

Comparison of Genetic, Protein Expression and Clinicopathological Parameters in Patients With Colorectal Cancer Having Vegetarian vs. Mixed Diet

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Abstract

Purpose: To assess the prevalence of KRAS codon 12/13 mutations and protein expression in patients with Colorectal Cancer (CRC). The outcomes were correlated with clinicopathological characteristics in vegetarian versus mixed diet population.

Methods: Dietary patterns were correlated with demographic and clinicopathological characteristics including KRAS codon 12/13 mutations and their protein expressions.

Results: Retrospective analysis was performed on 78 CRC patients, 55% of whom were men and had a median age of 56 years. Majority of the patients were vegetarian (73%) while remaining had a mixed diet. The association between dietary patterns and clinicopathological characteristics was observed and vegetarianism was more prevalent in patient with early and advanced cancer stages. The final histopathology revealed a statistically significant correlation between KRAS codon 12 mutations (23%) and necrosis ($p=0.025$) as well as a higher death rate ($p=0.021$). In the subgroup of rectal cancer patients, positive K-ras protein expression was linked to a higher risk of disease relapse ($p=0.045$). Vegetarians had higher levels of B-raf protein expression and was associated with poor prognostic factors, such as perineural invasion (86%) ($p=0.051$), positive lymph nodes ($p=0.020$) and high circulating carcinoembryonic antigen levels (>5.0 ng/ml) ($p=0.006$).

Conclusion: In our study, we found a higher prevalence of vegetarianism in patients with both early and advanced cancer stages. Among vegetarian patients, a link between B-raf protein expression with unfavorable prognostic factors was also observed. KRAS codon 12/13 mutations and K-ras/B-raf protein expression may act as vital prognostic and predictive indicators in CRC patients.

Keywords: Colorectal cancer; BRAF; KRAS; Dietary pattern

therapeutic options. They include endoscopic and surgical local excision, down staging pre-operative radiotherapy and systemic therapy, extensive surgery for locoregional and metastatic disease, local ablative therapies for metastases, palliative chemotherapy, targeted therapy and immunotherapy [1]. These treatments effectively and gradually reduce cancer progression and improve overall survival [2]. Additionally, proper enforcement of screening methods including fecal occult blood testing and endoscopy increases the survival rate and likelihood of CRC cure [3].

In 2020, GLOBOCAN estimated over 1.9 million new CRC cases with 0.93 million fatalities in 185 countries. The incidence and mortality rates were highest for males in European nations and lowest for females in African and South Asian nations. By 2040, it is estimated that 3.2 million new cases and 1.6 million deaths would occur from CRC. In developing countries such as India, CRC incidences were reported in a total of 0.065 million cases comprising 61.83% males and 38.17% females [4].

The rising incidence of CRC is mainly attributed to changes in lifestyle, aging and genetic aberrations. Numerous other risk factors, including poor diet, physical inactivity, smoking, chronic inflammatory bowel disease, advancing age and accumulation of genetic aberrations, have a substantial impact on the incidence and mortality of CRC [5]. Diet can influence the risk of CRC either through dietary components or indirectly through changes in gut flora or body weight. Healthy dietary patterns include high consumption of insoluble fiber, vegetables, fruits and low-fat milk. High consumption of processed foods, red meat, refined carbohydrates, high calories and a low-calcium diet promotes an inflammatory response and increases the risk of CRC [6]. Individuals who have a sedentary lifestyle or are obese have a 25%-50% higher risk of developing colon cancer compared to those who are physically active [7].

In addition to lifestyle factors, multiple genetic pathways and mechanisms contribute to CRC pathogenesis. The two important oncogenes that promote the progression of CRC are KRAS and

Introduction

Recent advancements in our understanding of the pathophysiology of Colorectal Cancer (CRC) have led to increased

BRAF, which are key players in the RAS/RAF/MEK/MAP kinase signaling pathways [8]. K-ras is a small GTPase transducer protein with intrinsic GTPase activity that controls regulatory pathways and promotes cell growth. B-raf is a serine-threonine protein kinase that acts as a direct effector of Ras and promotes tumor growth and survival by activating the Mitogen-Activated Protein Kinase (MAPK) pathway. It has been established that KRAS and BRAF mutations are present in approximately 40% and 10% of CRC cases, respectively [9,10]. The most frequent mutation in KRAS is an amino acid change at codon 12 or 13 in exon 2, whereas the most prevalent mutation in BRAF, V600E, occurs at codon 600 in exon 15 and accounts for 5%–10% of all cases of colon cancer [11].

Limited studies have examined genome-wide transcriptional alterations in KRAS and BRAF-transfected cell lines, despite their common significance in carcinogenesis [12]. Both dietary and genetic modifications play a significant role in the pathogenesis of CRC. Therefore, exploring the correlation between dietary patterns and K-ras and B-raf protein expression may be beneficial for the management of patients with CRC. In light of these factors, the objective of the present study was to investigate the prevalence of KRAS codon 12 and 13 mutations along with K-ras and B-raf protein expression in CRC patients to further correlate the findings with various clinicopathological parameters and to compare them with either vegetarian or mixed diets.

Materials and Methods

Patients

A total of 78 untreated and histologically confirmed colorectal cancer patients registered at 'The Gujarat Cancer and Research Institute (GCRI)', Ahmedabad, India, from 2013 to 2015 were enrolled in this study. The tissue acquired during the surgery was stored in the immunochemistry and cancer biology laboratory. The tests for KRAS 12/13 codons and K-ras and B-raf proteins were performed in 2017. The demographic characteristics of the patients including age, sex, family history, dietary habits, alcohol consumption, tobacco smoking and other potential confounding factors were collected. Detailed clinical and pathological characteristics of the patients including tumor size, lymph node status, histological grade, disease stage and treatment given, etc. were captured from the case files maintained at the medical record department of the institute and recorded in the registers kept at the tumor biology, GCRI. The key inclusion criteria were as follows: Untreated patients with histopathologically confirmed adenocarcinoma; patients with pre-op colonoscopy and biopsy; CT scan/MRI; patients with pre-op Carcinoembryonic Antigen (CEA) levels; and patients who had not previously received anticancer treatment at GCRI or any other hospital. The exclusion criteria included patients who underwent treatment prior to surgery (including previous surgery, chemotherapy and radiation therapy), as well as those who tested positive for HBsAg, HIV and HCV.

Sample collection

Prior to sample collection, written informed consent was obtained from each patient. Primary tumor tissue samples from

78 CRC patients were obtained on ice right from the operation theater for the analysis of K-ras codon 12 and 13 mutations. The tumor tissue portion from the collected specimen was selected by a pathologist, which was then snap frozen in liquid nitrogen and preserved at -80°C. Paraffin-embedded tumor tissue blocks from all patients were retrieved from the histopathology department of the institute for the analysis of K-ras and B-raf protein expression.

Methodology

The dietary patterns of the patients were analyzed and correlations were made with different demographic and clinicopathological details including K-ras codon 12 and 13 mutations and expression of K-ras and B-raf proteins.

Statistical analysis

Statistical analysis was performed using the Statistical Package for the Social Sciences software (SPSS). Relapse-free survival and overall survival were calculated using the Kaplan-Meier method and Log-rank tests. The correlation between the two parameters was calculated using Spearman's correlation coefficient (r). P -value ≤ 0.05 was considered significant.

Results

Demographic characteristics

A retrospective analysis was performed on 78 CRC patients, comprising 55% males and 45% females, with a median age of 56 years (range 20–80 years). Demographic details are provided in **Table 1**. A total of 73% of the patients were vegetarian, while the remaining patients had a mixed diet including a non-vegetarian or eggetarian diet. In addition, 47% of patients were tobacco consumers, smokers and alcoholics. The majority (59%) of patients had tumors in the colon while the remaining (41%) had a tumor in the rectum. The patients were at different stages of cancer; 74% had T3 tumors, 18% had T2 tumors and 8% had T4 tumors. According to the AJCC clinical staging system, 47% and 41% of the patients had stage II and III tumors, respectively. The colorectal cancer staging criteria used by Duke identified, 59% of the patients had Duke B and 41% had Duke C. Patients with Duke D were excluded from the study. In 74% of the cases, a moderately differentiated tumor was present while a poorly differentiated tumor was found in 11% of cases. Histological examination revealed adenocarcinoma in 73% and mucinous adenocarcinoma in 27% of cases. Nodal status was negative in 63% of the patients, while 37% had nodal-positive disease. In the analysis of tumor characteristics, 91% of patients did not show necrosis. Lymphatic permeation was absent in 80% of patients. Vascular permeation and perineural invasion were absent in 95% and 82% of the patients, respectively. Preoperative CEA levels were ≥ 5 ng/ml in 55% of the cases and ≤ 5 ng/ml in 45% of the cases. In 18% of the cases, the only form of treatment was surgery, in 58% of cases chemotherapy was added to surgery and radiotherapy added to surgery in 19% of cases. All three modalities were used in 5% of cases. Of the 63 patients studied, 57 (90%) exhibited no recurrence, while six experienced recurrence. Overall Survival (OS) was observed in 80% of patients ($n=62$), whereas 20% of patients ($n=16$) died during the follow-up period.

Table 1: Demographic details of CRC patients.

Characteristics		N=78	Percentage (%)
Age (Range: 20-80 years) (Median age: 56 years)	<56	34	44
	>56	44	56
Sex	Female	35	45
	Male	43	55
Diet	Vegetarian	57	73
	Mixed	21	27
Habit	No	41	53
	Yes	37	47
Tumor site	Colon	46	59
	Rectum	32	41
Tumor size	T2	14	18
	T3	58	74
	T4	6	8
AJCC staging	I	9	11
	II	37	47
	III	32	41
	Early stage (Stage I+Stage II)	46	59
	Advanced stage (Stage III+Stage IV)	32	41
		46	59
Dukes' stage	B	46	59
	C	32	41
Tumor differentiation	Well differentiated	11	14
	Moderately differentiated	58	74
	Poorly differentiated	9	11
Histological type	Adenocarcinoma	57	73
	Mucinous adenocarcinoma	21	27
Nodal status	Negative	49	63
	Positive	29	37
Necrosis	Absent	71	91
	Present	7	9
Lymphatic permeation	Absent	62	80
	Present	16	20
Vascular permeation	Absent	74	95
	Present	4	5
Perineural invasion	Absent	64	82
	Present	14	18
Lymphocytic stromal response	Absent	63	81
	Present	15	19
Pre-operative serum CEA levels (ng/ml) (N=81)	<5	35	45
	≥ 5	43	55
Treatment given	Surgery alone	14	18
	S+CT	45	58
	S+RT	15	19
	S+CT+RT	4	5
Recurrence (N=63)	Absent	57	90
	Present	6	10
Survival (N=78)	Alive	62	80
	Mortality	16	20

Clinical and pathological parameters

When the patient's dietary patterns were correlated with various clinical parameters such as age, sex, habit, family history and tumor site, no statistically significant correlation was established. A similar trend was observed for various pathological parameters as shown in **Table 2**. In patients who presented at an early or advanced stage, the prevalence of vegetarianism was higher

than in patients who had a mixed dietary pattern ($p=0.063$). In comparison to 33% ($n=15$) of the patients with a mixed dietary pattern, 67% ($n=31$) of the vegetarian patients developed early CRC. Furthermore, compared to 19% ($n=06$) of patients on a mixed diet, 81% ($n=26$) of the vegetarian patients also experienced advanced CRC.

Table 2: Correlation of diet with clinical and pathological parameters in CRC patients.

Characteristics	N	Diet		χ^2	r	P
		Vegetarian N (%)	Mixed N (%)			
Age (years)						
<55	34	25 (73)	09 (27)	0.006	0.009	0.938
>55	44	32 (73)	12 (27)			
Sex						
Female	35	28 (80)	07 (20)	1.547	0.141	0.219
Male	43	29 (67)	14 (33)			
Habit						
No	41	32 (78)	09 (22)	1.086	0.118	0.304
Yes	37	25 (68)	12 (32)			
Family history						
No	76	57 (75)	19 (25)	2.411	0.267	0.12
Yes	2	00 (00)	02 (100)			
Tumor site						
Colon	46	33 (72)	13 (28)	0.102	-0.036	0.753
Rectum	32	24 (75)	08 (25)			
TNM stage						
I	9	04 (44)	05 (56)			
II	37	27 (73)	10 (27)	4.837	-0.212	0.063
III	32	26 (81)	06 (19)			
Early (I+II)	46	31 (67)	15 (33)	1.842	-0.154	0.179
Advanced (III)	32	26 (81)	06 (19)			

Genetic mutation and clinicopathological correlation

Genetic mutations in KRAS codons 12 and 13 were analyzed using PCR-RFLP in primary tumor samples from patients with CRC. A total of 23% ($n=18$) of the patients showed the mutant genotype while 77% ($n=60$) of the patients had the wild-type

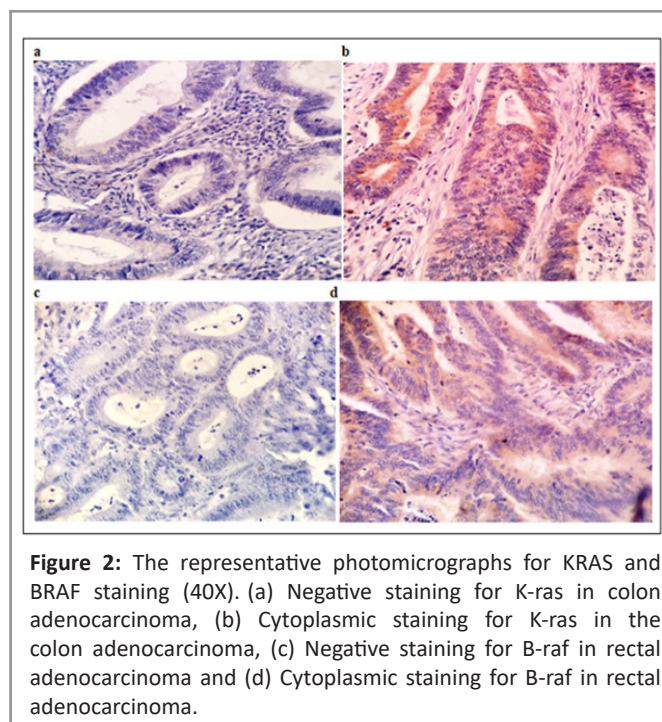
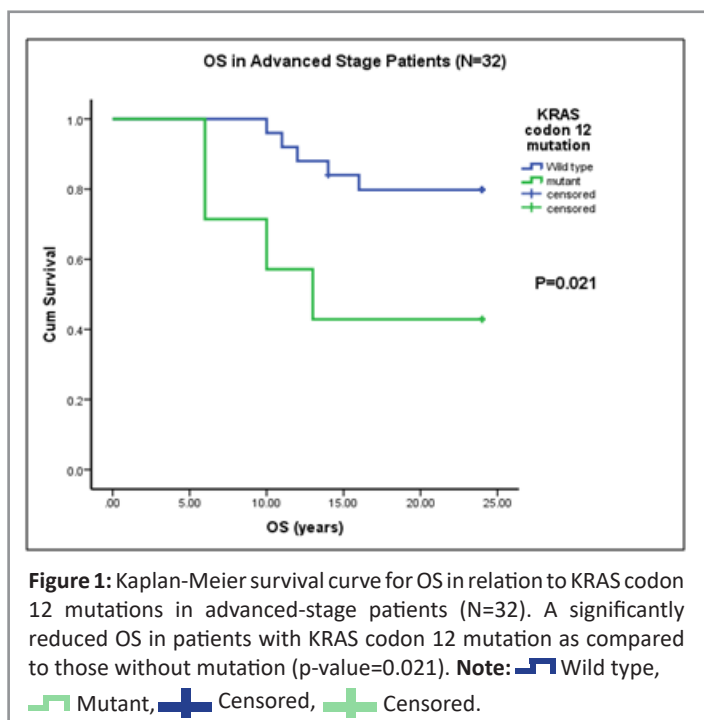
genotype of KRAS codon 12. For KRAS codon 13, all patients had wild-type genotype. The correlation between these mutations and clinical and pathological parameters was studied in patients with CRC. A statistically significant correlation was found between KRAS codon 12 mutations and necrosis ($p=0.025$). In addition, a noticeable trend was observed for tumor differentiation ($p=0.06$) and pre-operative CEA level ($p=0.087$) (**Table 3**).

Table 3: Correlation of KRAS codon 12 and 13 mutation with clinical and pathological parameters in CRC patients.

Characteristics	N	KRAS codon 12 mutation		χ^2	r	P
		Wild type N (%)	Mutant type N (%)			
Diet						
Vegetarian	57	44 (77)	13 (23)	0.009	0.011	0.927
Mixed	21	16 (76)	05 (24)			
Tumor site						
Colon	46	36 (78)	10 (22)	0.113	0.038	0.741
Rectum	32	24 (75)	08 (25)			
Tumor differentiation						
Well	11	06 (55)	05 (46)	4.015	-0.214	0.06
Moderate	58	46 (79)	12 (21)			
Poor	9	08 (89)	01 (11)			

Necrosis						
Absent	71	57 (80)	14 (20)	5.027	0.254	0.025
Present	7	03 (43)	04 (57)			
Pre-op circulating CEA (ng/ml) (N=131)						
<5.0	35	30 (86)	05 (14)	2.961	0.196	0.087
≥ 5	42	29 (69)	13 (31)			

Univariate analysis was performed to study Recurrence-Free Survival (RFS) and Overall Survival (OS) of patients with KRAS codon 12 mutations. In our study, 80% (n=62) of the 78 patients were alive while 20% (n=16) died at the time of follow-up. Of the 63 patients studied, only six had a recurrence of CRC (Table 1). In patients with advanced disease stage, Kaplan-Meier univariate survival analysis for OS revealed a statistically higher (p=0.02) incidence of death (Figure 1). However, no significant trend was observed for RFS in patients with the KRAS codon 12 mutations.



Protein mutation and expression

Histopathological analysis was performed to examine the cytoplasmic expression of the K-ras and B-raf proteins. Figure 2 shows the immunoreactivity-based detection of K-ras (Figures 2a and 2b) and B-raf (Figures 2c and 2d) protein expression in 35% (n=27) and 63% (n=49) of patients, respectively. The correlation of K-ras and B-raf protein expression with clinical and pathological parameters in patients with CRC was investigated. As shown in Table 4, there was no correlation between K-ras protein expression and clinical or pathological parameters, including variations between the vegetarian and mixed diets. In a subgroup of rectal cancer patients, Kaplan-Meier univariate survival analysis for RFS and OS revealed a substantially higher incidence of disease relapse in patients with positive K-ras protein expression than in those with negative K-ras protein expression (p=0.045), (Figure 3). However, no significant relationship was observed for OS.

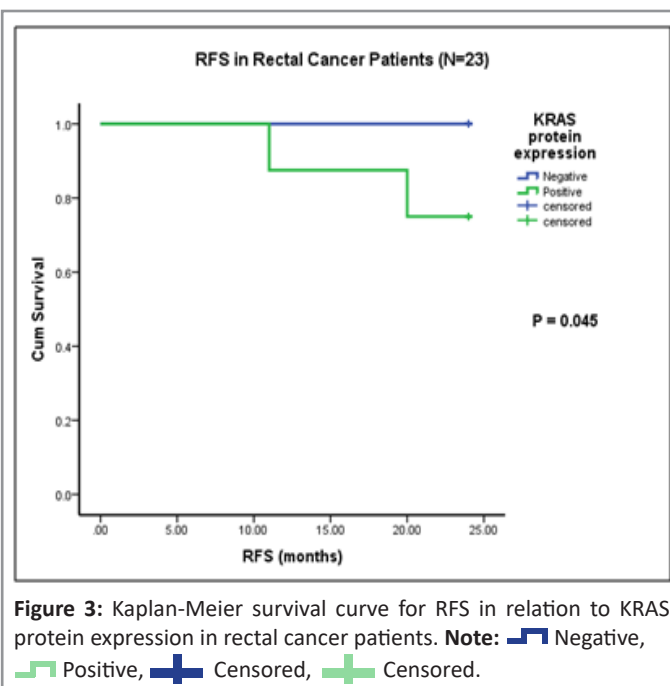


Table 4: Correlation of KRAS protein expression with clinicopathological parameters in CRC patients.

Characteristics	N	KRAS protein expression		χ^2	r	P
		Negative N (%)	Positive N (%)			
Diet						
Vegetarian	57	38 (67)	19 (33)	0.154	0.044	0.7
Mixed	21	13 (62)	08 (38)			
Tumor site						
Colon	46	28 (61)	18 (39)	1.01	-0.114	0.321
Rectum	32	23 (72)	09 (28)			
Nodal status						
Negative	49	30 (61)	19 (39)	1.008	-0.114	0.322
Positive	29	21 (72)	08 (28)			
TNM stage						
I	9	05 (56)	04 (44)			
II	37	23 (62)	14 (38)	1.15	-0.121	0.29
III	32	23 (72)	09 (28)			
Early (I+II)	46	28 (61)	18 (39)	1.01	-0.114	0.321
Advanced (III)	32	23 (72)	09 (28)			
Dukes' stage						
B	46	28 (61)	18 (39)	1.01	-0.114	0.321
C	32	23 (72)	09 (28)			
Tumor differentiation						
Well	11	07 (64)	04 (36)	0.691	-0.069	0.549
Moderate	58	37 (64)	21 (36)			
Poor	9	07 (78)	02 (22)			
Histologic type						
Adenocarcinoma	57	36 (63)	21 (37)	0.464	-0.077	0.502
Mucinous/Signet ring cell	21	15 (71)	06 (29)			
Lymphatic permeation						
Absent	62	40 (65)	22 (36)	0.101	-0.036	0.755
Present	16	11 (69)	05 (31)			
Vascular permeation						
Absent	74	49(66)	25(34)	0.441	0.075	0.513
Present	4	02(50)	02(50)			
Lymphocytic stromal response						
Absent	63	41 (65)	22 (35)	0.013	-0.013	0.909
Present	15	10 (67)	05 (33)			
Perineural invasion						
Absent	64	42 (66)	22 (34)	0.009	0.011	0.925
Present	14	09 (64)	05 (36)			
Necrosis						
Absent	71	46 (65)	25 (35)	0.124	-0.04	0.729
Present	7	05 (71)	02 (29)			
Pre-op circulating CEA (ng/ml) (N=77)						
<5.0	35	24 (69)	11 (31)	0.157	0.045	0.697
≥ 5.0	42	27 (64)	15 (36)			

The correlation of B-raf protein expression with clinical parameters in CRC patients revealed that vegetarians had higher B-raf protein expression than those with a mixed diet. B-raf protein expression was detected in 68% (n=39) of 57 vegetarian patients and 48% (n=10) of 21 patients on a mixed diet. Compared to pathological parameters, B-raf protein expression was significantly associated with positive nodal disease (p=0.020). Furthermore, the immune reactivity of B-raf protein was significantly higher

in patients with perineural invasion (86%) than in those without B-raf protein 58%, (p=0.051), as well as in patients with ≥ 5.0 ng/ml pre-operative CEA levels (76%) as compared to those with <5.0 ng/ml pre-operative CEA levels 46%, (p=0.006). Thus, higher B-raf positivity was more frequently associated with poor prognostic factors, such as perineural invasion, positive lymph nodes and high circulating CEA levels (>5.0 ng/ml) (**Table 5**).

Table 5: Correlation of BRAF protein expression with clinical and pathological parameters in CRC patients.

Characteristics	N	BRAF protein expression		χ^2	r	P
		Negative N (%)	Positive N (%)			
Diet						
Vegetarian	57	18 (32)	39 (68)	2.843	-0.191	0.094
Mixed	21	11 (52)	10 (48)			
Tumor site						
Colon	46	18 (39)	28 (61)	0.183	0.048	0.674
Rectum	32	11 (34)	21 (66)			
Nodal status						
Negative	49	23 (47)	26 (53)	5.374	0.262	0.02
Positive	29	06 (21)	23 (79)			
TNM stage						
I	9	03 (33)	06 (67)			
II	37	18 (49)	19 (51)	4.173	0.164	0.151
III	32	08 (25)	24 (75)			
Early (I+II)	46	21 (46)	25 (54)	3.446	0.21	0.065
Advanced (III+IV)	32	08 (25)	24 (75)			
Dukes' stage						
B	46	21 (46)	25 (54)	3.446	0.21	0.065
C	32	08 (25)	24 (75)			
Tumor differentiation						
Well	11	03 (27)	08 (72)	3.993	-0.195	0.087
Moderate	58	20 (34)	38 (66)			
Poor	9	06 (67)	03 (33)			
Perineural invasion						
Absent	64	27 (42)	37 (58)	3.829	0.222	0.051
Present	14	02 (14)	12 (86)			
Pre-op circulating CEA (ng/ml) (N=131)						
<5.0	35	19 (54)	16 (46)	7.552	0.313	0.006
≥ 5.0	42	10 (24)	32 (76)			

Discussion

The prevalence of colorectal cancer is rising worldwide and is strongly linked to changing lifestyles and food habits. According to epidemiological research, modifiable risk factors, such as food, can prevent approximately half of the risk of colon cancer [13]. Researchers have discovered a positive link between the risk of colorectal cancer and processed meat and preserved foods, whereas consumption of vitamins, minerals and general vegetable/fruit/fiber, reduces the risk of colorectal cancer [14]. Such vegetarian eating patterns are often associated with a lower body mass index, which in turn results in reduced fat accumulation and there is substantial evidence linking higher adiposity to an increased risk of colon cancer [15]. However, in our study, no significant correlation between diet and CRC was found with any clinicopathological or demographic parameter. Further, among vegetarian patients, the majority presented in the early and advanced stages of CRC in contrast to patients with mixed dietary patterns.

Colorectal cancer develops across several stages because of the accumulation of various genetic mutations. Ras and Raf, which are crucial intermediates in the RAS-mediated signaling cascade, are the most significant. Ras proteins are proto-oncogenes that are frequently mutated in human cancer. They are encoded by

three ubiquitously expressed genes: HRAS, KRAS and NRAS [16].

The present study examined KRAS codon 12 and 13 mutations using PCR-RFLP and indicated that 23% (n=18) of patients had KRAS codon 12 mutations, while all patients had only wild-type KRAS codon 13. Similar studies have reported the prevalence of KRAS mutations in 42.8% of patients with metastatic CRC in an Indian population [17]. Of all KRAS mutations, a mutation in codon 12 was the most common, followed by mutations in codons 13 and 61 [18]. These variations in KRAS mutation patterns may be due to racial differences and etiological factors. Given the results of several clinical trials, screening for KRAS mutations in codons 12 and 13 for metastatic colorectal cancer treatment has been recommended.

With regard to clinicopathological parameters, the present study showed a significant correlation between KRAS codon 12 mutation and necrosis. Furthermore, a trend of a higher incidence of KRAS codon 12 mutations was found in patients with well-differentiated tumors than in those with moderately and poorly differentiated tumors. Studies have shown that well-differentiated metastatic CRC tumors have significantly more mutation positivity than moderately and poorly differentiated tumors. In contrast, KRAS mutations are significantly associated with moderately and poorly differentiated colorectal adenocarcinoma [19]. Another study found that peritoneal metastasis, liver-peritoneum

metastases and multi-organ metastases were notably related to KRAS codon 12 mutations, but not codon 13 mutations, as compared to all wild types. These findings may help in predicting prognosis and selecting better treatment strategies [20].

Hence, the present study focused on the prognostic and predictive value of KRAS codon 12 and 13 mutation status in CRC patients. Additionally, we examined the role of KRAS codon 12 and 13 mutations in early and advanced-stage patients as well as in the subgroups of colon and rectal cancer patients. The study showed a significant reduction in OS with KRAS codon 12 mutations compared to KRAS wild-type in the advanced stage and prominently in advanced-stage rectal cancer patients. Also, a trend of reduced OS was noted with KRAS codon 12 mutant type in advanced-stage patients treated with adjuvant therapy. Accordingly, patients with KRAS mutations had significantly decreased median survival times compared to patients with wild-type KRAS genes in CRC.

Cytoplasmic expression of K-ras and B-raf proteins was examined using immunohistology. None of the clinicopathological parameters was found to be significantly correlated with K-ras protein expression. However, B-raf protein expression was significantly higher in vegetarians as compared to those with a mixed diet. B-raf immunopositivity was significantly higher in patients with perineural invasion, high CEA levels and positive nodal disease.

The expression levels of K-ras and its prognostic evaluation in patients with CRC remain unknown. To the best of our knowledge, this is the first study to demonstrate the prognostic and predictive roles of K-ras immunohistopathological localization in CRC. We observed that the subgroup of rectal cancer patients showed significantly reduced RFS with K-ras positive expression compared to K-ras negative expression ($p=0.045$). A similar trend of reduced RFS was observed with positive K-ras expression in the subgroup of patients with rectal cancer treated with adjuvant therapy ($p=0.073$).

Hence, the present study demonstrated that positive cytoplasmic K-ras protein expression could be a useful biomarker for predicting unfavorable prognosis and treatment response in patients with rectal cancer.

Limitations

Further studies with larger patient groups including non-vegetarian populations and longer follow-up periods are needed to better understand the prognostic and predictive role of KRAS and BRAF mutations and K-ras and B-raf protein expression in CRC cases. This would aid in the determination of CRC prevention, therapeutic response and prognosis of CRC patients.

Conclusion

Colorectal cancer is the most prevalent gastrointestinal malignancy worldwide, affecting both males and females. Dietary pattern has been postulated to affect the demographics of CRC patients. Unfortunately, most CRC tumors are benign, grow slowly, and do not manifest symptoms until they grow larger. Most genetic alterations observed in colorectal tumorigenesis are associated with K-ras and B-raf. These mutations most commonly

occur in KRAS codons 12 and 13 and render the therapeutic modulation of epidermal growth factor receptor irrelevant.

The present study did not show any significant clinicopathological variations among vegetarian and mixed-diet patients. However, a trend of an advanced stage among vegetarian patients was observed in this study. The study also demonstrated a correlation between patient prognostic factors and B-raf expression as well as a slight positive association of B-raf expression in vegetarian patients. KRAS codon 12 and 13 mutations as well as K-ras and B-raf protein expression could be useful prognostic and predictive markers in CRC patients. These data need to be validated in a larger cohort to potentially influence treatment decisions.

Author Contributions

Dr. Nikhil Garg was responsible for the conception and design, analysis and interpretation of the data. Dr. Chamanjot Singh assisted in the drafting of the manuscript or revising it critically for intellectual content. The author has approved the final version of the manuscript to be published.

Data Availability

The data that support the findings of this study are available on request from the corresponding author Dr. Nikhil Garg upon reasonable request.

Declarations

Ethical approval

The ethical approval was taken from the institutional ethics committee of 'The Gujarat Cancer & Research Institute (GCRI)' which is registered under the Central Licensing Authority (Central Drugs Standard Control Organization) with registration No ECR/41/Inst/GJ/2013/RR-19.GCRI/GCS.

Conflict of Interest

The author declares no conflict of interest.

Competing Interest

The authors declare there are no financial conflicts of interest to disclose.

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