

*Research Article***Study of Immunohistochemical Expression of Ezrin and Icam-1 in Colorectal Adenocarcinoma**

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Abstract

Background: Colorectal cancer is one of the most common cancers worldwide. Colorectal carcinogenesis is a complex process resulting from the cumulative effect of sequential genetic alterations. Ezrin is involved in the regulation of cellular architecture, controlling cellular adhesion and motility which are essential for the entire process of tumorigenesis and distant metastasis. Intercellular adhesion molecule-1 (ICAM-1) is a member of the immunoglobulin superfamily and is expressed on many cell types including cancer cells where it can be upregulated in response to various cytokines. **Aim:** The present work was conducted to study the immunohistochemical expression of Ezrin and ICAM-1 in different types of colorectal carcinoma (CRC), to evaluate their expression patterns and examine their association with various clinicopathological features and finally to investigate the presence of a possible relationship between both markers in CRC. **Material and Methods:** The present study comprised one hundred and two (102) cases of primary colorectal carcinomas including; 89 (87.2 %) cases of adenocarcinoma, 7 (6.8%) cases of mucoid carcinoma and 6 (5.9%) cases of signet ring cell carcinoma. Sections from 20 tissue blocks of adjacent normal colorectal epithelium were also immunohistochemically stained as the control group. Thirty four lymph node metastases were also evaluated for Ezrin and ICAM-1 expression. Immunohistochemistry for Ezrin and ICAM-1 antibodies was performed using the avidin biotin-peroxidase complex method. **Results:** The present study identified Ezrin expression in 97.1% of colorectal carcinoma cases Ezrin expression was significantly higher in carcinoma cases compared to adjacent normal colorectal epithelium with different Ezrin localization reflecting different functions in both locations. Cytoplasmic Ezrin expression was significantly associated with lymph node spread ($p < 0.0001$) lymphovascular invasion ($p = 0.038$), advanced TNM ($p < 0.0001$) and Dukes stages ($p < 0.0001$). Regarding membranous Ezrin expression, statistically significant positive associations were detected between membranous expression of Ezrin and nodal metastasis ($p = 0.02$), advanced TNM stage ($p = 0.03$) and Dukes stage ($p = 0.02$). Regarding ICAM-1 expression, the present study identified ICAM-1 expression in 35.3% of carcinoma cases while it was undetectable in normal epithelium. Cytoplasmic ICAM-1 expression was significantly higher in adenocarcinoma compared to mucoid and signet ring cell carcinoma subtypes ($p < 0.0001$). Among adenocarcinoma cases, a significant inverse association was found between cytoplasmic ICAM-1 expression and tumor grade ($p < 0.0001$). Also, statistically significant inverse associations were found between cytoplasmic ICAM-1 expression and both TNM stage ($p = 0.038$), Dukes stage ($p = 0.004$) and lymphovascular invasion ($p = 0.007$). However, statistically significant positive associations were found between cytoplasmic ICAM-1 expression and both tumor infiltrating lymphocytes ($p = 0.007$) and tumor budding ($p = 0.001$). A significant positive relationship was found between membranous ICAM-1 expression and tumor grade ($p = 0.02$). No significant correlation was observed between Ezrin and ICAM-1 expression in the tumor cells ($r = -0.03$ and $p = 0.8$ by Spearman's rho test, $p = 0.2$ by Chi-Square test). **Conclusion:** Ezrin expression is significantly associated with poor prognostic factors and thus can be used as a poor prognosticator for colorectal carcinoma patients. Cytoplasmic expression of ICAM-1 is significantly associated with good prognostic features except for tumor budding reflecting contradictory functions of ICAM-1 in CRC. In spite of their complementary functions in carcinogenesis, Ezrin and ICAM-1 are not

significantly correlated in this series of colorectal carcinoma. Altogether, this study suggests the important role of Ezrin and ICAM-1 in colorectal carcinogenesis and progression

Key Words: Ezrin, ICAM-1, Colorectal Carcinoma, Adenocarcinoma, Mucoïd Carcinoma, Signet Ring Cell Carcinoma.

Introduction

Colorectal cancer is the most common malignancy of gastrointestinal tract and is a common leading cause of mortality in many countries around the world. Colorectal carcinogenesis is a multifactorial process, in which genetic factors, environmental exposures (including diet), and inflammatory conditions of the digestive tract are involved⁽¹⁾.

The main prognostic factors in colorectal cancer are clinicopathological characteristics of the disease, including tumor stage, and grade. Although these parameters do reflect biological features of the tumor, they do not fully predict individual clinical outcome and thus, there is a strong need for better markers to identify patients with poor prognosis. Recent researches have focused on the potential role of new biological factors involved in the carcinogenic process as prognostic markers to aid accurate prediction of clinical outcome of patients with colorectal cancer⁽¹⁾.

Much attention has been focused on the involvement of Ezrin in tumor development and progression. Ezrin is a cytoskeleton associated protein that has traditionally been known as a molecule involved in maintaining the integrity and morphology of cells. It also has been widely reported to regulate the migration of the cells being suggested to be a candidate molecule in directional movement of cancer cells helping the processes of cancer invasion and metastasis. The main functions of Ezrin are thought to be acting as protein linker who links the cell membrane to the cytoskeletal proteins, cell surface molecules (cell adhesion molecules) to the cytoskeleton and, by doing so regulating cell adhesion and migration).

Among the many intriguing molecular features of cancer is the expression of intercellular adhesion molecule-1 (ICAM-

1), which is widely known for its ability to stimulate immune activity. ICAM-1 is a member of the immunoglobulin superfamily, which belongs to cell adhesion molecules. The interaction, mediated by ICAM-1, between cell-cell and between cells and extracellular matrix, is a form of communication between cells, which plays an important role in cell differentiation, movement, and immunity. ICAM-1 is also expressed in many tumor types where it is associated with good prognosis and improved survival⁽¹⁾.

Material and Methods

Subject description: The present study comprised 102 randomly selected cases of primary colorectal carcinomas which were obtained from the archives of histopathological laboratories of Minia University Hospital and National Cancer Institute of Cairo (In the period between 2004 and 2012). The cases included; 89(87.2%) cases of adenocarcinoma, 21 (20.6%) cases of mucoïd carcinoma and 12 (11.7%) cases of signet ring cell carcinoma. Another 40 tissue blocks of adjacent normal colorectal epithelium (available blocks) were immunostained as the control group. In addition, 34 of metastatic malignant lymph nodes for lymph node positive cases were also included in the study.

Drugs and reagents: Monoclonal Mouse anti-human Ezrin and ICAM-1 Abs (Lab vision Laboratories) were used and DAB (Lab vision Laboratories) was used as a detection kit. Immunostaining was performed following the instructions of the manufacturers. PBS (0.01 mmol/L) was used to replace the first antibody as a negative control, while the normal colorectal mucosa was a positive control for Ezrin and normal human tonsillar tissue was a positive control for ICAM-1.

Assessment of staining: To assess positive staining for Ezrin, immunoreactivity was

scored on a scale of 0-3 depending on the intensity of staining and the percentage of positive cells. Staining intensity was graded as 0 (negative), 1 (weak), 2 (moderate) and 3 (strong). The percentage of positive cells was scored as 0 (<1%), 1 (1-25%), 2 (26-50%), 3 (51-75%) and 4 (76-100%). The two scores were multiplied and then the total immunohistochemical score (IH score) values (0-12) were classified as follows: score 0 as negative, score 1 (multiplication values 1 and 2) as weakly positive, score 2 (multiplication values 3, 4 and 6) as moderately positive and score 3 (multiplication values 8, 9, 12) as strongly positive. Cytoplasmic and membranous staining, when observed, were evaluated separately.

For scoring of ICAM-1 staining, staining intensity was graded as 0 (negative), 1 (weak), 2 (moderate) and 3 (strong). The numbers of positive cells were expressed as the percentage of the total number of epithelial. The two scores were multiplied and then immunohistochemical score (IH score) for each tumor specimen was calculated. Cytoplasmic and membranous staining, when observed, was evaluated separately. No definite cut-off point for identification of positivity and grade of ICAM-1 expression was reported in literature. Therefore, we preferred to use continuous scores to study the association of ICAM-1 in relation to different clinicopathologic features.

Statistical analysis

Data were checked, coded, entered and analyzed by using SPSS (The Statistical Package for Social Sciences) version 22 software. Association between immunoreactivity and different clinicopathological data were done by Chi-Square test, McNemar test, Mann-Whitney test and Kruskal-wallis test. Spearman's rho coefficient Wilcoxon test and Friedman test were used for continuous variables. Statistical significance was determined at p value of ≤ 0.05 for all tests except Friedman test where significance level is set as 0.02.

Results

Information regarding different clinical and histopathological features for all colorectal

carcinoma (CRC) patients is illustrated in Table (1).

On studying the cytoplasmic expression of Ezrin (table 2), the expression was significantly higher in colorectal carcinoma compared to normal tissues.

Cytoplasmic Ezrin expression was significantly higher in lymph node positive than lymph node negative cases and in cases with lymphovascular invasion than cases without invasion. Similarly, there were significant variations in cytoplasmic Ezrin expression among different stages of colorectal carcinoma being markedly higher in tumors with advanced TNM and Dukes stages compared to lower stage tumors.

As for adenocarcinoma cases, grade III tumors showed more frequent strong cytoplasmic Ezrin expression than grade I and II however, such differences did not reach the significant statistical level (p value= 0.2). No significant associations were also observed between cytoplasmic Ezrin expression and patients' age, gender, tumor size, location, tumor type, perineural invasion, tumor budding or tumor infiltrating lymphocytes.

On comparing cytoplasmic and membranous Ezrin expression in primary tumors and their corresponding metastatic lymph nodes, no significant differences were observed in Ezrin expression between both locations.

Concerning cytoplasmic ICAM-1 expression (table 3), this work identified significant differences in expression between different types of colorectal carcinoma being markedly higher in adenocarcinoma compared to mucoid and signet ring subtypes. Among adenocarcinoma cases, a significant inverse association was found between cytoplasmic ICAM-1 expression and tumor grade where ICAM-1 expression scores were significantly higher in grade I than grade II and III tumors.

Statistically significant inverse associations were found between cytoplasmic ICAM-1 expression and both TNM and Dukes staging systems and lymphovascular invasion. However, statistically significant

positive associations were found between cytoplasmic ICAM-1 expression and both and tumor infiltrating lymphocytes and tumor budding. No significant associations were found between cytoplasmic ICAM-1 expression and other clinicopathologic features.

Comparing ICAM-1 expression in primary tumors and their corresponding metastatic

lymph nodes revealed no significant differences in ICAM-1 expression scores between primary tumors and their matched lymph node metastasis.

Despite having apparently complementary functions, Ezrin and ICAM-1 immunoreactivity results did not display significant association between both markers in this study.

Table 1: Clinicopathological features for patients with CRC (n=102)

Clinicopathological features	No. (%)
Age at Surgery, y	
- ≤ 40	21 (20.6%)
- > 40	81 (79.4%)
Gender	
- Male	53 (52%)
- Female	49 (48%)
Location	
1- Caecum	2 (2%)
2- Ascending colon	36 (35.3%)
3- Transverse colon	9 (8.8%)
4- Descending colon	17 (16.7%)
5- Sigmoid colon	19 (18.7%)
6- Rectum	19 (18.7%)
Histological subtypes	
1- Adenocarcinoma	59 (57.8%)
2- Mucooid carcinoma	28 (27.5%)
3- Signet ring cell carcinoma	15 (14.7%)
Tumor grade (for adenocarcinoma only)*	
1- Grade I	4 (3.9%)
2- Grade II	32 (31.3%)
3- Grade III	23 (22.6%)
Regional lymph node involvement	
Absent	58 (56.9%)
Present	44 (43.1%)
TNM stage groupings	
1-I	18 (17.7%)
2-II	37 (36.3%)
3-III	36 (35.3%)
4-IV	11 (10.8%)
Dukes classification	
1-A	0 (0%)
2-B1	13 (12.7%)
3-B2	41 (40.2%)
4-C1	3 (2.9%)
5-C2	34 (33.3%)
6-D	11 (10.8%)
Perineural invasion	
Absent	82 (80.4%)
Present	20 (19.6%)
Lymphovascular invasion	
Absent	72 (70.6%)
Present	30 (29.4%)
Tumor budding	
Absent	79 (77.5%)
Present	23 (22.5%)
Tumor infiltrating lymphocytes	
1-Absent	46 (45.1%)
2-Mild	22 (21.6%)
3-Moderate	13 (12.7%)
4-Marked	21 (20.6%)

Table (2): Association between cytoplasmic Ezrin expression and clinicopathological features for patients with CRC

Clinicopathological features	Total n (%)	Negative expression No. (%)	Weakly positive No. (%)	Moderately positive No. (%)	Strongly positive No. (%)	P value
Age at Surgery, y						
≤ 40	21 (20.7)	0 (0)	2 (9.5)	9 (42.9)	10 (47.6)	0.16
> 40	81 (79.3)	3 (3.7)	18 (22.2)	29 (35.8)	31 (38.3)	
Gender						
-Male	53 (52)	2 (3.8)	10 (18.9)	19 (35.8)	22 (41.5)	0.8
-Female	29 (28)	1 (3.4)	10 (34.5)	19 (65.5)	19 (65.5)	
Location						
1-Caecum	2 (2)	0 (0)	0 (0)	1 (50)	1 (50)	0.23
2-Ascending colon	37 (35.3)	1 (2.7)	10 (27.1)	18 (48.6)	18 (48.6)	
3-Transverse colon	9 (8.8)	1 (11.1)	1 (11.1)	2 (22.2)	5 (55.6)	
4-Descending colon	17 (16.7)	0 (0)	1 (5.9)	11 (64.1)	5 (29.4)	
5-Sigmoid colon	19 (18.6)	1 (5.3)	4 (21.1)	10 (52.6)	4 (21.1)	
6-Rectum	19 (18.6)	0 (0)	4 (21.1)	9 (47.4)	6 (31.5)	
Histological subtypes						
1-Adenocarcinoma	59 (57.8)	0 (0)	9 (15.3)	20 (33.9)	20 (33.9)	0.20
2-Mucoid carcinoma	28 (27.5)	1 (3.6)	8 (28.6)	10 (35.7)	9 (32.1)	
3-Signet ring cell carcinoma	10 (9.7)	2 (20)	3 (30)	3 (30)	2 (20)	
Tumor grade (adenocarcinoma) *						
1-Grade I	4 (3.8)	0 (0)	1 (25)	2 (50)	1 (25)	0.2
2-Grade II	32 (31.2)	0 (0)	6 (18.8)	16 (50)	10 (31.2)	
3-Grade III	23 (22.5)	0 (0)	2 (8.7)	7 (30.4)	14 (60.9)	
Regional lymph node involvement						
Present	44 (43.1)	2 (4.5)	0 (0)	14 (31.8)	28 (63.7)	< .0001*
Absent	58 (56.9)	1 (1.7)	20 (34.5)	24 (41.5)	13 (22.5)	
TNM stage groupings						
1-I	18 (17.6)	0 (0)	12 (66.7)	6 (33.3)	0 (0)	< .0001*
2-II	37 (36.3)	1 (2.7)	8 (21.6)	18 (48.6)	10 (27)	
3-III	37 (36.3)	2 (5.4)	0 (0)	14 (37.9)	20 (54.7)	
4-IV	11 (10.8)	0 (0)	0 (0)	0 (0)	11 (100)	
Dukes classification						
1-A	0 (0)	0 (0)	0 (0)	0 (0)	0 (0)	< .0001*
2-B1	13 (12.8)	0 (0)	11 (85.4)	2 (15.4)	0 (0)	
3-B2	41 (40.2)	1 (2.4)	9 (22)	21 (51.2)	10 (24.4)	
4-C1	3 (2.9)	0 (0)	0 (0)	2 (66.7)	1 (33.3)	
5-C2	34 (33.3)	2 (5.9)	0 (0)	13 (38.2)	19 (55.9)	
6-D	11 (10.8)	0 (0)	0 (0)	0 (0)	11 (100)	
Perineural invasion						
Present	20 (19.6)	1 (5)	1 (5)	11 (55)	7 (35)	0.86
Absent	82 (80.4)	2 (2.4)	19 (23.2)	27 (32.9)	34 (41.5)	
Lymphovascular invasion						
Present	30 (29.4)	0 (0)	3 (10)	11 (36.7)	16 (53.3)	0.38*
Absent	72 (70.6)	3 (4.2)	17 (23.6)	27 (37.5)	25 (34.7)	
Tumor budding						
Present	23 (22.5)	0 (0)	2 (8.7)	12 (52.2)	9 (39.1)	0.64
Absent	79 (77.5)	3 (3.8)	18 (22.8)	26 (32.9)	32 (40.5)	
Tumor infiltrating lymphocytes						
1-Absent	46 (45.1)	2 (4.3)	8 (17.4)	10 (21.7)	20 (43.6)	0.47
2-Mild	22 (21.6)	1 (4.5)	3 (13.6)	7 (31.8)	11 (50)	
3-Moderate	13 (12.7)	0 (0)	3 (23.1)	7 (53.8)	3 (23.1)	
4-Marked	21 (20.6)	0 (0)	0 (0)	9 (42.9)	12 (57.1)	

Table (3): Association between cytoplasmic ICAM-1 expression and clinicopathological features for patients with CRC (n=102)

Clinico-pathological features	Total (%)	Cases with no expression	cases showing expression	(Min-Max) Expression scores	Median	M ± SD	P value
Age at Surgery, y - ≤ 40 - > 40	21 (20.7) 81 (79.3)	16 (76.2) 86 (71.7)	5 (23.8) 31 (38.3)	(0-180) (0-270)	0 0	14.02 ± 43.8 27.9 ± 59.7	0.19
Gender -Male -Female	53 (52) 49 (48)	33 (62.3) 33 (67.3)	20 (37.7) 16 (32.7)	(0-270) (0-210)	0 0	20.8 ± 56.3 29.9 ± 57.7	0.27
Location 1-Caecum 2-Ascending colon 3-Transverse colon 4-Descending colon 5-Sigmoid colon 6-Rectum	2 (2) 37 (35.3) 9 (8.8) 17 (16.7) 19 (18.7) 19 (18.7)	2 (100) 27 (70) 7 (77.8) 10 (58.8) 9 (47.4) 12 (63.2)	0 (0) 9 (20) 3 (33.3) 7 (41.2) 10 (52.6) 7 (37.8)	(0-0) (0-180) (0-210) (0-100) (0-270) (0-160)	0 0 0 0 0 0	0 ± 0 13.3 ± 39.7 26.1 ± 69.3 22.7 ± 38.3 53.4 ± 89 24 ± 49.7	0.94
subtypes 1-Adenocarcinoma 2-Mucoid 3-Signet ring	59 (57.8) 28 (27.5) 15 (14.7)	27 (45.8) 20 (89.3) 14 (93.3)	32 (54.2) 3 (10.7) 1 (6.7)	(0-270) (0-0) (0-3)	0 0 0	43.3 ± 79.4 0.32 ± 1 0.2 ± 0.77	<0.0001*
Tumor grade (adenocarcinoma)* 1-Grade I 2-Grade II 3-Grade III	4 (6.8) 32 (54.2) 23 (39)	0 (0) 8 (20) 19 (82.6)	4 (100) 24 (70) 4 (17.4)	(80-270) (0-240) (0-20)	180 20 0	177.0 ± 11.4 56.7 ± 76.0 1.3 ± 4.3	<0.0001*
lymph node involvement Present Absent	44 (43.1) 58 (56.9)	32 (72.7) 34 (58.7)	12 (27.3) 24 (41.3)	(0-270) (0-240)	0 0	23.7 ± 59.7 26.3 ± 50.1	0.2
TNM stage 1-I 2-II 3-III 4-IV	18 (17.7) 37 (36.3) 36 (35.3) 11 (10.8)	11 (61.1) 20 (54.1) 26 (72.2) 9 (81.8)	7 (38.9) 17 (45.9) 10 (27.8) 2 (18.2)	(0-240) (0-210) (0-270) (0-210)	0 0 0 0	43.4 ± 74.8 20.2 ± 43.8 22.0 ± 57.2 20.9 ± 73	0.38*
Dukes classification 1-A 2-B1 3-B2 4-C1 5-C2 6-D	0 (0) 13 (12.8) 41 (40.2) 3 (2.9) 34 (33.3) 11 (10.8)	0 (0) 11 (84.6) 20 (48.8) 0 (0) 26 (76.5) 9 (81.1)	0 (0) 2 (15.4) 21 (51.2) 3 (100) 8 (23.5) 2 (18.2)	0-0 (0-240) (0-210) (60-270) (0-150) (0-210)	0 0 1 60 0 0	0 ± 0 24.6 ± 78.4 28 ± 53.3 130 ± 121 14.1 ± 38.7 20.9 ± 73	0.004*
Perineural invasion Present Absent	20 (19.6) 82 (80.4)	10 (50) 76 (78.3)	10 (50) 26 (19.6)	(0-160) (0-270)	20 0	39 ± 54.7 21.8 ± 57.1	0.08
Lymphovascular invasion Present Absent	30 (29.4) 72 (70.6)	20 (83.3) 41 (56.9)	5 (16.7) 31 (43.1)	(0-160) (0-270)	0 0	7.0 ± 30.2 32.7 ± 73.0	0.007*
Tumor budding Present Absent	23 (22.5) 79 (77.5)	8 (34.8) 71 (73.4)	15 (65.2) 21 (26.6)	(0-270) (0-240)	10 0	44.6 ± 78.9 19.0 ± 51.9	0.001*
TILs 1-Absent 2-Mild 3-Moderate 4-Marked	46 (45.1) 22 (21.6) 13 (12.7) 21 (20.6)	37 (80.4) 14 (63.7) 4 (30.8) 12 (57.1)	9 (19.6) 8 (16.3) 9 (69.2) 9 (42.9)	(0-150) (0-180) (0-210) (0-270)	0 0 20 0	9.7 ± 30.9 21.7 ± 50 41.3 ± 70.1 42 ± 77.9	0.007*

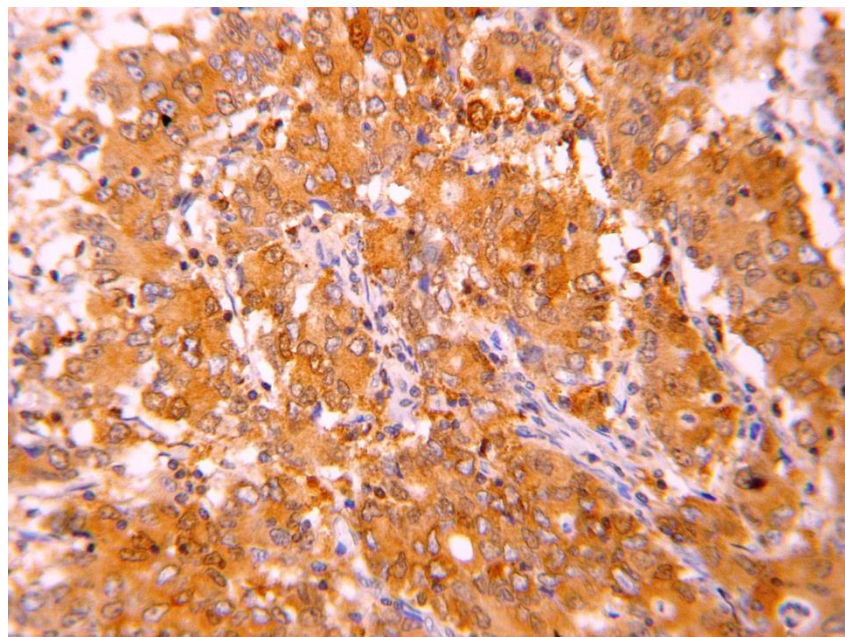


Figure 1: Immunohistochemical appearance of grade III colorectal adenocarcinoma stained with Ezrin, showing positive cytoplasmic staining of strong intensity (original magnification 400).

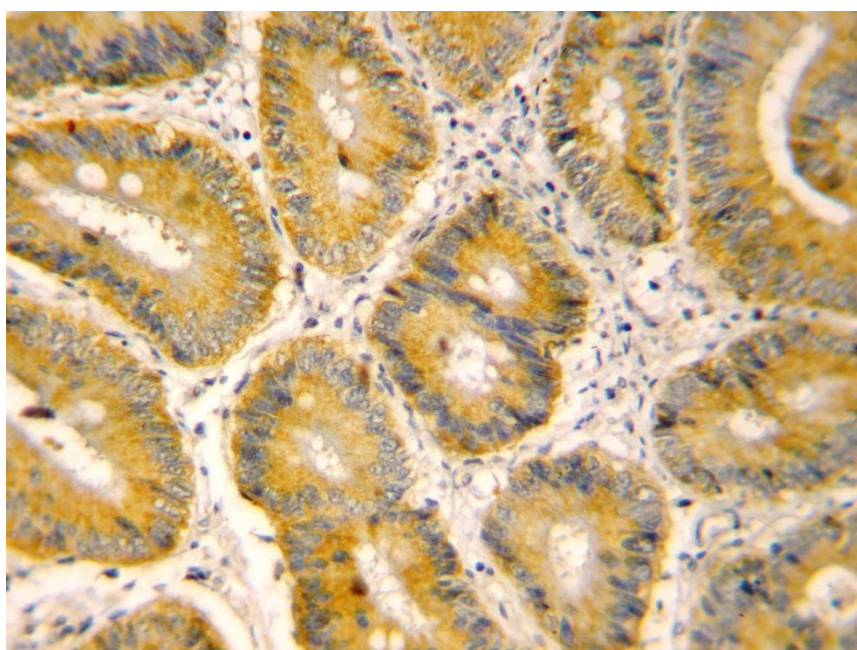


Figure 2: Immunohistochemical appearance of grade I colorectal adenocarcinoma with ICAM-1 immunostaining, showing positive cytoplasmic staining of strong intensity (original magnification 400).

Discussion

Colorectal cancer is one of the most common cancers worldwide. It affects men and women of all racial and ethnic groups, and is most often found in people aged 60 years or older in developed countries⁽⁵⁾.

Ezrin plays specific roles in the adherent junction between trans-membranous proteins and cytoskeleton within cell and between cells, and acts as a signal transducer. In this way, Ezrin is involved in the regulation of phenotypical change of cellular architecture, controlling cellular motility, tethering action between cell to cell and cell to matrix, recognizing and conducting internal and/or external signal. All of these cellular functions are essential for maintaining homeostasis of normal cell as well as entire process of tumorigenesis and distant metastasis^(6,7). Intercellular adhesion molecule (ICAM)-1 (CD54) is a member of the immunoglobulin superfamily and expressed on many cell types including endothelial cells, leukocytes and cancer cells where it can be upregulated in response to various cytokines⁽⁸⁾.

Regarding subcellular localization of Ezrin, the current study detected Ezrin expression in the intercellular junctions and cytoplasm of normal colorectal epithelial cells which explains an important function of Ezrin in normal cells, i.e. maintaining intercellular adhesion. In tumor cells, Ezrin was found either in the cytoplasm or in both cytoplasm and membrane. This finding was reported by several studies performed on Ezrin expression in normal colorectal mucosa and colorectal carcinoma⁽⁹⁾. In the current study, a statistically significant difference in expression of Ezrin was detected between normal colorectal mucosa and colorectal carcinoma and this was in agreement with other studies performed by^(10,11). 63.6% of cases positive for lymph node metastases showed strongly positive cytoplasmic Ezrin expression as compared to those negative for lymph node metastases which showed only 22.4% strongly positive expression. This was in accordance with⁽¹²⁾ who demonstrated significant association between Ezrin expression and nodal metastases. A significant association was

observed between cytoplasmic Ezrin expression and advanced TNM stage. The frequency of cytoplasmic Ezrin expression was much higher in both stage IV and stage III when compared to stage II and stage I. This was in line with findings of⁽³⁾. Our findings together with others suggest that Ezrin appears to be an important prognostic factor for patients with colorectal cancer as it is more frequently expressed in carcinoma than in normal epithelium and its expression correlates with aggressive tumor features such as lymph node involvement, lymphovascular invasion and advanced tumor stage. It may also allow therapeutic options such as miRNA and siRNA to be developed. No drugs are yet reported to specifically inhibit Ezrin; however, some of the epigenetic drugs and estrogen have been shown to upregulate Ezrin. Rational designed compounds that have impact on the activity of Ezrin have also been recently reported⁽¹³⁾.

The broad expression of Ezrin together with the association of Ezrin expression with poor prognostic features in colorectal carcinoma confirms the important role of Ezrin in pathogenesis of colorectal carcinogenesis and suggests its role as an adjuvant therapy for high risk patients.

ICAM-1 expression was totally absent in normal colorectal epithelium while in colorectal carcinoma, the positive expression rate was 30.3%. ICAM-1 was found in the cytoplasm, membrane or in both cytoplasm and membrane. This was consistent with several studies that mentioned absent ICAM-1 expression in normal mucosa and reported positive expression rates ranging from 33% up to 60% in colorectal carcinoma⁽¹⁴⁾.

Concerning the cytoplasmic expression of ICAM-1 in different subtypes of CRC, the current study showed highly significant differences in expression between adenocarcinoma and other histopathological subtypes. ICAM-1 positive expression rates were significantly much higher in adenocarcinoma cases compared to mucoid and signet ring cell carcinomas, a finding that was not previously reported in literature as

ICAM-1 expression was not previously investigated in mucoid or signet ring colorectal carcinoma cases. In a study performed on gastric carcinoma,^(7,11) found significant differences in ICAM-1 expression between gastric adenocarcinoma and other subtypes including mucoid and signet ring cell carcinomas. The current study demonstrated a significant inverse association between ICAM-1 expression and grade of colorectal adenocarcinoma, where positive expression rates were much higher in grades I and II compared to grade III tumors and this was consistent with previous studies^(1,3&6). Likewise, a significant inverse association between ICAM-1 expression and tumor stage was noticed in this study. The rate of ICAM-1 expression was significantly higher in stages I and II tumors compared to stage III and IV tumors. This was in line with⁽¹⁾ who reported significant association between loss of ICAM-1 expression and advanced stage. ICAM-1 positive expression was significantly lower in tumors showing lymphovascular invasion. Interestingly, ICAM-1 expression was significantly associated with tumor budding where the positive expression rates were markedly higher in tumors with budding than those without budding. Actually, multiple examples for adhesion molecules are known to display seemingly controversial roles, such as EpCAM or Maspin. In this respect, in breast cancer cells, silencing of ICAM-1 expression by siRNA resulted in decreased tumorigenicity in vitro⁽¹¹⁾. Another study performed on hepatocellular carcinoma suggested that, ICAM-1 can trigger the calcium influx and take a part in disintegration of the cells' local adhesion, contraction of cells and the reconstruction of cytoskeleton, so as to promote the transfer of cancer cells and that the high expression of ICAM-1 helps cancer cells to separate from each other becoming detached and transfer to other sites⁽¹²⁾, a mechanism that could explain the significant association between ICAM-1 expression and tumor budding. Infiltration of TILs was more frequently observed in the ICAM-1-positive tumors than in the ICAM-1-negative tumors, a finding that was also supported by previous studies performed on colorectal carcinoma^(1,3).

Taken together, our previous findings suggest that positive cytoplasmic ICAM-1 expression in colorectal carcinoma is a useful important biologic marker predicting good prognosis.

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