Turmeric Is Therapeutic in Vivo on Patient-Derived Colorectal Cancer Xenografts: Inhibition of Growth, Metastasis, and Tumor Recurrence

Mingyue Li1,2†, Grace Gar-Lee Yue2†, Lianxiang Luo3, Stephen Kwok-Wing Tsui1, Kwok-Pui Fung1,2, Simon Siu-Man Ng4 and Clara Bik-San Lau2*

1 School of Biomedical Sciences, The Chinese University of Hong Kong, Hong Kong, China, 2 Institute of Chinese Medicine and State Key Laboratory of Research on Bioactivities and Clinical Applications of Medicinal Plants, The Chinese University of Hong Kong, Hong Kong, China, 3 The Marine Biomedical Research Institute, Guangdong Medical University, Zhanjiang, China, 4 Department of Surgery, Faculty of Medicine, The Chinese University of Hong Kong, Hong Kong, China

Colorectal cancer is the third most frequently diagnosed cancer worldwide. Clinically, chemotherapeutic agents such as FOLFOX are the mainstay of colorectal cancer treatment. However, the side effects including toxicity of FOLFOX stimulated the enthusiasm for developing adjuvants, which exhibit better safety profile. Turmeric extract (TE), which has been previously shown to suppress the growth of human and murine colon xenografts, was further demonstrated here for its inhibitory effects on colon cancer patient-derived xenografts (PDX). PDX models were successfully established from tissues of colon cancer patients and the PDX preserved the heterogeneous architecture through passages. NOD/SCID mice bearing PDX were treated either with TE or FOLFOX and differential responses toward these treatments were observed. The growth of PDX, metastasis and tumor recurrence in PDX-bearing mice were suppressed after TE treatments with 60% anti-tumor response rate and 83.3% anti-metastasis rate. Mechanistic studies showed that TE reduced tumor cell proliferation, induced cell apoptosis, inhibited metastasis via modulating multiple targets, such as molecules involved in Wnt and Src pathways, EMT and EGFR-related pathways. Nevertheless, FOLFOX treatments inhibited the PDX growth with sharp decreases of mice body weight and only mild anti-metastasis activities were observed. Furthermore, in order to have a better understanding of the underlying mechanisms, network pharmacology was utilized to predict potential targets and mechanism. In conclusion, the present study demonstrated for the first time that oral TE treatment was effective to suppress the growth of colon PDX and the recurrence of colon tumors in mice. The findings obtained from this clinically relevant PDX model would certainly provide valuable information for the potential clinical use of TE in colorectal cancer patients. The application of PDX model was well illustrated here as a good platform to verify the efficacy of multi-targeted herbal extracts.

Keywords: colorectal cancer, patient-derived xenografts, turmeric, tumor recurrence, herbal medicines, network pharmacology
INTRODUCTION

Colorectal cancer (CRC) is the third most frequently diagnosed malignancy and is the second leading cause of cancer death worldwide (1). The incidence and death rates of CRC decreased among individuals aged ≥ 50 years, but increased by 13% in those aged less than 50 years (2). Metastasis results in nearly 90% of all cancer deaths (3) and it occurs in 20%–30% of CRC patients (2, 4). Poor prognosis of patients with non-resectable stage III–IV metastatic CRC (mCRC) provoked the development of preclinical models with novel mechanisms of action. Nonetheless, adverse events were reported in some targeted therapies (5, 6). Hence, there remains high unmet needs for safer adjuvant and/or therapeutic agents for mCRC patients, which could be searched from natural sources.

In the progress of development of anti-tumor and anti-metastatic agents, preclinical models play vital roles in translating preclinical data to clinical efficacy. However, high failure ratio of novel anti-tumor therapies in clinical trials partly results from the inability of conventional cell line xenograft models in reliably predicting the clinical efficacy (7, 8). In fact, cell lines used in xenograft models have adapted to the passaging on plastic outside of a natural tumor environment, whereas cancer cell-stromal cell interactions should have taken place in the presence of human extracellular matrix component. Hence, the cell line xenograft models may not be accurate enough to reflect interaction with the tumor microenvironment and tumor heterogeneity (8). In contrast, patient-derived xenograft (PDX) models, which have been established and adapted in recent years, are shown to preserve the intratumoral cell heterogeneity, histopathological and genetic alterations (9, 10). The PDX models are now regarded as effective, reproducible and clinically relevant preclinical models for testing the efficacy of new target-directed therapies, validating biomarkers of drug response, and evaluating preclinical drugs or therapeutic agents (11–13).

Along this line, natural product curcumin has been proven to have multiple molecular targets in cancer (14), and other components such as turmerones and polysaccharides have also been shown to possess anti-proliferative and immune-stimulatory effects (15, 16). However, poor bioavailability and low solubility limit curcumin to achieve its satisfactory therapeutic outcome in clinical trials. Previous studies suggested that turmeric (main ingredient in curry and a source of curcumin) may contribute to the lower CRC incidence in Indians (17, 18). Our previous studies also demonstrated that the whole complex of turmeric extract (TE) rather than curcumin alone could exert better inhibitory effects on colon cancer cell line generated xenografts growth in mice (19). Xenografts derived from cell lines generally exhibit poorly differentiated carcinomas that lack resemblance to the original patient tumor in terms of molecular characteristics (20). In order to confirm the benefits arising from the use of turmeric in clinical-relevant animal models, which would be a crucial step before launching clinical trial for this health supplement, we established a panel of PDX of CRC. The efficacies of TE on tumor growth, metastasis and tumor recurrence were then investigated in this panel of PDX for the first time.

Systems biology approach, in particular network pharmacology enable the paradigm shift from “one-target, one-drug” to a “network-target, multiple-component-therapeutics” mode, which is a powerful tool for the analysis of drug combinations, especially for traditional Chinese medicines (21, 22). Network pharmacology is a new method based on “disease-gene-drug” network which provides a new way to explore the mechanism of drug action at molecular, cellular, tissue and biological levels (23). The multi-dimensional mechanism of drug action is evaluated by target prediction, pharmacokinetic measurement and network analysis. More importantly, network pharmacology has been successfully applied to predict/identify the molecular mechanisms of traditional Chinese medicine in the treatment of various diseases (21), including cardiovascular diseases (24), neurodegenerative diseases (25), cancer (26), etc. Here, we have identified potential active components of TE against CRC targets, and predicted their network, suggested the potential pathways and key regulatory genes.

METHOD AND MATERIALS

Reagents

Turmeric ethanolic extract (TE) was prepared as previously reported (19, 27). In brief, the dried rhizome of Curcuma longa was extracted under reflux using 95% ethanol and evaporated under reduced pressure at 60°C to dryness. Quantification of two commercially available chemical markers, curcumin and Ar-turmerone (Sigma-Aldrich, MO, USA), in turmeric ethanolic extract was performed using UPLC and the chemical profiles were registered (27). The contents of curcumin and Ar-turmerone present in turmeric ethanolic extract were 18.7% (w/w) and 5.3% (w/w), respectively. Standardization of turmeric ethanolic extract used in the present study was same as the method reported previously, in which the 3-D chromatograms of the extract as well as the extracted ion chromatograms (EIC) and MS spectra of the marker compounds were also well reported previously (27).

Chemotherapeutics FOLFOX consisted of fluorouracil, folinic acid and oxaliplatin, were obtained from Sigma-Aldrich (MO, USA). Primary antibodies against RhoA, β-catenin, Src, p-Src, and GAPDH were purchased from Cell Signaling (MA, USA); Ki-67 from Abcam (MA, USA); CD31 from Dako (CA, USA); mouse/rabbit specific HRP/DAB (ABC) Detection IHC kit from Abcam Inc (Cambridge, UK). The in situ Cell Death Detection kit for TUNEL staining was from Roche (IN, USA). The Trizol reagent was from Thermo Fisher Scientific (MA, USA) and QuantiFast® SYBR® Green PCR mini kit was from Qiagen (CA, USA).

Establishment of Colorectal Cancer Patient Derived Xenograft Mouse Model

Experimental protocols and procedures were reviewed and approved by the Joint Chinese University of Hong Kong – New Territories East Cluster Clinical Research Ethics Committee (CREC Ref No.: 2016.444) and the Animal Experimentation Ethics Committee of The Chinese University of Hong Kong (Ref No. 15-164-MIS).
Freshly resected tissues (tumor and adjacent normal tissues) from consented CRC patients were collected after colon surgery. Half portion of resected tumor tissues were stored in –80°C freezer for further histological and molecular analysis and another half portion of tumors were transferred at 4°C into RPMI-1640 medium with antibiotics. Within 24 h of surgical resection, tumor tissues were trimmed, cut into 3- to 5-mm sizes and inoculated subcutaneously on the back of 6- to 8-week-old NOD/SCID mice, which were supplied by the Laboratory Animal Services Centre of The Chinese University of Hong Kong. Mice were maintained in pathogen-free conditions in specifically designed air-controlled rooms with a 12-h light/dark cycle. When the tumors reached 1,000–1,500 mm³, mice with the first generation of xenografts (P1) were sacrificed and then the xenografts were isolated and expanded for two more generations (P2 and P3). When P3 xenografts reached an average volume of 50 mm³, mice were then subjected to TE or FOLFOX treatments (Figure 1A).

**Administration of Turmeric Extract or FOLFOX to PDX-Bearing Mice**

In total, seven batches of PDX samples were established for the following pharmacological experiments. Six batches of PDX-bearing mice were treated with TE (200 mg/kg, orally administered) or vehicle control 6 days a week for 2–8 weeks depending on the tumor sizes and the well-being of mice. FOLFOX (15 mg/kg fluorouracil and 5 mg/kg folinic acid daily plus 5 mg/kg oxaliplatin once a week) was administered intraperitoneally for 2 or 4 weeks depending on the well-being of mice. The treatments and assessment details of the PDX samples were listed in Table 1. In our previous study, TE at 400 mg/kg has been shown to inhibit growth of human colon HT29 subcutaneous xenografts (27). Thus, both 200 and 400 mg/kg TE have been tested in our pilot studies in PDX-bearing mice. Results showed that the inhibitory activities of these two doses of TE were comparable and without significant difference. Hence, the minimum effective dose (200 mg/kg) was chosen for the subsequent PDX experiments in the present study.

Tumor volumes and body weights were measured once a week until the end of experiments. Tumor volumes were calculated using the formula: length × width × depth/2 (mm³) as described previously (28). Tumor growth inhibition was calculated using the formula: (TGI= Average tumor weight in control group – Average tumor weight in treatment group)/ Average tumor weight in control group × 100%. Relative tumor volume was equaled to tumor volume (on the measuring day) divided by tumor volume (on day 0).

**Histological Assessments**

PDX samples were collected at the end of experiments and separated into two halves, one half was snap frozen at –80°C for further Western blot/RT-PCR analysis and the other half was fixed in 10% formalin. Lungs and livers were also fixed in 10% formalin, embedded in paraffin and then sectioned for hematoxylin and eosin (H&E) staining. Tumor area in liver and lung H&E sections were used to assess the level of metastasis. The metastatic area was calculated as percentage of metastatic area in whole view of photo. Embedded PDX tissues from each group were sectioned and stained with H&E or subjected to IHC staining against β-catenin and Ki 67 antibodies. The TUNEL assay was performed to assess apoptosis in tumor sections as per kit’s instruction. H&E and IHC staining were performed as previously described (28). Immunoreactivity was analyzed in 5 random areas for each tumor tissue and was scored as 0 (no staining), 1 (weak staining), 2 (moderate staining), 3 (strong staining), and 4 (very strong staining) (29).

**Western Blotting**

Western blot was performed using the lysates of PDX samples. Twenty to sixty micrograms of protein were loaded to denaturing gel electrophoresis. After transfer to PVDF membrane, membranes were blocked with 5% BSA and incubated overnight with primary antibodies, RhoA, p-Src, Src, β-catenin, cleaved caspase-3, and GAPDH in 1:1,000 dilutions. Blots were then incubated with secondary antibodies (1:2,000). Quantification was performed using Image J software as described previously (28).

**RNA Extraction and RT-PCR**

Total RNA was extracted from three batches (CC15OT, CC15PT, and CC15ZT) of PDX samples using Trizol reagent. The reverse transcription and quantification were performed as previously described (30). The gene expression of Wnt signaling elements (AXIN2, DKK1, and SMAD7), EMT pathway related molecules (N-cadherin, Snail and Vimentin, and EpCAM), growth factor receptors [platelet-derived growth factor receptor B (PDGFRB), epidermal growth factor receptor (EGFR)], asparagine synthetase (ASNS), Rho signaling (RTKN2), ATP-binding cassette transporters (ABCA13), and iNOS were evaluated using qPCR. Besides, total RNA of CC15OT primary tumor and adjacent normal tissue samples were extracted and subjected to qPCR for the expressions of KRAS and EGFR. In addition, total RNA of lungs CC16AET were extracted and the expressions of human-specific 850-bp fragment of the α-satellite DNA on human chromosome 17 (Ch17) were detected by qPCR (31, 32). In brief, real time semi-quantitative PCR of cDNA samples were performed in Bio-Rad CFX96™ Real-time system C1000 Thermal cycler using the QuantiFast SYBR Green RT-PCR kit (Qiagen, USA). The primers sequences were listed in Table S1 (in supplementary information), which were synthesized by Thermo Fisher Scientific (Hong Kong). The specific gene mRNA levels (including Ch17 expression level) were normalized to GAPDH and expressed as a fold change compared to the vehicle control group.

**Data Preparation and Analysis of Network Pharmacology**

The potential active compounds and putative targets of turmeric were searched using Traditional Chinese Medicine Systems Pharmacology Database (TCMSP, http://tcmspw.com/tcmsp.php), and The Encyclopedia of Traditional Chinese Medicine (ETCM, http://www.tcmip.cn/ETCM/index.php/Home/Index/index.html). The optimal cutoff values of OB and DL in
FIGURE 1 | Schematic diagram showing the establishment of PDX samples and histological assessment of PDX samples. (A) Patients’ tumors were obtained, inoculated subcutaneously on the back of NOD-SCID mice (P1). When the size of PDX samples reached 1,000 mm³, PDX samples were sub-cultured into the next batch of mice (P2), which subsequently expanded to the third passage (P3). Treatments of turmeric extract or FOLFOX or vehicle control were applied in the mice bearing P3 PDX samples. (B) The histology of representative patients’ adjacent normal tissues, patients’ primary tumors and PDX were determined using H&E staining. Bar = 50 μm.
TCMSP were set to 30% and 0.18. Information on associated target genes of CRC was obtained from GeneCards (https://www.genecards.org/), and the top 1,000 genes were reserved for further analysis. The String database (http://string-db.org/) was utilized to obtain the data of protein-protein interaction (PPI) with the species limited to "Homo sapiens". The Cytoscape 3.7.2 software was applied for constructing the Compound-target Network, Compound-candidate target Network, PPI Network, and Hub Genes Network. GO Slim of candidate targets was analyzed by WebGestalt (http://www.webgestalt.org), and the biological processes and KEGG pathway were analyzed by DAVID 6.8 (https://david.ncifcrf.gov).

Statistical Analysis
All quantitative data were expressed as mean ± standard error of the mean (SEM). The unpaired Student’s t-test was conducted to determine statistical significant differences. p < 0.05 was considered as statistically significance.

RESULTS
Generation of PDX From Colon Cancer Patients’ Tumor Samples
PDX samples for colon cancer were established from seven patients’ tumor samples. PDX were successfully grown through three serial passages in mice. The latency time from initial inoculation time to post-treatment varied from 6 to 8 months. Clinical characteristics including tumor size, origin of samples, tumor/node/metastasis stage and survival time, etc. were listed in Table 2. All patients’ tumors included in this study were moderately differentiated without metastasis. Histological examination of the H&E stained PDX sections was conducted to compare the similarity of PDX with the original patients’ tumors. Both the primary tumors and PDX show features of adenocarcinoma (Figure 1B). PDX could represent diverse clinical characteristics typically observed in patients according to pathologist’s assessment. Meanwhile, the adjacent normal tissues were also stained by H&E to illustrate the differences between normal tissues and tumor tissues. Most of the adjacent tissues have well differentiated histomorphology, which were apparently different from tumor tissues.

Turmeric Extract Inhibited Tumor Growth in Colon PDX
Mice bearing 4 batches of PDX samples (CC15OT, CC15PT, CC16AGT, and CC16AET) were received TE treatments. Since the size of the original patients’ tumors varied, the number of mice in each batch of PDX samples were different. Besides, the duration of treatment was determined by the tumor size and/or the well-being of mice. Some of the experiments were terminated when the tumor size of untreated control group was large (~600–700 mm³) or the PDX-bearing mice became very weak (even the PDX size was not large), as it was unethical to continue the experiments.
Nevertheless, our results could show that oral TE treatment decreased the size of PDX in three batches (except CC16AET) with different extent. As shown in Figure 2, all tumors in TE-treated groups were smaller than those in vehicle control groups, with tumor growth inhibition for CC15OT, CC15PT, and CC16AGT at 79.1%, 36.7%, and 51.5%, respectively. The inhibitory effects of TE were shown to be the most effective in CC15OT sample, the final tumor weight of TE-treated group was 54.9 ± 7.43 g versus that of control group 263.2 ± 67.3 g (p < 0.01, Figure 2A). Body weights of mice did not significantly differ between treatment and control groups in these three tested batches, indicating TE did not cause systemic toxicity in mice.

Anti-Metastatic Effect of TE Treatment in Colon PDX Models

The efficacy of TE treatment on tumor metastasis in PDX models was reflected by metastasis area in livers and lungs. Levels of metastasis varied among batches, metastatic area ranged from 7.8% to 20.9% in lungs, and from 6.3% to 17.7% in livers. The highest tumor burden was observed in CC16AET sample, with the metastatic area as 17.7% and 20.9% in liver and lung, respectively. The liver metastasis in four batches of PDX samples (CC15OT, CC15PT, CC16AGT, and CC16AET) could be significantly decreased after TE treatment (Figure 3A). Whereas TE treatment significantly reduced the lung

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**TABLE 2 | Characteristics of the patients for CRC PDX model.**

| Sample codes | CC15OT | CC15PT | CC16AGT | CC15AGT | CC15ZT | CC16ANT | CC16AET |
|--------------|-------|-------|--------|--------|-------|--------|--------|
| Range of age            | 70-74 | 80-84 | 50-54  | 65-69  | 80-84 | 65-69  | 60-64  |
| Metastases at presentation (Y/N) | N     | N     | N      | N      | N     | N      | N      |
| Tumor length (cm)        | 2.5   | 4     | 6      | 3.5    | 9     | 4      | 6      |
| Origin of sample         | sigmoid | rectum | ascending colon | rectum | ascending colon | splenic flexure | transverse colon |
| Number of lymph nodes removed  | 12    | 13    | 14     | 32     | 20    | 29     | 31     |
| Tumor/node (TN) staging  | T3N0  | T2N0  | T3N0   | T3N0   | T2N0  | T3N0   | T3N0   |
| Tumor differentiation (well/moderate/poor) | moderate | moderate | moderate | moderate | moderate | moderate | moderate |
| Lymphovascular permeation (Y/N) | N     | N     | N      | N      | N     | N      | N      |
| Adjuvant chemotherapy (Y/N) | N     | N     | N      | N      | N     | N      | N      |
| Adjuvant radiotherapy (Y/N) | N     | N     | N      | N      | N     | N      | N      |
| Follow up time from surgery (months) | 42    | 41    | 30     | 35     | 37    | 23     | 32     |
| Recurrence (Y/N)         | N     | N     | N      | N      | N     | N      | N      |

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**FIGURE 2 | Effects of TE treatment in 3 PDX samples CC15OT, CC15PT, and CC16AGT.** The tumor volume and weight in the control group and TE treatment groups were compared using (A) CC15OT (n = 4-6), (B) CC15PT (n = 6-7), and (C) CC16AGT (n = 7). Tumor volume was measured once a week while the final tumor weight was measured at the end of the experiment. Representative photos for tumors from 4 to 7 mice in each PDX sample. Data were presented as mean ± SEM, *p < 0.05, **p < 0.01 vs. vehicle control at the same time point.
metastatic area in CC15OT, CC16AGT, and CC16AET samples (Figure 3B). Importantly, most of the metastasis occurred near blood vessels where the tumor cells permeated from and then colonized. As the highest inhibition rate (51.0%) of lung metastasis by TE treatment was observed in CC16AET sample, the expression of the human-specific 850-bp fragment of the α-satellite DNA on human Ch17 (31, 32) was determined in lung tissues. Results showed that the relative Ch17 expression (after normalization with housekeeping gene GAPDH) in TE-treated tissue was 1.04 ± 0.70, while that in control group was 2.80 ± 1.56. There was significantly decreased of the Ch17 expression in TE-treated mice (p < 0.05), suggesting that the inhibitory effect of TE treatment on the metastasis and/or colonization of human xenograft cells in mice lungs.

**Underlying Mechanisms of Anti-Tumor and Anti-Metastatic Effects of TE**

As the tumor sizes were reduced after TE treatment, the tumor sections were subjected to TUNEL assay for evaluating apoptosis in tumors. As shown in Figure 4A, TE treatment could significantly induce apoptosis in tumors in CC15PT and CC16AGT samples as shown by TUNEL assay and the expression of cleaved caspase-3. Besides, TE treatment decreased the expression of Ki 67 (proliferation marker, Figure 4B) in tumor sections. On the other hand, β-catenin, involved in tumor cell proliferation (33) and EMT process (34) which is considered as the first step in tumor metastasis, was also assessed in tumor sections. In CC15PT and CC16AGT samples (Figure 4C), β-catenin expression was shown to be significantly decreased in tumors. As the sizes of TE-treated tumors in CC15OT batch were small, tumors were subjected to the extraction of proteins and RNA instead of immunohistochemical analysis. The protein expression of β-catenin in CC15OT samples was examined using Western blot as shown in Figure 5A, and it was apparently reduced after TE treatment.

Furthermore, TE treatment significantly reduced the protein expressions of β-catenin, Src and RhoA in CC15OT (p < 0.05) and slightly reduced the protein expressions of RhoA in CC15PT samples (p = 0.074, Figure 5A). As the RNA of CC15OT and CC15PT samples were extracted, the gene expression of Wnt signaling elements (AXIN2, DKK1, and SMAD7), EMT pathway related molecules (N-cadherin, Snail and Vimentin, and EpCAM), growth factor receptors (PDGFRB), asparagine synthetase (ASNS), Rho signaling (RTKN2), ATP-binding cassette transporters (ABCA13), and iNOS were evaluated using qPCR. As shown in Figure 5B, mRNA expressions of Wnt signaling elements (AXIN 2), EMT pathway related molecules (Vimentin and EpCAM), EGFR, and RTKN2 were all down-regulated in TE-treated tumors of CC15OT samples, whereas DKK1 was significantly up-regulated by TE treatment. On the other hand, the expressions of KRAS and EGFR in primary tumor tissues were shown to be significantly higher than those in adjacent normal tissues (p < 0.001, Figure 5C, left panel). While the CC15OT PDX-bearing mice treated with TE, the expressions of these two genes in the PDX samples were significantly decreased when compared with untreated control (p < 0.05, Figure 5C, right panel).

Furthermore, in the samples of CC15PT, the expressions of AXIN2, SMAD7, N-cadherin, vimentin, ASNS, EGFR, PDGFRB, ABCA13, and DKK1 were all significantly down-regulated after TE treatment (Figure 5D). These results suggested the multi-targeted activities of TE treatment on the colon PDX.

**Novel Finding of TE in Reducing Colon Tumor Recurrence**

Since the anti-tumor and anti-metastatic efficacies were observed in PDX-bearing mice treated with TE, the potential inhibitory effect of TE treatment on tumor recurrence was further investigated in another 2 batches of PDX samples. Mice bearing PDX samples CC15ZT and CC16ANT were treated with TE for
7 and 4 weeks, respectively. Then, the xenografts were removed by surgery. The primary tumor volumes of TE-treated groups were smaller than control group, and the average tumor weight in control group was 1.47 ± 0.39 g versus 0.74 ± 0.23 g in TE-treated group (Figure 6A). After the mice recovered from surgery, they were maintained without any treatment in the following 4 weeks. The recurrent tumors could be observed in mice of vehicle control group. The mice were sacrificed when the largest recurrent tumors reached the size of 1,000 mm³, and the recurrent tumors, livers and lungs were collected for analysis. The tumor incidence after surgery and recurrent tumor weight could indicate the inhibitory effect of TE on tumor recurrence. The incidence of tumor in vehicle control group was 100% (4/4), while that in TE treatment group was only 50% (2/4) (Figure 6A). The average weight of recurrent tumors in TE treatment group (0.018 ± 0.012 g) were lower than that in control group (0.393 ± 0.207 g). TUNEL assay for the primary xenografts and recurrent tumors revealed that TE treatment induced apoptosis in tumors (Figure 6B). The protein expressions of β-catenin, Src and pSrc were suppressed in the tumors in TE-treated group (Figure 6C). The gene expressions of EMT pathway related molecules (N-cadherin, EpCAM, EGFR, ASNS, and RTKN2) were also down-regulated after TE-treatment (Figure 6D). All these molecular alterations in the primary xenografts might contribute to the inhibitory effects of TE on the growth of primary and recurrent tumors. Lastly, lung metastasis was found to be significantly reduced after TE treatment (Figure 6E).

The inhibitory effect of TE on tumor recurrence was also demonstrated in mice bearing PDX sample CC16ANT, which were treated with TE for 4 weeks, and the primary xenografts were removed by surgery. The average tumor weight in vehicle control group was 1.82 ± 0.32 g while that in TE-treated group was 0.95 ± 0.19 g, which was significantly decreased (Figure 6F, left panel). Five weeks after the removal of the primary xenografts, recurrent tumor was found in 1 out of 3 mice treated with TE (i.e., tumor incidence: 1/3). While in vehicle control group, five mice out of seven were found to have recurrent tumors (Figure 6F, right panel). Similarly, the significant reduction of metastasis in liver and lung could be observed in this set of experiment (Figure 6G).

**Efficacy of FOLFOX in Colon PDX Model**

Beyond surgery or in metastatic phase, FOLFIRI (fluorouracil, folinic acid, and irinotecan) or FOLFOX (plus oxaliplatin) are utilized as system treatment (35). Previous studies demonstrated the side effect of FOLFOX including neurotoxicity, immune suppression, hair loss, nausea, etc. (28, 36). Therefore, there is a
strong desire for developing potent and safe anti-tumor and anti-metastatic therapeutics. In order to show the potential superior efficacy of TE over the conventional therapeutics, the efficacy of FOLFOX was also examined in the present study. Chemotherapeutics FOLFOX was administered to mice bearing CC16AGT and CC15AGT PDX samples. Surprisingly, the mice cannot even bear long time FOLFOX treatment. After 2 and 4 weeks of FOLFOX treatments, gradually decreases (>20%) of body weights (Figures 7A, D), the treatments were observed and the treatments were terminated at 2 and 4 weeks in CC16AGT and
CC15AGT PDX samples, respectively. The mice were maintained without any treatment until the end of experiments.

In the CC16AGT PDX samples, the tumor volume was smaller in FOLFOX treatment group when compared with vehicle control group (days 14–21). However, the final tumor volume and weight were similar in FOLFOX-treated group and the vehicle control group (Figure 7B), which might due to the termination of FOLFOX treatment on day 15. Nonetheless, the anti-tumor activities of FOLFOX could still sustain as the apoptotic area in tumors was higher in FOLFOX-treated group than that in vehicle control group (Figure 7C). The metastasis in liver and lung were also examined after FOLFOX treatment, and results showed that the metastatic area was slightly increased in FOLFOX-treated group (Figure 7C).

For the CC15AGT PDX samples, which should be sensitive to FOLFOX treatment because the tumor growth was suppressed by

FIGURE 6 | Effects of TE treatment on the growth of PDX and the recurrence of tumor. Mice bearing PDX samples CC15ZT and CC16ANT were treated with TE and the primary xenografts were removed by surgery. For the CC15ZT PDX sample, (A) tumor volume changes, final tumor weight and photo of primary xenografts were shown on the left panel. The final tumor weight and photo of recurrent tumors were shown on the right panel. The tumor incidence was listed in the brackets next to the photo. (B) Representative images of TUNEL in primary xenografts and recurrent tumor sections and quantitative results of positive-stained cells in the tumor sections were shown. Bar = 50 μm. Data were presented as mean ± SEM. n = 20 sections in each group. (C) Western blot of primary xenograft protein against β-catenin, p-Src, Src, and GAPDH and the quantitative data presented as fold of vehicle control. (D) Real-time PCR of selected gene expressions in the primary xenografts. GAPDH was used for normalization. n = 4 tumors in each group. (E) Liver and lung sections were stained with H&E and the metastatic area (shown by yellow arrows) was assessed. n = 20 sections in each group. Quantitative results of liver and lung metastasis were shown on the right panel. For the CC16ANT PDX sample, (F) tumor volume changes, final tumor weight and photo of primary xenografts were shown on the left panel. The final tumor weight and photo of recurrent tumors were shown on the right panel. The tumor incidence was listed in the brackets next to the photo. (G) Liver and lung sections were stained with H&E and the metastatic area (shown by yellow arrows) was assessed. n = 15–25 sections in each group. Data were shown as mean ± SEM. *p < 0.05, **p < 0.01, ***p < 0.001 vs. vehicle control.
FOLFOX treatment as shown in the tumor growth curve (Figure 7E), meanwhile the body weight was decreased gradually (Figure 7D). After termination of FOLFOX treatment, the tumor volume of this group increased sharply (Figure 7E), resulting in comparable tumor size with vehicle control group. The apoptotic area of tumors were slightly higher in FOLFOX treatment group than that in vehicle control group (Figure 7F). Nevertheless, the metastasis in lung was found to be suppressed by FOLFOX treatment in this batch of PDX-bearing mice, further suggesting the diverse responses of PDX samples toward chemotherapeutics.

**Network Pharmacology Predicts Potential Mechanism of Turmeric Extract Treatment in Colorectal Cancer**

A total of 18 candidate targets for treating CRC were collected from 14 potential active compounds (Table S2) of TE. The PPI network of candidate targets contained 38 nodes and 202 edges, in which average node degree is 10.6 (Figure 8A). The cytoHubba plugin of the Cytoscape software was used to analyze the 38 nodes and found that 3 targets belonged to the hub genes in candidate targets according to rank by Degree, Closeness, and Betweenness (Figure 8B). Hence, we suspected that the following three genes encode proteins in pivotal roles: AKT1, ESR1, and PTGS2. We then performed a GO Slim analysis by WebGestalt, and it was suggested that the candidate targets participated in protein binding, ion binding, and lipid binding in membrane, protein-containing complex, and nucleus (Figure 8C). To further examine the signaling pathways and functions of these candidate targets, functional enrichment analysis was carried out using DAVID 6.8. Candidate targets were found involved in biological processes such as signal transduction, inflammatory response, and positive
regulation of nitric oxide biosynthetic process (Figure 8D). The results of KEGG analysis indicated that candidate targets were intensively associated with PI3K-Akt signaling pathway and thyroid hormone signaling pathway (Figure 8E). In fact, previous studies have shown that these signaling pathways are related to regulating cancer cell growth and proliferation (33, 34), which indicates a possible direction for further cancer related research. Except three hub genes and pathways in cancer, other targets of TE which could be potential targets in treating other diseases were also proposed.

**DISCUSSION**

The in vivo efficacies of turmeric ethanolic extract, which contains absorbable curcumin, were demonstrated in mice bearing colon PDX for the first time in the present study. Seven PDX samples were established from the moderately differentiated colorectal tumors from consented patients and the comprehensive evaluation on anti-tumor (including tumor recurrence) and anti-metastatic effects of TE and chemotherapeutics FOLFOX have been conducted. In our established PDX samples, histological features were well preserved through passages, despite the high degree of heterogeneous architecture and molecular characteristics, they could recapitulate in vivo tumor biology and be employed in the accurate preclinical drug evaluation (8). The responses of these PDX samples toward therapeutic agents (natural or synthetic) would be more clinically relevant than those from cell-derived xenograft models.

In view of the multiple entities of pathology of CRC, PDX model would be the most appropriate model to allow accurate testing of potential multi-targeted therapeutics. Recent studies demonstrated the use of combination of targeted drugs, e.g., ABT-263 plus YM-155 (37), panitumumab plus TAS-102 (38) in colon PDX models. In fact, apart from those novel synthetic small molecules and biomolecules (antibodies) for cancer treatment, herbal medicines are certainly alternative source of therapeutic agents since herbal medicines have long been used for chronic disorders management. The multi-components properties of herbal medicines would match the multiple combination treatment approaches in many diseases, such as cancer.

To prove this concept, a well-known Chinese and Ayurvedic herbal medicine, turmeric, the dried rhizome of the plant *Curcuma longa* Linn. was chosen to be verified of its anti-tumor effects in colon cancer using PDX models. One of the major components of turmeric, curcumin, has been shown to be anti-tumor in various cancer types in preclinical studies (39, 40). However, low water solubility, poor bioavailability, rapid metabolism, and systemic elimination of curcumin hinder its pharmacological application (41). Nevertheless, our previous studies demonstrated that turmeric ethanolic extract, which contains curcuminoids, turmerones, and other components, exerted potent anti-tumor and anti-metastatic effects as well as augmented anti-tumor activities of bevacizumab in colon cancer...
xenograft-bearing mice (19, 27, 42). Despite the mice models we used which covered both immuno-deficient nude mice and immuno-competent Balb/c mice, the xenografts were formed by homogeneous cell lines. Hence, the efficacy of TE treatment in colon cancer has been further verified using the PDX model in the present study, which would reveal the actual potential responders toward the treatments.

The present study demonstrated the anti-tumor and anti-metastatic effects of TE in six PDX samples. The responsive rate of anti-tumor effect toward TE was 50.0% (significantly reduced tumor weight in three out of six samples), and that of anti-metastatic effect was 83.3% responsive rate (five out of six) in both lungs and livers. Although the direct anti-tumor efficacy of TE was not as potent as that observed in cell-derived tumor-bearing model, the present findings further confirmed the potent anti-metastatic activities of TE treatment in PDX models, in which colon cancer cells from different patients metastasized to distal organs from the original xenografts. A recent study reported the inhibition of curcumin on tumor growth and aggressiveness using xenograft model with patient-derived oral squamous cell carcinoma cells (43). Nonetheless, the inhibitory effects of TE treatment toward lung and liver metastasis of colon PDX have not yet been reported. Interestingly, in our project, we did see all of moderately differentiated CRC tumors developed metastasis in PDX model. In fact, in clinical situation, around 70% of CRC tumor are moderately differentiated (44). For the hypothesis that moderately differentiated CRC tumors result in liver and lung metastasis, in our PDX model, we inoculated primary tumors into SCID mice, in which the immune surveillance to tumor metastasis is lacking. Meanwhile, not all primary tumors from patients could develop into PDX. Hence, the successful established PDX needs to be aggressive and may have potential to develop metastasis in patients as well. So, it is understandable for all models here turned out to be metastatic.

The samples of PDX after treatments were subjected to protein and molecular analysis in order to determine the mechanisms of action. Wnt signaling pathway controls cell fate, regulates homeostasis and is closely associated with malignant transformation (45). β-catenin is a key element in Wnt pathway and the loss of degradation of β-catenin in cytoplasm results in the transcription of cell proliferative, survival genes such as MYC, cyclin D (46). DKK1 encodes a Wnt suppressor gene which exerts pro-apoptotic and strong anti-proliferative activities in CRC cell lines (47, 48). TE treatment significantly up-regulated DKK1 expression in CC15OT sample (Figure 5B) and CC16ANT (data not shown) which might account for its anti-tumor effects. AXIN2 encoding a Wnt signaling component and promotes colon carcinoma oncogenic activity (49, 50). SMAD7 expression can affect β-catenin levels and lead to increased Wnt signaling (51). Results from IHC, Western blot and q-PCR showed the down-regulating effects of TE on Wnt signaling pathway, which involves in the CRC progression, including tumor initiation, growth and metastasis (52). In addition, KRAS and EGFR gene expressions were down-regulated by TE treatment in all tested PDX samples. EGFR signal pathway involves in angiogenesis, cancer cell proliferation, migration, invasion as well as inhibition to apoptosis (53). Furthermore, protein expression of Src was also down-regulated after TE treatment, indicating that TE also exerted anti-tumor and anti-metastatic effects through influencing cell growth, adhesion and migration (54). Vimentin, EpcAM, and N-cadherin (belong to EMT pathway) were all down-regulated in TE treatment group in tested PDX samples. These alternations of gene or protein expressions strongly suggested the multiple targets of TE treatment in colon cancer, such targets are also consistent with other studies regarding curcumin (55, 56).

On the other hand, the potential of TE treatment on inhibiting tumor recurrence in colon PDX model was firstly revealed in this study. Although the direct anti-tumor effect of TE treatment may not be as good as the first-line chemotherapeutics, the results of reduced numbers and sizes of recurrent tumors shown here should provide evidences on the “chemopreventive” efficacy of TE treatment. The molecular alterations after TE treatment may also explain the inhibition of tumor recurrence observed in the present PDX model. The results from the present study are of clinical significance as PDX model mimics clinical situation and predict the tumor metastasis. To the best of our knowledge, this is possibly the first study of herbal extract on PDX model of colon cancer.

The first-line chemotherapy for CRC in clinics is based on fluorouracil (5-FU) alone or in combination as FOLFOX (5-FU, folic acid and oxaliplatin)/FOLFIRI (5-FU, folic acid and irinotecan) for long time (35, 57). Unfortunately, despite effectiveness of the chemotherapeutics, the incidence of side effects is as high as 98% (58), including neurotoxicity, immune system suppression, hair loss, nausea and vomiting, severely affected the quality of life of patients (59). In the present study, two batches of mice bearing PDX samples were used to evaluate the efficacy of FOLFOX, which has seldom been reported. Similar to the side effects observed in patients, the body weight of mice bearing PDX received human equivalent dosages of FOLFOX decreased gradually (Figure 7). Meanwhile, the tumor growth was suppressed by FOLFOX treatment. The termination of treatment resulted in tumor volume increases. The overall outcomes of the incomplete FOLFOX treatment were not that positive, although the apoptotic activities retained (shown by TUNEL+ stained cells in Figure 7C). FOLFOX treatment did not effectively suppress liver metastasis in the mice-bearing PDX (Figures 7C, F) and it did not affect the Ki 67 and β-catenin expressions in tumor sections (data not shown). Taken together, the observed differential responses toward TE and FOLFOX treatments were as expected because the study subjects (i.e., each batch of PDX samples) were heterogeneous. In general, TE treatment might exert superior anti-metastasis efficacy than FOLFOX treatment in the PDX samples tested in this study.

Beyond the molecular mechanisms examined here, we utilized system biology method especially network pharmacology approach to predict the potential targets of TE based on its main components. Collectively, we constructed the protein-protein network of the 18 candidate targets of TE treated CRC in which the hub genes were revealed: AKT1, ESR1, and...
PTGS2. Furthermore, we discovered that the candidate targets may involve in signal transduction, inflammatory response, and positive regulation of nitric oxide biosynthetic process. Moreover, as the result of KEGG analysis, multiple pathways like PI3K-Akt signaling pathway and thyroid hormone signaling pathway, may be the potential pathways of turmeric in the treatment of CRC. We postulated that patients may benefit from the combination use of PI3K-Akt and/or thyroid hormone signaling pathway inhibitors together with TE treatment.

In conclusion, we have successfully established PDX model from a panel of CRC patients’ tumors which was a good platform for evaluating the multi-targeted herbal medicines. The preclinical anti-tumor and anti-metastatic activities of turmeric ethanolic extract were firstly demonstrated in individual colon PDX. To make this study more informative, we utilized network pharmacology to figure out more potential drug targets and working mechanisms. This precision medicine approach can potentially provide crucial information regarding preclinical translational evidence to clinical trials and together with network pharmacology to illustrate the potential therapeutic values of TE in CRC prevention and treatment.

DATA AVAILABILITY STATEMENT
The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

ETHICS STATEMENT
The studies involving human participants were reviewed and approved by Joint Chinese University of Hong Kong–New Territories East Cluster Clinical Research Ethics Committee. The patients/participants provided their written informed consent to participate in this study. The animal study was reviewed and approved by Animal Experimentation Ethics Committee of the Chinese University of Hong Kong. Written informed consent was obtained from the individual(s) for the publication of any potentially identifiable images or data included in this article.

REFERENCES
1. Cassidy S, Syed BA. Colorectal cancer drugs market. Nat Rev Drug Discov (2017) 16(8):525–6. doi: 10.1038/nrd.2017.59
2. Siegel RL, Miller KD, Fedewa SA, Ahnen DJ, Meester RGS, Barzi A, et al. Colorectal cancer statistics, 2017. CA Cancer J Clin (2017) 67(3):177–93. doi: 10.3322/caac.21395
3. Christofori G. New signals from the invasive front. Nature (2006) 441 (7092):444–50. doi: 10.1038/nature04872
4. Engstrand J, Nilsson H, Stromberg C, Jonas E, Freedman J. Colorectal cancer liver metastases - a population-based study on incidence, management and survival. BMC Cancer (2018) 18(1):78. doi: 10.1186/s12885-017-3925-x
5. Ohhara Y, Fukuda N, Takeuchi S, Homma R, Shimizu Y, Kinoshita I, et al. Role of targeted therapy in metastatic colorectal cancer. World J Gastrointest Oncol (2016) 8(9):642–55. doi: 10.4251/wjgo.v8.i9.642

AUTHOR CONTRIBUTIONS
CL and GY conceived the study. ML and GY performed the in vitro and in vitro studies and analyzed the data. LL performed the network pharmacology analysis. SN provided colon cancer patients’ tissue samples. ST, SN, and K-PF provided advices on study design and data interpretation. CL and K-PF contributed essential reagents and tools, ML, GY, and CL wrote the manuscript. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL
The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fonc.2020.574827/full#supplementary-material

6. Kim BJ, Jeong JH, Kim JH, Kim HS, Jang HJ. The role of targeted agents in the adjuvant treatment of colon cancer: a meta-analysis of randomized phase III studies and review. Oncotarget (2017) 8(19):31112–8. doi: 10.18632/oncotarget.16091
7. Ellis LM, Fidler II. Finding the tumor copycat. Therapy fails, patients don’t. Nat Med (2010) 16(9):974–5. doi: 10.1038/nm0910-974
8. Mattie M, Christensen A, Chang MS, Yeh W, Said S, Shostak Y, et al. Molecular characterization of patient-derived human pancreatic tumor xenograft models for preclinical and translational development of cancer therapeutics. Neoplasia (2013) 15(10):1138–50. doi: 10.1593/neo.13922
9. Bertotti A, Migliardi G, Galimi F, Sassi F, Torti D, Isella C, et al. A molecularly annotated platform of patient-derived xenografts ("xenopatients") identifies HER2 as an effective therapeutic target in cetuximab-resistant colorectal cancer. Cancer Discov (2011) 1(6):508–23. doi: 10.1158/2159-8290.CD-11-0109
28. Li M, Merino-Trigo A, Lacroix L, Pocard M, Goede D, Mariani P, et al. Characterization of a large panel of patient-derived tumor xenografts representing the clinical heterogeneity of human colorectal cancer. *Clin Cancer Res* (2012) 18(19):5314–28. doi: 10.1158/1078-0432.CCR-12-0372

26. Zhang Z, Li B, Huang J, Huang S, He D, Peng W, et al. A Network Approach to Explore the Mechanisms of Uncaria Alkaloids in Treating Induced Vascular Smooth Muscle Cells. *Drug Des Devel Ther* (2017) 11:476–89. doi: 10.2147/DDDT.S220719

27. Yue GG, Lee JK, Kwok HF, Cheng L, Wong EC, Jiang L, et al. Novel PI3K/AKT targeting anti-angiogenic activities of 4-vinylphenol, a new therapeutic potential of a well-known styrene metabolite. *Sci Rep* (2015) 5:11149. doi: 10.1038/srep11149

25. Berger SI, Iyengar R. Network analyses in systems pharmacology. *Nat Rev Genet* (2015) 16(1):56–64. doi: 10.1038/nrg.2013.11.010

23. Hidalgo M, Amant F, Blankin AV, Budinska E, Byrne AT, Caldas C, et al. Patient-derived xenograft models: an emerging platform for translational cancer research. *Cancer Discov* (2014) 4(9):998–1013. doi: 10.1158/2159-2890.CD-14-0001

24. Pari L, Tewas D, Eckel J. Role of curcumin in health and disease. *Arch Physiol Biochem* (2008) 114(2):117–29. doi: 10.3109/15419061.2011.636465

22. Rowan BG, Gimble JM, Sheng M, Anbalagan M, Jones RK, Frazier TP, et al. Human adipose tissue-derived stromal/stem cells promote migration and early metastasis of triple negative breast cancer xenografts. *PLoS One* (2014) 9(2):e89595. doi: 10.1371/journal.pone.089595

21. Becker M, Nitsche A, Neumann C, Aumann J, Junghahn I, Fichtner I. Sensitive PCR method for the detection and real-time quantification of human cells in xenotransplantation systems. *Br J Cancer* (2002) 87(11):1328–35. doi: 10.1038/sj.bjc.6600573

20. Wu JQ, Niu T, Yang H, Wang Y, Li G, Su Q, et al. Human adipose tissue-derived stromal/stem cells promote migration and early metastasis of triple negative breast cancer xenografts. *PloS One* (2014) 9(2):e89595. doi: 10.1371/journal.pone.089595

19. Liu J, Li Q, Fan Z, Li Q. Cellular signaling pathways implicated in metastasis of colorectal cancer and the associated targeted agents. *Future Oncol* (2015) 11(21):2911–22. doi: 10.2217/fon.15.235

18. Liu SY, Lange A, Frenzler A, Manms MP, Kubicka S, Vogel A. Targeted therapies in metastatic colorectal cancer: a systematic review and assessment of currently available data. *Onco Targets and Therapy* (2014) 19(11):1156–68. doi: 10.1056/thoracolengineering-2014-0032

17. Rahman KM, Desai DC. Epidemiology of digestive tract cancers in India. *V. Large and small bowel*. Indian *J Gastroenterol* (1999) 18(3):118–21.

16. Yue GG, Chan BC, Hon PM, Kennelly EJ, Yeung SK, Cassileth BR, et al. Immunostimulatory activities of polysaccharide extract isolated from *Curcuma longa*. *Int J Biol Macromol* (2010) 47(3):342–7. doi: 10.1016/j.ijbiomac.2010.05.019

15. Yue GG, Chan BC, Hon PM, Fung KP, Leung PC, et al. Evaluation of in vitro anti-proliferative and immunomodulatory activities of compounds isolated from *Curcuma longa*. *Food Chem Toxicol* (2010) 48(8–9):2011–20. doi: 10.1016/j.fct.2010.04.039

14. Yue GG, Chan BC, Hon PM, Kennelly EJ, Yeung SK, Cassileth BR, et al. Immunostimulatory activities of polysaccharide extract isolated from *Curcuma longa*. *Int J Med Macromol* (2010) 47(3):342–7. doi: 10.1016/j.ijjjibio.2010.05.019

13. Mohandas KM, Desai DC. Epidemiology of digestive tract cancers in India. V. Large and small bowel. *Indian J Gastroenterol* (1999) 18(3):118–21.

12. Chauhan DP. Chemotherapeutic potential of curcumin for colorectal cancer. *Curr Pharm Des* (2002) 8(9):1695–706. doi: 10.2174/1381612023394016

11. Yue GG, Li Q, Fan Z, Li Q. Cellular signaling pathways implicated in metastasis of colorectal cancer and the associated targeted agents. *Future Oncol* (2015) 11(21):2911–22. doi: 10.2217/fon.15.235

10. Liu J, Liu Y, Fan Z, Li Q. Cellular signaling pathways implicated in metastasis of colorectal cancer and the associated targeted agents. *Future Oncol* (2015) 11(21):2911–22. doi: 10.2217/fon.15.235

9. Srivastava S, Roy NK, Prasad S, et al. Curcumin, the golden nutraceutical: multitargeting for multiple chronic diseases. *Br J Pharmacol* (2017) 174(11):1325–48. doi: 10.1111/bph.13621

8. Rahimi HR, Naneinina R, Sepahni Shamloo A, Nikdoust S, Kazemi Oskuee R. Novel delivery system for natural products: Nano-curcumin formulations. *Asian J Phytomed* (2016) 6(4):383–98.

7. Li M, Yue GG, Tsui SK, Fung KP, Lau CB. Turmeric extract, with absorbable curcumin, has potent anti-metastatic effect in vitro and in vivo. *Phytochemistry* (2018) 146:131–41. doi: 10.1016/j.phytochemical.2018.03.065

6. de Campos PS, Matte BF, Diel LF, Jesus LH, Bernardi L, Alves LM, et al. Low Doses of Curcuma longa Modules Cell Migration and Cell-Cell Adhesion. *Phytother Res* (2017) 31(9):1433–40. doi: 10.1002/ptr.5872

5. Fleming M, Ravula S, Tattishchev SF, Wang HL. Colorectal carcinoma: Pathologic aspects. *J Gastrointest Oncol* (2012) 3(3):153–73. doi: 10.3978/j.issn.2078-6891.2012.030

4. Gregoire A, Clevers H. Wnt signaling in the intestinal epithelium: from endoderm to cancer. *Genes Dev* (2005) 19(8):877–90. doi: 10.1101/gad.1295405

3. Baylin SB, Ohm JE. Epigenetic gene silencing in cancer - a mechanism for early oncogenic pathway addiction? *Nat Rev Cancer* (2006) 6(2):107–16. doi: 10.1038/nrc1799

2. Sato H, Suzuki H, Toyota M, Nojima M, Maruyama R, Sasaki S, et al. Frequent epigenetic inactivation of DKK1-KOFP family genes in human gastrointestinal tumors. *Carcinogenesis* (2007) 28(12):2459–66. doi: 10.1093/carcin/bgm178

1. Aguilera G, Fraga MF, Ballestar E, Paz MF, Herranz M, Espada J, et al. Epigenetic inactivation of the Wnt antagonist DKK1-KOFP-1 (DKK1-) gene in human colorectal cancer. *Oncogene* (2006) 25(29):4116–21. doi: 10.1038/sj.onc.1209439
49. Thorvaldsen TE, Pedersen NM, Wenzel EM, Stenmark H. Differential Roles of AXIN1 and AXIN2 in Tankyrase Inhibitor-Induced Formation of Degradasomes and beta-Catenin Degradation. *PloS One* (2017) 12(1): e0170508. doi: 10.1371/journal.pone.0170508

50. Wu ZQ, Brabletz T, Fearon E, Willis AL, Hu CY, Li XY, et al. Canonical Wnt suppressor, Axin2, promotes colon carcinoma oncogenic activity. *Proc Natl Acad Sci U S A* (2012) 109(28):11312–7. doi: 10.1073/pnas.1203015109

51. Broderick P, Carvajal-Carmona I, Pittman AM, Webb E, Howarth K, Rowan A, et al. A genome-wide association study shows that common alleles of SMAD7 influence colorectal cancer risk. *Nat Genet* (2007) 39(11):1315–7. doi: 10.1038/ng.2007.18

52. Anastas JN, Moon RT. WNT signalling pathways as therapeutic targets in cancer. *Nat Rev Cancer* (2013) 13(1):11–26. doi: 10.1038/nrc3419

53. Chen A, Xu J, Johnson AC. Curcumin inhibits human colon cancer cell growth by suppressing gene expression of epidermal growth factor receptor through reducing the activity of the transcription factor Egr-1. *Oncogene* (2006) 25(2):278–87. doi: 10.1038/sj.onc.1209019

54. Frame MC. Src in cancer: deregulation and consequences for cell behaviour. *Biochim Biophys Acta* (2002) 1602(2):114–30. doi: 10.1016/s0304-419x(02)00040-9

55. Villegas I, Sanchez-Fidalgo S, Alarcon de la Lastra C. New mechanisms and therapeutic potential of curcumin for colorectal cancer. *Mol Nutr Food Res* (2008) 52(9):1040–61. doi: 10.1002/mnfr.200700280

56. Teiten MH, Eifes S, Dicato M, Diederich M. Curcumin—the paradigm of a multi-target natural compound with applications in cancer prevention and treatment. *Toxins* (Basel) (2010) 2(1):128–62. doi: 10.3390/toxins2010128

57. Yu Y, Kanwar SS, Patel BB, Nautiyal J, Sarkar FH, Majumdar AP. Elimination of Colon Cancer Stem-Like Cells by the Combination of Curcumin and FOLFOX. *Transl Oncol* (2009) 2(4):321–8. doi: 10.1593/tno.09193

58. Schuell B, Gruenberger T, Kornek GV, Dworan N, Depisch D, Lang F, et al. Side effects during chemotherapy predict tumour response in advanced colorectal cancer. *Br J Cancer* (2005) 93(7):744–8. doi: 10.1038/sj.bjc.6602783

59. Wiel-Ahoenska A, Kowalska T, Filipczyk-Cisarz E, Lapinski L, Nartowski K. Evaluation of the toxicity of anticancer chemotherapy in patients with colon cancer. *Adv Clin Exp Med* (2013) 24(1):103–11. doi: 10.17219/acecm/38154

**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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