

Exendin-4 inhibits the survival and invasiveness of two colorectal cancer cell lines via suppressing GS3K β / β -catenin/NF- κ B axis through activating SIRT1

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Abstract: This study examined if the anti-tumorigenesis effect of Exendin-4 in HT29 and HCT116 colorectal cancer (CRC) involves modulation of SIRT1 and Akt/GSR3K/ β -catenin/NF- κ B axis. HT29 and HCT116 cells were treated either with increasing levels of Exendin-4 (0.0-200 μ M) or with Exendin-4 (at its IC₅₀) in the presence or absence of EX-527 (10 μ M/a selective SIRT1 inhibitor) or Exendin-4 (9-39) amide (E (9-39) A) (1 μ M/an Exendin-4 antagonist). In a dose-dependent manner, Exendin-4 inhibited cell survival, but enhanced levels of lactate dehydrogenase (LDH) and single-stranded DNA (ssDNA) in both HT29 and HCT116. In both cell lines and at its IC₅₀ (45 μ M for HT29 and 35 μ M for HCT116), Exendin-4 also significantly reduced cell survival, migration, and invasion of both cell types, with no effect on the expression of GLP-1 receptors (GLPRs) nor of the activity of Akt. At these doses, Exendin-4 also increased the expression of SIRT1 but reduced the acetylation of NF- κ B and the expression of Bax and cleaved caspase-3 and in both cell lines. Concomitantly, protein levels of p-GS3K β (Ser⁹), total and acetylated β -catenin, and Anix2 were significantly decreased, but levels of p-GS3K β (Ser⁹) and p- β -catenin (Ser^{33/37}/Thr⁴¹) were significantly increased in both HT29 and HCT116-exendin-4 treated cells. All the effects exerted by Exendin-4 were completely prevented by Ex527 or E (9-39) A. In conclusion, Exendin-4 suppresses the tumorigenesis of HT29 and HCT116 CRC cell activation of GS3K β -induced inhibition of β -catenin and NF- κ B in a SIRT1-dependent mechanism.

Introduction

Colorectal cancer (CRC) is one of the most common solid tumors among male and female patients, which develops due to environmental, nutritional, and genetic factors (Thanikachalam and Khan, 2019). Despite the large improvement in the screening tools and therapeutic options, the mortality rates among the affected patients

remained significantly high and are expected to increase to 60% by 2030, with a projection that 90% of those patients will be among the young whose age is between 24–34 years (Siegel *et al.*, 2019).

The highly conserved canonical Wnt/ β -catenin signaling pathway is a leading pathway that mainly involved in embryonic development and adult homeostasis, including wound healing, maintenance and renewal of stem cell, and cell proliferation, survival, and motility (El-Sahli *et al.*, 2019; Logan and Nusse, 2004). Under normal conditions, the cytoplasmic levels of β -catenin, a transcription co-activator, remain very low due

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to their rapid ubiquitination and proteasome degradation as they are always phosphorylated by the β -catenin destruction complex (Krishnamurthy and Kurzrock, 2018). This complex is composed of several proteins, including axin, casein kinase I (CKI), adenoma polyposis coli (APC), and glycogen-synthase kinase-3 β (GS3K β), where the activation of the GS3K β is indispensable for β -catenin-induced degradation (Metcalfe and Bienz, 2011). However, the unphosphorylated β -catenin will escape this degradation and accumulates in the cytoplasm and nucleus to trigger the transcriptional program (Metcalfe and Bienz, 2011). Within this view, the binding of the Wnt protein to its specific transmembrane receptor (Fzd and LRP) triggers the recruitment of Dishevelled proteins, which leads to several events which end up with phosphorylation (inhibition) and/or internalization of GS3K β , thus blocking the GS3K β -induced β -catenin degradation (Metcalfe and Bienz, 2011; Rajkumar, 2016).

Currently, accumulating evidence is showing that sustained activation of Wnt/ β -catenin pathway underlies multiple hematological and solid tumors, including chronic myelogenous leukemia (CML), CRC, and breast, ovaries, prostate, stomach, liver, lung, and thyroid cancers (Beurel and Jope, 2006; El-Sahli et al., 2019; Polakis, 2012; Taciak et al., 2018). Accordingly, the aberrant activation of β -catenin triggers the transcription of numerous genes involved tumor proliferation (e.g., Cyc D1, and Myc), differentiation (e.g., SOX9), stemness (e.g., Nanog and OCT4), and migration (CXCR4, CCL18, ICAM-1, and VCAM-1) (El-Sahli et al., 2019; Polakis, 2012; Rajkumar, 2016). Besides, β -catenin stimulates tumor growth and development by stimulating cancer metabolism, including glycolysis, glutaminolysis, and lipogenesis (El-Sahli et al., 2019). Furthermore, β -catenin can stimulate the tumor inflammatory microenvironment, which plays a significant role in the tumor cells, proliferation, invasiveness, and resistance to drugs by increasing the transcription of TNFSF9 and the nuclear factor- κ B (NF- κ B) (Hanahan and Weinberg, 2011; Polakis, 2012; Rajkumar, 2016).

However, the process of β -catenin activation seems to be very complicated and involved not Wnt1-dependent mediators. Although it is a major mechanism to induce cancer, inflammation, the components of the NF- κ B signaling pathways, including IKK α , RelA (p65), p50, and other inflammatory cytokines, such as IL-6, can upregulate β -catenin transcriptional activity (Albanese et al., 2003; Carayol and Wang, 2006; Cho et al., 2008; Li et al., 2014; Rajkumar, 2016; Schwitalla et al., 2013; Yang et al., 2012). Besides, β -catenin activity is negatively regulated by deacetylation by the silent information regulator 1 (SIRT1) (Cho et al., 2012; Lin and Fang, 2013; Salaroli et al., 2008). However, the contribution of these pathways in β -catenin activation in the different tumor types is still not completely understood.

On the other hand, accumulating line of evidence is currently showing that genetic deletion and silencing or pharmacological inhibition of β -catenin is considered a highly valuable clinical therapeutic option to prevent and slow the progression of CRC (Cong et al., 2003; Green et al., 2001; Mogni et al., 2010; Morin et al., 1996; Polakis, 2012; Tetsu and McCormick, 1999; van de Wetering et al., 2002; Verma et al., 2003). The glucagon-like peptide-1 analog,

Exendin-4, has received much attention during the last years as a therapeutic and anti-tumor drug due to its potent antioxidant and anti-inflammatory potentials, *in vivo* and *in vitro* (Defronzo et al., 2005; Lonborg et al., 2014; Yamamoto et al., 2013). Indeed, Exendin-4 anti-tumor effects were demonstrated against breast, prostate, colon, and ovarian cancers through suppressing inflammation and cell survival by modulating several cellular pathways, including NF- κ B, PI3K/Akt, AMPK, and MAPKs (Iwaya et al., 2017; Nomiya et al., 2014; Tsutsumi et al., 2015; Tzotzas et al., 2017). Of note, Exendin-4 also induced cell apoptosis and decreased colony formation CT26 colon cancer cells by increasing intracellular levels of cAMP and inhibiting GS3K β and ERK1/2 signaling (Koehler et al., 2011).

Given the complexity of β -catenin activation and its role in CRC, in this study, we hypothesized that Exendin-4 may induce cell death and inhibit cell proliferation and invasiveness of two colorectal cell lines by attenuating the aberrant β -catenin activation through affecting the expression/activities of Akt, GS3K β , NF- κ B, and SIRT1.

Materials and Methods

Cell culture

The Human HT29 & HCT116 CRC cell lines were supplied from the American Type Culture (USA) and always cultured in a humidified atmosphere (5% CO₂/37°C) in DMEM medium (ThermoFisher Scientific) supplemented with 10% heat-inactivated FBS, 1% penicillin, and 0.1% streptomycin (ThermoFisher Scientific). In all the parts of this study, the cells were always used when they reveal a confluent density of <85%. Accordingly, both cell lines 5 × 10⁴ cells (150 μ L cell suspension in the media) were seeded in 96-well tissue culture plates and incubated under standard culture conditions (5% CO₂/37°C) for 24 h. each part of any experiment performed in this study was conducted in three experiments each done in triplicate.

Cell treatment

Cells were treated with increasing levels of Exendin-4 (10, 25, 50, 100, and 200 μ M) (Cat. No. E7144, Sigma Aldrich, MA, USA) for 24 h at 37°C and 5% CO₂. Control cells were treated only with 0.05% DMSO (dimethyl sulfoxide) (Cat. No. 276855, Sigma Aldrich, MA, USA). Cell survival, medium levels of lactate dehydrogenase (LDH), and single-stranded DNA (ssDNA) were measured as described below. Based on these data, the inhibitory concentration (IC₅₀) of Exendin-4 in both cell lines was calculated from the survival data and then used for the rest of the experiments. In these experiments, the cells were seeded in the same manner but were incubated with the IC₅₀ of Exendin-4 (45 μ M for HT29 and 35 μ M for HCT116). In some cases, cells of both cell lines were also pre-incubated for 1 h with 10 μ M EX-527 (a SIRT1 inhibitor (Cat. No. E7144, Sigma Aldrich, MA, USA) or 1 μ M Exendin (9-39) amide (a GLP1 receptor antagonist (Cat. No. ab141101, Abcam, Cambridge, UK). The doses of EX-527 and Exendin (9-39) amide were pre-determined in our preliminary data, where they completely inhibited SIRT1 activity and cAMP generation, respectively, by more than 90%. All drugs were always

prepared in DMSO and diluted with PBS (pH = 7.4). In all experiments, the DMSO final concentration was always 0.05%.

Determination of cell viability

The viability of cells of all treatments was determined using a colorimetric kit (cell counting kit-8/CCK-8) (Cat No. CK04-13, Dojindo, Japan). Briefly, part of the culture medium (100 μ L) was discarded and replaced with an equal volume of fresh medium containing 10% of CCK8 solution. The plate containing the cultured cells was incubated at 37°C for 3 h, and the absorbance was read at 450 nm. Cell viability was calculated as a percent using the following formula ($[\text{Abs of the sample} - \text{Abs of the blank}] / [\text{Abs of the control} - \text{Abs of the blank}]$). The assay followed the manufacturer's instructions.

Determination of LDH activity in the supernatants

The cells of all treatments were centrifuged for 5 min at 200 x g at 4°C. The pellets were discarded, and the resulting supernatants were collected for analysis of LDH using an assay kit (Cat. No. ab102526; Abcam, Cambridge, UK), which is based on the LDH-dependent reduction of NAD⁺ to NADPH which can be detected using a special probe at 450 nm. In brief, 50 μ L of the prepared reaction mixture was added to 20 μ L the pre-adjusted, tested sample, and the color was read immediately at 450 nm. The activity of LDH was calculated from a standard curve.

Determination of apoptosis (ssDNA assay)

The levels of cell apoptosis in all treated groups were calculated using the ssDNA assay ELISA kit (Cat No. APT225, Millipore, USA). All steps followed the manufacturer's instructions. In the beginning, cells were pelleted (200 x g, 5 min, 4°C) and fixed for 30 min at room temperature with 200 μ L of the provided fixative. Then, the fixative was removed followed by the addition of 100 μ L formamide solution. The plate was incubated at room temperature for 10 min and then heated in an oven for another 10 min at 75°C to denature DNA. After that, the plate was allowed to cool in the fridge, and the formamide solution was discarded. After that, all wells were blocked by the addition of 3% skim milk (prepared in distilled water) and incubated at 37°C for another 1 h. Then, the milk was discarded, and the cells of all wells were incubated with 100 μ L of the provided mouse monoclonal anti-ssDNA antibody. The plate was incubated again at room temperature for an extra 30 min. All wells were then washed (3X/2 min) with 250 μ L of the provided washing. Following, the membranes were incubated at room temperature with the and HRP-secondary anti-mouse secondary antibody (200 μ L) for 30 min followed by the addition of 100 μ L ABTS solution. Finally, 100 μ L of the supplied stop solution was added to each well and the absorbance of all wells was read at 405 nm.

Migration and invasion assays

Rates of cell migration and invasiveness of both HT29 and HCT1165 were evaluated with or without Exendin-4 treatment as described in our laboratories previously (El-Kott *et al.*, 2019). To evaluate the cell migration, 5×10^4 cells (100 μ L) and 200 μ L serum-free medium (SFM) were plated on the top of the 8- μ m pore size Corning Transwell chambers (diameter of 6.5 mm) (USA) and either treated Exendin-4 (45 μ M for HT29 and 35 μ M for HCT1165) in the presence

or absence of the inhibitors (i.e., 10 μ M EX-527 or 1 μ M Exendin (9-39) amide). Control cells were incubated with 0.05% DMSO. In all treatments, 500 μ L 10% FBS was added to the bottom of the chamber. The cells were incubated in this setting for 24 h. Then, the cells presented in the lower chamber were fixed at room temperature with methanol for 10 min. To visualize the cells, they were stained with 0.2% crystal violet and left for 10 min at 25°C and then counted under 10X of a light microscope. A similar procedure, but with 5-mm pore size Corning Transwell chambers that is percolated with 50 Matrigel μ L (Cat. No. DLW354263, Sigma Aldrich, USA) was used for the invasion test. However, cells first starved in the SFM for 3 h and then loaded onto the upper chamber. Cell count was done as described in the migration test.

Cell fractionation and homogenates preparation

The cytoplasmic/nuclear fractions of all treated cells were prepared using a commercially available isolation kit (Cat No. 78835 and ThermoFisher Scientific). To prepare total cell homogenates for the western blotting protocol, the cell pellets were homogenized in radioimmunoprecipitation (RIPA) buffer (250 μ L), centrifuged at 11000 x g for 10 min at 4°C to isolate the supernatants. Protein levels in the cytoplasmic, nuclear, and total fractions were evaluated using a Bradford assay (Cat. No. 23300, ThermoFisher Scientific, MA, USA). The levels of target protein expression were calculated relative to the expression of β -actin.

Determination of SIRT1 and NF- κ B p65 in the nuclear fractions
SIRT1 deacetylase activity and the nuclear activity of NF- κ B p65 were determined using assay kits (Cat. No. CY-1151V, Nagano, Japan and TransAM, Cat. No. 40596, Active Motif, Tokyo, Japan) as per the manufacturers' instructions.

Immunoprecipitation (IP)

This procedure was done to detect the acetylated protein levels of both β -catenin and NF- κ B p65 by western blotting (Cat. No. 206996, Abcam, Cambridge, UK). In brief, 4 μ g of the antibodies against β -catenin or NF- κ B p65 or normal rabbit IgG antibodies were added to each sample (containing 100 μ g proteins), and the total volume was adjusted to 500 μ L using the lysis buffer containing the protease inhibitor cocktail. The whole mixture was kept in rotation for 4 h at 4°C. Four different samples, then 30 μ L of protein A/G Sepharose beads were added to every reaction and incubated, and the mixture was further incubated for 1 h at 4°C. All reactions were then centrifuged (2000 x g, 3 min) to collect the protein A/G Sepharose beads. These were then washed (3x) with the provided washing buffer. The beads were allowed to dry at room temperature for 20 min and then were loaded with 40 μ L 2x Laemmli buffer, boiled for 5 min and then stored at -80°C for western blotting protocol.

Western blotting analysis

Western blotting was conducted in accordance with the methods established by others (Shan *et al.*, 2020; Su *et al.*, 2020). Equal protein concentrations of the all-cell fractions (40 μ g/well) were separated using an SDS-polyacrylamide gel electrophoresis (PAGE) for 1 h at 100 V. The bands in every gel were then transferred onto nitrocellulose membranes

for another 1.5 h at 80 V. Membranes were then blocked by 5% skimmed milk prepared in Tris-buffered saline-tween-20 (TBST buffer) and blotted using each target antibody (Tab. 1) (prepared in TBST buffer) or with acetylated lysine antibody (Cat. No. 9441, 1:500) for 2 h at room temperature. Membranes were then washed with the TBST buffer (3X/10 min) and then incubated for 1.5 h at room temperature with the HRP-2nd antibodies. After three times washing with the TBST buffer, the bands were developed using a Pierce ECL kit (ThermoFisher, USA) and scanned and analyzed using the C-Di Git blot scanner (LI-COR, NE, USA).

Statistical analysis

Data were analyzed on Graph Pad Prism statistical software package (version Differences using the one-way ANOVA and Tukey's test as *post hoc*. The level of significance was considered at $P < 0.05$.

Results

Exendin-4 inhibits cell survival and induces cell death in both HT29 and HCT116 CRC cell lines in a dose-dependent manner
As shown in the Suppl. Figs. S1A–S1D and S2A–S2D, Exendin-4 significantly and in a dose-dependent manner (10–200 μ M) reduced cell survival and increased the release of LDH and levels of ssDNA in both HT29 and HCT119 CRC cell lines as compared to their controls treated with diluted DMSO. The calculated IC₅₀ levels of Exendin-4 in HT29 and HCT116 CRC cells were 44.19 μ M and 35.45 μ M, respectively (Suppl. Figs. S1A–S1D). These doses were used in the rest of the experiments.

Exendin-4 induces cell death and inhibits cell survival, migration, and invasions in both HT29 and HCT116 CRC cell lines in SIRT1 and GLP-1R-dependent mechanism

In the same line, treating the CRC cells with Exendin-4 at its IC₅₀ (45 μ M was for HT29 and 35 μ M for HCT116)

significantly reduced cell survival, invasion, and migration and increased the release of LDH and levels of ssDNA in both cell lines as compared to DMSO treated cells (Figs. 1A–1F and 2A–2D). However, these effects were completely reversed and restored to their basal control levels in both cell lines when Exendin-4 treated cells were pre-treated with Ex-527, a selective SIRT1 inhibitor, or with Exendin (9-39) amide, a GLP-1R antagonist (Figs. 1A–1F and 2A–2D).

Exendin-4 has no effects on the expression of GLP-1Rs but significantly stimulates the activity and protein levels of SIRT1 in both HT29 and HCT116 CRC cell lines

The protein levels of GLP-1Rs remained significantly unaltered in both CRC cell types of any treatment (Figs. 3A–3B). However, the nuclear activity of SIRT1 (Suppl. Fig. S3), as well as total protein levels of SIRT1 (Figs. 3C–3D), was significantly increased in Exendin-4-treated cells of both HT29 and HCT116 cells as compared to their corresponding control cells treated with DMSO. However, as compared to Exendin-4 treated cells, the activity and protein levels of SIRT1 were significantly reduced to their normal levels in Exendin-4-treated HT29, and HCT116 cells were pre-incubated with either Ex-527 or Exendin (9-39) amide (Suppl. Figs. S3A–S3B and Figs. 3C–3D).

Exendin-4 enhances the activity of GS3K β , independent of Akt, in both HT29 and HCT116 CRC cell lines in a SIRT1 and GLP-1Rs-dependent mechanism

As depicted in Figs. 4A and 4B, there was no significant variation in total I protein levels of Akt, as well in the levels of p-Akt (Thr³⁰⁸) between all study groups of both HT29 and HCT116 CRC cell lines as compared to DMSO-treated cells. These data indicate no effect of Exendin-4 on Akt activity in both cell lines. However, the catalytic activity of GS3K β is inhibited by phosphorylation at Ser⁹, whereas its apoptotic activity is increased by Tyr²¹⁶ phosphorylation (Jain *et al.*, 2017). Herein, the protein level p-GS3K (Ser⁹) were

TABLE 1

Primary antibodies used in the western blotting protocol

Target	Cat. No./Supplier	kDa
GLP-1Rs	sc-390774/Santacruz biotechnology	56
SIRT1	2496/Cell Signalling Technology	120
Bax	2772/Cell Signalling Technology	20
cleaved caspase-3	9661/Cell Signalling Technology	17/19
Ak	9272/Cell Signalling Technology	60
p-Akt (Thr308)	3038/Cell Signalling Technology	60
GS3K β	9315/Cell Signalling Technology	46
p-GS3K β (Ser ⁹)	9322/Cell Signalling Technology	46
p-GS3K β (Tyr ²¹⁶)	b75741/Abcam	47
β -catenin	8480/Cell Signalling Technology	92
p- β -catenin (Ser ^{33/37} /Thr ⁴¹)	9561 Cell Signalling Technology	92
p-NF- κ B (Ser ⁴⁶⁸)	3039/Cell Signalling Technology	65
Axin	2151/Cell Signalling Technology	95
β -actin	3700/Cell Signalling Technology	45

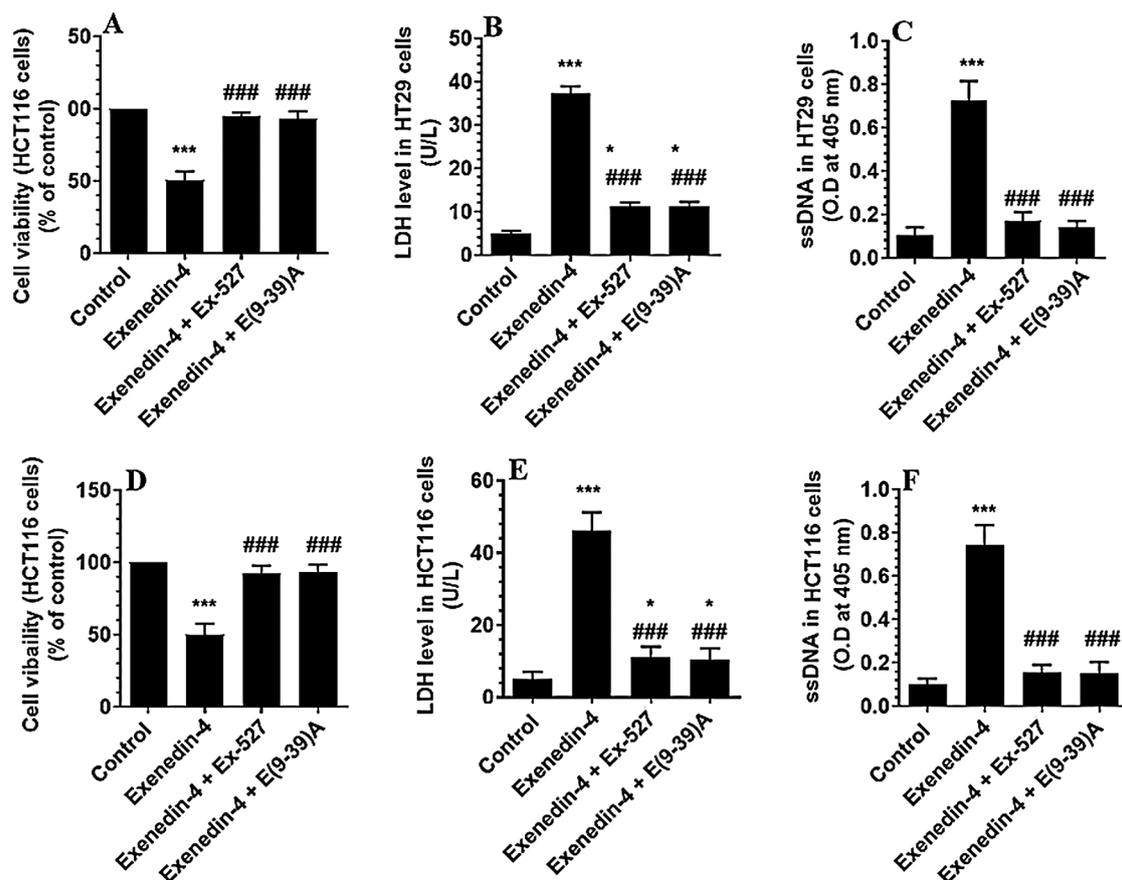


FIGURE 1. Cell viability and levels of Lactate Dehydrogenase (LDH) and single-stranded DNA (ssDNA) in HT29 (A–C, respectively) and HCT116 (D–F, respectively) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exendin-4 at its IC_{50} (45 μ M for HT29 and 35 μ M for HCT116) for 24 h (37°C and 5% CO_2) with or without 1 h pre-incubation with EX-527 (10 μ M), A selective SIRT1 inhibitor or Exendin (9-39) amide (E (9-39) A) (1 μ M), a GLP-1 antagonist. ***, ***: vs. control (DMSO-treated) at $P < 0.05$ and $P < 0.001$, respectively. ###: vs. Exendin-4-treated cells at $P < 0.001$.

significantly decreased, whereas protein levels of p-GS3K β (Tyr²¹⁶) were significantly increased in Exendin-4-treated HT29 and HCT116 cells as compared to their control groups (Figs. 5A–5D), thus suggesting activation of the catalytic and apoptotic activity of GS3K β . Also, levels of Bax and cleaved caspase-3 were significantly increased in both cell lines which were treated with Exendin-4 as compared to their control untreated cells (Figs. 6A–6D). On the contrary, all these effects afforded by Exendin-4 on the phosphorylation of GS3K β and total protein levels of Bax and cleaved caspase-3 were completely reversed in both Exendin-4-treated HT29 and HCT116 cells when they were pre-incubated with Ex-527 or Exendin (9-39) amide (Figs. 5A–5D and 6A–6D).

Exendin-4 suppressed total protein level of β -catenin, Acetylated β -catenin, and p- β -catenin (Ser^{33/37}/Thr⁴¹) in both HT29 and HCT116 CRC cell lines in a SIRT1 and GLP-1Rs-dependent mechanism

GS3K β phosphorylates β -catenin at Ser^{33/37} and Thr⁴¹ to enhance its ubiquitination and proteasome degradation (Rajkumar, 2016). Also, SIRT1 inhibits β -catenin activity by deacetylation at Lys³⁴⁵ (Levy et al., 2004; Wolf et al., 2002). Associated with the increased activity of GS3K β and SIRT1 and levels of SIRT, Exendin-4 significantly decreased total and acetylated protein levels of β -catenin and increased

protein levels of p- β -catenin (Ser^{33/37}/Thr⁴¹) in both cell lines as compared to control cells treated with DMSO (Figs. 7A–7D). Also, it reduced levels of Axin-2, a major downstream target of β -catenin in both HT29 and HCT116 CRC cells (Figs. 8A and 8B). These data suggest that Exendin-4 inhibits β -catenin signaling in both HT29 and HCT116 CRC cells. On the other hand, total and acetylated protein levels of β -catenin, as well as levels of p- β -catenin (Ser^{33/37}/Thr⁴¹), were not significantly different when Exendin-4 + Ex-527 and Exendin-4 + Exendin (9-39) amide-treated cells of both cell lines were compared to their control groups (Figs. 7A–7D and 8A–8B), thus suggesting that the suppressive effect of Exendin-4 on β -catenin activation is SIRT1 and GLP-1R-dependent.

Exendin-4 inhibits the activation of NF- κ B in both HT29 and HCT116 CRC cell lines in a SIRT1 and GLP-1Rs-dependent mechanism

The activity of NF- κ B p65 is negatively controlled by GS3K β (phosphorylation at Ser⁴⁶⁸) and by deacetylation by SIRT1 (Buss et al., 2004; Yang et al., 2012). The activity of NF- κ B was significantly decreased in both cell lines which were treated with Exendin4 as compared to their control cells (Suppl. Figs. S3C–S3D). In addition, the protein levels of p-NF κ B p65 (Ser⁴⁶⁸) were significantly increased, and its acetylated levels were significantly decreased in both HT29

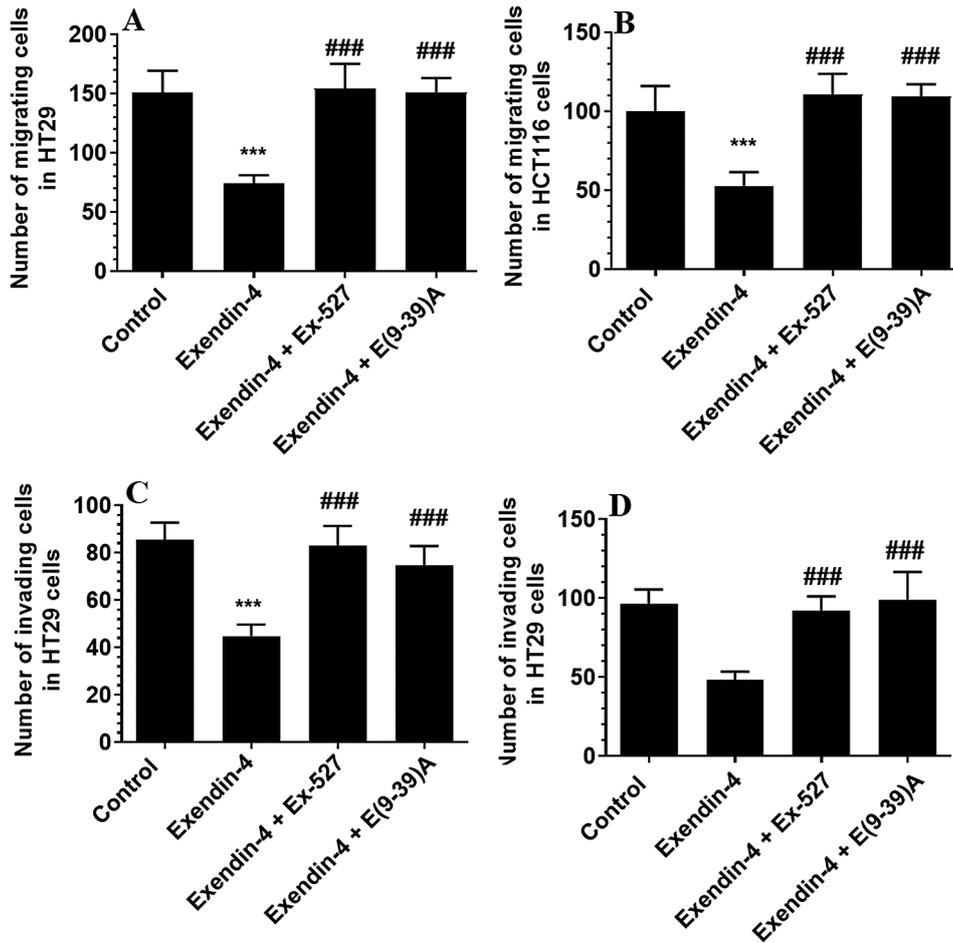


FIGURE 2. Number of migrating (A–B) and invading cells (C–D) in HT29 and HCT116 colorectal cells, respectively) and their calculated half inhibitory concentration (IC_{50}) (B–D, respectively). The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exendin-4 at its IC_{50} (45 μ M for HT29 and 35 μ M for HCT116) for 24 h (37°C and 5% CO_2) with or without 1 h pre-incubation with EX-527 (10 μ M), A selective SIRT1 inhibitor or Exendin (9-39) amide (E(9-39)A) (1 μ M), a GLP-1 antagonist. Data are presented as mean \pm SD of three independent experiments, each performed in triplicate. ***: vs. control (DMSO-treated) at $P < 0.001$. ###: vs. Exendin-4-treated cells at $P < 0.001$.

and HCT116 cells treated with Exendin-4 (Figs. 9A–9B). However, there was no significant variation in total protein levels in all groups of treatments for both cell lines (Figs. 9A–9B). All these effects were completely prevented in both cells, which were treated with Exendin-4 but pre-treated with Ex-527 or E (9-39) A (Suppl. Figs. S3C–S3D; Figs. 9A–9D).

Discussion

The salient findings of this study show that Exendin-4 inhibited the survival and invasiveness of two colorectal cancer (CRC) cell lines, HT29 and HCT116, by suppressing β -catenin activation. These effects were associated with increasing the activity GS3K β , inhibiting NF- κ B activity, and upregulating and activating SIRT1. Further investigations have revealed that the anti-tumorigenesis effect of Exendin-4 is GLP-1 and SIRT1-dependent as pre-incubating the cells with Exendin-4 (9-39) amid, a GLP-1 antagonist or EX-527, a selective SIRT1 inhibitor, completely prevented all the effects afforded by Exendin-4. A summary of the protective effect of these pathways is shown in graphical abstract.

Although the tumor suppressor and the inhibitory effect of Exendin-4 have been confirmed in many solid tumors, few studies have examined its role in a CRC, and contradictory results exist. Koehler *et al.* (2011) have shown that Exendin-4 induced cell apoptosis and reduced colony formation in

murine CT26 cells line by increasing intracellular levels of cyclic adenosine monophosphate (cAMP) and inhibiting GS3K β and ERK1/2 signaling (Koehler *et al.*, 2011). On the opposite, He *et al.* (2017) have shown the Exendin-4 does not modify the growth and survival of human colon cancer cells, including Colo320, Caco2, SW480, and LOVO. This variation was attributed to the presence or absence of GLP-1Rs between these cells where the expression of GLP-1Rs was abundant in the CT26 cell lines but absent in all the above-mentioned tested human cell lines. In our preliminary data, we first tested the expression of GLP-1Rs in several cell lines, including HT29 and HCT116, DLD-1, SW480, CaCo2, and CT26, using real-time PCR and western blotting techniques with a very specific monoclonal antibody. Among all, we have found that GLP-1Rs are abundantly expressed in HT29 and HCT116 and to a less extent in CT26 cells, but completely absent in all other tested cells (data not shown except for HT29 and HCT116). Hence, we have decided to continue our experiments using HT29 and HCT116 cell lines.

While the lack of GLP-1Rs on SW480 and CaCo2 supports the data presented by He *et al.* (2017), our findings contradict their results from the perspective of GLP-1Rs expression in the HT29 cells. This variation could be explained by the age of the cells used, freezing and thawing, culturing conditions, and the specificity of the antibody used. Indeed, several commercial antibodies are not specific to GLP-1Rs (Ussher and Drucker, 2014). While we have

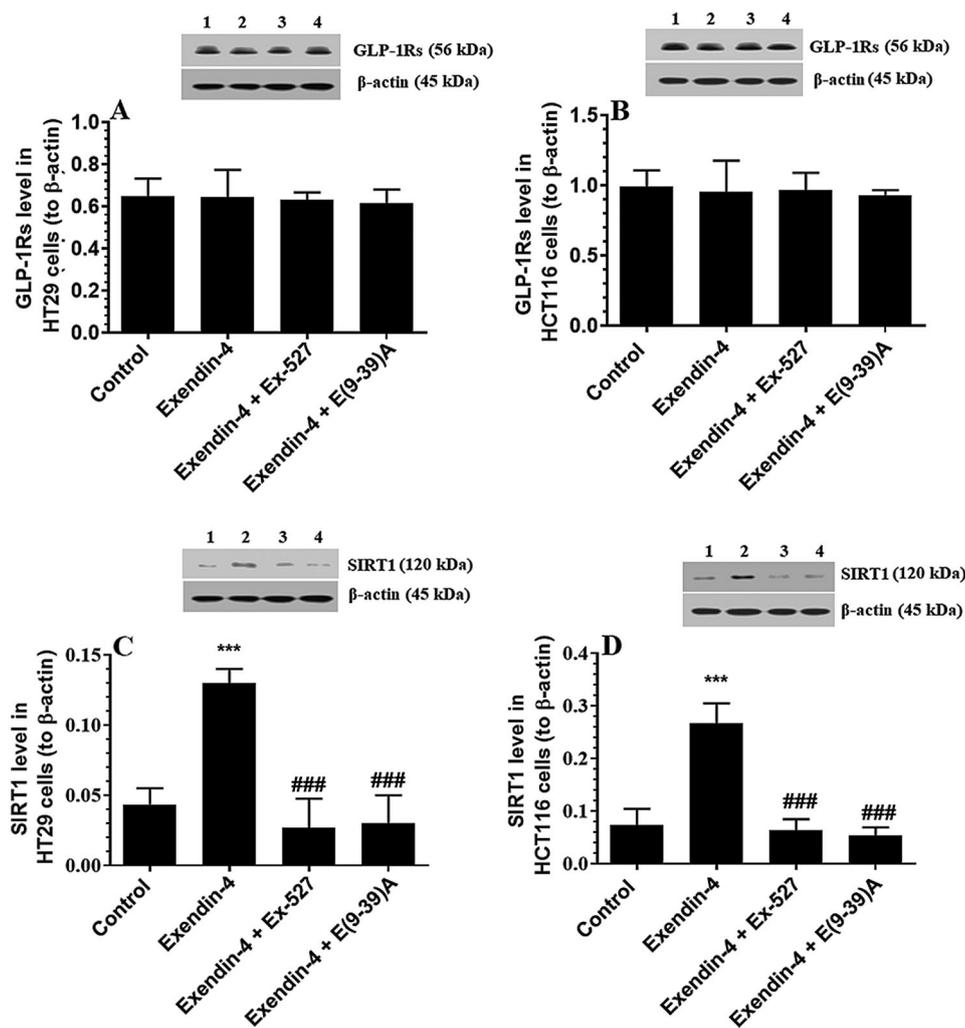


FIGURE 3. Protein levels of GLP1 and SIRT1 in HT29 (A–C, respectively) and HCT116 (B–D, respectively) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exendin-4 at its IC₅₀ (45 μ M for HT29 and 35 μ M for HCT1165) for 24 h (37°C and 5% CO₂) with or without 1 h pre-incubation with EX-527 (10 μ M), A selective SIRT1 inhibitor or Exendin (9-39) amide (E(9-39)A) (1 μ M), a GLP-1 antagonist. Data are presented as mean \pm SD of three independent experiments, each performed in triplicate. ***: vs. control (DMSO-treated) at $P < 0.001$. ###: vs. Exendin-4-treated cells at $P < 0.001$. Lane 1: control cells; Lane 2: Exendin-4-treated cells; Lane 3; Exendin-4 + Ex-527-treated cells. Lane 4: Exendin-4 + E(9-39)A-treated cells, respectively.

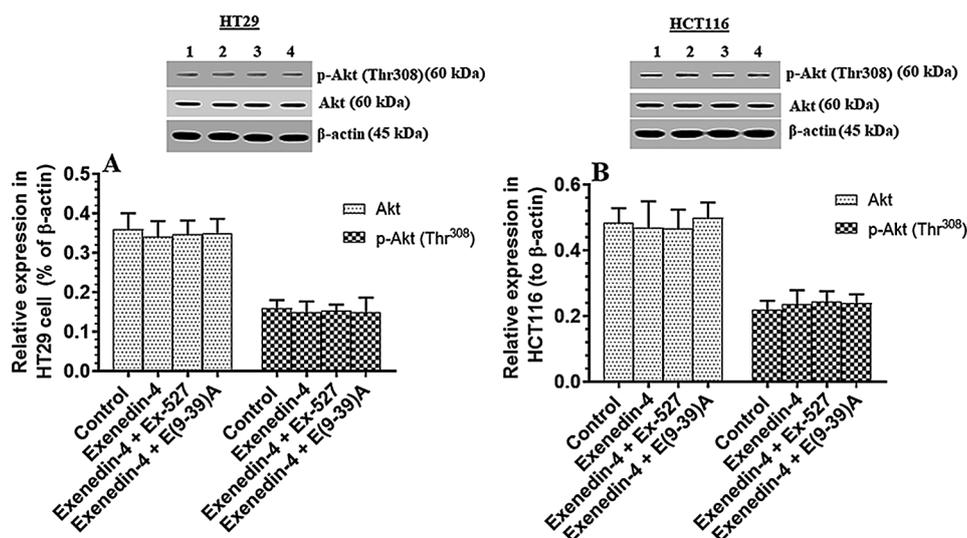


FIGURE 4. Protein levels of Akt and phosphor-Akt (Thr³⁰⁸) in HT29 (A) and HCT116 (B) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exendin-4 at its IC₅₀ (45 μ M for HT29 and 35 μ M for HCT1165) for 24 h (37°C & 5% CO₂) with or without 1 h pre-incubation with EX-527 (10 μ M), A selective SIRT1 inhibitor or Exendin (9-39) amide (E(9-39)A) (1 μ M), a GLP-1 antagonist. Data are presented as mean \pm SD of three independent experiments, each performed in triplicate. No significant variation in the levels of Akt or p-Akt (Thr³⁰⁸) was detected between all groups of the study of both cell lines. Lane 1: control cells; Lane 2: Exendin-4-treated cells; Lane 3; Exendin-4 + Ex-527-treated cells. Lane 4: Exendin-4 + E(9-39)A-treated cells, respectively.

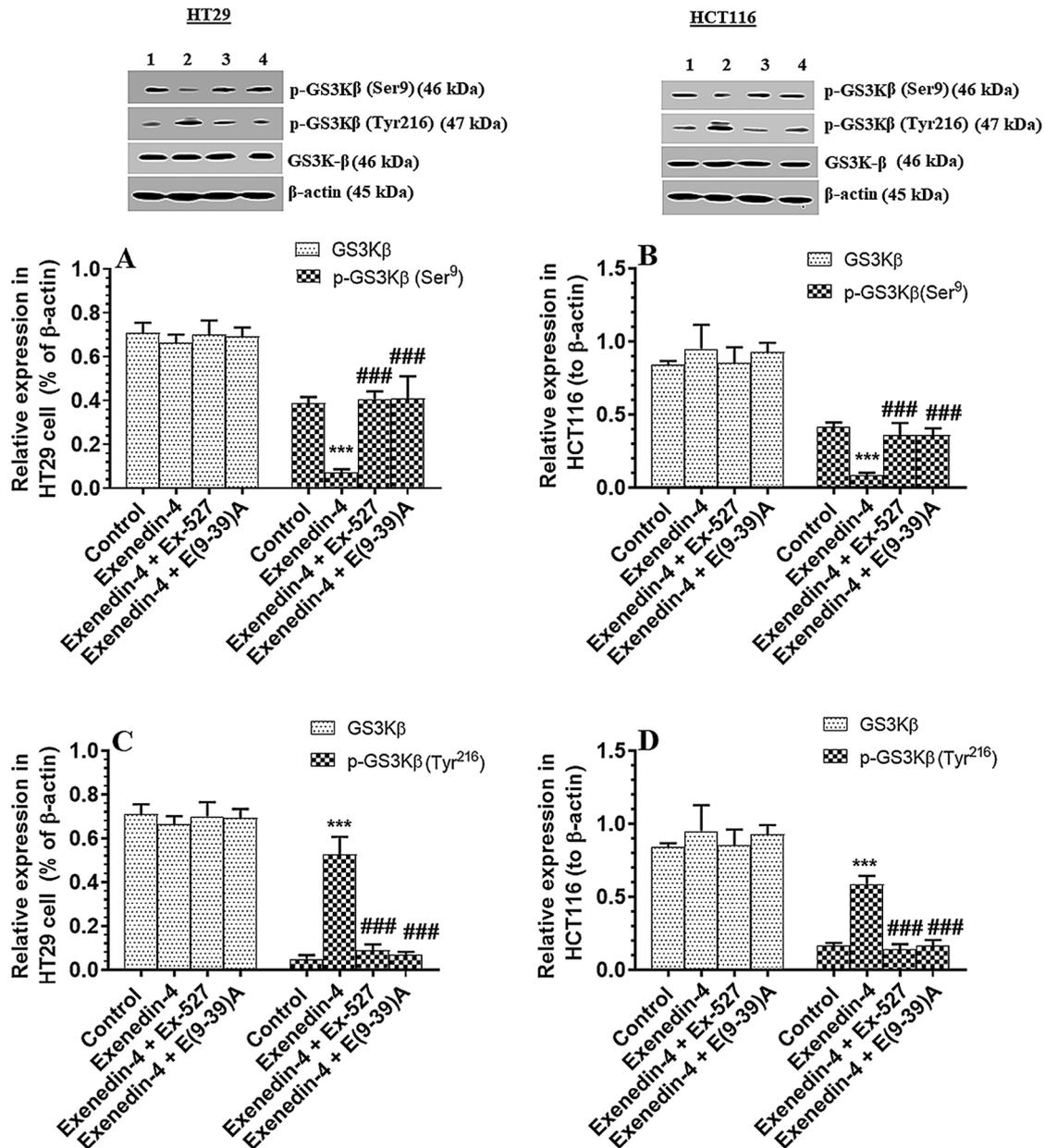


FIGURE 5. Protein levels of total and p-GS3Kβ (Ser⁹ and Tyr⁶¹²) in HT29 (A&C, respectively) and HCT116 (B&D, respectively) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exenedin-4 at its IC₅₀ (45 μM for HT29 and 35 μM for HCT116) for 24 h (37°C & 5% CO₂) with or without 1 h pre-incubation with EX-527 (10 μM), A selective SIRT1 inhibitor or Exenedin (9-39) amide (E(9-39)A) (1 μM), a GLP-1 antagonist. Data are presented as mean ± SD of three independent experiments, each performed in triplicate. ***: vs. control (DMSO-treated) at $P < 0.001$. ###: vs. Exenedin-4-treated cells at $P < 0.001$. Lane 1: control cells; Lane 2: Exenedin-4-treated cells; Lane 3: Exenedin-4 + Ex-527-treated cells. Lane 4: Exenedin-4 + E(9-39)A-treated cells, respectively.

used monoclonal specific antibodies, He *et al.* (2017) have used they have used two polyclonal antibodies. Supporting our data, Saber-Ayad *et al.* (2018) have shown that both HT29 and HCT116 cell lines express GLP-1Rsinn in abundant quantities. Further evidence shown in this study is the all the anti-tumorigenesis effects exerted by Exednin-4 were significantly attenuated by the use of Exenedin (9-39), the antagonist of the GLP-1R.

However, the aberrant activation of the β-catenin is believed to be a major mechanism underlying the development and progression of several solid tumors, including CRC (Bennecib *et al.*, 2000). The activation of this

pathway triggers the transcription of several targets involved in cell inflammation, survival, division, proliferation, and migration (Hanahan and Weinberg, 2011; Polakis, 2012). Of interest, genetic silencing or pharmacological inhibition of this cellular molecule afforded potent anti-tumorigenesis effects on Serval cell lines of CRC (Cong *et al.*, 2003; Green *et al.*, 2001; Mologni *et al.*, 2010; Morin *et al.*, 1996; Polakis, 2012; Tetsu and McCormick, 1999; van de Wetering *et al.*, 2002; Verma *et al.*, 2003). Supporting this evidence, we are demonstrating here that Exenedin-4, and at least, by inhibiting β-catenin activation, can induce cell death and attenuate the proliferation and invasiveness of both T29 and

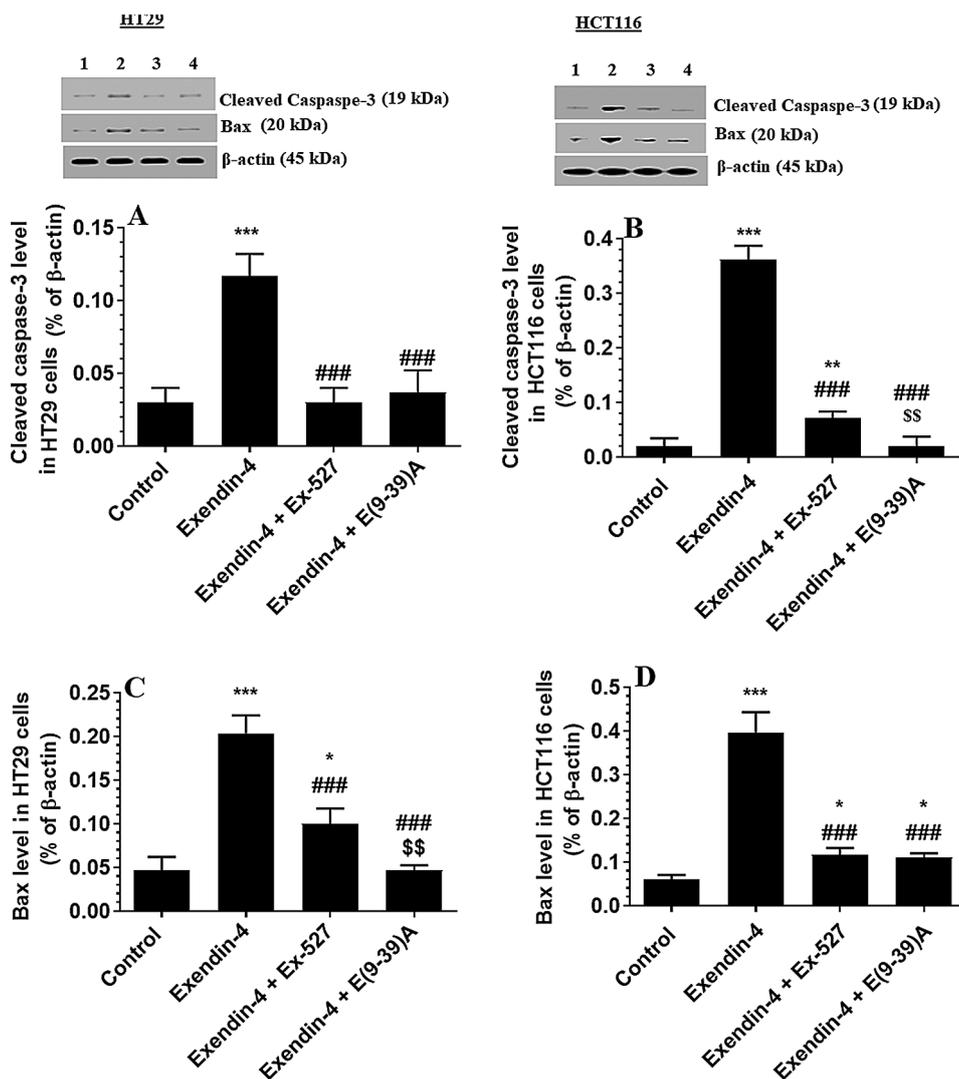


FIGURE 6. Protein levels of Cleaved caspase-3 and Bax in HT29 (A&C, respectively) and HCT116 (B&D, respectively) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exendin-4 at its IC₅₀ (45 μM for HT29 and 35 μM for HCT116) for 24 h (37°C & 5% CO₂) with or without 1 h pre-incubation with EX-527 (10 μM), A selective SIRT1 inhibitor or Exendin (9-39) amide (E(9-39)A) (1 μM), a GLP-1 antagonist. Data are presented as mean ± SD of three independent experiments, each performed in triplicate. ***,***: vs. control (DMSO-treated) at P < 0.05; 0.01, and 0.001, respectively. ###: vs. Exendin-4-treated cells at P < 0.001. \$\$\$: vs. Exendin-4 + Ex527 at P < 0.001. Lane 1: control cells; Lane 2: Exendin-4-treated cells; Lane 3: Exendin-4 + Ex-527-treated cells. Lane 4: Exendin-4 + E(9-39)A-treated cells, respectively.

HCT116 cell lines, *in vitro*. Indeed, Exendin-4 at an IC₅₀ dose of 36.6 μM increased the phosphorylation of β-catenin (Ser³³/Ser³⁷/Thr⁴¹), which are the main site for the action of the GS3Kβ (Metcalf and Bienz, 2011; Rajkumar, 2016). Besides, it significantly reduced protein levels of Axin2, which is considered a major marker of β-catenin activation (Lustig *et al.*, 2002; Mccubrey *et al.*, 2014; Rajkumar, 2016). Of note, the inhibition of this β-catenin is mediated by a cytoplasmic proteome degradation due to phosphorylation GS3Kβ or by deacetylation by SIRT1 (Cho *et al.*, 2012; Lin and Fang, 2013; Metcalfe and Bienz, 2011; Salaroli *et al.*, 2008). In the first scenario, it is well-reported that activation of GS3Kβ is indispensable for cytoplasmic degradation and inhibition of β-catenin (Metcalf and Bienz, 2011; Rajkumar, 2016). Of note, phosphorylation of the Ser⁹ residue inactivates GS3Kβ, whereas the phosphorylation of Tyr²¹⁶ residue activates it (Jacobs *et al.*, 2012). Therefore, we have examined the effect of Exendin-4, in both cell lines, on the activity of GS3Kβ and the level/activity of SIRT1. Interestingly, Exendin-4 significantly suppressed the phosphorylation of GS3Kβ at its Ser⁹ and concomitantly increased its phosphorylation at Tyr²¹⁶, thus implicated activation. These effects occurred without altering the activity of Akt, a major upstream target of GS3Kβ. At the

same time, it increased the levels and activity of SIRT1. Taken together, these data suggest the anti-tumorigenesis of Exendin-4 is mediated by its ability to inhibit β-catenin, at least by activation of GS3Kβ and SIRT1.

On the other hand, inflammation is a significant hallmark in the majority of cancers, which stimulate tumor proliferation, survival, metastasis, and angiogenesis and reprogramming its energy metabolism toward glycolysis and lipid synthesis (Greten and Grivennikov, 2019). The activity of NF-κB is mainly regulated by phosphorylation (Lin and Fang, 2013). GS3Kβ is known to inhibit NF-κB p65 by direct phosphorylation at its Ser⁴⁶⁸ residue and by deacetylation by SIRT1 (Buss *et al.*, 2004). Besides, SIRT1 can inhibit NF-κB by deacetylation (Lin and Fang, 2013; Yang *et al.*, 2012). However, NF-κB and activation of β-catenin can positively affect each other (Rajkumar, 2016). Indeed, overexpression of β-catenin upregulated βTrCP, which in turn activates NF-κB by stimulating the degradation of IκB-α (Winston *et al.*, 1999). Also, NF-κB can directly activate β-catenin by either downregulating LZTS2 and/or reducing the activation of GS3Kβ (Cho *et al.*, 2008; Rajkumar, 2016). Also, IKKα upregulated and activated β-catenin signaling by inhibiting the GS3Kβ (Carayol and Wang, 2006). Besides, TNF-α is a potent

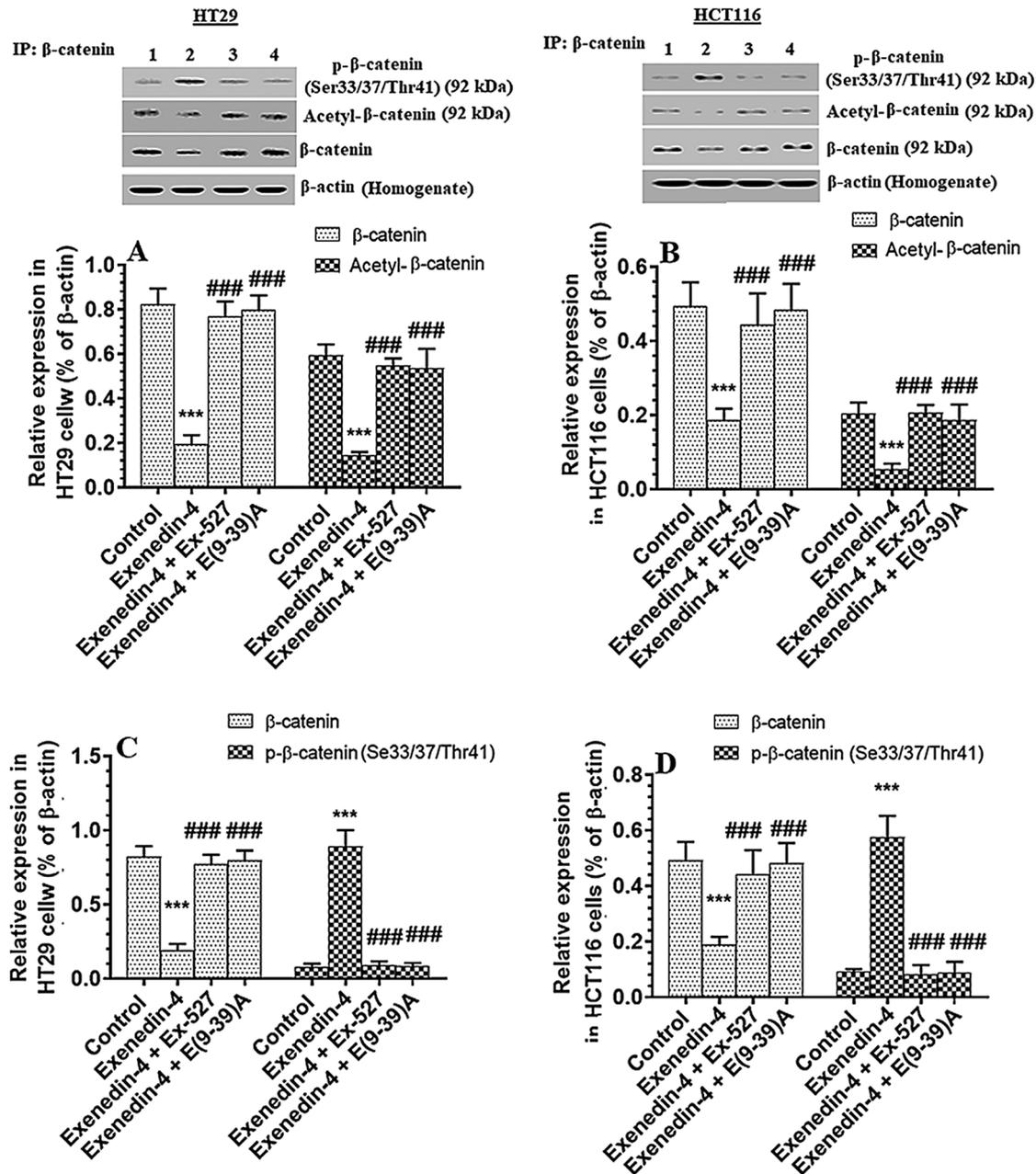


FIGURE 7. Protein levels of total, p-(Ser^{33/37}/Thr⁴¹), and acetylated β -catenin in HT29 (A&C, respectively) and HCT116 (B&D, respectively) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exendin-4 at its IC₅₀ (45 μ M for HT29 and 35 μ M for HCT1165) for 24 h (37°C & 5% CO₂) with or without 1 h pre-incubation with EX-527 (10 μ M), A selective SIRT1 inhibitor or Exendin (9-39) amide (E(9-39)A) (1 μ M), a GLP-1 antagonist. Data are presented as mean \pm SD of three independent experiments, each performed in triplicate. ***: vs. control (DMSO-treated) at $P < 0.001$, respectively. ###: vs. Exenedin-4-treated cells at $P < 0.001$. Lane 1: control cells; Lane 2: Exendin-4-treated cells; Lane 3: Exendin-4 + Ex-527-treated cells. Lane 4: Exendin-4 + E(9-39)A-treated cells, respectively.

activator of β -catenin signaling through the inhibition of GS3K β (Li *et al.*, 2014; Oguma *et al.*, 2008).

In this study, we have also found that Exendin-4 can ameliorate the inflammatory microenvironment in both cell lines by attenuating the activation of NF- κ B. In this regard, Exendin-4 not only reduced the phosphorylation of NF- κ B at Ser⁴⁶⁸ but also decreased its acetylation. These data support many other studies which have shown that Exendin-4 is capable of attenuating inflammation in breast cancer as well as in obesity, myocardial infarction, atherosclerosis, and stroke by inhibiting NF- κ B (Athauda and Foltynie, 2016;

Guo *et al.*, 2016; Iwaya *et al.*, 2017; Lee *et al.*, 2012; Ma *et al.*, 2014). Although it could be possible that Exendin-4 suppresses β -catenin by inhibiting cell inflammation inflammatory or vice versa, our data also suggest that Exendin-4-induced inhibition of NF- κ B is mediated by direct phosphorylation of GS3K β and upregulation of SIRT1. In support, Eid *et al.* (2020) have recently shown that Exendin-4 attenuated cardiac remodeling after myocardial infarction and protects the heart from the damaging effect post-ischemia/reperfusion by activation/upregulation of SIRT1-induced inhibition of NF- κ B (Eid *et al.*, 2020).

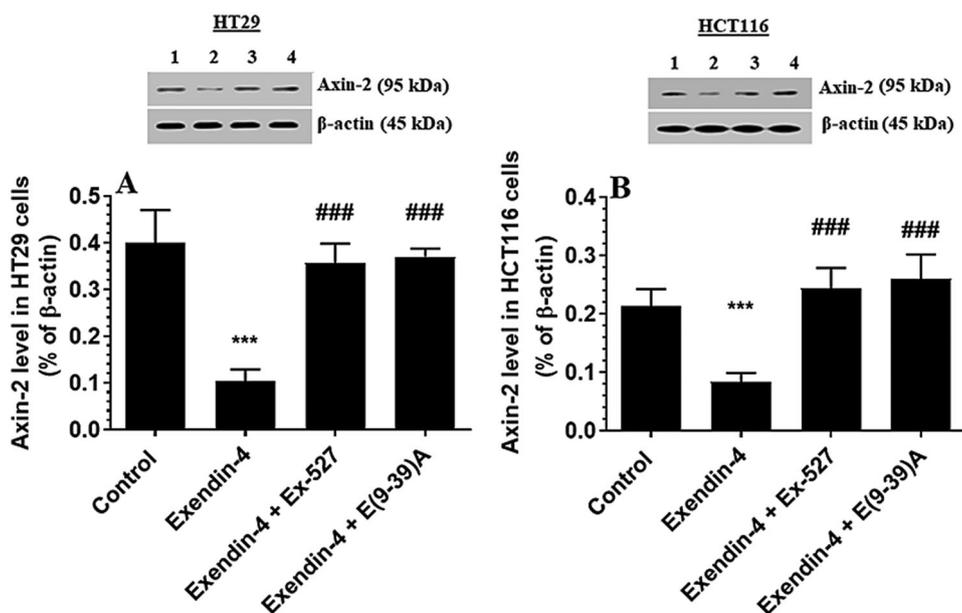


FIGURE 8. Protein levels Axin2 in HT29 (A) and HCT116 (B) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exendin-4 at its IC_{50} (45 μ M for HT29 and 35 μ M for HCT116) for 24 h (37°C & 5% CO_2) with or without 1 h pre-incubation with EX-527 (10 μ M), A selective SIRT1 inhibitor or Exendin (9-39) amide (E(9-39)A) (1 μ M), a GLP-1 antagonist. Data are presented as mean \pm SD of three independent experiments, each performed in triplicate. ***: vs. control (DMSO-treated) at $P < 0.001$, respectively. ###: vs. Exendin-4-treated cells at $P < 0.001$. Lane 1: control cells; Lane 2: Exendin-4-treated cells; Lane 3: Exendin-4 + Ex-527-treated cells. Lane 4: Exendin-4 + E(9-39)A-treated cells, respectively.

Another important observation in this study is that pre-incubating the cells with EX-527, a selective SIRT1 inhibitor, or Exendin (9-39), a GLP-1 antagonist, completely prevented Exendin-4-induced cell death and increased cell proliferation, survival, and migration. Besides, they prevented the inhibitory effects of Exendin-4 on NF- κ B and β -catenin and increased the phosphorylation of GS3K β . These data confirm that the activation of SIRT1 by Exendin-4 underlies its anti-tumorigenesis activity in both HT29 and HCT116 cell lines, an effect that is mediated through GLP-1Rs.

Generally, SIRT1 has a dual effect on cancer, of either acting as a tumor promoter or a suppressor. This matter is still under debate. In some studies, the tumor-suppressive effect has been suggested to its ability to inhibit NF- κ B as evident in this study (Kong *et al.*, 2011; Lin and Fang, 2013; Yeung *et al.*, 2004). On the other hand, SIRT1 can stimulate tumor growth and metastasis by several other mechanisms, including the deacetylation of p53 and FOXO-1, as well as promoting autophagy (Rajkumar, 2016). P53 mediated intrinsic cell apoptosis directly by upregulation of Bax (Chipuk *et al.*, 2004). Upon acetylation by SIRT1, p53 is unable to induce cell apoptosis due to the downregulation of Bax. However, this contradicts our results which have shown increased expression of Bax and cleaved caspase-3 even with higher levels of SIRT1. Of note, besides acetylation, the activation of p53 is mediated by a wide array of post-translational mechanisms, including phosphorylation, ubiquitination, methylation, glycosylation, simulation, and neddylation (Jacobs *et al.*, 2012). A balance between these various mechanisms determines the fate of p53. During inflammation, NF- κ B antagonizes cell death by decreasing the transcriptional activity of p53 (Xia *et al.*, 2014). On the contrary, the activation of GS3K β entails an apoptotic signal in various cells by inhibiting the proapoptotic transcription factor such as heat shock protein-1 and CREB and boosting the activity of other transcription factors such as p53 (Jacobs *et al.*, 2012). Within this view, GS3K β can also bind to the c-terminal of

p53 to directly phosphorylate it at multiple serine residues to promote its nuclear translocation and transcription activity (Beurel *et al.*, 2004; Jacobs *et al.*, 2012; Qu *et al.*, 2004; Turenne and Price, 2001; Watcharasit *et al.*, 2003). Also, several studies have shown that the phosphorylation of GS3K β at Tyr²¹⁶ is essential for its apoptotic activity (Jacobs *et al.*, 2012; Koehler *et al.*, 2011; Yang *et al.*, 2012). Therefore, the decrease in the activation of NF- κ B and the concomitant activation of GS3K β could explain the apoptotic effect of Exendin-4 and the associated increase in Bax and cleaved caspase-3 levels the treated cell lines.

However, despite this interesting finding, we still have some limitations. Most importantly, although our data have shown that Exendin-4 anti-tumorigenic effect is mediated mainly by a SIRT1 mechanism and through GLP-1Rs, by the use of inhibitors, knocking down these targets is indispensable to strengthen these findings. Therefore, more investigations using transgenic animals or gene silencing *in vitro* could support our findings. Besides, given the complicated positive and negative interactions between GS3K β , inflammatory cytokines, NF- κ B, and β -catenin, the data of this study are not sufficient to determine the initial mechanism that triggers all the remaining events. Hence, further studies by targeting each of these markers individually are needed to explain these data. Furthermore, more studies to explain the precise mechanism by which Exendin-4 activates SIRT1 and the subsequent selective activation of GS3K β are highly recommended. Of note, the protein phosphatase PP2A is the best-known activator of GS3K β by removing the phosphorylation at Ser⁹ residue (Bennecib *et al.*, 2000). On the other hand, the proline-rich tyrosine kinase 2 (PYK2) and the Fyn tyrosine kinase are the most common kinases that phosphorylate GS3K β at Tyr²¹⁶ (Lesort *et al.*, 1999). Besides, the GLP-1Rs are of G-coupled protein receptors (GPCRs) type, which normally responds to the ligand-binding in a biased agonism (i.e., some signaling is activated while others are inhibited, at the same time to regulate cell signaling). G protein and

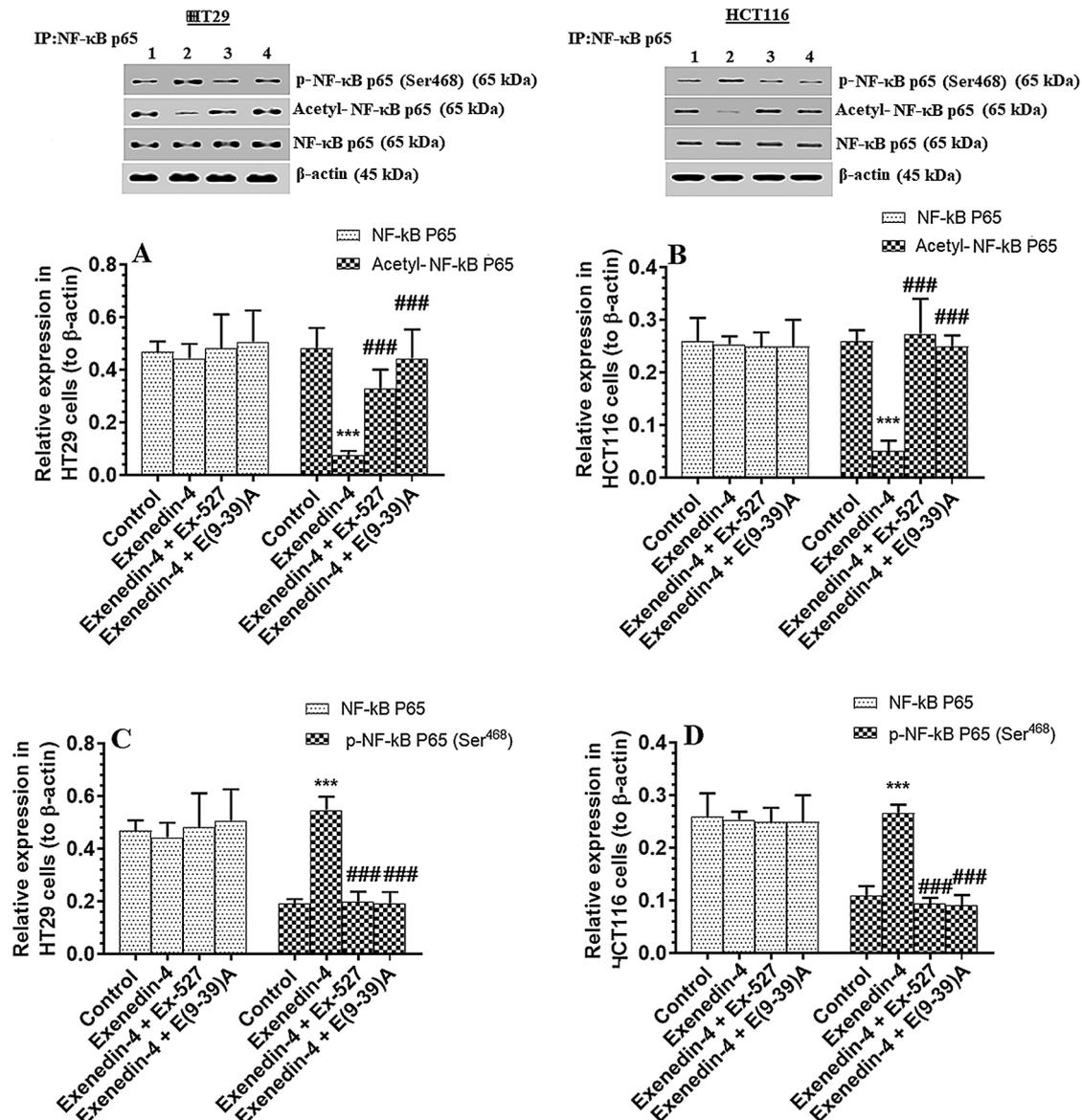


FIGURE 9. Protein levels of total, p-NF-κB p65 (Ser⁴⁶⁸) and acetylated NF-κB p65 in HT29 (A&C, respectively) and HCT116 (B&D, respectively) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exenidin-4 at its IC₅₀ (45 μM for HT29 and 35 μM for HCT1165) for 24 h (37°C & 5% CO₂) with or without 1 h pre-incubation with EX-527 (10 μM), A selective SIRT1 inhibitor or Exendin (9-39) amide (E(9-39)A) (1 μM), a GLP-1 antagonist. Data are presented as mean ± SD of three independent experiments, each performed in triplicate. ***: vs. control (DMSO-treated) at $P < 0.001$. ###: vs. Exenidin-4-treated cells at $P < 0.001$. Lane 1: control cells; Lane 2: Exenidin-4-treated cells; Lane 3; Exenidin-4 + Ex-527-treated cells. Lane 4: Exenidin-4 + E(9-39)A-treated cells, respectively.

β-arrestin1/2 are major substrates activated after GPCRs activation. β-arrestin-2 is an important inducer of PP2A and an inhibitor of NF-κB (Beaulieu *et al.*, 2005; Lei *et al.*, 2018; Tang *et al.*, 2019). Therefore, further studies targeting β-arrestin may help to explain our data.

Finally, the Cmax of exenidin-4 after SC injection (10 μg) is approximately 250 pg/mL (approximately 60 pM). Therefore, the selected dose of Exenidin-4 used in this study seems to be higher than the physiological levels in both humans and animals. However, given that the use of EX-527 reversed the apoptotic and anti-tumorigenic of Exenidin-4. These data suggest that the observed effects are not due to non-specific

toxicity. Although this dose is much lower than the doses used in other studies of different cell cancer types, further studies examining the anti-tumorigenic effect of Exenidin-4 with lower physiological doses on these cell lines are needed.

In conclusion, the data between our hands still interesting and shows for the first time that Exenidin-4, a common GLP-1 agonist that is used to treat diabetes, could be a potent suppressor of CRC. At molecular levels, Exenidin-4 acts in several interconnected mechanisms, including activation of GS3Kβ and inhibiting β-catenin and NF-κB. However, these effects are mediated by the activation of SIRT1 and through the GLP-1R.

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Author Contribution: The study conception and design: Attalla El-kott, Ayman E. El-Kenawy and Mashael Mohammed Bin-Meferij; data collection: Eman R. Elbealy and Ali S. Alshehri; analysis and interpretation of results: Heba S. Khalifa, Ehab E. Massoud and Amira M. Alramlawy; draft manuscript preparation: Attalla El-kott, Ayman E. El-Kenawy and Amira M. Alramlawy; All authors reviewed the results and approved the final version of the manuscript.

Availability of Data and Materials: The data used to support the findings of this study are available from the corresponding author upon request.

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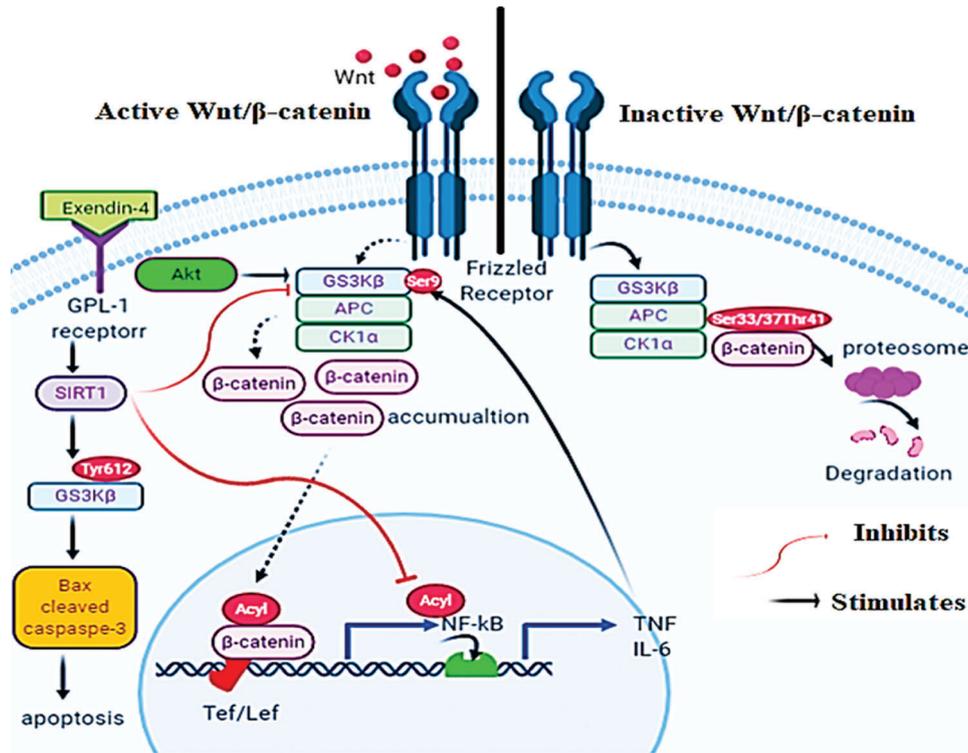
Conflicts of Interest: The authors have declared no conflict of interest.

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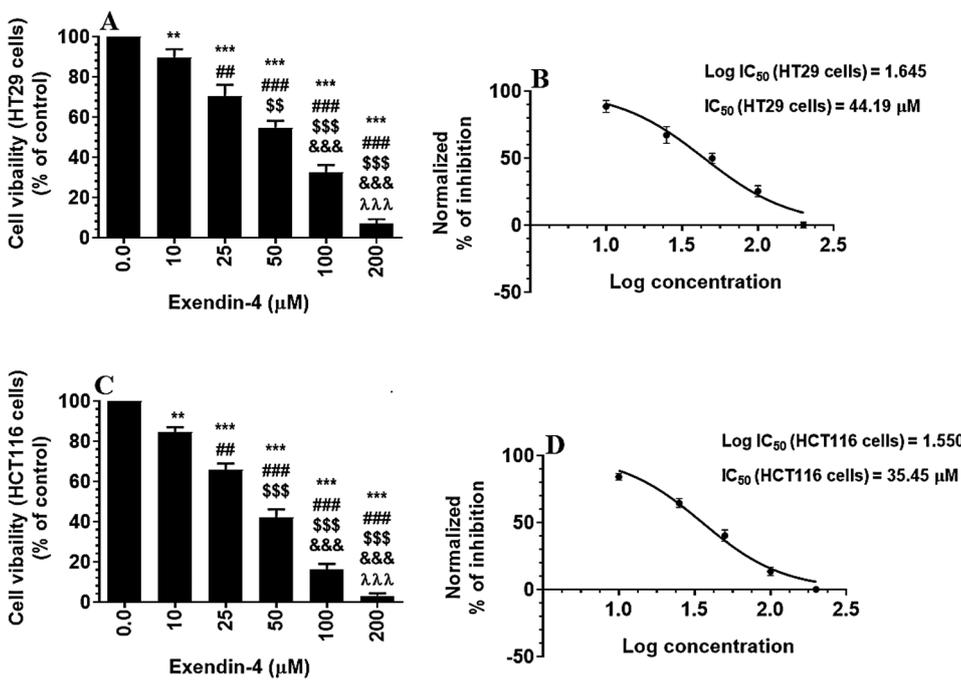
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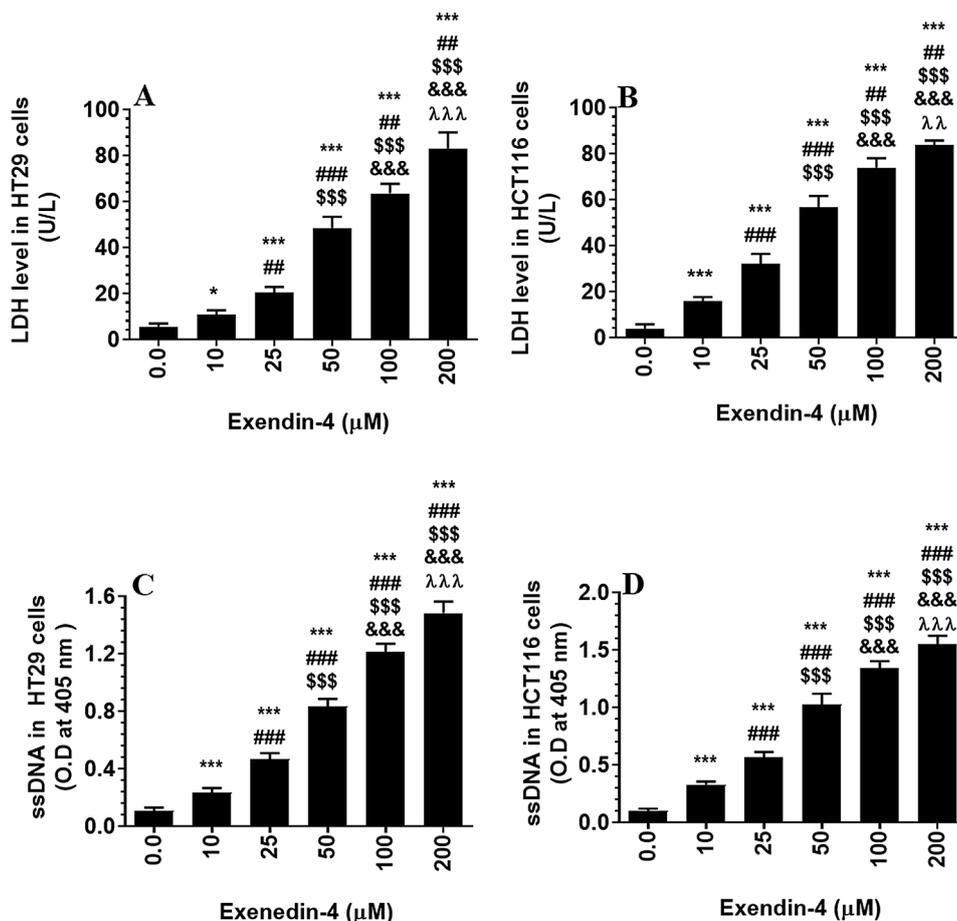


Graphical Abstract

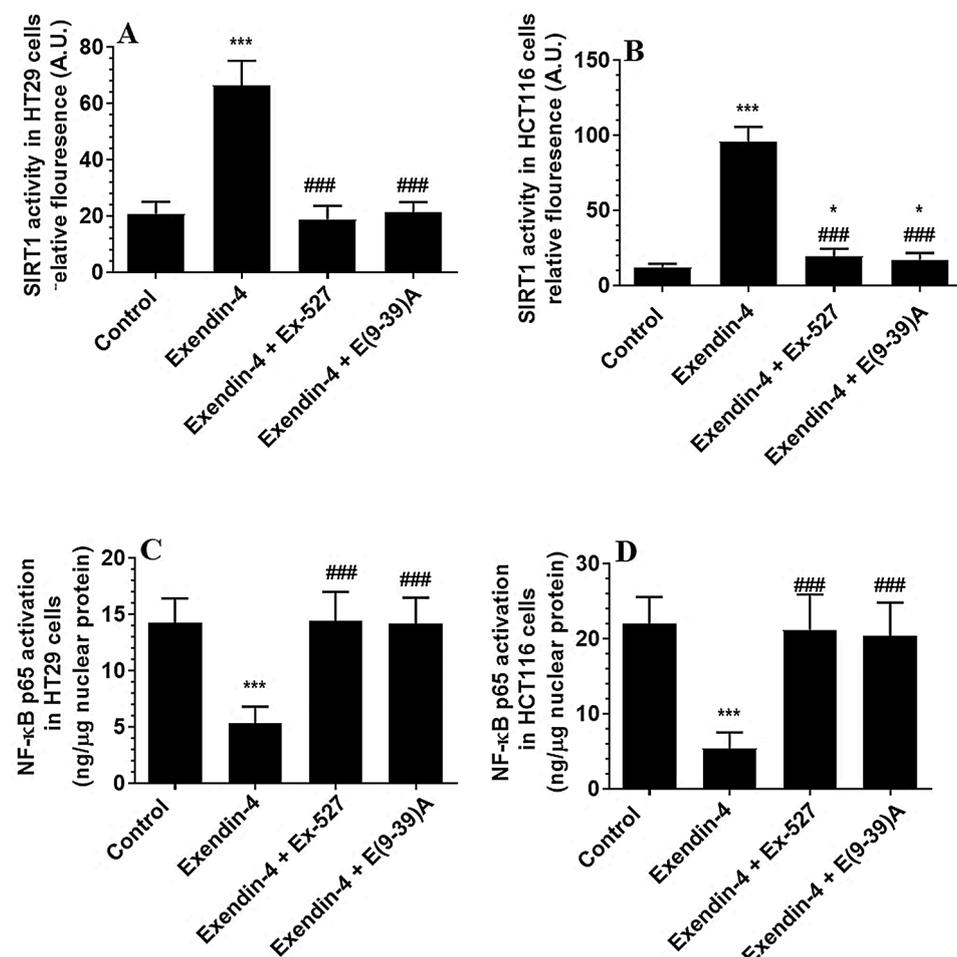
A schematic diagram for presenting the mechanisms by which Exendin-4 induces apoptosis in HT29 and HCT116 colorectal (CRC) cells. In the absence of Wnt, GS3K is active and stimulates the proteasome degradation of β-catenin by phosphorylating it at Ser^{33/37}/Thr⁴¹ (Right side). On the other hand, Wnt stimulates the phosphorylation of Dishevelled protein that inhibits the degradation complex by phosphorylating GS3Kβ at Ser⁹ (Left side). Also, Akt can inhibit GS3Kβ by increasing its phosphorylation at Ser⁹. These events allow the nuclear translocation of β-catenin that stimulates the transcription activity of NF-κB. Furthermore, the transcriptional activity of NF-κB and β-catenin is activated by acetylation. Exendin-4, and through GLP-1 receptors, increases SIRT1 levels in both HT29 and HCT116 CRC cell lines. In SIRT1-dependent mechanisms (still unknown), Exendin-4 inhibits the phosphorylation of GS3Kβ at Ser⁹ and thus stimulates β-catenin degradation and decreases its transcriptional activity.



SUPPLEMENTARY FIGURE S1. Cell Viability in HT29 and HCT116 colorectal cells (A–C, respectively) and their calculated half inhibitory concentration (IC₅₀) (B–D, respectively). The cells were treated with DMSO (0.05%) (control, 0.0 μM of Exendin-4) or with increasing concentration of Exendin-4 (prepared in DMSO) (10–200 μM) for 24 h (37°C and 5% CO₂). Data are presented as mean ± SD of three independent experiments, each performed in triplicate. **, ***: vs. control (DMSO-treated) at P < 0.01 and P < 0.001, respectively. #, ##: vs. 10 μM at P < 0.01 and P < 0.001, respectively. \$\$\$: vs. 25 μM at P < 0.001. &&&: vs. 50 μM at P < 0.001. λλλ: vs. 50 μM at P < 0.001.



SUPPLEMENTARY FIGURE S2. Levels of Lactate dehydrogenase (A–B) and single stranded DNA (ssDNA) (B–C) in HT29 and HCT116 colorectal cells, respectively) and their calculated half inhibitory concentration (IC₅₀) (B and D, respectively). The cells were treated with DMSO (0.05%) (control, 0.0 μM of Exendin-4) for 24 h (37°C and 5% CO₂) or with increasing concentration of Exendin-4 (prepared in DMSO) (10–200 μM) for 24 h. Data are presented as mean ± SD of three independent experiments, each performed in triplicate. *, ***, vs. control (DMSO-treated) at *P* < 0.05 and *P* < 0.001, respectively. ##, ###: vs. 10 μM at *P* < 0.01 and *P* < 0.001, respectively. \$\$\$, vs. 25 μM at *P* < 0.001, respectively. &&&, vs. 50 μM at *P* < 0.001, respectively. &&&, vs. 50 μM at *P* < 0.001. λλλ: vs. 50 μM at *P* < 0.001.



SUPPLEMENTARY FIGURE S3. The activity of SIRT1 and NF-κB in HT29 (A–B, respectively) and HCT116 (C–D, respectively) colorectal cells of all treated groups. The cells were treated with DMSO (0.05%) (control) in the presence or absence of Exendin-4 at its IC₅₀ (45 μM for HT29 and 35 μM for HCT116) for 24 h (37°C and 5% CO₂) with or without 1 h reintubation with EX-527 (10 μM), A selective SIRT1 inhibitor or Exendin (9-39) amide (1 μM), a GLP-1 antagonist. Data are presented as mean ± SD of three independent experiments, each performed in triplicate. ***, vs. control (DMSO-treated) at *P* < 0.05 and *P* < 0.001, respectively. ###: vs. Exendin-4-treated cells at *P* < 0.001.