

# Significance of Irisin (FNDC5) Expression in Colorectal Cancer

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**Abstract.** *Background/Aim: The role of irisin, the extracellular part of fibronectin type III domain containing 5 (FNDC5), in colorectal cancer (CRC) is unclear. The aim of this study was to investigate immunohistochemical (IHC) expression level of irisin and correlations with clinicopathological factors in patients with CRC. Materials and Methods: We collected 222 archived CRC samples and 26 control samples from autopsies conducted at the Department of Forensic Medicine. They were used to perform IHC reactions detecting irisin, Ki-67, minichromosome maintenance protein complex component 3 (MCM3), and urine diphosphate-galactose ceramide galactosyltransferase (UGT3) expression. The correlations with Ki-67, MCM3, and UGT3 were analyzed. Irisin expression was also evaluated in cancer cell lines by immunofluorescence reaction and western blot. Results: Irisin expression was higher in cancer cells compared to the control tissues ( $p < 0.0001$ ). Irisin expression was significantly higher in stage I than in stage III ( $p = 0.013$ ) and IV CRC ( $p = 0.05$ ). Conclusion: The correlation between higher expression of irisin and cancer stages indicates its potential usefulness as a marker in CRC.*

Currently, colorectal cancer (CRC) is the world's fourth most deadly cancer, causing almost 900,000 deaths annually (1). It is the second most commonly diagnosed cancer annually and the second most common cause of death among all types of cancer in Europe – 500,000 new cases and 243,000 deaths were reported in 2018 (2). The morbidity and mortality in the

USA are similar – approximately 135,000 new cases and more than 50,000 deaths in 2017 (3). In Poland, between 1980 and 2013, there was a 4.2-fold increase in CRC morbidity in men (from 2,288 to 9,548) and a 3.2-fold increase in women (from 2,432 to 7,902), with 75% of new cases being diagnosed in patients over 60 years of age in both sexes. In 2017, CRC was diagnosed in 10,905 patients from the Polish population (5,073 women and 5,832 men) (4). The etiopathology of this cancer is multifactorial, and the main causes, which are well known, include an aging population, obesity (due to dietary habits), lack of physical exercise, smoking, and immune system pathology (1, 5, 6). The response of the immune system to pathological signs may lead to carcinogenesis (7, 8).

The extracellular peptide known as irisin is released from skeletal muscle by an unknown protease acting on fibronectin type III domain containing 5 (FNDC5), and plays an essential role in regulating energy homeostasis. It was detected for the first time by Boström *et al.* in 2012 (9). Apart from converting different adipose tissues, it is involved in other metabolic activities, *e.g.*, inflammatory processes and aging (10). Moreover, increased irisin expression was reported in some digestive tract malignancies (*e.g.*, hepatocellular, colon, esophageal, and gastric cancer) (11-13). Its role in pancreatic cancer is unclear (14, 15). Therefore, we were interested in understanding the role of associated factors (including irisin) in the progression of CRC.

Our study is the continuation of our research on irisin expression in lung cancer (16). We observed increased irisin expression in cancer cells and stromal cells of lung cancer. Higher irisin expression in stromal cells of lung cancer was observed in patients with shorter overall survival. Moreover, irisin expression correlated with Ki-67 expression in stromal cells of lung cancer. Therefore, we investigated irisin expression in CRC. The Ki-67 antigen is routinely used in the diagnostic process to check the proliferative ability of different cancer types. Minichromosome maintenance protein complex component 3 (MCM3) is also a proliferation

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marker. This protein is a part of the MCM complex, which participates in replication. Its increased expression has been observed in various cancer types (17, 18).

Additionally, urine diphosphate-galactose ceramide galactosyltransferase (UGT8) influences sulfatide biosynthesis and is also related to cell proliferation. Increased sulfatide activates integrin  $\alpha V\beta 5$ -mediated signaling, which contributes to breast cancer progression (19). Research has also demonstrated that  $\alpha V$  integrins are receptors for irisin in osteocytes (20). Therefore, we investigated irisin expression in CRC and its correlation with the markers Ki-67, MCM3, and UGT8. Furthermore, the aim of this study was to evaluate irisin expression in association with clinicopathological parameters in CRC.

## Materials and Methods

**Patient characteristics.** This was a retrospective study conducted on 222 archived paraffin blocks from patients with CRC (women n=90; men n=132) (Table I). The specimens were collected between 2012 and 2016 during surgical removal of tumor. The surgical procedures were conducted at the Wrocław Military Hospital. Patients whose specimens were analyzed had a sedentary lifestyle. They had not undergone chemo- or radiotherapy before surgery. Patients were excluded from the study if they were diagnosed with any other cancer. All the patients in the analyzed group were diagnosed with CRC. The histopathological evaluation was made in accordance with the World Health Organization criteria, and the pathological staging was standardized according to the eighth TNM edition (21).

The control group (26 specimens) included samples from autopsies conducted at the Department of Forensic Medicine in Wrocław. Tissues were collected from every segment of the colon. Only patients with sudden death were included. The exclusion criteria were as follows: Other neoplasms, gastrointestinal tract surgery, intestinal inflammation, or intoxication. Samples of poor quality were excluded from further analysis. The project was approved by the Bioethics Committee at Wrocław Medical University (no. 604/2018)

**Cell line and culture conditions.** Three intestinal cancer cell lines were used in *in vitro* studies, namely CaCo-2, LoVo and HT-29, and the normal intestinal cell line CCD-18Co. Appropriate culture media were used to culture the cell lines. CaCo-2 cells were grown in minimum essential medium (Lonza, Switzerland, Basel), LoVo in F-12K medium (American Type Culture Collection, Manassas, VA, USA), HT-29 in McCoy's 5A medium (Merck KGaA, Darmstadt, Germany), and CCD-18Co in Eagle's minimum essential medium (Lonza). All culture media were supplemented with 10% fetal bovine serum (Merck KGaA) except for the CaCo-2 cell culture medium to which 20% fetal bovine serum (Merck KGaA) and 1% non-essential amino acids (Merck KGaA) were added. All culture media were also supplemented with 1% penicillin/streptomycin (Merck KGaA). Cell cultures were maintained in a humid atmosphere at 37°C with 5% CO<sub>2</sub>. Cells were trypsinized with 0.25% trypsin (Merck KGaA).

**Tissue microarray (TMA) preparation.** The material for the study consisted of 222 archived paraffin-embedded samples of colon

Table I. Clinicopathological characteristics of patients (n=222) related to irisin expression in colorectal cancer cells.

Clinicopathological parameter	Frequency, n (%)	Irisin expression in CRC cells		p-Value*
		Low (<6)	High (≥6)	
Age				0.6268
≤60 Years	51 (22.96)	21	30	
>60 Years	171 (77.04)	77	94	
Gender				0.3680
Male	132 (59.47)	55	77	
Female	90 (40.53)	43	47	
Tumor size				0.5439
T1	1 (0.89)	0	1	
T2	28 (24.56)	7	21	
T3	79 (69.29)	27	52	
T4	6 (5.26)	3	3	
Lymph node status				0.1776
N0	70 (61.40)	26	44	
N1-N3	44 (38.60)	11	33	
Metastasis				0.1283
M0	95 (83.34)	28	67	
M1	19 (16.66)	9	10	
Stage				0.3340
I	22 (19.30)	5	17	
II	40 (35.08)	14	26	
III	33 (28.94)	9	24	
IV	19 (16.66)	9	10	
Grade of malignancy				0.2844
1	44 (19.81)	16	28	
2	156 (70.27)	68	88	
3	19 (8.56)	11	8	
Ki67 score index				<b>0.0352</b>
1-2	107 (48.27)	51	56	
3-4	115 (51.73)	71	44	
MCM3 score index				<b>0.0189</b>
1-2	84 (37.93)	59	25	
3-4	138 (62.07)	75	63	
UGT8 score index				0.1045
1-6	128 (57.5)	50	78	
7-12	94 (42.5)	47	47	

MCM3: Minichromosome maintenance protein complex component 3; UGT8: urine diphosphate-galactose ceramide galactosyltransferase. \*By chi-squared test. Statistically significant p-values are shown in bold.

adenocarcinoma. Sections 7-μm-thick were stained with hematoxylin and eosin and later scanned using a histology scanner Panoramic MIDI (3DHitech, Budapest, Hungary) under 20× magnification to create virtual slides. Scans were examined by two independent pathologists, and representative spots were selected to create microarrays (three spots from each block, 1.5 mm diameter each). TMAs were created using TMA Grand Master automatic system (3DHitech).

**Immunohistochemistry (IHC) reactions.** IHC was performed on 4-μm-thick sections obtained from the TMA blocks using a Link48 Autostainer (Dako, Glostrup, Denmark). To deparaffinize, rehydrate

and unmask the epitope, the slides were boiled in EnVision FLEX Target Retrieval Solution (97°C, 20 min; pH 9) in PT-Link. The activity of endogenous peroxidase was blocked by 5-min incubation with EnVision FLEX Peroxidase-Blocking Reagent (Dako). Next, primary rabbit polyclonal anti-irisin/FNDC5 antibody (NBP2-14024; Novus Biologicals, Littleton, CO, USA) was applied at 1:300 for 1 h. Next, the slides were incubated with EnVision FLEX/horseradish peroxidase (HRP) for 20 min. Finally, a substrate for HRP (3,3'-diaminobenzidine) was added for 10-min incubation. All sections were counterstained with EnVision FLEX Hematoxylin (Dako) for 5 min. After dehydration in graded ethanol concentrations (70%, 96%, 99.8%) and xylene, the slides were sealed with coverslips in Dako Mounting Medium (Dako). After IHC, TMAs were analyzed to evaluate irisin expression using a BX41 Optical Microscope (Olympus Hamburg, Germany) at  $\times 200$  magnification. The semiquantitative immunoreactive score by Remmele and Stegner (22) was used for the assessment of cytoplasmic irisin and UGT8 reaction. The final result was the product of the scores obtained by the estimation of the intensity of the color reaction (*i.e.*, 1 point for weak, 2 point for moderate, 3 point for strong reaction) and the percentage of positively stained cancer cells (0 points: no expression, 1 point: 1-10%, 2 points: 11-50%, 3 points: 51-80%, 4 points: >80%) (22). Nuclear IHC reaction of Ki-67 and MCM3 were determined with the use of the five-point evaluation scale (0-no expression, 1 point—1%-10%, 2 points—11%–25%, 3 points—26%–50%, 4 points >50%). The assessment was performed by two independent pathologists. Discrepancies were re-evaluated until consensus.

**Immunofluorescence.** Caco-2, Lovo, HT-29, and CCD-18Co cell lines were set up at  $2 \times 10^4$  cells per well in Millicell EZ 8-well glass slides (Merck KGa) in appropriate medium overnight.

The cells were then fixed with 4% paraformaldehyde for 12 min at room temperature and permeabilized using 0.2% Triton X-100 for 10 min. Non-specific binding was blocked using 3% bovine serum albumin (BSA) in 0.1% phosphate-buffered saline buffer (1 h at room temperature). The cells were incubated overnight at 4°C with primary anti-Irisin/FNDC5 (dilution 1:200; NBP2-14024; Novus Biologicals). Subsequently, secondary anti-rabbit Alexa Fluor 568 was applied at 1:2,000 (ab175470; Abcam, Cambridge, UK), for 1 h at room temperature. Negative controls were performed with 1% BSA in phosphate-buffered saline instead of the specific antibody. The preparations were mounted in the ProLong Gold Antifade Mountant with 4',6-diamidino-2-phenylindole (Thermo Fisher Scientific, Waltham, MA, USA). A Fluoview FV3000 (Olympus) confocal laser scanning microscope coupled with CellSense software (Olympus) was used for evaluating fluorescence at a magnification of  $\times 600$ .

**Protein isolation and western blot.** The whole-cell protein lysates from CCD-18Co, HT-29, CaCo-2, and LoVo cell lines were extracted using CellLytic™ MT Cell Lysis Reagent (Sigma-Aldrich, Saint Louis, MO, USA) with the addition of Halt™ Protease Inhibitor Cocktail (Thermo Fisher Scientific, Waltham, MA, USA) and 0.2 mM PMSF Protease Inhibitor (Sigma-Aldrich). Protein concentrations were quantified with a Pierce BCA Protein Assay Kit (Thermo Fisher Scientific) and a NanoDrop™ 1000 (Thermo Fisher Scientific) spectrophotometer. Equal amounts of total protein (30  $\mu$ g) were mixed with the Laemmli sample buffer (23) and resolved on 10% acrylamide gel by sodium dodecyl sulphate-polyacrylamide gel

electrophoresis. After electrophoresis, the samples were transferred to Immobilon-P polyvinylidene difluoride membranes (Merck Darmstadt, Germany) in the XCell SureLock™ Mini-Cell Electrophoresis System (Thermo Fisher Scientific). Next, the membranes were blocked in 4% BSA (Merck KGaA) solution in TBST buffer (0.2 M Tris; 1.5 M NaCl; 0.1% Tween-20). After blocking, the membranes were incubated overnight at 4°C with primary polyclonal rabbit antibody to irisin/FNDC5 (NBP2-14024; Novus Biologicals, Centennial, CO, USA) diluted at 1:200. The membranes were also incubated for 1 h at room temperature with secondary HRP-conjugated donkey anti-rabbit antibody (711-035-152; Jackson ImmunoResearch, West Grove, PA, USA) diluted at 1:3,000. Finally, the membranes were rinsed and treated with Luminata Classico (Merck KGaA) chemiluminescent substrate. The data were collected using the ChemiDoc Imaging System (Bio-Rad, Hercules, CA, USA).  $\beta$ -Tubulin, detected with primary rabbit anti-human  $\beta$ -tubulin (ab6046; Abcam) diluted at 1:1,000, and the secondary HRP-conjugated donkey anti-rabbit antibody (711-035-152; Jackson ImmunoResearch) diluted at 1:3,000, was used as an internal control to normalize the amounts of irisin/FNDC5. The densitometric analysis of the results was performed using Image Lab software suite (Bio-Rad). The experiment was performed in triplicate.

**Statistical analysis.** The associations between the IHC expression of irisin/FNDC5 and clinical parameters were analyzed statistically with GraphPad Prism 5.0 (La Jolla, CA, USA). The data distribution was evaluated with the Kolmogorov-Smirnov test. All the quantitative variables were described as medians and ranges. The Mann-Whitney *U*-test, the Kruskal-Wallis test, and the  $\chi^2$  test were used to compare the analyzed groups. Spearman's rank correlation was used to evaluate the association between the expression of the Ki-67 antigen, MCM3 and UGT8 and that of irisin in cancer cells. Kaplan-Meier analysis and the log-rank test were used to verify the relationship between the intensity of irisin expression and patient overall (OS) and event-free (EFS) survival. The results were considered statistically significant when  $p < 0.05$ .

## Results

**IHC evaluation of irisin expression levels in CRC TMAs.** We found weak or no expression of irisin in normal colorectal cells in 26 cases (Figure 1A); strong irisin expression was present only in goblet cells (Figure 1B). In CRC cells, the IHC reaction showed irisin in the cytoplasm (Figure 1C-E). In CRC, we also observed the presence of irisin in the apical part of cells and the extracellular space (Figure 1D). The mean irisin expression level in CRC cells was significantly higher than in normal colorectal cells [ $6.51 \pm 2.46$  (SD) vs.  $4.86 \pm 1.48$  (SD), respectively;  $p = 0.0016$ , Mann-Whitney *U*-test].

**Associations between irisin expression in CRC cells and clinicopathological parameters.** We compared the mean values of irisin expression and their relationship with the clinicopathological parameters (Table I and Table II). We observed changes in irisin levels according to tumor size, grade of malignancy, lymph node status, and distant metastasis. Irisin levels decreased with higher T-status.

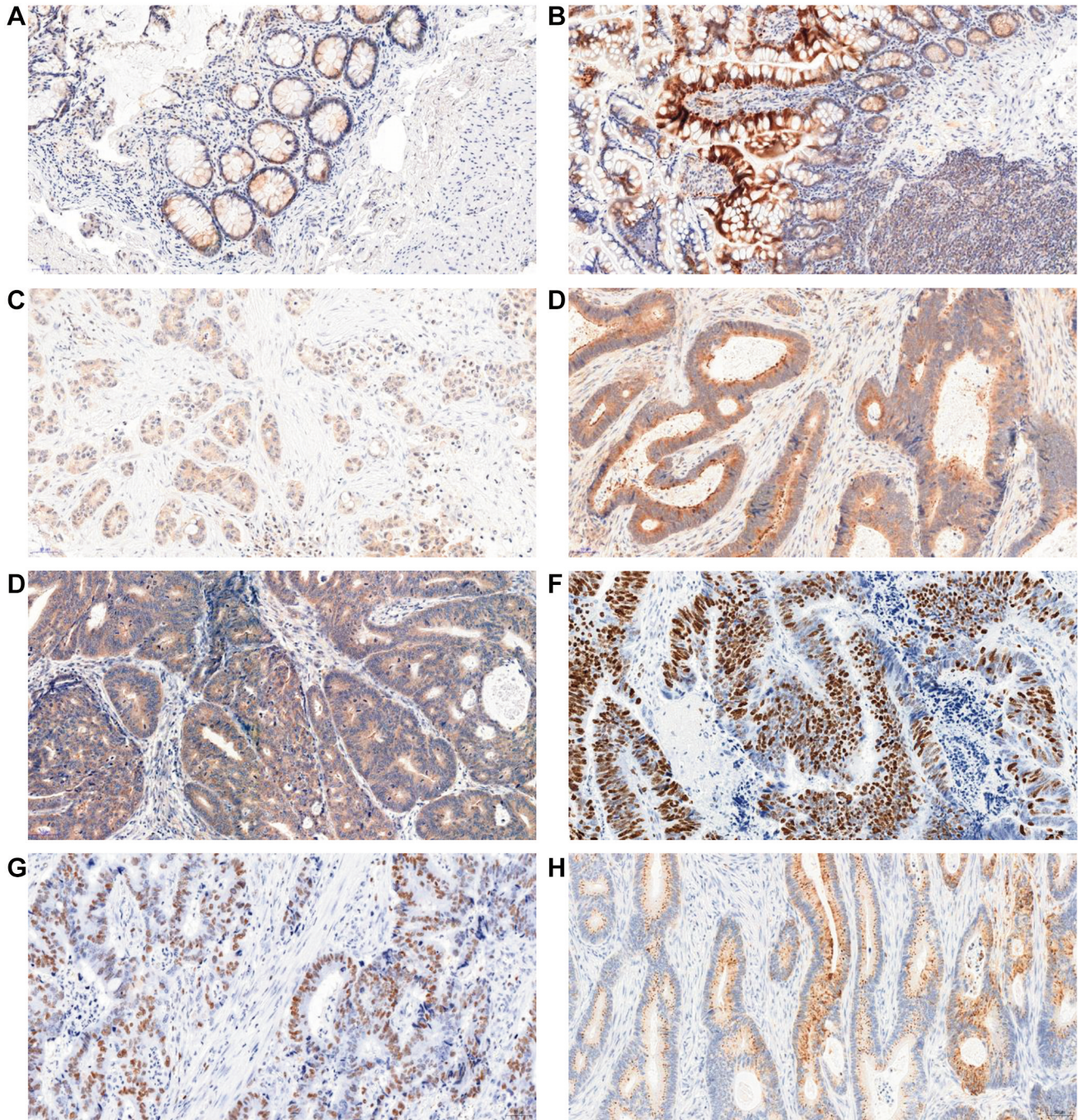


Figure 1. Immunohistochemical reactions (brown color) for irisin, Ki-67, minichromosome maintenance protein complex component 3 (MCM3) and urine diphosphate-galactose ceramide galactosyltransferase (UGT8) expression in colorectal cancer (CRC). Irisin cytoplasm expression in healthy colorectal tissue (A: no expression, B: expression in goblet cells) and in different grades of malignancy of CRC (C: grade 1, D: grade 2, E: grade 3) as evidenced by brown-stained cytoplasm. CRC tissue was also positive for expression of Ki-67 (F), MCM3 (G) and UGT8 (H); magnification, ×200.

Differences between T1-T2 and T3-T4 were of borderline significance (Mann-Whitney *U*-test,  $p=0.052$ ). The differences in irisin expression levels between groups of patients according to the pN or pM status were not

statistically significant. We analyzed irisin expression in different CRC stages and observed significant differences between stages I and III (Mann-Whitney *U*-test,  $p=0.01$ ) and between stages I and IV (Mann-Whitney *U*-test,  $p=0.05$ ).

Table II. Associations of irisin expression level with clinicopathological characteristics in patients with colorectal cancer.

Parameter	Subgroup	Mean value±SD	Comparison	p-Value*
Lymph nodes	N0	6.046±2.23	N0 vs. N1-N3	0.346
	N1-N3	5.830±2.07		
Grade of malignancy	1	6.206±2.53	1 vs. 2	0.6345
	2	5.964±2.03	1 vs. 3	0.3020
	3	5.438±2.44	2 vs. 3	0.2416
Tumor size	T1	8.00±0.0	T1-T2 vs. T3-T4	0.052
	T2	7.49±2.24	T2 vs. T3	0.085
	T3	6.23±2.54	T2 vs. T4	0.163
	T4	5.93±1.90	T3 vs. T4	0.652
Stage	I	6.75±2.0	I vs. II	0.077
	II	5.82±2.2	I vs. III	<b>0.013</b>
	III	5.73±2.2	I vs. IV	0.050
	IV	5.67±1.6	II vs. III	0.438
			II vs. IV	0.618
Metastasis			III vs. IV	0.981
	M0	5.98±2.21	M0 vs. M1	0.796
	M1	5.83±1.73		

CRC: Colorectal cancer Statistically significant p-values are shown in bold. \*Mann-Whitney U-test.

*Associations between the expression of irisin and OS and EFS.* We did not observe any differences in OS according to irisin expression in cancer cells in patients with CRC (log-rank Mantel-Cox test;  $p=0.46$ ) (Figure 2A). However, longer EFS was observed in patients with higher irisin expression (Figure 2B). This relationship was close to the statistical significance (log-rank Mantel-Cox test;  $p=0.09$ ).

The results of the univariate and multivariate analyses are given in Table III. The multivariate analysis showed that a high grade of malignancy was related to shorter OS. Additionally, advanced pT and pN status and the stage of CRC were related to shorter OS.

*Associations between the proliferation of cancer cells and irisin expression.* We investigated the relationship between irisin levels and CRC cell proliferation. For this purpose, we compared the level of irisin with the proliferation markers such as the Ki-67 antigen, UGT8, and the MCM3 protein. Irisin expression in CRC cells weakly positively correlated with the expression of the Ki-67 antigen ( $r=0.20$ ,  $p=0.0395$ ). Moreover, we observed a weakly positive correlation between irisin and MCM3 expression ( $r=0.26$ ,  $p=0.0091$ ) and a medium positive correlation with UGT-8 ( $r=0.34$ ,  $p=0.0005$ ). Figure 3 shows plots of correlations with the markers of proliferation.

*Irisin expression in CRC cell lines.* We analyzed irisin expression of the normal colon cell line (CCD-18Co) compared to cancer cell lines (LoVo, HT29 and CaCo-2) by

western blot and immunofluorescence reaction. The comparison of irisin levels is given in Figure 4. We observed higher expression of irisin in cancer cells compared to CCD-18Co cells. The highest irisin level was found in LoVo cells, which are cells derived from a metastatic site.

## Discussion

We performed our study on a large group of 222 patients to investigate irisin expression in CRC tissue specimens in association with different clinical parameters describing CRC. We also correlated the irisin expression by IHC with Ki-67, MCM3, and UGT8 in CRC. Additionally, we used an *in vitro* model and analyzed the differences in immunofluorescence between normal colon cells (CCD-18Co) and selected colon cancer lines (CaCo-2, LoVo and HT-29). Increased irisin concentration is responsible for a decrease in adenosine triphosphate (ATP) production. However, increased heat production is observed (24). We decided to investigate this molecule due to its importance – an increase in local heat can kill cancer cells by coagulating their proteins and destroys their small blood vessels (25).

Aydin *et al.* found that irisin was significantly increased in colon cancer cells (26). They suggested two explanations for this finding. Firstly, that it might be related to adaptation to changes in cell conditions (adaptation to inhibit ATP generation). Secondly, an increased irisin level raises the conversion of white adipose tissue to brown adipose tissue, which leads to the release of more heat through mitochondria that contain uncoupling protein 1 (the molecule that controls adaptive thermogenesis by the release of heat) rather than ATP (9, 26, 27).

We confirmed the above finding, obtained on tissues by IHC method, using immunofluorescence reaction in the selected colon cancer cells lines (CaCo, LoVo and HT29) compared to normal colon cells. Additionally, we found higher expression of irisin by immunofluorescence in CRC cell lines. Irisin is useful to differentiate between stage I (initial colon cancer) and more advanced stages of CRC. The levels of many molecules are increased in tumors. We revealed statistically significant correlations of irisin with Ki-67, MCM3, and UGT3.

According to Shi *et al.*, irisin stimulated cell proliferation, migration, and invasion of hepatocellular carcinoma. Furthermore, it reduced the cytotoxicity of doxorubicin in HepG2 cells, which suggests that an increased expression of irisin might have a protective role in liver cancer (13). These findings are not in line with those revealed by Aydin *et al.* (26), who did not report increased irisin levels in liver cancer (28). Zhu *et al.* showed that a reduced serum irisin level was detected in patients with CRC. Their research suggested irisin might be a new important diagnostic indicator for CRC (29). Zhang *et al.* found that irisin inhibited cell

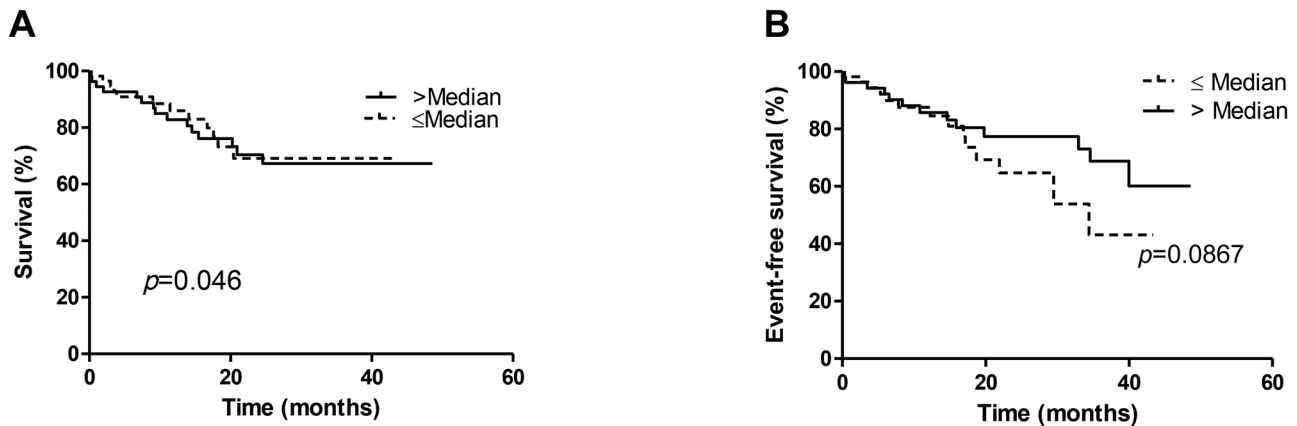


Figure 2. Kaplan-Meier survival curves presenting the prognostic impact of irisin expression levels detected by immunohistochemistry in cancer cells of patients with colorectal cancer on overall survival (A) and event-free survival (B). Patient groups were compared according to the median value of the immunohistochemical expression of irisin in colorectal cancer cells.

Table III. Univariate and multivariate Cox proportional hazards analyses in patients with colorectal cancer.

Parameter	Subgroup comparison	Univariate analysis		Multivariate analysis	
		HR (95% CI)	p-Value	HR (95% CI)	p-Value
Age	≤60 vs. >60 Years	0.46 (0.17-1.16)	0.1050		
Gender	Male vs. female	0.98 (0.46-2.03)	0.9548		
pT	1-2 vs. 3-4	2.58 (1.17-5.73)	<b>0.0192</b>	2.00 (0.87-4.63)	0.2016
pN	N0 vs. N+	1.93 (0.90-4.10)	0.0891		
Grade	1 vs. 2-3	3.07 (1.37-6.87)	<b>0.0062</b>	4.86 (1.14-20.70)	<b>0.0320</b>
Stage	I-II vs. III-IV	2.92 (1.40-6.12)	<b>0.0043</b>	2.28 (0.63-8.06)	0.1040
Irisin	<25% vs. ≥25%	1.41 (0.61-3.32)	0.4570		
Ki-67	<25% vs. ≥25%	0.97 (0.51-1.85)	0.9626		
MCM3	<25% vs. ≥25%	0.91 (0.47-1.81)	0.8158		

CI: Confidence interval; HR: hazard ratio; MCM3: minichromosome maintenance protein complex component 3. Statistically significant *p*-values are shown in bold.

proliferation, induced apoptosis, and reduced the migration and invasion of pancreatic cancer cells (15). These findings were the basis for our study on the role of irisin in CRC. Liu *et al.* showed that irisin suppressed the migration and invasion of selected pancreatic cancer cell lines (14). Moon *et al.* found that irisin had no effect on cell proliferation, adhesion, or malignant potential of obesity-related cancer cell lines (12).

In our opinion, the IHC analysis of irisin expression of cancer tissues offers considerable advantages over serum-based techniques. This protein is mostly produced in the muscles after exercise, and various tissues have different irisin concentrations. The serum level of this peptide shows fluctuation, which is related to discrepancies when the serum irisin level is correlated with clinical cancer parameters. We are of the opinion that irisin is a useful protein that can be used in further analysis. Irisin has been found in different

cells (*e.g.*, in muscles, pancreas, liver, brain, adipose tissue). It plays a role in regulating fat metabolism but is not related to specific cell physiology. Therefore, this protein may be responsible for energy balance of all cancer cells, thereby releasing the additional local heat.

We can support the findings for CRC clinical staging. We observed no correlation between IHC irisin expression and cancer cell malignancy or a clinical stage.

Altay *et al.* observed increased expression of FNDC5 in adipose tissue in mice with induced gastric cancer, but no expression in the gastric cancer group (30). We revealed the increased expression of irisin in CRC cells, which is not in line with the findings of the above authors.

We speculate that irisin may be responsible for thermal conditions inside CRC cells, thus leading to the progression of weight loss in cachexia. If so, further analysis is warranted to improve the treatment of cancer cachexia. This process is

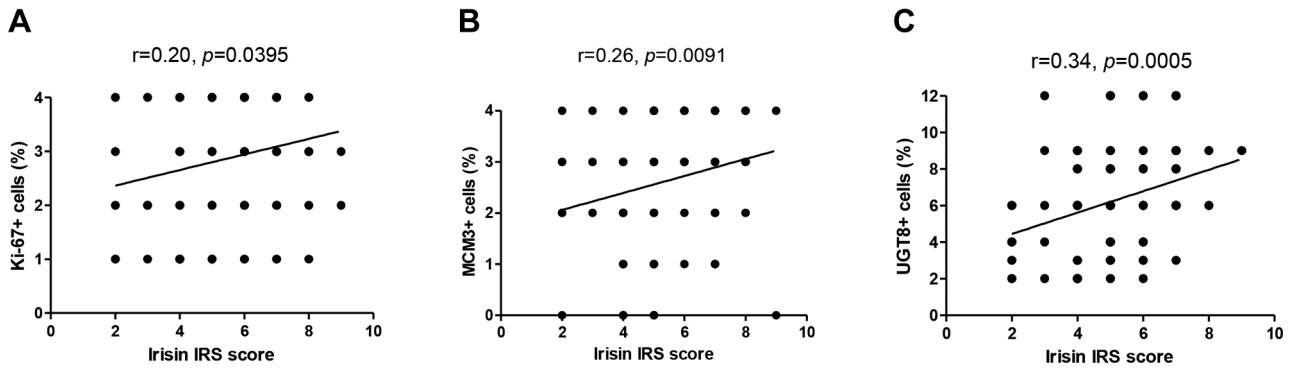


Figure 3. Correlations between irisin expression level and proliferation markers Ki-67 (A), minichromosome maintenance protein complex component 3 (MCM3) (B) and urine diphosphate-galactose ceramide galactosyltransferase (UGT8) (C) in colorectal cancer cells from patients. IRS: Immunoreactive score.

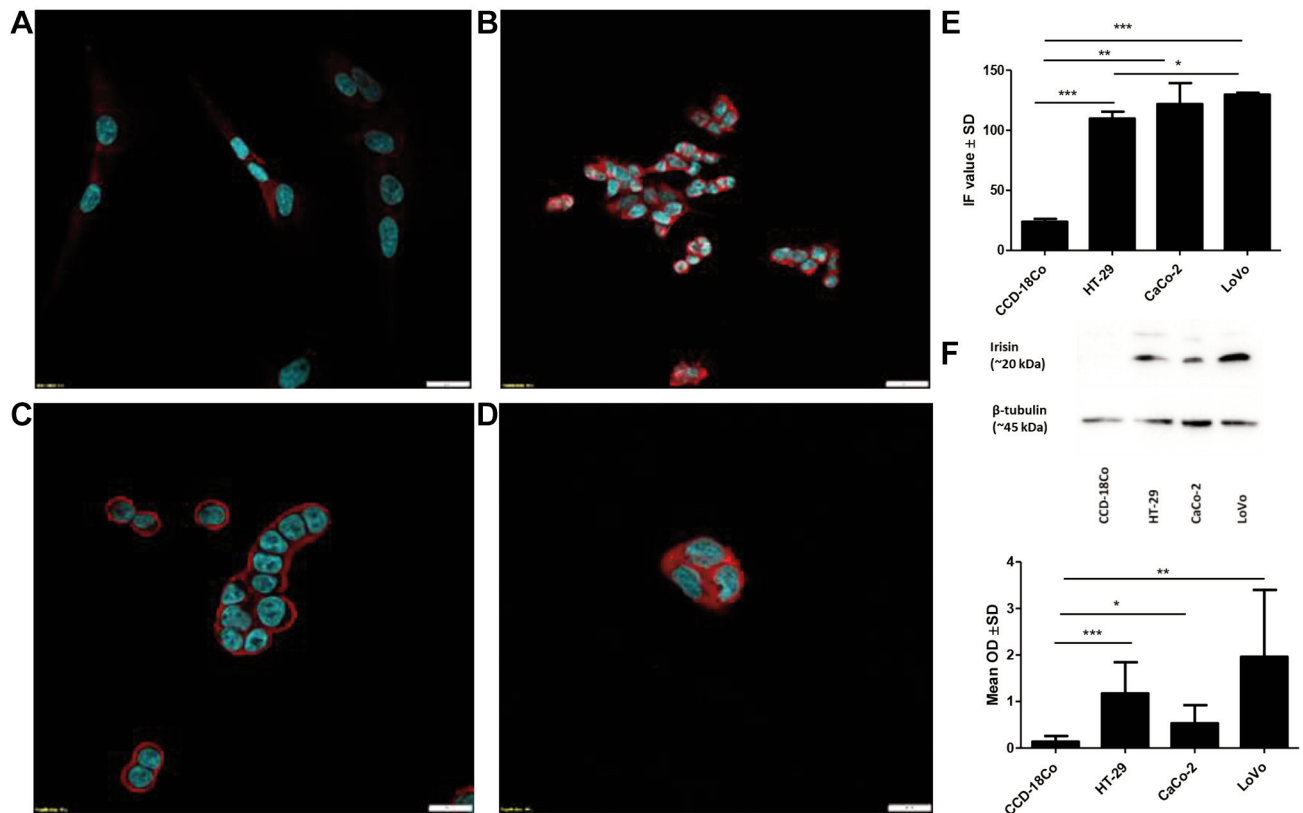


Figure 4. Comparison of irisin expression levels detected by the immunofluorescence (IF) method in normal colon CCD-18Co cells (A) and in different colorectal cancer cell lines (B: HT-29; C: CaCo-2; D: LoVo). Bars=20  $\mu$ m. The level of irisin expression was quantified by IF value (E) and western-blot densitometry (F) in normal and cancer cells. Significantly different at \* $p \leq 0.05$ , \*\* $p \leq 0.005$  and \*\*\* $p \leq 0.001$ .

a multifactorial disease and can be found in 50-80% of patients with cancer (31). We speculate that the change in irisin production is observed during cancer progression, and is reflected by the differences in its level when stage I is compared to stage III or IV.

We speculate that different results of intracellular irisin levels can mirror the differences in the number or functions of irisin in different types of gastrointestinal cancer, and hence reflects the different results in gastric, liver, and colorectal cancer. There is a possibility that the situation in

the gastrointestinal tract is similar to that of adipose tissue, in which the level of irisin expression is different in brown, white and beige tissues. Therefore, the final results in various types of cancer are also divergent. However, the data from our study may be useful for understanding the role of this peptide in CRC progression.

## Conflicts of Interest

The Authors declare no conflicts of interest.

## Authors' Contributions

KN: methodology, formal analysis and writing; SW: investigation, resources formal analysis, validation and writing; MC: data curation; PD: review and editing.

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