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CCK₂ RECEPTOR EXPRESSION TRANSFORMS NON-TUMORIGENIC HUMAN NCM356 COLONIC EPITHELIAL CELLS INTO TUMOR FORMING CELLS

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Abstract

Expression of gastrin and cholecystokinin 2 (CCK₂) receptor splice variants (CCK₂R and CCK_{2i4sv}R) are upregulated in human colonic adenomas where they are thought to contribute to tumor growth and progression. To determine the effects of ectopic CCK₂ receptor variant expression on colonic epithelial cell growth *in vitro* and *in vivo*, we employed the non-tumorigenic colonic epithelial cell line, NCM356. Receptor expression was induced using a retroviral expression vector containing cDNAs for either CCK_{2i4sv}R or CCK₂R. RT-PCR and intracellular Ca²⁺ ([Ca²⁺]_i) imaging of RIE/CCK₂R cells treated with conditioned media (CM) from NCM356 revealed that NCM356 cells express gastrin mRNA and secrete endogenous, biologically active peptide. NCM356 cells expressing either CCK₂R or CCK_{2i4sv}R (71 and 81 fmol/mg, respectively) grew faster *in vitro*, and exhibited an increase in basal levels of phosphorylated ERK (pERK), compared to vector. CCK₂ receptor selective antagonist, YM022, partially inhibited the growth of both receptor-expressing NCM356 cells, but not the control cells. Inhibitors of mitogen activated protein kinase pathway (MEK/ERK) or protein kinase C (PKC) isozymes partially inhibited the elevated levels of basal pERK and *in vitro* growth of receptor-expressing cells. Vector-NCM356 cells did not form tumors in nude mice, whereas, either CCK₂ receptor-expressing cells formed large tumors. Autocrine activation CCK₂ receptor variants are sufficient to increase *in vitro* growth and tumorigenicity of non-transformed NCM356 colon epithelial cells through a pathway involving PKC and the MEK/ERK axis. These findings support the hypothesis that expression of gastrin and its receptors in human colonic adenomas contributes to tumor growth and progression.

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Ectopic expression of CCK₂ receptor splice variants in the non-tumorigenic colonic epithelial cell line, NCM356, is sufficient to promote enhanced cell adhesion, cell growth, and tumorigenesis, in part, through autocrine activation of protein kinase C (PKC) and mitogen-activated protein kinase (MAPK) pathways. This data supports the hypothesis that CCK₂ receptors and gastrin contribute to colorectal tumor development and growth.

Keywords

Gastrin; CCK₂ receptors; colorectal tumorigenesis; NCM356 cells

INTRODUCTION

The adenoma- to carcinoma-multistage sequence of colorectal cancer development is characterized by specific histopathologic criteria as well as defined genetic mutations, which result in the activation of oncogenes (e.g., *K-ras*) and inactivation of tumor suppressors (e.g., adenomatous polyposis coli [APC] and p53).¹ It is now well recognized that the key initiating events that underlie most cases of colorectal tumorigenesis, whether familial or sporadic, are the mutations within the Wnt/APC/ β -catenin signaling pathway.² However, increasing evidence also indicate that epigenetic changes in DNA and/or chromatin structure, causing aberrant mRNA splicing and/or inappropriate expression of normal genes, can interact with genetic mutations to contribute to the development of the malignant phenotype.³ Aberrant expression of the gastrin/cholecystokinin 2 (CCK₂) receptor, gastrin and its biosynthetic precursors, in a majority of pre-malignant adenomatous polyps strongly implicate a relevant role for this signaling axis in the adenoma-carcinoma sequence.⁴⁻⁶

Carboxyl-terminus-amidated gastrin (i.e., mature gastrin) is produced from the cleavage and post-translational processing of a preprohormone protein. Although both gastrin precursors and mature gastrin are prevalent during the early stages of malignant transformation, as well as in established colon cancers, their exact roles in each context are controversial (summarized in a recent review⁷). Several lines of evidence suggest a potentially important function for the peptide hormone particularly in the early stages of colorectal cancer development. First, patients with hypergastrinemia, associated either with Zollinger-Ellison syndrome or chronic autoimmune gastritis, exhibited increased rates of colonic mucosal cell proliferation,^{8, 9} whereas hypergastrinemia due to pernicious anemia, *Helicobacter pylori* infection or other causes has been associated with an increased risk of developing colonic polyps and/or cancers.¹⁰⁻¹² Second, a mechanistic relationship has been established between mutations in the Wnt/APC/ β -catenin pathway, aberrant gastrin gene expression, and gastrin-mediated signal transduction using both *in vivo* and *in vitro* rodent models of early colorectal carcinogenesis. Specifically, Koh et al.¹³ have shown that mice derived from a cross between the APC^{min-/+} mouse, a model of familial adenomatous polyposis, and a gastrin gene knockout mouse developed fewer intestinal polyps. Additionally, Watson et al.¹⁴ showed in the APC^{min-/+} mouse model, that proton pump inhibitor-induced hypergastrinemia increased mucosal proliferation, polyp development, and decreased survival. Treatment with anti-gastrin antibodies inhibited the effects of hypergastrinemia on mucosal proliferation and animal survival. Finally, *in vitro* experiments demonstrated that induction of the wild-type APC decreased gastrin mRNA expression, while transfection of constitutively active β -catenin increased gastrin promoter activity.¹³ These data suggest that aberrant gastrin expression is mechanistically linked to initiating genetic mutations within the Wnt/APC/ β -catenin pathway and contributes to the development of the malignant phenotype.

Although it is well known that in normal tissues and cells the biologic actions of mature gastrin are mediated by CCK₂ receptors, members of the rhodopsin β subclass of G protein-coupled receptors,¹⁵ whether CCK₂ receptor is the only mediator the tumorigenic activities of gastrin in colorectal cancer remains an unresolved issue. Central to the controversy is the contention that since the biosynthetic precursors of gastrin (i.e., progastrin and glycine-extended gastrin [Gly-G]) bind the CCK₂ receptor with significantly lower affinities than that for amidated gastrin, there must also exist unique high-affinity receptor(s) for these

precursors. The molecular identity and characteristics of these other receptors, however, currently are not completely elucidated.^{16, 17} Nonetheless, several studies suggest a potential role for CCK₂ receptor variants in the early stages of colorectal carcinogenesis. As the result of alternative splicing at intron 4, at least two functionally distinct CCK₂ receptor variants (CCK₂R and CCK_{2i4sv}R) have been identified in a subset of clinical specimens from human pre-malignant adenomatous polyps when compared to normal colonic mucosa.^{4, 6} Furthermore, genetic ablation of the CCK₂ receptor reduced azoxymethane-induced tumor size and burden compared to wild-type controls and mice overexpressing progastrin,¹⁸ suggesting a relevant function for the CCK₂ receptor, even in the context of gastrin precursors. Finally, intestinal polyps from hypergastrinemic APC^{min-/+} mice exhibited a 6-fold increase in CCK₂ receptor mRNA expression compared to normgastrinemic control animals,¹⁴ supporting an endocrine mechanism of CCK₂ receptor activation. However, human colon adenomas from patient samples demonstrate contemporaneous overexpression of both gastrin peptides and their cognate receptors, implying that a more local or cell-autonomous mechanism of tumor progression may be present. The lack of a non-tumorigenic human colonic epithelial cell model to study potential autocrine/paracrine activation of the CCK₂ receptor signaling pathways has limited our understanding of how aberrant expression of gastrin and CCK₂ receptor variants lead to the development and the promotion of colorectal carcinogenesis.

To address whether acquisition of the CCK₂ receptor is sufficient to convert a non-tumorigenic, gastrin-producing colorectal cell into a tumorigenic cell, we employed the NCM356 epithelial cell line, derived from the normal colon mucosa wide margin resection of a patient with rectal adenocarcinoma.¹⁹ The NCM356 cells express colon epithelial antigens, including cytokeratin, villin and mucin. They do not grow in soft agar, are non-tumorigenic in nude mice, and yet, are immortal in cell culture and as we show herein, inappropriately expresses and secretes functional gastrin peptides. We report that ectopic expression of CCK₂ receptor splice variants is sufficient to promote enhanced cell adhesion, cell growth, and tumorigenesis, in part, through the activation of protein kinase C (PKC) and mitogen-activated protein kinase (MAPK) pathways.

MATERIALS AND METHODS

Materials

The parental NCM356 cell line was acquired under an MTA from InCell Corp. (San Antonio, TX). Antibodies for immunoblotting of phospho-ERK (pERK) and β -actin were purchased from Santa Cruz Biotechnologies (Santa Cruz, CA). Chemical inhibitors PD98059 and GF109203X were purchased from Calbiochem (San Diego, CA). Receptor-mediated signaling was antagonized with YM022 (Sigma; St. Louis, MO) and JB93182, a gift from the James Black Foundation. Gastrin 1-17 (G17) was purchased from Biomol (Plymouth Meeting, PA), and dimethyl sulfoxide (DMSO) was purchased from Sigma (St. Louis, MO).

Retroviral expression constructs and transduced cell lines

CCK₂ receptor variant cDNAs were cloned into a bicistronic packaging murine oncoretroviral vectors based on pFB (Stratagene). The vector contains the murine leukemia retrovirus (MLV) packaging sequence and a multiple cloning site (MCS), flanked by the MLV long terminal repeat (LTR) regions. The 5' LTR functions as a strong promoter upon chromosomal integration of proviral DNA. The pFB plasmid was modified to contain a cassette comprising an ECMV internal ribosome entry site (IRES) followed by a gene encoding β -galactosidase (modified with a nuclear localization signal), which enabled retrovirus titer and transcript expression levels to be determined by staining for β -

galactosidase.²⁰ Retroviruses were made by simultaneous transfection of HEK293FT cells (Invitrogen; Carlsbad, CA) with the CCK₂ receptor expression plasmid and plasmids encoding MLV gag-pol and vesicular stomatitis virus envelope protein. Cell supernatants were used to transduce NCM356 cells (MOI=10). The level of receptor expression in transduced cell lines was quantified using [¹²⁵I]labeled-G17 competition binding to isolated cell membranes as previously described.²¹ NCM356-CCK₂R and -CCK_{2i4sv}R cells express 71 and 81 fmol receptor/mg of membrane protein, respectively. All experiments were performed using the original transduced cell line; there was no additional subcloning of the cultures.

Cell Culture

The parental and transduced NCM356 cell lines were cultured at 37°C in a humidified atmosphere of 95% air and 5% CO₂ in culture media obtained from Incell Corp. (San Antonio, TX). Serum free media M3™ (Incell Corp.) was mixed with M3:10™ containing 10% fetal bovine serum (FBS) to obtain 1% FBS media for the *in vitro* cell proliferation experiments.

Reverse-transcription-polymerase chain reaction (PCR)

Total RNA was extracted using Ultraspec reagent (Biotecx; Houston, TX) and treated with 1 unit of RNase-free DNase I at 37°C for 30 min (Promega, Madison, WI). Messenger RNA was converted to cDNA using Retroscript (Ambion). PCR was performed using the primers for gastrin: sense, 5'-CTTAGGTACAGGGGCCAACA-3' and anti-sense 5'-TCCATCCATCCATAGGCTTC-3'. The PCR conditions were 94°C for 5 min followed by 40 cycles of 94°C for 30 s, 45°C for 30 s, and 72°C for 1 min. A 1-kb DNA ladder (10 µl) (Life Technologies, Inc.) was used to determine the relative size of the PCR products.

Intracellular Ca²⁺ ([Ca²⁺]_i) measurements

[Ca²⁺]_i imaging of rat intestinal epithelial (RIE) cells expressing recombinant CCK₂R (RIE/CCK₂R) was used as a bioassay, to detect gastrin in conditioned culture media collected from parental NCM356 cells. Briefly, RIE/CCK₂R cells were plated on 25 mm glass coverslips, washed with a physiological medium (KRH) containing NaCl (125 mM), KCl (5 mM), KH₂PO₄ (1.2 mM), MgSO₄ (1.2 mM), CaCl₂ (2 mM), glucose (6 mM), HEPES (25 mM; pH 7.4), and loaded with 2 µM Fura-2AM (Molecular Probes, Eugene, OR) at 25°C for 50 min. The cells were stimulated with NCM356 conditioned media (CM) and single cell changes in the concentration of free [Ca²⁺]_i were recorded with a Nikon Diaphot inverted microscope (Garden City, NY) and a CCD camera (Dage-MITI, Inc., Michigan City, IN). Data points were collected every 1-8 s from approximately 35 cells/coverslip and processed using ImageMaster software.

In vitro cell proliferation assay

To study the effects of receptor variant expression on NCM356 cell growth *in vitro* with and without chemical inhibitors, single cell suspensions of 2 × 10⁴ cells were seeded into 24-well plates in triplicate and cultured in 1.0 ml M3:10 media (10% FBS) mixed with M3 (serum free) to a final concentration of 1% FCS with or without G17 (10 nM). Cells were trypsinized and counted by a Coulter counter (Beckman Coulter, Inc., Fullerton, CA) daily for 9 days. The doubling time was calculated using the GraphPad Prism program, Version 4.0 (GraphPad Software Inc, San Diego, CA). Each experiment was performed at least twice.

In vivo tumor growth assay

The athymic nude mouse xenograft model was used to assess the effects of receptor variant expression on NCM356 cell growth *in vivo*. Replicates (n=6 mice for each group) of nu/nu Balb/C female mice (age 8-10 weeks, weight approximately 20 g each) were injected subcutaneously with 5×10^6 cells per cell line in the left dorsum. Tumor diameters were measured transcutaneously using calipers over a 3-week period. Tumor volumes for the generally spherical tumors were calculated using the formula: $\text{volume} = 4/3\pi r^3$. At harvest, the wet weights of the tumors were recorded. T-test comparisons of the tumor weights for each CCK₂ receptor group was performed using Graphpad Prism with significance determined at $p < 0.05$. This experiment was repeated three times.

Western blot

Cells were plated into 12-well plates at a density of approximately 10^5 cells/well in media. After 2 days, cells were washed with ice-cold PBS and solubilized in lysis buffer containing 1% Triton X-100, 150 mM NaCl, 20 mM Tris pH 7.5, 1 mM EGTA, 1 mM EDTA, 2.5 mM sodium pyrophosphate, 1 mM β -glycerophosphate, 1 mM Na₃VO₄, 1 mM phenylmethylsulfonyl fluoride and 5 $\mu\text{g}/\text{ml}$ each of chymostatin, pepstatin A, leupeptin and antipain at 4°C for 15 min. Triton X-100 insoluble cellular material was removed by centrifugation at 14,000 rpm for 15 min and the protein concentrations of the supernatant were determined using the Bio-Rad DC protein assay kit (Bio-Rad Laboratories; Hercules, CA). Protein (20 μg) from each sample was resolved on 8% sodium dodecyl sulfate-polyacrylamide gel electrophoresis, transferred to nitrocellulose membrane and probed with anti-pERK1,2 antibody. Immunoreactive proteins were visualized using the ECL Western blotting detection system (Amersham Biosciences, Piscataway, NY). The blots were stripped and reprobed for β -actin to insure the equal protein loading in all lanes. Densitometry analyses were performed using Un-Scan-It (Silk Scientific; Orem, UT).

RESULTS

The human colonic epithelial cell line NCM356 secretes gastrin peptide

Gastrin binding to the CCK₂ receptor leads to the G_q-mediated activation of phospholipase C- β , which cleave membrane phosphatidylinositol 4,5-bisphosphate to produce the second messengers 1,4,5-inositol triphosphate (IP₃) and 1,2-diacylglycerol. IP₃, in turn, induces Ca²⁺ release from the endoplasmic reticulum by activating inositol-1,4,5-trisphosphate receptors (IP₃R), resulting in a transient increase in the concentration of free cytosolic Ca²⁺ [Ca²⁺]_i.

To assess whether the NCM356 cells secrete gastrin peptides, we developed an [Ca²⁺]_i imaging bioassay, taking advantage of the fact that CCK₂ receptors are coupled to IP₃R. We generated the rat intestinal epithelial (RIE) cell line which expresses recombinant human CCK₂R (RIE/CCK₂R). We treated RIE/CCK₂R loaded with the Ca²⁺ indicator dye Fura-2, with known concentrations of agonists to establish the sensitivity, efficacy and specificity of the bioassay. We determined that the RIE/CCK₂R cells did not induce a calcium response to 10 nM of Gly-G, but did respond robustly to 1 nM of purified G17 (Fig. 1A). Increasing the Gly-G dose to 100 nM, or a 100-fold excess in comparison to 1 nM G17, did cause an increase in [Ca²⁺]_i, but to a lesser extent as compared to that of 1 nM G17 (Fig. 1B). As expected, pretreatment with the selective CCK₂ receptor antagonist L 365,260 (100 nM) abrogated the calcium response to both 100 nM Gly-G and 1 nM G17 in the RIE/CCK₂R cells; however, L 365,260 did not affect the increase in [Ca²⁺]_i due to activation of the bradykinin (BK) receptor after application of 100 nM BK (Fig 1C). As previously reported by Hellmich et al.,⁴ Gly-G binds to the CCK₂ receptors with significantly lower affinity

than the amidated form of gastrin, G17, and this binding can be blocked with specific inhibitors to CCK₂ receptors.

We then applied CM from cultures of parental NCM356 to the RIE/CCK₂R cells in our bioassay and showed that the CM produced a transient increase in [Ca²⁺]_i (Fig. 1D). To demonstrate that the NCM356 CM-induced Ca²⁺ response was mediated by the CCK₂R expressed on RIE cells, the CM was supplemented with 1 μM JB93182, a CCK₂ receptor-specific antagonist. The antagonist blocked the transient increase in [Ca²⁺]_i induced by CM (Fig. 1E). To verify that the cells were loaded with Fura-2 and capable of responding, we applied the muscarinic receptor agonist carbachol, which induced an increase in [Ca²⁺]_i (Fig. 1E). Treating RIE/CCK₂R cells with 1 nM purified G17 induced an increase in [Ca²⁺]_i of similar amplitude and duration (Fig. 1A and B) as cells treated with NCM356 CM (Fig. 1D). Finally, addition of exogenous G17 (100 nM) to parental NCM356 did not induce an increase in [Ca²⁺]_i, indicating the absence of expression of CCK₂ receptors by these cells (Fig. 1D). Together, these data demonstrate that, like many pre-malignant polyps, NCM356 secrete forms of gastrin that activate CCK₂ receptors, such as Gly-G and G17. Although other precursor forms of gastrin may also be present, the receptors mediating their actions are either not known or well defined, and therefore, were not assessed with our bioassay. Finally, since the NCM356 cells do not express functional CCK₂ receptors, this colorectal cell line represents a good model system to address the effects of receptor acquisition, autocrine signaling on cell proliferation, and malignant potential.

Expression of human recombinant CCK₂ receptor splice variants in NCM356 cells increases cellular adhesion and proliferation in vitro

Concomitant with the upregulation of gastrin expression, which has been reported to have a prevalence of 78% in polyps, is expression of CCK₂ receptor variants in up to 81% of colon adenomas,⁵ suggesting the existence of a potential autocrine signaling axis. To model the observations from adenomatous polyps, we transduced parental NCM356 cells with retrovirus (pFB-IRES) expression constructs containing the cDNAs for either CCK₂R or CCK_{2i4sv}R. Cells also were generated containing the pFB-IRES vector alone to control for non-specific effects of transduction. Reverse transcriptase/polymerase chain reaction with primer sequences to human gastrin verified mRNA expression by all three transduced cell lines (Fig. 2A) and the RIE/CCK₂R cell bioassay confirmed the presence of secreted gastrin in CM (data not shown).

In culture, the parental NCM356 cells grow predominantly (>98%) as a suspension of loosely aggregated round cells with a small (<2%) population of adherent (A) cells.¹⁹ The NCM356 vector-transduced cells were phenotypically indistinguishable from parental NCM356 cells (Fig. 2B). However, in contrast to the vector transduced cells, both CCK₂R- and CCK_{2i4sv}R-expressing NCM356 cell lines exhibited predominantly (>99%) an A and spread phenotype in culture with <1% non-adherent (NA) cells (Fig. 2C and D, respectively).

To assess the effects of CCK₂ receptor variant expression on NCM356 cell proliferation, cells were plated in multi-well plates and counted using a Coulter cell counter. The doubling time for vector-transduced NCM356 cells was 3.9 days (95% confidence intervals of 2.5, 6.2). In contrast, both CCK₂R- and CCK_{2i4sv}R-expressing cells grew faster, with a shorter doubling times of 2.5 days (2.2, 2.8); and 2.7 (2.4, 3.1), respectively (Fig. 2D). Comparison of the best fit curve of exponential growth between vector- versus either CCK₂R- or CCK_{2i4sv}R-expressing cell was statistically significant (p=0.0001). However, there were no differences in doubling times between each the two CCK₂ receptor variants (p=0.48; comparison of best-fit values).

YM022, a CCK₂ receptor-specific antagonist/inverse agonist, partially inhibits the proliferation of CCK₂R and CCK_{2i4sv}R cells *in vitro*—To determine whether the enhanced proliferation of receptor-expressing NCM356 cells was due to autocrine activation of the CCK₂ receptor variants by endogenous gastrin, the cells were treated at a 48-h interval beginning two days after plating either with vehicle (DMSO) or the CCK₂R antagonist/inverse agonist YM022^{22, 23} over a 9-day time course (Fig. 3). YM022 partially inhibited the proliferation of both CCK₂R- and CCK_{2i4sv}R-expressing cells when compared to cells treated with vehicle (Fig. 3A and B, respectively). Vector-transduced cells grew slower than receptor-expressing cells and were not inhibited by YM022 treatment (Fig. 3A and B). The p-values for best-fit linear regression growth curve between DMSO- versus YM022-treated were statistically significant for both CCK₂ receptor-expressing cell lines (p<0.001), but not for the vector-expressing cells (p=0.38). These data suggest that autocrine activation of the CCK₂ receptor signaling axis is, in part, responsible of the enhanced proliferation rate observed with receptor-transduced NCM356 cells. Additionally, YM022 inhibited the growth response of both CCK₂R- and CCK_{2i4sv}R-expressing cells to the same extent, suggesting that there was no significant constitutive activity associated with CCK_{2i4sv}R in this cell system. If CCK_{2i4sv}R had significant constitutive growth promoting effects, YM022 would have suppressed the growth of these cells to a greater extent when compared to the CCK₂R-expressing cells.

CCK₂ receptor expression is sufficient to transform NCM356 cells

Having established the presence of an *in vitro* autocrine growth loop, we next investigated whether acquisition of either CCK₂ receptor variant affected the NCM356 cell tumorigenic potential. Transduced cells were injected into the flanks of athymic nude mice (5×10⁶ cells per mouse, 6 replicates per condition). This experiment was performed three times and a representative experiment is shown in Fig. 4A. Neither parental NCM356 cells (data not shown) nor the vector-transduced cells formed tumors in athymic nude mice (Fig. 4B). However, both CCK₂R- and CCK_{2i4sv}R-expressing cells formed large tumor masses at the subcutaneous injection site (Fig. 4B). At sacrifice on day 33, no statistically significant differences (p=0.6, t-test) were noted between the mean weights of CCK₂R- (0.68 ± 0.05 g) and CCK_{2i4sv}R-expressing tumors (0.63 ± 0.09 g). Hematoxylin and eosin (H&E)-stained tissue sections from both CCK₂R- (Fig. 4C) and CCK_{2i4sv}R-expressing (Fig. 4D) tumors showed similar histopathology. The tumors appeared to be moderately differentiated adenocarcinoma, with clusters of tumor cells exhibiting marked nuclear atypia surrounded by scant intervening stromal tissue including murine blood vessels. Together, these data demonstrate that acquisition of either CCK₂ receptor variant is sufficient to transform the non-tumorigenic NCM356 cells into tumor-forming cells.

Basal phosphorylation of extracellular signal-regulated kinase (ERK)-1 and ERK2 are elevated in NCM356 cells expressing CCK₂ receptor variants

Extracellular signal-regulated kinases are members of the MAPK family that includes c-Jun N-terminal kinase (JNK), and p38^{MAPK}. The ERK subfamily is comprised of multiple isoforms, of which ERK1 and ERK2 are the most extensively characterized. Both are activated by phosphorylation of the threonine and tyrosine residues with the T-E/D-Y consensus motif by the dual-specificity kinases, MAP/ERK kinase (MEK)1 and/or MEK2. Previous studies have shown that CCK₂R and CCK_{2i4sv}R can couple gastrin stimulation to the activation (phosphorylation) of ERK1 and ERK2.^{24, 25}

To begin to elucidate potential signaling mechanisms for the enhanced proliferation and tumorigenicity of CCK₂ receptor variant-expressing NCM356 cells, we compared the basal phosphorylation state of ERK1 and ERK2 in protein extracts from CCK₂R-, CCK_{2i4sv}R- and vector-transduced NCM356 cells. Since the vector-transduced cells contain both A and

NA cells, we separately evaluated the levels of phosphorylated ERK1 and ERK2 (p-ERK1,2) in each subpopulation of cells. Compared to either NA or A vector-transduced cells, both CCK₂ receptor variants exhibited a marked increase in basal levels of activated ERK1 and ERK2 (Fig. 5A). No differences were observed between the basal levels of p-ERK1 and p-ERK2 in the NA and A subpopulations of vector-transduced cells (Fig. 5A). Densitometric analyses of immunoblot data from 6 independent experiments, after normalization using β actin as a protein loading control, demonstrated a significant difference in the basal p-ERK1 and p-ERK2 levels comparing vector-transduced cells to either CCK₂R- ($p=0.0006$) or CCK_{2i4sv}R-expressing lines ($p=0.005$) (Fig. 4B). Additionally, we evaluated whether the Src or PI3K/Akt pathways were activated by CCK₂ receptor expression. Western blots revealed that neither phospho-Src nor phospho-Akt levels increased in CCK₂R- and CCK_{2i4sv}R-expressing cells when compared to vector-transduced control cultures (data not shown).

The MEK inhibitor PD98059 and the PKC inhibitor GF109203X partially inhibit basal ERK phosphorylation in CCK₂R- and CCK_{2i4sv}R-expressing NCM356 cells

Todisco et al.²⁶ reported that gastrin-stimulation of CCK₂R expressed on the rat pancreatic acinar cell line AR42J activated ERK by both MEK- and PKC-dependent mechanisms. To further define the mechanisms involved in enhanced basal activation of ERK1 and ERK2 in NCM356 cells, we treated CCK₂R- and CCK_{2i4sv}R-expressing cells either with the MEK inhibitor PD98059 or the PKC inhibitor, GF109203X. Immunoblot analyses of protein extracts from vector-transduced as well as CCK₂R- and CCK_{2i4sv}R-expressing cells, showed a partial but dose-dependent inhibition of p-ERK1 and p-ERK2 in cells treated with PD98059 (Fig. 6A). Densitometric analyses from three independent experiments comparing p-ERK1 and p-ERK2 levels between vehicle- and PD98059-treated groups for CCK₂R- and CCK_{2i4sv}R-expressing cells demonstrated a $54 \pm 9.9\%$ ($p=0.006$) and $63 \pm 10\%$ ($p=0.003$) inhibition of ERK activity, respectively, when treated with 10 μ M PD98059 for 30 min (Fig. 6B). The basal p-ERK1 and p-ERK2 levels were also partially decreased when receptor-expressing cells were treated with the PKC inhibitor, GF109203X (5 μ M), for 30 min (Fig. 6C). Comparison of p-ERK1 and p-ERK2 levels between vehicle- and GF109203X-treated groups for CCK₂R- and CCK_{2i4sv}R-expressing cells demonstrated a $72 \pm 14\%$ ($p=0.036$) and $55 \pm 10\%$ ($p=0.035$) inhibition of ERK activity, respectively, when treated with 5 μ M GF109203X for 30 min ($n=3$) (Fig. 6B). Together these data indicate that receptor expression enhances basal ERK activation in NCM356 cells, in part, through a pathway involving GF109203X-sensitive PKC isozymes.

Inhibition of MEK and PKC slow the proliferation of CCK₂R- and CCK_{2i4sv}R-expressing NCM356 cells

To assess the involvement of the MEK/ERK pathway and GF109203X-sensitive PKC isozymes in receptor-stimulated NCM356 cell proliferation, 2 days after plating (20,000 cells/well), cells were treated at 48-h intervals either with vehicle (DMSO), PD98059 (10 μ M) or GF109203X (5 μ M) over an 8-day time course. Cell proliferation was determined by counting the total number of attached cells per well on day 6 and 8 using a Coulter Cell Counter. Both PD98059- and GF109203X-treatment inhibited the proliferation of receptor-transduced NCM356 cells. GF109203X was more effective inhibiting 86% and 88% of CCK₂R-associated (Fig. 6A) and 90% and 96% of CCK_{2i4sv}R-associated (Fig. 6B) cell proliferation at days 6 and 8, respectively. By comparison, PD98059 inhibited only 45% and 54% of CCK₂R-associated (Fig. 6A) and 41% and 47% of CCK_{2i4sv}R-associated cell proliferation (Fig. 6B) at days 6 and 8.

DISCUSSION

The role of gastrin peptides in colorectal cancer is complex and multifactorial. Experimental models using the hypergastremic APC^{min-/+} mice¹⁴ have demonstrated that both CCK₂ receptors and gastrin are upregulated in intestinal epithelial cells early in the development of colorectal carcinogenesis, indicating a positive role for endocrine gastrin. However, clinical data have shown that most adenomatous polyp specimens coexpress gastrin peptides and CCK₂ receptors in the absence of hypergastrinemia,⁶ suggesting that an autocrine mechanisms may be more important to tumor progression. Here we have provided evidence demonstrating neoplastic transformation of a human colorectal epithelial cell line by autocrine activation of two CCK₂ receptor variants. Additionally, we have shown that gastrin precursors such as Gly-G may also contribute to the initiation of colorectal tumorigenesis through calcium imaging of the RIE/CCK₂R cell line, a useful bioassay for the activation of the CCK₂ receptor. Although a 100-fold concentration of Gly-G (compared to the amidated G17) is required to generate an increase in [Ca²⁺]_i through the CCK₂ receptor, these incompletely processed forms of gastrin peptide are abundantly expressed in human colonic polyps. Smith et al.,⁵ using immunohistochemistry to identify the expression of gastrin peptides, reported that among 55 human polyps that were examined, the prevalence for progastrin, Gly-G, and amidated G17 was 91, 80, and 47%, respectively. Our study does not specifically examine, and does not preclude, the potential contributions of progastrin or other receptors to the precursor forms of gastrin. Most importantly, we demonstrate that expression of either CCK₂R or CCK_{2i4sv}R in the gastrin-producing, non-malignant, human colonic mucosa NCM356 cells, in distinct contrast to the vector-expressing cells, is sufficient to cause the cells to spread, adhere, and proliferate faster in cell culture, upregulate basal p-ERK 1,2 protein expression, and also, confer a tumorigenic phenotype in athymic nude mice.

Acquisition of the CCK₂ receptor has afforded the non-tumorigenic NCM356 cell line a means to transmit gastrin-mediated, growth-stimulatory signals and to activate the MEK/ERK pathway. Perhaps, as postulated by Hanahan and Weinberg,²⁷ the functional consequence of CCK₂ receptor overexpression into the context of the NCM356 cell, is to phenocopy the actions of a mutated *K-ras*, an early oncogene implicated in the progression of colorectal adenoma to carcinoma in about 40% of cases.¹ Thus, CCK₂ receptor activation, early in the development of an adenoma, in effect, may substitute for one of the multiple effector pathways downstream of *K-ras*, such as activation of the Raf/MEK/ERK pathway.²⁸ As a non-malignant polyp transitions to carcinoma, additional genetic mutations are gained, some of which may also sustain the MEK/ERK pathway, decreasing the selective pressure to maintain the gastrin-CCK₂ receptor autocrine loop for cellular proliferation and survival. Carcinomas maintain genetic mutations, while epigenetic regulation of other genes, perhaps genes with redundant functions, such as the expression of CCK₂ receptors, may become less sustainable, as a carcinoma progresses toward greater chromosomal instability. This theory is consistent with the finding that while the majority (81%) of adenomatous polyps examined express the CCK₂ receptor,⁵ colorectal carcinomas only express the receptor in 30-40% cases examined.^{29, 30}

We and others³¹⁻³⁵ have shown that CCK₂R and CCK_{2i4sv}R regulate intracellular pathways in an agonist-dependent manner involving MAPKs. Both MEK inhibitor PD98095 and PKC inhibitor GF109203X partially decreased the steady-state p-ERK levels to a similar extent in both the NCM356-CCK₂R and -CCK_{2i4sv}R cells, corresponding to the MEK-mediated partial decrease in the cell growth. Partial growth inhibition was also achieved by the specific antagonist for CCK₂ receptor, YM022, verifying that autocrine activation of the CCK₂ receptor in the transfected cells has a role in cell growth. The possibility that an endogenous receptor, not inhibited by YM022 and activated by precursor

gastrin peptides, also may be responsible for partial cell growth. The *in vitro* cell growth mediated by the CCK₂ receptor variants was almost entirely inhibited in a PKC-dependent manner, suggesting that the MEK/ERK pathway is only partially responsible for cell growth. Furthermore, the enhanced cell growth conferred by ectopic expression of either CCK₂R or CCK_{2i4sv}R, which can be almost fully attenuated with GF109203X, implicates additional growth mechanisms involving other PKC isozymes that do not utilize the MEK/ERK pathway. Future studies will delineate the specific PKC isozymes responsible for MEK/ERK-dependent and -independent mechanisms of G17-stimulated cell growth in CCK₂ receptor variant-expressing NCM356 cells.

In conclusion, our results support the concept that inhibition of CCK₂ receptor activation may be an important component in the prevention of adenoma to carcinoma progression in colorectal cancer. Indeed, G17-DT (Gastrimmune), an antibody therapy which neutralizes gastrin, was successful in preclinical murine studies, reducing the incidence of polyps and increasing survival.¹⁴ However, in a Phase I/II clinical trial of 50 patients with advanced colorectal cancer, Gastrimmune failed to reduce tumor burden in the later stages of disease, suggesting that Gastrimmune may be more relevant for use in prevention of high-risk patients. Additionally, several CCK₂R antagonists have been developed and used in human studies for diagnostic imaging of and therapeutic radiotherapy of receptor-expressing tumors (reviewed in Berna, et al.³⁷). In particular, one CCK₂ receptor antagonist, JB95008 (Gastrazole) has in two clinical trials for advanced pancreatic cancer, another tumor type that aberrantly overexpresses CCK₂ receptors.³⁸ Chau and colleagues³⁸ demonstrated a significant one-year survival benefit (log rank p=0.03) in a small trial of 18 patients. In a follow-up study of 98 patients with inoperable, advanced pancreatic cancer, however, the survival of patients treated with Gastrazole was equivalent to that of 5-fluorouracil, with a median survival of 3.6 and 4.2 months, respectively. These clinical trials were directed at late-stage colorectal and pancreatic patients, and therefore, it is not surprising that a survival benefit was not realized for either treatment. Our study supports the hypothesis that early targeting of the gastrin and the CCK₂ receptor autocrine loop, perhaps at the adenomatous polyp stage with CCK₂ receptor antagonists or antibodies, may be a successful therapeutic option for colorectal cancer prevention. Currently, chemoprevention with nonsteroidal anti-inflammatory drugs are very efficacious, but are not widely adopted for use due to risk of gastrointestinal bleeding or cardiovascular events.³⁹ In contrast, both the Gastrazole and Gastrimmune trials established acceptable toxicity profiles, which do not include cardiovascular side-effects. Blockade of CCK₂ receptors, therefore, may prove to be effective alternative chemopreventative agents.

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Abbreviations

(CCK₂)	cholecystokinin 2
(pERK)	phosphorylated ERK
(APC)	adenomatous polyposis coli
(PKC)	protein kinase C
(MAPK)	mitogen-activated protein kinase

(G17)	Gastrin 1-17
(MLV)	murine leukemia retrovirus
(MCS)	multiple cloning site
(LTR)	long terminal repeat
(DMSO)	dimethyl sulfoxide
(IRES)	internal ribosome entry site
(PCR)	polymerase chain reaction
(CM)	conditioned media
(H&E)	hematoxylin and eosin
(RIE)	rat intestinal epithelial
(JNK)	Jun N-terminal kinase
(Gly-G)	glycine-extended gastrin
(BK)	bradykinin
(FBS)	fetal bovine serum
(A)	adherent
(NA)	non-adherent

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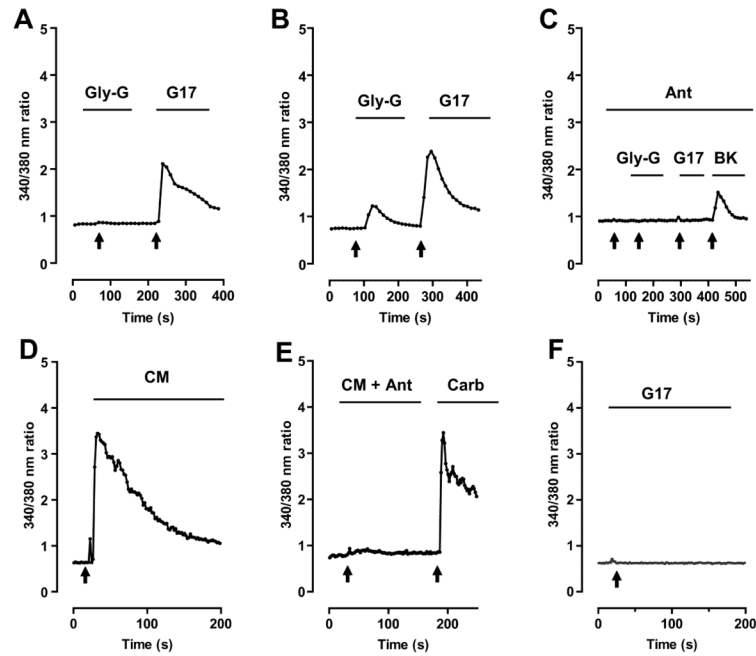


Figure 1. Secretion of functional gastrin-like peptide by non-tumorigenic human colonic epithelial cell line NCM356

Intracellular Ca^{2+} ($[\text{Ca}^{2+}]_i$) imaging of rat intestinal epithelial (RIE) cells expressing recombinant human CCK_2 receptor (RIE/ CCK_2R) loaded with the Ca^{2+} indicator dye Fura-2 was used as a bioassay to test the sensitivity, efficacy, and specificity of gastrin peptides. Black arrows indicate the time of addition of each treatment. Black bars indicate duration of exposure to each treatment. (A) Treatment of the RIE/ CCK_2R cells with 10 nM glycine-extended gastrin (Gly-G) failed to induce an increase in $[\text{Ca}^{2+}]_i$, whereas 1 nM of gastrin 1-17 (G17) to RIE/ CCK_2R cells does. (B) A 100-fold increase in concentration of Gly-G (100 nM) results in calcium response less robust than that of 1 nM of G17. (C) Pretreatment with CCK_2 receptor antagonist 100 nM of L365,260 abolishes the Ca^{2+} response by 100 nM of Gly-G and 1 nM of G17 but not 100 nM of bradykinin (BK). (D) Conditioned media (CM) from NCM356 cultures was applied to the RIE/ CCK_2R cells, inducing a transient increase in $[\text{Ca}^{2+}]_i$ in the cells of similar amplitude and duration as treating cells with 1 nM G17 (A,B). (E) Pretreatment with selective CCK_2 receptor antagonist 1 μM of JB93182 (JB) abrogates CM-induced Ca^{2+} response. Cells were treated with 10 μM carbachol (Carb) to insure that the cells were loaded with Fura-2. (F) Although NCM356 cells secrete gastrin, addition of exogenous G17 (100 nM) does not induce an increase in $[\text{Ca}^{2+}]_i$ indicating the absence of functional CCK_2 receptor.

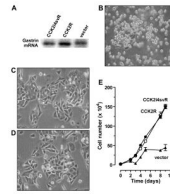


Figure 2. Phenotypic Effects of CCK₂ receptor variants expression on NCM356 cell growth *in vitro*

(A) All three transduced cell lines express gastrin mRNA by reverse transcriptase/polymerase chain reaction assay. (B) Photomicrograph of NCM356 cells stably transduced with the pFB-IRES retroviral expression vectors. Vector cells exhibited a small round, non-adherent (NA) phenotype indistinguishable from the parental line. In contrast, NCM356 cells transduced either with pFB-IRES containing recombinant human CCK₂R (C) or CCK_{2i4sv}R (D) exhibit a spread, adherent (A) phenotype with few floating cells. (E) Change in cell number in cell culture over time. NCM356-CCK₂R (open circle) or -CCK_{2i4sv}R (filled circle) grew faster compared to -vector control cells (filled triangle) in 10% FCS media conditions.

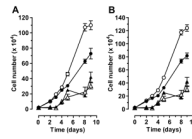


Figure 3. Effect of YM022 on cell growth in NCM356 cells expressing CCK₂ receptor variants
 Change in cell number in cell culture over time with application of either vehicle (DMSO) or YM022 (1 μ M) to the growth media. Receptor expressing cells were treated at a 48 h interval beginning on day 2 either with vehicle [DMSO (open circle)] or 1 μ M YM022 (filled circle) over a 9-day time course. Vector transduced NCM356 cells were also treated with either vehicle (open triangle) or 1 μ M YM022 (filled triangle) over the same time course. Data from the vector transduced cells are plotted on both graphs for comparison. **(A)** The growth curves for NCM356-CCK₂R compared to -vector cells. **(B)** The growth curves for NCM356-CCK_{2i4sv}R cells compared to -vector cells.

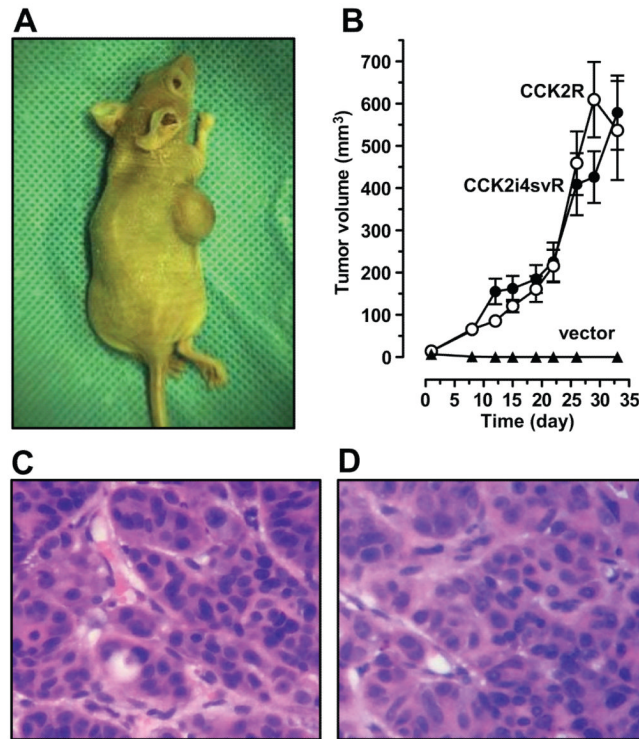


Figure 4. Phenotypic effects of CCK₂ receptor variants expression on NCM356 cell *in vivo*
 The athymic nude mouse model was used to assess whether acquisition of either CCK₂ receptor variant was sufficient to transform non-tumorigenic NCM356 cells into tumor forming cells. (A) Photograph of representative nude mouse with subcutaneous xenograft at harvest on day 33. (B) Change in tumor volume over time. NCM356 cells infected with the control vector (filled triangle) did not form tumors in nude mice; whereas, both -CCK₂R (open circle) and -CCK_{2i4sv}R cells (filled circle) formed subcutaneous tumors (~600 mm³) over a 33-day time course. The graph represents summary data from 6 mice /group. (C, D) Photomicrograph of formalin-fixed, hematoxylin-eosin stained section of NCM356-CCK₂R and -CCK_{2i4sv}R tumors, respectively (400X magnification). Tumors, at day 33, showed similar moderately-differentiated adenocarcinomas with marked nuclear atypia are surrounded by scant stroma.

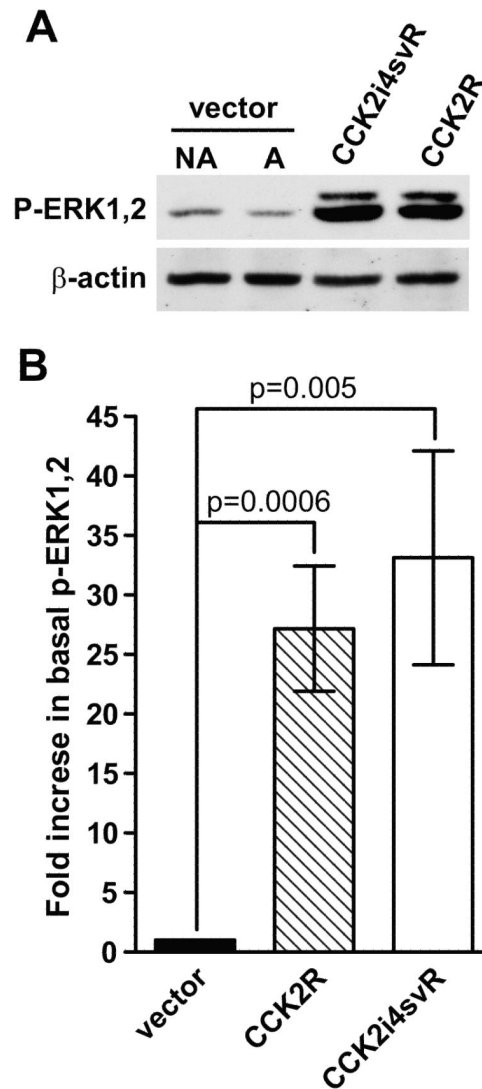


Figure 5. Effect of basal levels of phosphorylated (activated) ERK1 and ERK2 in NCM356 cells expressing CCK2 receptor variants

(A) Western blot comparing the basal levels of phospho-ERK1 and 2 (p-ERK1,2) levels in the non-adherent (NA) cell population NCM356-vector cells, the adherent (A) -vector cells, -CCK₂R pooled cells, and the -CCK_{2i4svR} pooled cells. Blots were reprobed with an antibody to β-actin to insure the equal loading and transfer of proteins in each lane. (B) Graph representing densitometric analyses of immunoblot data from 6 independent experiments comparing the fold change in basal p-ERK1,2 levels between NCM 356-vector, -CCK₂R and -CCK_{2i4svR} transduced cell lines. Statistical analysis using the t-test demonstrated significance between the basal p-ERK1,2 levels from vector and NCM356-CCK₂R (p=0.0006) and between vector and -CCK_{2i4svR} cells (p=0.005).

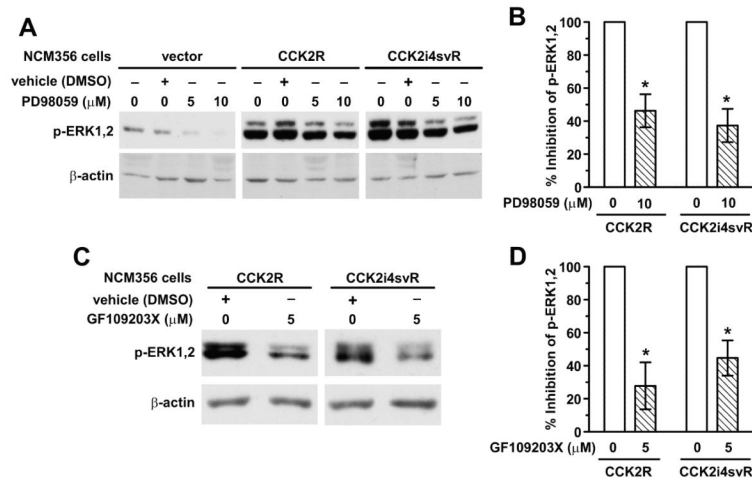


Figure 6. Effects of the MEK inhibitor PD98059 and the PKC inhibitor GF109203X on basal p-ERK1,2 levels

(A) Western blot of basal pERK levels in NCM356-vector, -CCK₂R, or -CCK_{2i4svR} expressing cells in the presence of vehicle (DMSO) or MEK inhibitor PD98059 (5 or 10 μM) for 30 min. Blots were reprobated with an antibody to β-actin to insure the equal loading and transfer of proteins in each lane. (B) Graph representing densitometric analyses of immunoblot data from 3 independent experiments comparing the % inhibition of basal p-ERK1,2 levels between treatment with vehicle or 10 μM PD98059 on NCM 356 -CCK₂R and -CCK_{2i4svR} transduced cell lines. T-test demonstrates statistical significance comparing the % inhibition of basal p-ERK1,2 levels between vehicle-treated and PD98059-treated NCM356-CCK₂R (p=0.006) and -CCK_{2i4svR} cells (p=0.003). (C) Western blot of basal pERK levels in NCM356-vector, -CCK₂R, or -CCK_{2i4svR} expressing cells in the presence of vehicle (DMSO) or PKC inhibitor GF109203X (5 μM) for 30 min. An antibody to β-actin was used to insure the equal loading and transfer of proteins in each lane. (D) Graphical representation of immunoblot data from 2 independent experiments comparing the % inhibition of basal p-ERK1,2 levels between treatment with vehicle or 5 μM GF109203X on NCM 356 -CCK₂R and -CCK_{2i4svR} transduced cell lines. Statistically significant difference between basal p-ERK1,2 levels of vehicle-treated compared to inhibitor-treated cells were noted in the NCM356-CCK₂R (p=0.036) and -CCK_{2i4svR} cells (p=0.035) using t-test analyses.

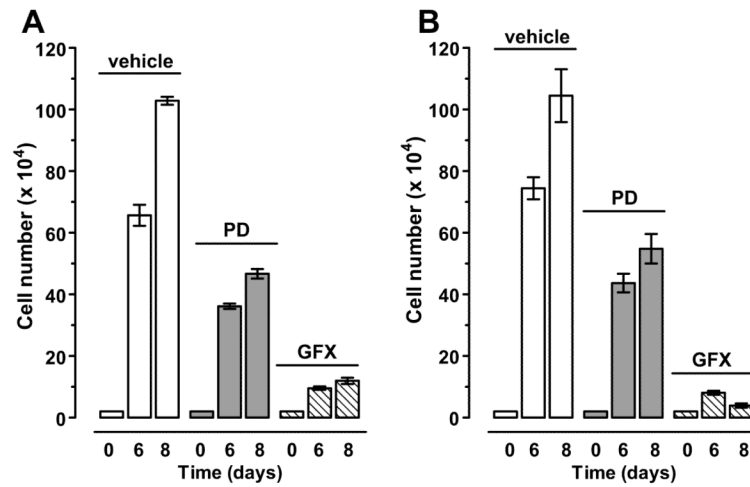


Figure 7. Effects of the MEK inhibitor PD98059 (PD) and the PKC inhibitor GF109203X (GFX) on the proliferation of NCM356-CCK₂R (A) and -CCK_{2i4sv}R (B) cells

Bar graphs comparing the growth of cells in 3 different conditions are shown. 2×10^5 cells were plated in triplicate on day 0 and starting on day 2, treated every 48 h with inhibitor. The cells were harvested and counted by Coulter Counter on days 6 and 8: vehicle-treated media (open bars), addition of PD98059 (5 μ M) (bars filled with dots), and GF109302X (5 μ M) (bars filled with slashes). Data represent the average cell numbers on days 6 and 8 \pm S.E.M for 3 separate experiments.