaglandin dehydrogenase (PGDH), allowing PGE2 to accumulate and activate the pathway repeatedly in a positive feedback loop. They show that EGFR signaling induces Snail, which binds to the conserved E-box elements in the PGDH promoter and represses transcription [31]. It is likely that the repression of the CysLT2R in the presence of EGF is a result of EGFR signaling via the EGFR, thereby activating the Snail, which then functions as a transcriptional repressor when bound to the E-boxes present in the CysLT2R promoter region.

Earlier, we have shown that LTC4 is able to induce the differentiation of Caco-2 cells [17]. In this study, we show that IFN-α is also able to induce differentiation in colon cancer cells. This was demonstrated by the induction of mucin-2 after IFN-α stimulation. Mucins are highly glycosylated proteins that build up the mucus layer protecting the underlying epithelial cells in the intestine. The colon epithelium mainly secretes mucin-2 [31], which is known to play a protective role against colorectal diseases, such as colon cancer [52] and colitis [53]. Mucin-2-deficient mice frequently develop adenomas, which progress into invasive adenocarcinomas, demonstrating that mucin-2 is involved in colon cancer suppression [32]. Interestingly, we demonstrate that both LTC4 and IFN-α significantly induced the mRNA expression of mucin-2. Taken together, this implicates that both LTC4 and IFN-α are involved in the stimulation of mucus synthesis and are important in protecting against mucosal damage in the colon. We have previously shown that endogenous signaling by overexpressed CysLT2R is enough to suppress cell migration in breast cancer cells [18]. Here, we show that LTC4 signaling mediates a suppression of cell migration and reduces the cell migration induced by EGF signaling. The ability of CysLT2R signaling to suppress cell migration is probably one of the reasons why high CysLT2R expression is connected to a better prognosis for colorectal cancer patients. It is possible that the balance between positive and negative gene regulators, such as repressors and transcription factors and their co-factors, are disrupted in the tumor microenvironment. This could explain the manner by which the down-regulation of CysLT2R plays a role in colon cancer progression.

We observed a decrease of IFNα/βR1 in tumors in colon cancer patients. Considering the protective role that IFNs play in the innate immune system (i.e. participating in the cell defense against infections and tumor cells), we speculate that tumor cells have an impaired innate immune response. There are epidemiological associations between pathogen invasion leading to chronic inflammation and malignant transformations [2]. For example, virus infections have been connected to increased incidences of cancer. Infection with hepatitis B or C viruses can result in chronic hepatitis and liver cirrhosis, which are believed to be a major cause
of hepatocellular carcinoma [2]. In addition, the IFN-α can achieve anti-tumorigenic effects by affecting tumor cell differentiation [47]. The observation that IFN-α/βR1 had lower expression in more aggressive tumors agrees with an anti-tumorigenic role for IFN-α. Furthermore, the expression of IFNα/βR1 was reduced in tumor tissue compared to normal tissue from the surgical borders of the tumor in the patients, contributing to the notion of altered IFN signaling in tumor cells.

Figure 4. CysLT2R signaling mediates anti-tumorigenic effects. (A) An alkaline phosphatase activity assay was used to determine the differentiation of Caco-2 cells. Cells were treated with IFN-α (500–2000 U/ml) and/or LTC4 (40 nM) for 72 h. The alkaline phosphatase activity was determined by measuring the absorbance at 405 nm due to formation of p-nitrophenol. (B) QPCR quantification of mRNA expression of MUC2 with or without treatment with LTC4 (40 nM), IFN-α (1000 U/ml), pretreatment for 30 min with CysLT2R inhibitor AP-100984 (1 μM) or CysLT1R inhibitor Montelukast (1 μM), in Caco-2 cells. (C) Caco-2 cells were incubated with LTC4 (40 nM) and/or IFN-α (1000 U/ml) for 48 h in medium containing 1.5% serum. To determine proliferation by thymidine uptake, [methyl-3H] thymidine (0.5 μCi/well) was added to the wells during stimulation. (D) Cell migration was analyzed with Int 407 cells grown in the presence or absence of EGF (100 ng/ml) and/or LTC4 (40 nM). The cells were allowed to invade a collagen gel on top of a Boyden chamber for 18 hrs. The results are shown as means ± SD of at least three different experiments; *, P<0.05; **, P<0.01; ***, P<0.001.

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The same clinical material displayed overexpression of EGFR in tumor tissue. The expression pattern for CysLT2R [17] is similar to the expression of anti-tumorigenic IFNα/βR1 while the expression of CysLT1R is increased in colon cancer [16] similar to the expression of EGFR. Both CysLT1R and EGFR exhibit pro-tumorigenic effects by driving the processes of cell migration and proliferation [29,30,33,54]. The nuclear localization of EGFR has been linked to more aggressive tumor types [29] and increased nuclear localization of CysLT1R has also been observed in colon cancer cells [34]. We show that EGF signaling induces CysLT1R mRNA while suppressing CysLT2R. Interestingly, EGFR and CysLT2R display a significant inverse correlation in their expression pattern among the colon cancer patients in the array. In conclusion, we have located one response element for IRF-7 and four E-boxes in the promoter region of CysLT2R. We demonstrate that the anti-tumorigenic IFN-α is able to induce CysLT2R promoter activity and expression while EGF, a known inducer of mitogenic effects, is able to suppress it. These data support the hypothesis that CysLT2R might have a protective role in colon cancer.

Figure 5. Representative IFNα/βR1 and EGFR staining in normal human colon tissue and colorectal adenocarcinomas. (A) Top row, shows the degree of IFNα/βR1 staining of carcinomas. Bottom row shows the degree of EGFR staining of carcinomas (+/− to ++++, 10x objective). (B) Distribution of high (+, ++++) and low (+/−, +) IFNα/βR1 staining intensities of tumors in Duke’s A, B and C, and (C) of EGFR. Samples are assessed according to total IFNα/βR1 and EGFR staining. Statistics are based on tumors from 78 patients that were included in the array. (D) Representative pairs of control and tumor immunostaining from a Duke’s C patient stained with IFNα/βR1, EGFR and CysLT2R.

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Table 1. CysLT2R immunoreactivity vs. IFNα/βR1 and EGFR.

| Relative Staining | Frequency (% of total) | CysLT2R  | Spearman’s rank | p |
|-------------------|-----------------------|----------|-----------------|---|
|                   |          | +/− | + | ++ | +++ | n |                     |
| IFNα/βR1          |          | 2.8 | 0  | 1 | 1 | 0 | 2 | 0.062 | 0.608 |
|                   |          | 40.8| 7  | 7 | 13| 2 | 29| |
|                   |          | 35.2| 2  | 11| 10| 2 | 25| |
|                   |          | 21.1| 1  | 8 | 5 | 1 | 15| |
| EGFR              | 1.4      | 0   | 1  | 0 | 0 | 1 | −0.392 | 0.001 |
|                   | 36.6     | 9   | 14 | 3 | 26| |
|                   | 36.6     | 3   | 12 | 9 | 2 | 26| |
|                   | 25.4     | 7   | 5  | 6 | 0 | 18| |

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Author Contributions
Conceived and designed the experiments: CM RE ML YC AS. Performed the experiments: CM AMB MI JL RE. Analyzed the data: CM AMB ML.

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