Therapeutic Efficacy and Inhibitory Mechanism of Regorafenib Combined With Radiation in Colorectal Cancer

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Abstract. Background: Although both chemotherapy and radiotherapy (RT) can sufficiently maintain tumor suppression of colorectal cancer (CRC), these treatments may trigger the expression of nuclear factor kappa B (NF- κ B) and compromise patients' survival. Regorafenib suppresses NF- κ B activity in various tumor types. However, whether regorafenib may act as a suitable radiosensitizer to

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enhance therapeutic efficacy of RT remains unknown. Materials and Methods: Here, we established a CRC-bearing animal model to investigate the therapeutic efficacy of regorafenib in combination with RT, through measurement of tumor growth, body weight, whole-body computed tomography (CT) scan and immunohisto-chemistry staining. Results: Smallest tumor size and weight were found in the combination treatment group. In addition, RT-induced upregulation of NF-kB and downstream proteins were diminished by regorafenib. Moreover, the body weight and liver pathology in the treated group were similar to those of the non-treated control group. Conclusion: Regorafenib may enhance the anti-CRC efficacy of RT.

Recent innovative radiotherapy (RT) technologies, including intensity-modulated RT, image-guided RT, volumetric-modulated arc therapy, intraoperative RT, and stereotactic body RT (SBRT), are widely used to treat many types of cancer (1-3). RT as neoadjuvant strategy offers therapeutic benefits for patients with colorectal cancer (CRC). The combination of RT and surgical resection or chemotherapy has been shown to reduce the local recurrence rate and improve the survival of patients with CRC (4, 5). For increasing anti-CRC efficacy of RT, novel radiosensitizers that sensitize cancer cells to radiation have been developed (6).

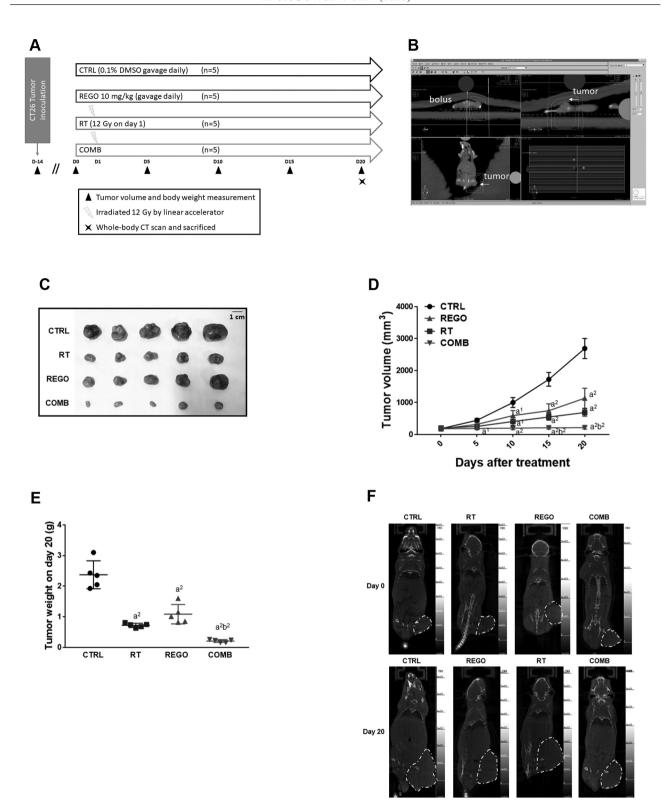


Figure 1. Regorafenib (REGO) enhanced radiotherapy (RT) inhibition of tumor growth in colorectal cancer-bearing mice. A: Experimental flow chart of animal experiment. B: Irradiation dosage and area definition are analyzed by XiO® treatment planning software. C: Photo of extracted tumors from each group of mice on day 20. D: Tumor size. E: Tumor weight. F: Whole-body computed tomographic (CT) scan images. CTRL: Control, 0.1% dimethyl sulfoxide; COMB: combination therapy. Significantly different at: a^1 : p<0.05, and a^2 : p<0.01 vs. control group; b^2 : p<0.01 vs. single treatment group.

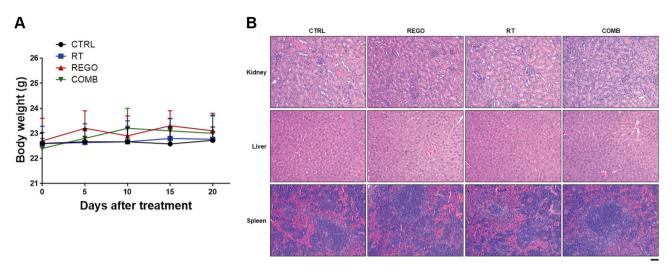


Figure 2. No general toxicity is found in mice treated with regorafenib (REGO) combined with radiotherapy (RT). A: Mouse body weight is measured every 5 days. B: Kidney, liver and spleen tissues from each group of mice on day 20 are displayed. CTRL: Control, 0.1% dimethyl sulfoxide; COMB: combination therapy. Scale bar: 100 µm.

Regorafenib, a Food and Drug Administration-approved oral multi-kinase inhibitor, as well as an analog of sorafenib, has been demonstrated to prolong the overall survival of patients with CRC. Regorafenib induces tumor regression through inactivation of angiogenic and oncogenic kinases such as vascular endothelial growth factor receptors 1-3, fibroblast growth factor receptor 1, epidermal growth factor receptor, RAF, tyrosine-protein kinase, and rearranged during transfection (7-10). The combination of regorafenib and SBRT was demonstrated to increase survival benefits in patient with metastatic CRC (11). However, the anti-CRC effect and mechanism of regorafenib when used along with RT is not fully understood. To harness the beneficial properties of regorafenib on CRC radiotherapy, in vivo investigative experiments were carried out to better understand its feasibility in radiosensitization and the underlying mechanism involved.

Materials and Methods

Cells, antibodies and reagents. Human CRC CT26 cells were purchased from the Bioresource Collection and Research Center (BCRC, Hsinchu, Taiwan, ROC). Cells were cultured in RPMI-1640 medium with additional 10% fetal bovine serum, 10 U/ml of penicillin and 100 mg/ml streptomycin sulfate at 37°C under a humidified atmosphere with 5% CO₂. Primary antibodies purchased from different companies as follow, Ki-67 (Elabscience, Houston, TX, USA), cyclin D1 (Elabscience), survivin, NF-kB p65 (Ser276) (Cell Signaling Technology, Danvers, MA, USA), cleaved caspase-3 (Elabscience), cleaved caspase-8 (Elabscience), cleaved caspase-9 (Proteintech, Rosemont, IL, USA), Fas cell surface death receptor (FAS) (Elabscience), Fas ligand (FAS-L) (Elabscience), myeloid cell leukemia-1 (MCL1) (Elabscience), X-linked inhibitor of apoptosis

protein (XIAP) (Elabscience), and cellular FADD-like interleukin- 1β -converting enzyme-inhibitory protein (C-FLIP) (Elabscience). Regorafenib and dimethyl sulfoxide (DMSO) were purchased from Sigma (St. Louis, MO, USA).

Animal experiment. Male BALB/c and CAnN.Cg-Foxn1nu/CrlNarl mice, approximately 22-28 g in weight and 4-6 weeks of age, were obtained from the National Laboratory Animal Center (Taipei, Taiwan, ROC). CT26 cells (5×106 cells/in 0.1-ml phosphatebuffered saline/mouse) were subcutaneously injected into the right flank of mice and developed into CRC in 2 weeks. A total of 20 mice were randomly divided into four groups (n=5 each) when tumor reached 100-120 mm³ and subsequently received different treatments: Control mice were treated with 0.1% DMSO in 100 µl double distilled water; regorafenib-treated mice were gavage-treated with 10 mg/kg/day regorafenib in 100 µl double distilled water; RTtreated mice were treated with 12-Gy radiation once on day 1 by linear accelerator; and the mice of the combination treatment group were treated with both regorafenib and radiation as described above (Figure 1A). Ethical approval was provided by the Institutional Animal Care and Use Committee of China Medical University (CMUIACUC-2019-288). Tumor volume was recorded every 5 days, evaluated by caliper and calculated by the equation, tumor volume=length × width² ×0.523. The mice were sacrificed with CO₂ for further studies on day 20 post-treatment.

Tumor, spleen, kidney and liver tissue collection. Tumor, spleen, kidney and liver samples were isolated from all mice after sacrificed and prepared for further processing of formalin fixing and paraffin embedding. Tumor tissues were then weighed individually.

Radiation exposure. Mice were anesthetized by 1-3% isoflurane during the irradiation procedure. For dose build up, the tumor was covered with 1.0 cm bolus and scanned with computed tomography (CT). After CT simulation, XiO[®] treatment planning software was used to evaluate the dose distribution in the mouse. The prescribed dose given by a

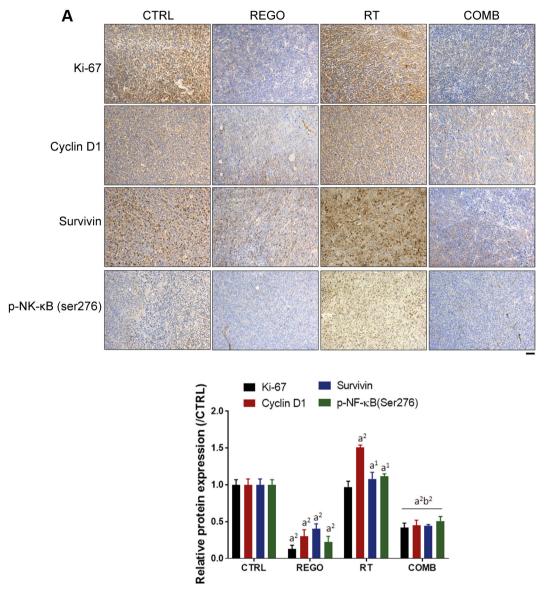


Figure 3. Continued

linear accelerator (6 MV photons; Elekta Synergy linear accelerator, Crawley, UK) to the gross tumor volume was 12 Gy (12).

Whole-body CT scan. One of the mice from each group was anesthetized by 1-3% isoflurane during whole-body CT scan on day 0 and 20. Whole-body CT scan was acquired by Mediso CT (Mediso Ltd., Budapest, Hungary). The scanning resolution was $145{\times}145{\times}145~\mu m$, the tube potential peak was 55 kVp and the tube current was $145~\mu A~(13)$.

Immunohistochemistry (IHC) staining. The protein expression level in tumor tissue including, Ki-67, cyclin D1, survivin, NF-kB p65 (Ser536), cleaved caspase-3, cleaved caspase-9, FAS, FAS-L, MCL1, XIAP, and C-FLIP were investigated in tumor

tissue sections by IHC staining as described in previous studies (9, 14). Stained slides were imaged under a Nikon microscope at ×100 (Nikon ECLIPSE Ti-U; Nikon, Minato City, Japan). Protein expression levels were measured by ImageJ software version 1.50 (National Institutes of Health, Bethesda, MD, USA).

Hematoxylin and eosin (H&E) staining. Tissue sections underwent H&E staining by Bio-Check Laboratories Ltd (New Taipei City, Taiwan, ROC) as described elsewhere (9). H&E-stained sections were then imaged under microscopy and quantified by ImageJ software version 1.50.

 $Statistical\ analysis.$ Data are presented as the mean±standard deviation (n=5). All data were analyzed using GraphPad Prism

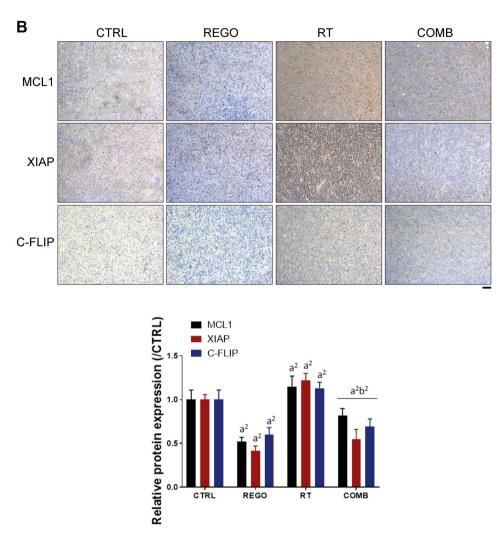


Figure 3. Regorafenib (REGO) reduced radiation (RT)-triggered nuclear factor kappa B (NF- κ B) activation and expression of anti-apoptosis proteins. Tumor immunostaining for expression of A: Ki-67, cyclin D1, survivin, phospho (p)-NF- κ B(ser276) and B: myeloid cell leukemia-1 (MCL1), X-linked inhibitor of apoptosis protein (XIAP), and cellular FADD-like interleukin-1 β -converting enzyme-inhibitory protein (C-FLIP) from each group of mice are displayed (left) and quantified (right). CTRL: Control, 0.1% dimethyl sulfoxide; COMB: combination therapy. Significantly different at: a^1 : p<0.05, and a^2 : p<0.01 vs. control group; b^2 : p<0.01 vs. single treatment group. Scale bar: 100 μ m.

version 7. Differences between the control and experimental groups were investigated by independent Student's t-test analysis when p<0.05 was considered as a statistically significant difference.

Results

Regorafenib effectively induced anti-CRC ability of RT. To investigate whether regorafenib may trigger toxicity of RT, we established CT26 tumor-bearing animal model. As shown in Figure 1A, mice were divided into four different groups: Control treated with 0.1% DMSO; RT alone (12 Gy/once); regorafenib alone (10 mg/kg/day); and combination treatment. RT treatment plan was analyzed XiO[®] treatment planning

software and irradiation was delivered by linear accelerator (Figure 1B). A 1-cm bolus for dose build up was placed on mouse tumors during irradiated procedure. Isolated tumors from each group at the endpoint (day 20) were photographed and displayed in Figure 1C. Smallest tumor size was found in the combination treatment group. Disparity in tumor volume between the combination treatment and control groups began at day 5 after treatment. After 15 days of treatment, the treatment efficacy of combination treatment group was better than that of the single treatment group (Figure 1D). Tumors extracted from mice on day 20 were weighed and the tumor weight of the combination treatment group was less than that of single treatment groups (Figure 1E). Whole-body CT scan

indicated marked tumor growth inhibition in the combination treatment group as compared to single treatment groups (Figure 1F). In summary, as found from the tumor growth pattern, regorafenib effectively sensitized CRC to RT.

Regorafenib combined with RT in CRC-bearing mice appeared not to induce general toxicity. To investigate whether regorafenib combined with RT influenced body weight alteration, we measured mouse body weight once every 5 days. As shown in Figure 2A, no significant difference change of body weight was found between untreated and treated groups. In addition, we also extracted liver, spleen, and kidney tissues from each group for pathology evaluation (Figure 2B). No noticeable pathological change was found in H&E staining between untreated and treated groups. Taken together, regorafenib combined with RT appeared not to induce any general toxicity effect.

Regorafenib diminished radiation-induced NF-κB and NF-κB-mediated protein expression on CRC-bearing mice. NF-κB phosphorylation and the expression of NF-κB-related proteins were detected by IHC staining of tumor sections. In Figure 3A, radiation-induced phosphorylation of NF-κB was reduced by combination with regorafenib. Tumor proliferation marker, Ki-67, was markedly reduced by regorafenib combined with RT. NF-κB-related proteins that regulate tumor proliferation, such as cyclin D1 and survivin were all reduced by combination therapy. Furthermore, NF-κB-related proteins that regulate apoptosis (12, 15), such as MCL1, XIAP and C-FLIP were also effectively reduced by regorafenib combined with RT (Figure 3B). Regorafenib successfully diminished RT-induced expression of NF-κB and its associated proteins.

Regorafenib enhanced RT-induced expression of extrinsic and intrinsic apoptosis-associated proteins in CRC-bearing mice. The levels of extrinsic and intrinsic apoptosis-related proteins, cleaved caspase-3, -8 and -9, were detected by IHC staining of tumor sections. As illustrated in Figure 4A, overall apoptosis-related marker cleaved caspase-3 was significantly expressed in the combination treatment group as compared to single treatment groups. Expression of FAS, FAS-L, and cleaved caspase-8, which are extrinsic apoptosis-related proteins, were induced by regorafenib combined with RT (Figure 4B). Moreover, cleavage of caspase-9, an intrinsic apoptosis-related protein was also triggered in the combination treatment group as compared to single treatment group. In conclusion, RT-induced expression of extrinsic and intrinsic apoptotic-related proteins were successfully enhanced by regorafenib.

Discussion

The combination of regorafenib and SBRT has been demonstrated to control disease progression and extend

progression-free survival to approximately 3 years in patients with late-stage metastatic CRC (11). Our results indicate that regorafenib, RT, and their combination significantly diminished tumor growth of CT-26 tumor-bearing mice (Figure 1C and D). The combination group exhibited small tumor volume and tumor weight compared to mice from individual single treatment groups (Figure 1E). We found that regorafenib, as a radiosensitizer, effectively diminished CRC tumor growth with RT *in vivo*.

Cleaved caspase-8 and -9 are required for apoptosis induced by radiation through extrinsic (*via* death receptor) and intrinsic (mitochondrial) apoptotic pathways. Caspase-3 can be activated or cleaved by caspase-8 or caspase-9, leading to initiation of apoptotic DNA fragmentation (16, 17). The increased protein expression of cleaved caspase-3 in tumor and tumor-associated stroma was associated with better prognosis in patients with CRC (18). Our results showed that expression of FAS, FAS-L, and cleaved caspase-3, -8, and -9 was significantly augmented with regorafenib, RT, and their combination (Figure 4). The combination treatment increased the expression of these proteins to a significantly higher level than that increased by regorafenib or radiation alone.

NF-κB, an oncogenic transcription factor composed of p50 and p65 dimers, can be activated with chemotherapeutic agents or radiation (19, 20). Increased NF-κB activity attenuates the anticancer efficacy of RT through promoting the expression of proliferative and anti-apoptotic proteins, such as cyclin-D1, survivin, MCL1, XIAP, and C-FLIP (12, 21). It has been found that worse 3- and 5-year overall survival of patients with CRC was correlated with overexpression of NF-κB activation (22). In addition, cell and animal models revealed suppression of NF-κB signaling enhanced radiosensitivity of CRC (12, 15, 21). Correspondingly, we found that radiation-triggered NF-κB activation, expression of proliferative, and anti-apoptotic proteins were significantly diminished by regorafenib concurrent treatment (Figure 3).

In conclusion, regorafenib was found to enhance the anti-CRC efficacy of RT. Induction of apoptosis and reduction of NF-κB signaling are involved in the radiosensitizing effect of regorafenib.

Conflicts of Interest

The Authors declare no competing financial interests regarding this study.

Authors' Contributions

YCL, JHH, MCW and FTH performed the experiments. YHL, ITC and CSC prepared the initial version of the article. YCL, JGC and SSL designed the study, performed the literature review, and prepared the final versions of the article.

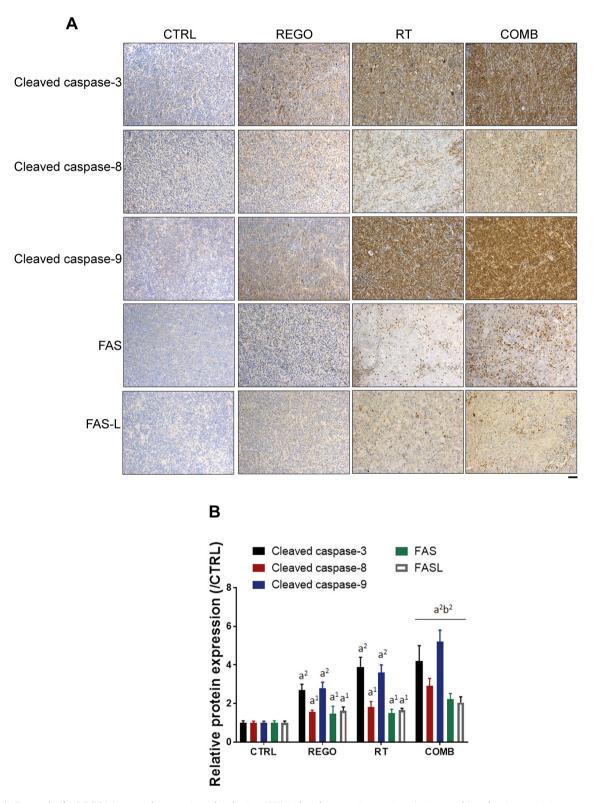


Figure 4. Regorafenib (REGO) increased expression of radiation (RT)-induced apoptotic proteins. A: Immunohistochemistry staining images and B: quantification of cleaved caspase-3, -8 and -9, Fas cell surface death receptor (FAS), and Fas ligand (FAS-L) expression in tumor from each group of mice are displayed. CTRL: Control, 0.1% dimethyl sulfoxide; COMB: combination therapy. Significantly different at: a^1 : p<0.05, and a^2 : p<0.01 vs. control group (0.1% dimethyl sulfoxide; CTRL); b^2 : p<0.01 vs. single treatment group. Scale bar: 100 μ m.

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