

**“THE IMMUNOHISTOCHEMICAL
EXPRESSION OF SYNDECAN 1(CD 138)
AND ITS CORRELATION WITH STAGING
AND GRADING OF COLORECTAL
CARCINOMA”**



BY

Dr. SUDARSHAN K, MBBS

DISSERTATION SUBMITTED TO
SRI DEVARAJ URS ACADEMY OF HIGHER EDUCATION &
RESEARCH

TAMAKA, KOLAR, KARNATAKA

IN PARTIAL FULFILLMENT
OF THE REQUIREMENTS FOR THE DEGREE OF
DOCTOR OF MEDICINE

IN
PATHOLOGY

UNDER THE GUIDANCE OF

Dr. HEMALATHA A, MD

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DEPARTMENT OF PATHOLOGY
SRI DEVARAJ URS MEDICAL COLLEGE,
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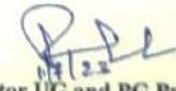

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ABSTRACT

In case of ascending adenocarcinoma, colorectal cancer (CRC) tends toward lymphatic and to the distal area frequent metastatic spread. A growing number of young adults are being diagnosed with colorectal cancer. Both cell-cell and cell-matrix interactions are known to be involved in CRC metastasis.

Syndecan-1 is expressed in colorectal cancer and is involved in CRC metastasis. Syndecan-1 is expressed in colorectal cancer and is involved in CRC metastasis. Syndecan-1 is expressed in colorectal cancer and is involved in CRC metastasis.

Objectives: To determine the expression of Syndecan-1 in the tumor tissue of colorectal adenocarcinoma and correlate the expression of Syndecan-1 with histopathological grading and staging of colorectal adenocarcinoma.

Results: The expression of Syndecan-1 was significantly higher in tumor with poor differentiation and lower in tumor with good differentiation. There was positive correlation between expression of Syndecan-1 and histopathological grading and staging of colorectal adenocarcinoma.

Conclusion: Poorly differentiated adenocarcinoma of the colon and rectum are linked to poor clinical outcomes, while well-differentiated adenocarcinoma express Syndecan-1 at high levels. As a result, elevated Syndecan-1 expression can be used routinely to evaluate prognosis in patients with colorectal cancer.

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LIST OF ABBREVIATIONS

CRC- Colo Rectal Carcinoma

APR- Abdomino-Perineal Resection

H&E- Hematoxylin and Eosin

HPF- High Power Fields

IHC- Immuno Histo Chemistry

AJCC- American Joint Committee on Cancer

ABSTRACT

BACKGROUND: In terms of mortality from cancer, colorectal cancer (CRC) ranks second and is the third most frequent malignancy worldwide. Syndecan-1 acts in both cell-cell and cell-matrix interactions. Controlling cell division, movement, and structure of the cell is one of its functions. Syndecan-1 expression was lower than surrounding normal epithelium in many types of malignancies and loss of its expression is associated with a poor prognosis in various malignancies.

AIMS & OBJECTIVES: To determine the proportion of Syndecan 1 in the tumor proper of colorectal carcinoma and its association with histopathological grading and staging.

MATERIALS AND METHODS: 95 colorectal cancer cases that underwent surgical resection in total were examined. All cases' H & E slides were examined, and immunohistochemistry was run against Syndecan-1. IHC expression levels were assessed, divided into groups based on high and low expression, and these values were compared to clinicopathological information about the cases, including age, sex, histological grading, lymph node status, and staging. IBM SPSS Statistics, Somers, NY, USA. Software was used in determining p value.

RESULTS: Based on 95 samples, Syndecan-1 expression was significantly low in tumours with poor differentiation and increased in well-differentiated CRCs and statistically significant with CRC malignancy grading (p 0.001). No correlation was found between age, sex, tumour site, tumour stage, vascular invasion, or perineural invasion.

CONCLUSION: Poorly differentiated colon and rectum adenocarcinomas express little Syndecan-1, but well-differentiated ones do. Thus, epithelial Syndecan-1 expression can be employed in all colorectal cancer cases to assess prognosis.

KEYWORDS: Colorectal carcinoma, Syndecan-1, Prognosis of CRCs

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INTRODUCTION

INTRODUCTION

One of the most common malignancies to be discovered is colorectal carcinoma (CRC), which also has a high mortality rate among cancer patients. Still, the risk of developing CRCs sporadically in Asian populations is low to minimal.^{1,2}

Numerous factors act at the molecular level, like DNA mismatch repair, microsatellite repeat regions, Mutations in the oncogenes KRAS and SMAD2 and SMAD4 as well as environmental variables all play critical roles in the progression to malignant colorectal tumours. CRCs are more common among the aged 60-70 years. Various other factors are considered risk factors for developing CRCs, including polyps, adenomas, diet (red meat, animal fat, alcohol), sedentary lifestyle, obesity, and positive family history. It is a slow-growing tumor and may remain asymptomatic for years.³

The tumor, nodes, and metastasis staging (TNM) give the prognosis of CRCs. The American Joint committee on cancer recommends additional prognostic factors that should be determined and reported to indicate the prognosis of CRCs.^{4,5}

Among the recommended parameters for CRC prognosis are serum carcinoembryonic antigen (CEA) levels, lymph nodes showing tumor cells within the area of its lymphatic drainage, and vascular and neural invasion, which shows a poorer prognosis.⁵

Multiple novel markers have emerged as important in determining CRC prognosis, according to recent studies. Syndecan-1 is one such marker; it facilitates communication between cells and between cells and their surrounding matrix. Syndecan-1 contributes to the regulation of cell division, migration, and morphology. Plasma cells and other

epithelial cell types in normal tissues express Syndecan-1.⁶

Expression of Syndecan-1 over the epithelial surface is deregulated in several cancers, while loss of its epithelial expression shows a bad prognosis in colorectal carcinoma.

Evaluation of Syndecan-1 by immunohistochemistry allows the identification of changing patterns of its expression which is involved in the progression and differentiation of the CRCs.^{7,8}

AIMS & OBJECTIVES

OBJECTIVES OF THE STUDY

1. To determine the proportion of Syndecan 1 in the tumor proper of colorectal carcinoma
2. To correlate the association of Syndecan 1 with histopathological grading and staging of colorectal carcinoma.

REVIEW OF LITERATURE

REVIEW OF LITERATURE

EMBRYOLOGY OF LARGE INTESTINE.^{9,10}

The primitive tube, which forms the foregut, midgut, and hindgut, includes the intestines from its posterior part. A relatively straight cylindrical primitive tube transforms into a folded complex of tubes that forms the characteristic adult intestinal tract.

The superior mesenteric artery, which comes from the aorta, sends blood through this tube to the midgut. Midgut forms the caecum & appendix. Along with continuation from caecum ascending colon, and most of the transverse colon are formed.

Around the sixth week of life inside the womb, the midgut loop forms. By the tenth week, it connects to the omphalomesenteric duct.

The midgut loop has a cranial and caudal limb, and mesentery suspends these loops in the abdominal cavity. The cranial loop multiplies and forms the intestinal loops. The caudal loops include the caecal swelling, which appears in the ante mesenteric border of the midgut loop, which further grows slowly in the apex, forming an appendix.

The remaining portion of the large intestine is the hindgut, which comprises of the left side of one-third of the transverse colon up to the anal canal and is fed by the inferior mesenteric artery. The cloaca that forms at the end of the hindgut contributes to the formation of the anal canal and rectum.

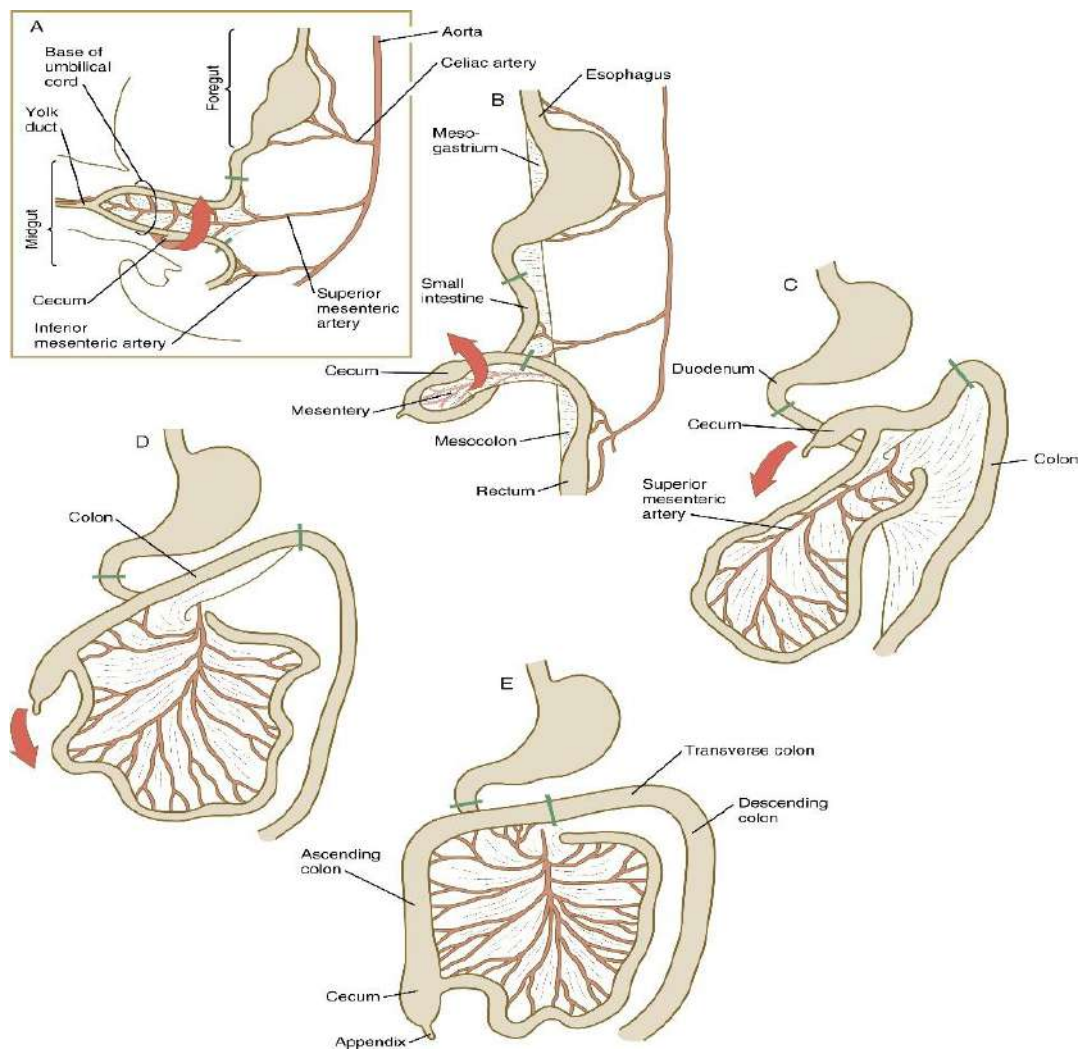


Figure 1. The stages of large intestine development and its rotation. (A) the fifth week of development. (B) the sixth week of intrauterine life. (C) the eleventh week of intrauterine life. (D) twelfth week of intrauterine life. (E) after twelve weeks of intrauterine life. (Image from Human Embryology and Developmental Biology Sixth Edition.⁹)

Anatomy of the large intestine^{11,12}

Colon starts from ileocecal junction and extends till anus. Colon begins in right iliac fossa and ascends superiorly in the lateral region till right hypochondriac region where it turns to left forming right colic flexure and continues as transverse colon. To produce the left colic flexure, the muscle twists to the left and curls on the abdomen's left side. At this point, the colon descends to form the descending colon, continues as the sigmoid colon in the pelvis, and transforms into the rectum, which is located at the level of the third sacral vertebrae in the smaller pelvis. At the level of the pelvic diaphragm, the rectus muscle forms the anal canal.

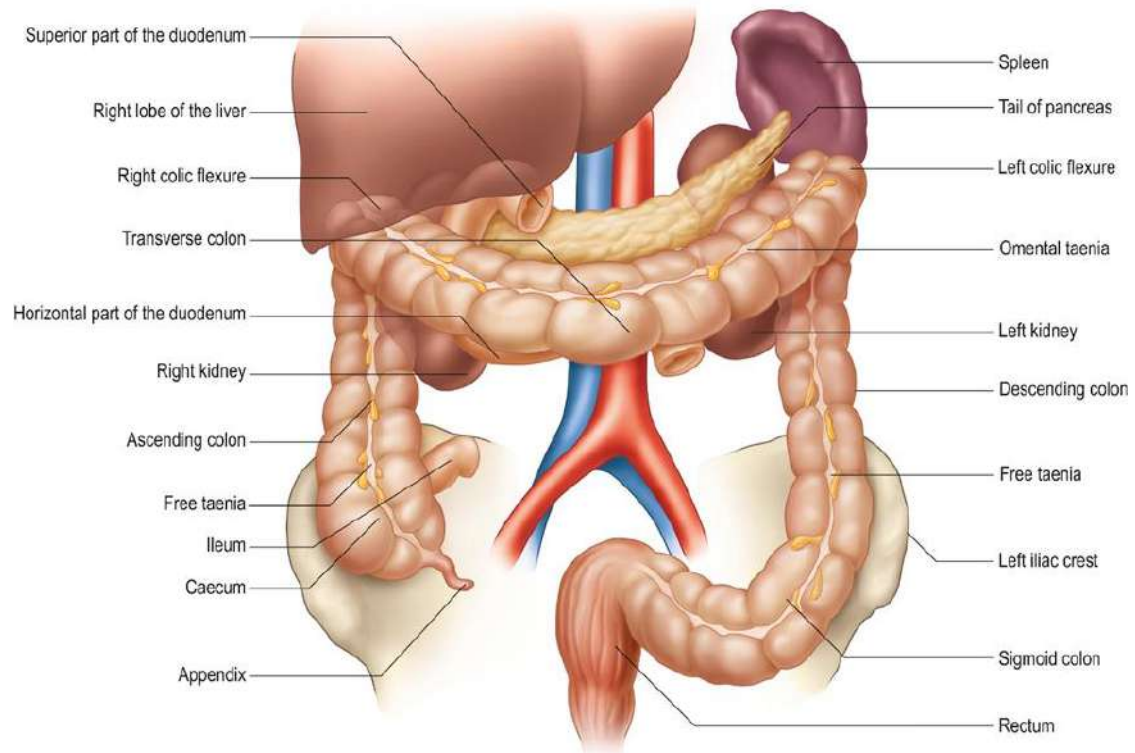


Figure 2. Large intestine in abdominal cavity (Image from Gray's Anatomy. 42nd edition¹¹)

HISTOLOGY OF LARGE INTESTINE^{14, 15}

The large intestine generally has four separate layers histologically, with minor alterations, like other regions of the alimentary canal. The simple to tall columnar epithelium that lines the large intestine digs into the surface to a depth of about 0.5 mm, creating crypts in the mucosa. The lining of the colon has a large number of goblet cells. The cellular lamina propria lies beneath the mucosal layer and may contain lymphoid clusters. Lamina propria is followed by the submucosa and muscularis propria. There is a significant amount of fully developed adipose tissue in the submucosa. The muscularis propria refers to the smooth muscle cells present in the colon. The large intestine is innervated by Meissner's plexus, which is between the submucosal layer and muscularis propria, and Auerbach's plexus, which lies between the layers of muscularis propria. Both the ascending and descending colon has adventitia due to its presence in retroperitoneal. The mesentery, which contains the remainder of the colon is enclosed by serosa. With a few of the following exceptions, the rectum and colon are comparable to one another. The rectum lacks taenia, lacks appendices epiploicae, has a continuous longitudinal muscle coat, and has serous layer covering on its lateral sides in the upper third of the rectum.



Figure 3. Histological layers of the colon (a) mucosa (b) submucosa (c) muscularis propria (d) serosa (e) lymphoid aggregates in submucosal layer (Image from Pathology Outlines – Anatomy & histology¹³)

Large intestines lymphatic drainage¹¹

Large intestine lymphatic drainage occurs after mesenteric artery blood flow. The result is that the superior mesenteric lymph nodes get lymph nodes from the ascending colon, the proximal section of the transverse colon, and the caecum. The lymph nodes that border the path of the inferior mesenteric artery drain the distal portion of the transverse colon until rectum.

Epidemiology of colorectal cancer¹⁴⁻¹⁶

CRCs are linked to a high rate of morbidity and deaths across North America and Europe, as well as the areas with similar dietary practices and lifestyles to those described above. CRCs account for 10% of malignancies which indicates it to be the third most common cancer next to breast and lung cancers accounting for 11.7% and 11.4% of overall cancers diagnosed and its related deaths worldwide.

In the year 2020, nearly 1.93 million of the newly diagnosed CRCs and around 0.9 million of CRC related deaths were registered. Incidence of CRC was 19.5 per million of CRC cases which include 23.4 per million in males and 16.2 per million in females. Mortality related to CRC was 9 per million of worldwide population in 2020. Highest incidence of mortality was seen in Northern European zone accounting for 33.6 per million and South-Central Asia with 5.5 per million as second highest rate for CRC related deaths.

CRC was among the fifth most common cancer of incidence in India, with record of 65358 newly diagnosed cases in 2020. Rate of incidence in Indian population was 4.8 per million and mortality rate was 2.8 per million.

In India, colorectal cancer is fifth among all cancers in incidence, accounting for 65358 new cases (40408 in males, 24950 in females) in 2020. The incidence rate is 4.8 per 100,000, and the mortality rate is 2.8 per 100,000 in India.

Risk factors for development of colorectal carcinoma^{14,16}

Increased Risk (Convincing or Probable Evidence)

- Intake of alcohol above 30 g per day
- Processed meat and meats preserved by smoking and adding up of chemicals for preservation
- Increased BMI

Increased Risk (Limited suggestive evidence)

- Tobacco smoking
- Mutagens and carcinogens in environment
- Foods that has heme iron
- Charbroiled and fried meat or fish which contain heterocyclic amines
- Reduced intake of fruits
- Inadequate intake of non-starchy vegetables
- Dysbiosis of intestinal microbiota

Decreased Risk (Convincing or probable evidence)

- Usage of drugs like calcium supplements, NSAIDs, Aspirin, COX-2 inhibitors and Oestrogen used in hormonal replacement therapy
- Consumption of dairy products
- Foods rich in fibers and whole grains
- Low body mass and physical activity

Decreased Risk (Limited suggestive evidence)

- Vitamin C and Vitamin D
- Fish
- Multivitamin supplements

Genetics of colorectal cancers.¹⁶

Many genetic factors are known to be involved in the development of CRCs, some of them are as follows,

- RAS gene mutations seen in over >50% of full-blown CRCs.
- Chromosome 17p deletion at TP53 region occurs in >75% cases.
- Chromosome 5 allele loss seen in 70 % of cases.
- 18q deletion seen in 50-70% cases of CRCs with invasion
- SMAD2 & SMAD4 deletions
- Loss of DCC gene in overt CRCs
- Accumulation of RAS and TP53 gene mutations, LOH on 5q & 18q mutations seen during the transition from benign adenoma to carcinomas
- CRC shows two to more of the above-mentioned alterations in 90% cases

Pathogenesis of colorectal carcinoma³

Colonic adenocarcinoma develops due to epigenetic and genetic alterations which are heterogeneous molecular changes. Most important mechanisms in development of CRCs are activation of APC/ β -catenin seen in transition of classic adenoma to carcinoma, and the association of repeated microsatellite regions caused by DNA mismatch repair mechanisms.

APC/ β -catenin pathway and DNA mismatch repair pathway occurs in a stepwise manner which causes several mutations that involve multiple genes. Epigenetic event in progression of both the pathways is methylation-induced gene silencing.

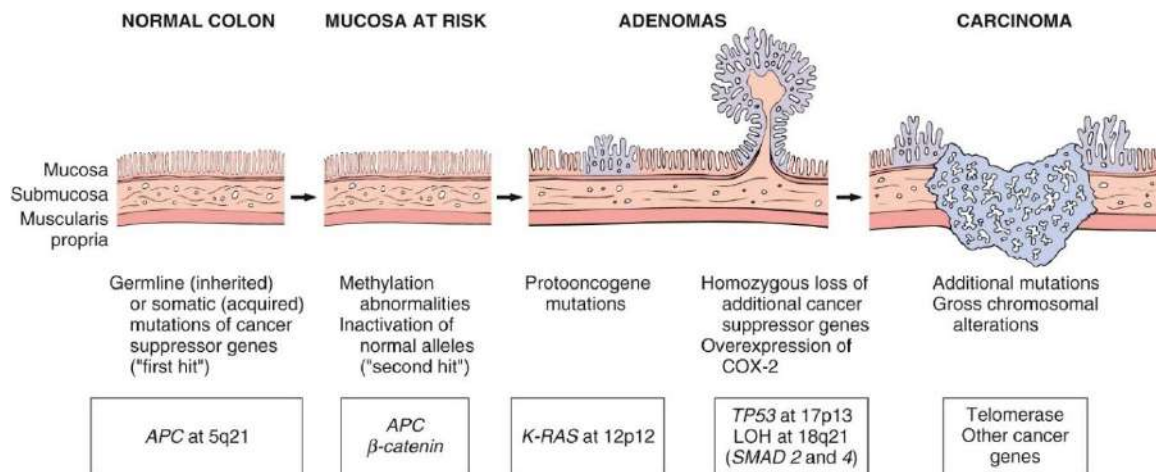


Figure 4. Progression of normal colon to adenomas and carcinomas through genetic alteration (Image taken from Robbins & Cotran Pathologic basis of the disease, 10th edition³)

Transition of classic adenomas into carcinomas occurs in nearly 80% of the cases. Inactivation of both alleles of APC gene is necessary for development of adenomas. Due to the loss of APC gene, β -catenin and TCF which are essential DNA-binding factors form a complex which promotes cell proliferation by activation MYC and cyclin D1 transcription.

In few cases there is KRAS mutations seen among tumors with <1cm size. In adenomas which are >1cm in size, KRAS mutations are seen in nearly 50% cases.

Tumor suppressor genes like SMAD2 and SMAD4 which affects TGF- β signaling is also known to cause CRCs by unrestricted cell growth and cell cycle inhibition.

CRCs show mutation in TP53 gene in form of chromosomal instability and chromosomal deletions. This causes instability of APC/ β -catenin pathway that act as a tumor suppressor.

Along with this methylation of CpG-rich zones, CpG islands and 5' regions which are promoter and transcriptional start sites that help in cell differentiation is mutated.

Telomerase expression is increased in cases with advanced disease.

Loss of BAX gene which helps in survival of mutated clones is seen in progressive CRCs.

Overall mutations in BRAF, MSI instability and methylation of MLH1 forms the carcinogenic pathway is a well-established phenomenon in CRCs.

Clinical features in colorectal carcinoma^{17,18}

CRC symptoms might manifest as sharp abdominal pain or as nonspecific, long-lasting symptoms. Anemia, which is a symptom of colon cancer on the right side, causes anemia-related symptoms like weakness, dullness, and decreased activity. Tumor on the left side of the colon presents with bleeding and tenesmus. Very rarely mass per abdomen will be the constitutional symptom in CRCs. Rarely hematuria and infection in urinary tract due to fistula in urinary bladder or fistula in gastrocolic region causing severe diarrhea.

Symptoms of the CRCs are non-specific. Stratification of patients as high-risk category for urgent investigations is based on the symptoms of changed bowel habits, mass per rectum and chronic anemia. Accordingly high-risk and low-risk symptoms are as follows

Higher risk

- Rectal bleed
- Consistent rectal bleeding without soreness, discomfort, itching, lumps, prolapse, or pain.
- Changes in bowel habits, such as hard stools or increased frequency of passing stools, that last for six weeks in individuals of all ages. (> 60 years)
- Right-sided abdominal lump that can be felt (all ages)
- Rectal lump that can be felt (not pelvic) (all ages)
- Unexplained anemia due to a lack of iron (all ages)

Low risk

- People who are not anemic
- Rectal bleeding with an evident external origin, such as anal fissure
- No lump per abdomen or rectum (all ages)
- Rectal bleeding with no evident change in bowel habits.
- Temporary changes in bowel habits brought on by passing firmer stools or less frequent urination
- Intestinal discomfort without blockage (all ages)

WHO HISTOLOGICAL CLASSIFICATION OF TUMORS OF THE COLON AND RECTUM⁴

BENIGN EPITHELIAL TUMOURS AND PRECURSORS:

Serrated dysplasia, low grade

Serrated dysplasia, high grade

Hyperplastic polyp, micro vesicular type

Hyperplastic polyp, goblet cell

Adenomatous polyp, low-grade dysplasia

Adenomatous polyp, high-grade dysplasia

Tubular adenoma, low grade

Tubular adenoma, high grade

Villous adenoma, low grade

Villous adenoma, high grade

Tubulovillous adenoma, low grade

Tubulovillous adenoma, high grade

Advanced adenoma

Glandular intraepithelial neoplasia, low grade

Glandular intraepithelial neoplasia, high grade

MALIGNANT EPITHELIAL TUMOURS:

Adenocarcinoma NOS

Serrated adenocarcinoma

Adenoma like adenocarcinoma

Micropapillary adenocarcinoma

Mucinous adenocarcinoma

Poorly cohesive carcinoma

Signet ring cell carcinoma

Medullary adenocarcinoma

Adenosquamous carcinoma

Carcinoma undifferentiated, NOS

Carcinoma with sarcomatoid component

Neuroendocrine tumors

Neuroendocrine tumor, grade 1

Neuroendocrine tumor, grade 2

Neuroendocrine tumor, grade 3

L cell tumor

Glucagon-like peptide-producing tumor PP/PYY-producing tumor Enterochromaffin –
cell carcinoid Serotonin-producing tumor

Neuroendocrine carcinoma NOS

Large cell neuroendocrine carcinoma

Small cell neuroendocrine carcinoma

Mixed neuroendocrine–non–neuroendocrine neoplasm (MiNEN)

COMMON COLORECTAL NEOPLASMS¹⁹

EPITHELIAL POLYPS

The majority of the colorectal polyps are those with epithelial genesis. Adenomatous polyps and serrated polyps are two major types into which they can be categorized. Juvenile (retention) polyp is the most frequent colonic polyp seen in children.

TUBULAR ADENOMA

Adenomatous polyps, sometimes referred to as tubular adenomas, are typically evenly distributed throughout the entire large intestine but less frequently in the rectum. These are typically asymptomatic and occasionally lead to changes in bowel habits. They can be sessile or pedunculated and are typically smaller than 1 cm in size. These adenomas are composed of tubular crypts that are closely spaced apart and only include 20% villous tissue. These exhibit cellular crowding and glandular hyperplasia, as well as possible abnormal nuclear characteristics. Increased positive is seen when Carcino embryonic antigen (CEA) expression is immunostained, especially in the atypical parts.

VILLOUS ADENOMA

These are frequently solitary and seen in older age range. Rectum and recto sigmoid areas are the most frequent sites, although due to the lesions' extremely soft consistency, even a digital inspection often misses them. More than 80% of the components in these adenomas are villous. They have a broad base from which finger-like villi emerge. Long papillary structures and a crown-like pattern may be visible under a light microscope. Treatment

varies depending on the size and severity of the lesion. A 29%–70% chance of malignant transformation exists.

SERRATED ADENOMA

These are typically sessile and tiny, not exceeding 5 mm. These adenomas are known as serrated because they exhibit saw-toothed architecture under a microscope. These are distinctive and consist of the glands folding into the lumen. Additional mitotic activity might be observed. Sessile, conventional, and hyperplastic serrated adenomas are the three types of serrated adenomas.

TUBULO VILLOUS ADENOMA

These often combine villous and tubular elements, with 20–80% of the villous element present.

Syndromes associated with colorectal carcinomas are Familial adenomatous polyposis (FAP; also known as polyposis coli), Gardner syndrome, Turcot syndrome and Cowden syndrome.

ADENOCARCINOMA

The tumour cells must completely penetrate the muscularis mucosae into the submucosa in order to be classified as a carcinoma.. They are usually asymptomatic; the most common presentation mode is a change in bowel habits, haematochezia, or anemia. Colonoscopy may aid in the early diagnosis. The growth pattern may be exophytic with intraluminal growth,

diffusely infiltrative/ linitis plastica type with endophytic change, or with complete circumferential involvement.

MUCINOUS CARCINOMA

Malignant cells with more than 50% extracellular mucin pools. Usually associated with microsatellite instability.

SIGNET RING CELL CARCINOMA

The cells should have eccentrically placed nuclei with intracellular mucin, and the cells should comprise more than 50 % of tumor cells.

ADENOSQUAMOUS CARCINOMA

The entity should contain a combination of squamous cell carcinoma and adenocarcinoma elements. There should be more than one component and convincing foci of squamous cell carcinoma.

MEDULLARY CARCINOMA

It is a rare tumor with a reasonably good prognosis and characterized by a solid pattern of cells with a vesicular nucleus, prominent nucleoli, and eosinophilic cytoplasm.

TNM CLASSIFICATION OF COLORECTAL TUMORS⁴

| Tumor | Regional lymph nodes | Distant metastasis |
|---|---|---|
| Tx- Primary tumor cannot be assessed | Nx- regional lymph nodes cannot be assessed | M0- no distant metastasis |
| T0- no evidence of primary tumor | N0- no regional lymph node metastasis | M1- distant metastasis to one or more sites or organs or peritoneal metastasis |
| Tis- carcinoma insitu, intramucosal carcinoma | N1- metastasis to one to three regional lymph nodes | M1a- metastasis to one site or organ without peritoneal metastasis |
| T1- tumor invades the submucosa | N1a- metastasis in one regional lymph node | M1b- metastasis to two or more sites or organs without peritoneal metastasis |
| T2- tumor invades the muscularis propria | N1b- metastasis in two or three regional lymph nodes | M1c- metastasis to peritoneal surface, alone or with other site or organ metastasis |
| T3- tumor invades through the muscularis propria into pericolic or perirectal tissues | N1c- tumor deposits in subserosa or in non-peritonealised pericolic or perirectal soft tissue without regional nodal metastasis | |
| T4- tumor invades visceral peritoneum or invades or adheres to adjacent organ | N2- metastasis in four or more regional lymph nodes | |
| T4a- tumor invades visceral peritoneum | | |
| T4b- tumor invades or adheres to adjacent organs | | |

Table 1: TNM classification of colorectal carcinomas

| | | | |
|------------|--------|---------|-----|
| Stage 0 | Tis | N0 | M0 |
| Stage I | T1, T2 | N0 | M0 |
| Stage IIA | T3 | N0 | M0 |
| Stage IIB | T4a | N0 | M0 |
| Stage IIC | T4b | N0 | M0 |
| Stage IIIA | T1, T2 | N1/ N1c | M0 |
| Stage IIIA | T1 | N2a | M0 |
| Stage IIIB | T3-T4a | N1/ N1c | M0 |
| Stage IIIB | T2, T3 | N2a | M0 |
| Stage IIIB | T1, T2 | N2b | M0 |
| Stage IIIC | T4a | N2a | M0 |
| Stage IIIC | T3-T4a | N2b | M0 |
| Stage IIIC | T4b | N1-N2 | M0 |
| Stage IVA | Any T | Any N | M1a |
| Stage IVB | Any T | Any N | M1b |
| Stage IVC | Any T | Any N | M1c |

Table 2: TNM staging of colorectal carcinoma.

MOLECULAR CLASSIFICATION OF COLORECTAL CARCINOMAS²⁰

Due to heterogenous molecular constituents in development of CRCs a single molecular classification for colorectal carcinomas is not prepared. Based on the proliferation index and tumor differentiation and survival rates associated with their expression CRCs can be classified into four groups.

| Molecular classes | Driving role of biomarkers | Median survival in months |
|-------------------|----------------------------------|---------------------------|
| CRC Novel Class-1 | Low Ki67, high CDX2 & low P53 | 30 |
| CRC Novel Class-2 | High ki67, low CDX2 & low P 53 | 25 |
| CRC Novel Class-4 | High ki67, high CDX-2 & high P53 | 26 |
| CRC Novel Class-4 | High ki67, high CDX-2 & low P53 | 23 |

Table 3: Molecular classification of colorectal carcinomas.

PROGNOSTIC FACTORS IN COLORECTAL CARCINOMA.^{21–43}

Factors associated with bad prognosis or poor outcomes are

- Stage of presentation
- Post treatment stage of carcinoma
- Depth of penetration of the tumor
- Local involvement of peritoneum
- Tumor size >4.5 cm
- Presence of tumor after definitive therapy
- The circumferential margin (CRM) and presence of tumor cell at ≤ 1 mm from CRM and the point of deepest penetration
- The presence of tumour cells in lymph nodes and the total number of lymph nodes that are affected
- Lymphovascular
- Perineural invasion
- CRCs with high-grade and its subtypes like poorly differentiated, undifferentiated carcinomas & signet ring cell variants
- Malignancy grade in the differentiation of carcinomas
- Expansile pattern of infiltration into the stroma
- Fibrosis due to desmoplastic changes in the stroma
- Involvement of the tumor with more of neuroendocrine cells
- Elevated levels of carcinoembryonic protein postoperatively within one year of tumor resection

-
- Lymphocyte infiltration into a tumour has been shown to be a significant prognostic indicator in a number of studies
 - Presentation with perforation or obstruction
 - Presence of genetic aberration in BRAF, RAS genes and mismatch repair deficiency

Factors associated with good prognosis

- Neoadjuvant chemoradiotherapy is associated with considerable tumour response and downstaging in suitably chosen patients with rectal cancer⁴¹
- Increased number of CD4+ and CD25+ T cells
- Presence of tumor beyond the splenic flexure

SYNDECAN 1(CD 138)

There is widespread expression of the transmembrane proteoglycan syndecan-1 (CD138) in both normal and malignant tissues.⁶

Syndecan-1 is encoded by the chromosome 2 gene SDC1.⁴⁴

The core protein of syndecan-1 has two intracellular and extracellular domains. Heparan sulphate and chondroitin sulphate glycosaminoglycans are used in place of the extracellular domain. It is possible to reduce the adhesion between cell-to-cell and cell-to-basement membrane adhesions that are present in the stroma of various tumours by cleaving a portion of this Syndecan-1 at the junction between the cell membrane and extracellular site. This can aid in tumour progression by suppressing, transforming, and migrating.^{7,45}

Reduced Syndecan-1 expression is related with tumour differentiation, progression, and clinical staging in colorectal cancers.⁷ Syndecan-1, E-cadherin, and beta-catenin complex expression are typically disrupted in CRCs.⁸

To evaluate syndecan-1, Western blot, ELISA, and immunohistochemistry techniques are utilized. Evaluation of Syndecan-1 is crucial as a predictive tool because of its significance and role as a target for Indatuximab, a monoclonal antibody combined with a cytotoxic agent, due to its differential expression in tumours.⁶

Together, integrins and FAK are activated more quickly when Sdc-1 is reduced, and this results in signals that promote cancer stem cell characteristics and invasiveness.⁴⁶

Syndecan-1 regulates the molecular mediators of tumour cell survival, proliferation, angiogenesis, and metastasis and is seen expressed by the surface of epithelial cells in mature tissues.⁸

MATERIALS AND METHODS

MATERIALS AND METHODS

STUDY DESIGN: Laboratory based observational Study

SOURCE OF DATA: The Department of Pathology at Sri Devaraj Urs Medical College, Tamaka, Kolar, received surgically resected colorectal cancer specimens from R.L. Jalappa Hospital and Research Center from October 2019 to November 2022. Additionally, the department also retrieved data and paraffin blocks for all colorectal cancer cases from the department's archives for the years 2008 to 2022.

DURATION OF STUDY: Two years

METHOD OF COLLECTION OF DATA: The Department of Pathology's archives were used to gather all Colorectal Cancer cases from 2008 to 2022, along with clinical information.

INCLUSION CRITERIA: All cases with histological diagnosis of colorectal carcinoma were admitted and underwent surgical resection in RLJH from 2008 to 2022.

EXCLUSION CRITERIA:

- Patients subjected to neoadjuvant radiotherapy/chemotherapy before excision of colorectal carcinoma.
- Patients who underwent chemotherapy for other cancers over the past five years.

SAMPLE SIZE: 91

Based on Antigony Mitselou et al.⁸ survey of Syndecan-1 expression in colorectal carcinomas (62.32%) with 95% confidence intervals and a 10% absolute error, the sample size for the current investigation has been estimated at 91.

Formula for calculating sample size

Sample size is equal to
$$\frac{Z^2 \cdot p(1-p)}{d^2}$$

Z 1- = Standard normal variation in this case.

P = Population Expected Proportion based on Prior Studies

10% absolute inaccuracy is given by d.

In the current investigation, 91 colorectal cancer patients were included using the aforementioned values at a 95% Confidence Interval.

METHODOLOGY

All the clinicopathological data of colorectal carcinoma cases, such as age, sex, histological grading, lymph node status, and staging, were collected. The resected specimens of all colorectal carcinoma, confirmed histopathologically, were included in the study. H & E To perform immunohistochemistry against Syndecan-1 (rabbit monoclonal antibody, prediluted, Biogenex) for all cases of colorectal cancer, slides from all cases were evaluated, tumour tissue was chosen, and the peroxidase and anti-peroxidase method was used. Positive and negative controls were carried out on each patient.

IMMUNOHISTOCHEMISTRY PROTOCOL:

- Sections are cut at 3-4 μm , floated on positively charged slides, incubated at 37 degrees 1 day, and further at 58 degrees overnight.
- Do not allow it to dry at any stage.
- Carry out steps of incubation with the antibody at 37 degrees.
- Deparaffinization carried out in 15-minute intervals with xylene-I and xylene-II.
- Dextylinisation with Absolute Alcohol I and II administered for a minute each
- Dealcoholisation for 1 minute
- Distilled water five min-Washing
- Antigen Retrieval using the microwave at power 10 for 6 minutes in citrate buffer pH 6.0
- Transfer to TBS buffer pH (7.6)-15 minutes three times, washing for 5 minutes
- Preparation of peroxidase block for 30 minutes.

-
- Power Block will be done for 10 minutes.
 - Drain and cover section with Primary Antibody
 - Wash with TBS buffer for 5 minutes three times to wash unbound antibodies
 - Secondary Antibody for 30 minutes
 - Super enhancer
 - TBS buffer wash 5 min three times
 - Colour development with a working color development solution for 5-8 minutes
 - Distilled water wash for 5 minutes.
 - Counterstain with Harris hematoxylin for one minute
 - Dehydration
 - Mount with DPX

POSITIVE CONTROL: Tonsil was taken as a positive control.

EVALUATION OF SYNDECAN-1 IMMUNOSTAINING⁴⁷

Syndecan-1 membranous/cytoplasmic staining was graded on a scale of 0 to 3, with 0 denoting no staining, 1 denoting faint staining, 2 denoting moderate staining, and 3 denoting high staining.

The following formula was used to determine the percentage of positively stained cells: (0, no stain; 1, 1-25%; 2, 26-50%; 3, > 50%).

A final score of 1-6 was obtained by combining the intensity and percentage scores.

A low-expression group (scores 0-2) and a high-expression group are created from the total score (scores 3-6).

STATISTICAL ANALYSIS:

The study's data were input into a Microsoft Excel data sheet and analyzed using SPSS 22 software. In order to depict the current variable and its values, Data was analyzed using frequency and proportional analysis. To determine if there was a statistically significant relationship between two sets of qualitative data, the chi-square test or Fischer's exact test (only for 2x2 tables) was used. Continuous data were used to illustrate the mean and standard deviation.

Data visualization: Several types of graphs were produced using Microsoft Word and Excel.

After considering all of the guidelines for statistical tests, a p-value (Probability that the result is accurate) of 0.05 was deemed to be statistically significant.

Statistical software utilized for analysis: MS Excel and IBM SPSS Statistics, Somers, NY, USA.

OBSERVATION AND RESULTS

RESULTS

Age distribution of subjects (n=95)

| | Frequency | Percent |
|----------|-----------|---------|
| 20-29yrs | 3 | 3.2 |
| 30-39yrs | 5 | 5.3 |
| 40-49yrs | 15 | 15.8 |
| 50-59yrs | 20 | 21.1 |
| 60-69yrs | 33 | 34.7 |
| 70-79yrs | 15 | 15.8 |
| 80-89yrs | 4 | 4.2 |
| Total | 95 | 100.0 |

Table 4: Categorical subjects distribution by to age.

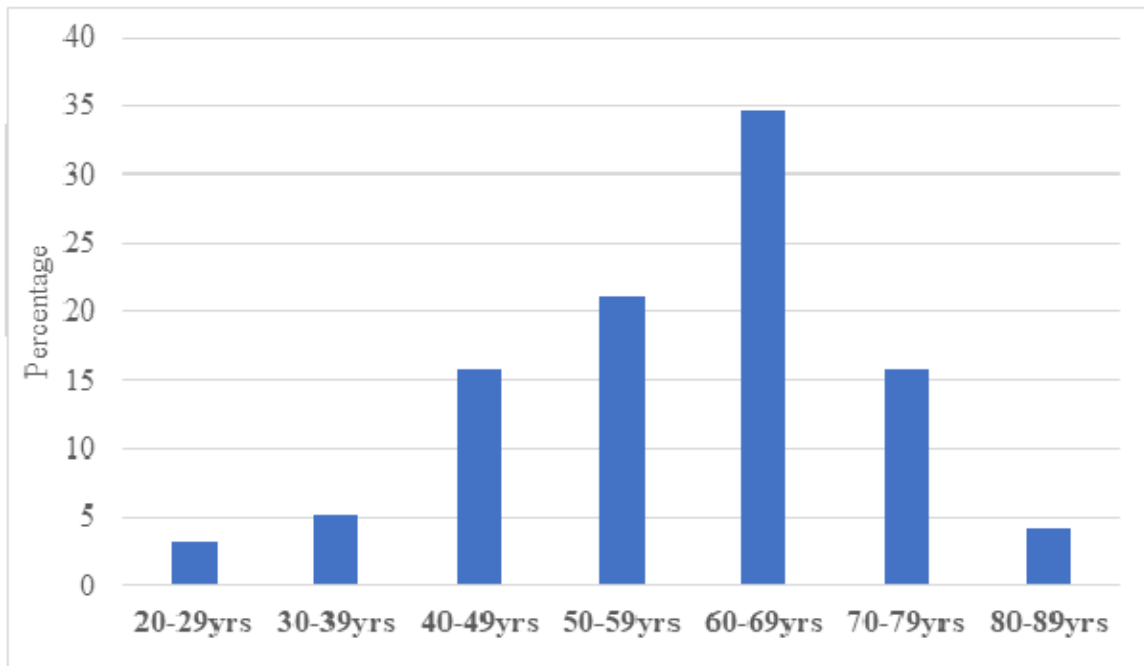


Chart 1: Categorical subjects distribution by to age.

Gender distribution of subjects (n=95):

| | Frequency | Percent |
|--------|-----------|---------|
| Female | 42 | 44.2 |
| Male | 53 | 55.8 |
| Total | 95 | 100.0 |

Table 5: Categorical subjects distribution by sex

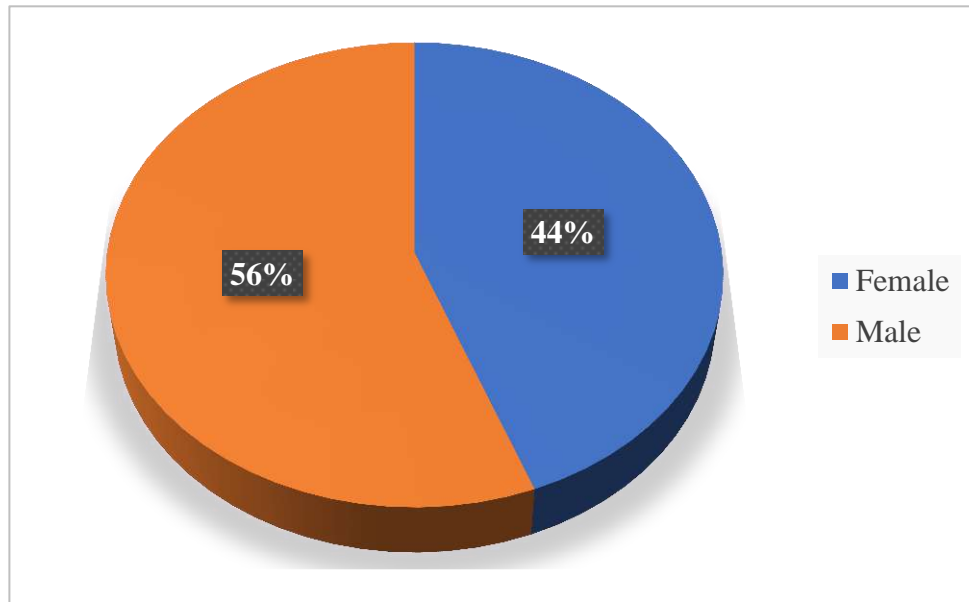


Chart 2: Categorical subjects distribution by sex

Categorical subjects distribution by site of tumor (n=95):

| | Frequency | Percent |
|------------------|-----------|---------|
| Ascending colon | 21 | 22.1 |
| Descending colon | 11 | 11.6 |
| Rectum | 41 | 43.2 |
| Sigmoid colon | 15 | 15.8 |
| Transverse colon | 7 | 7.4 |
| Total | 95 | 100.0 |

Table 6: Categorical subjects distribution by site of tumor

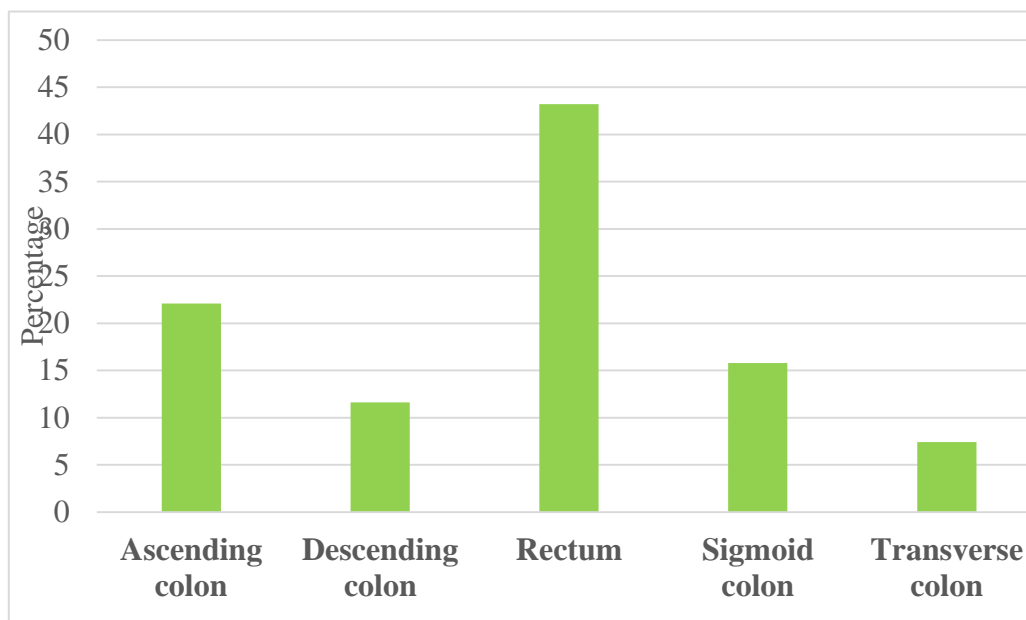


Chart 3: Categorical subjects distribution by site of tumor

Distribution of subjects according to malignancy grading (n=95):

| | Frequency | Percent |
|---------------------------|-----------|---------|
| Well-differentiated | 36 | 37.9 |
| Moderately differentiated | 40 | 42.1 |
| Poorly differentiated | 19 | 20.0 |

Table 7: Distribution of subjects according to malignancy grading

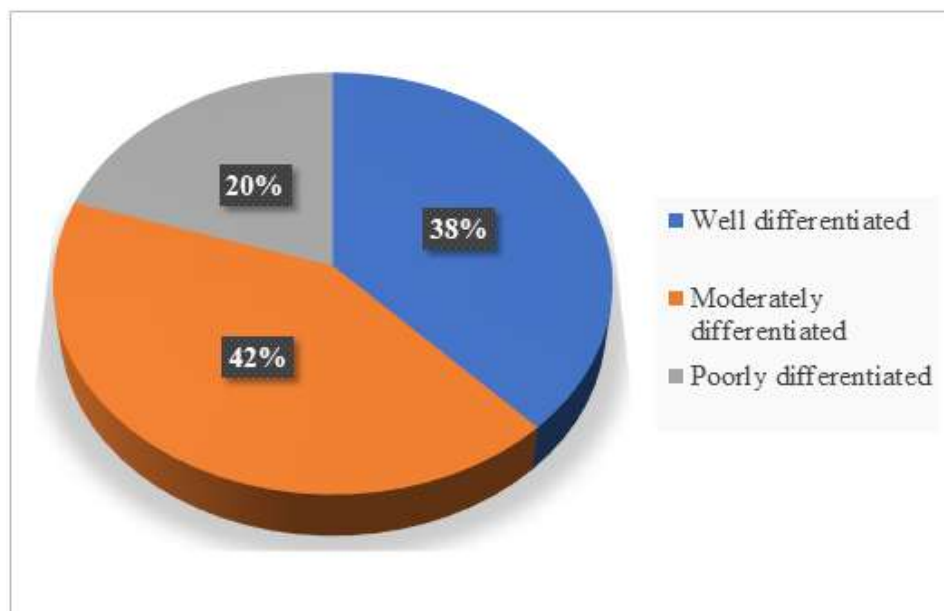


Chart 4: Distribution of subjects according to malignancy grading

Categorical subjects distribution by stage of the tumor (n=95):

| | Frequency | Percentage |
|-----------|-----------|------------|
| Stage I | 25 | 26.31 |
| Stage II | 29 | 30.52 |
| Stage III | 41 | 43.15 |

Table 8: Categorical subjects distribution by stage of the tumor

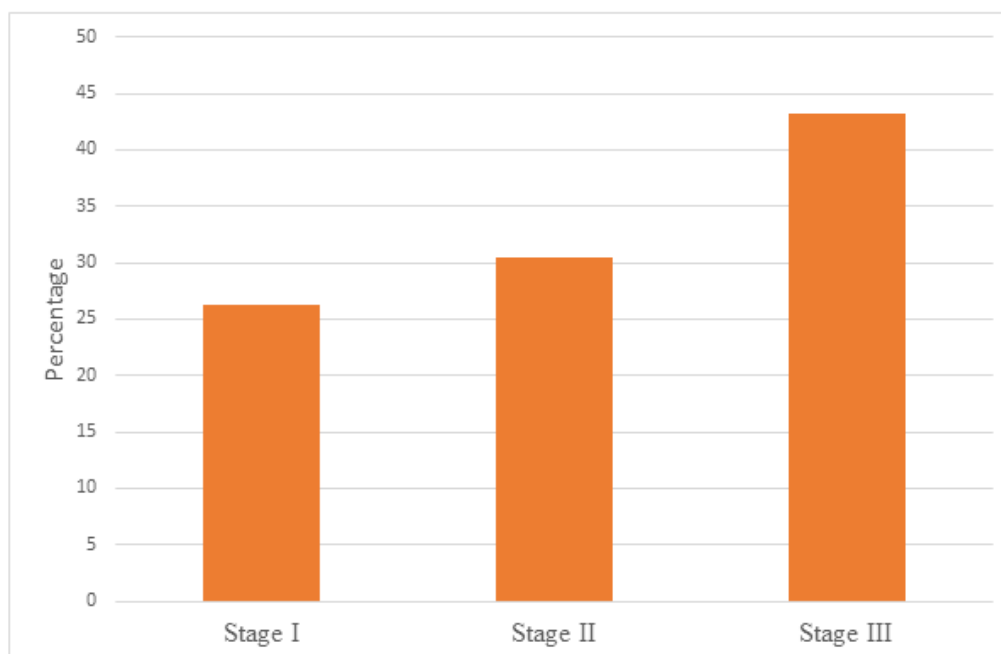


Chart 5: Categorical subjects distribution by stage of the tumor

Categorical subjects distribution by tumor size (n=95):

| | Frequency | Percentage |
|----|-----------|------------|
| T1 | 4 | 4.21 |
| T2 | 26 | 27.36 |
| T3 | 53 | 55.78 |
| T4 | 12 | 12.63 |

Table 9: Categorical subjects distribution by tumor size

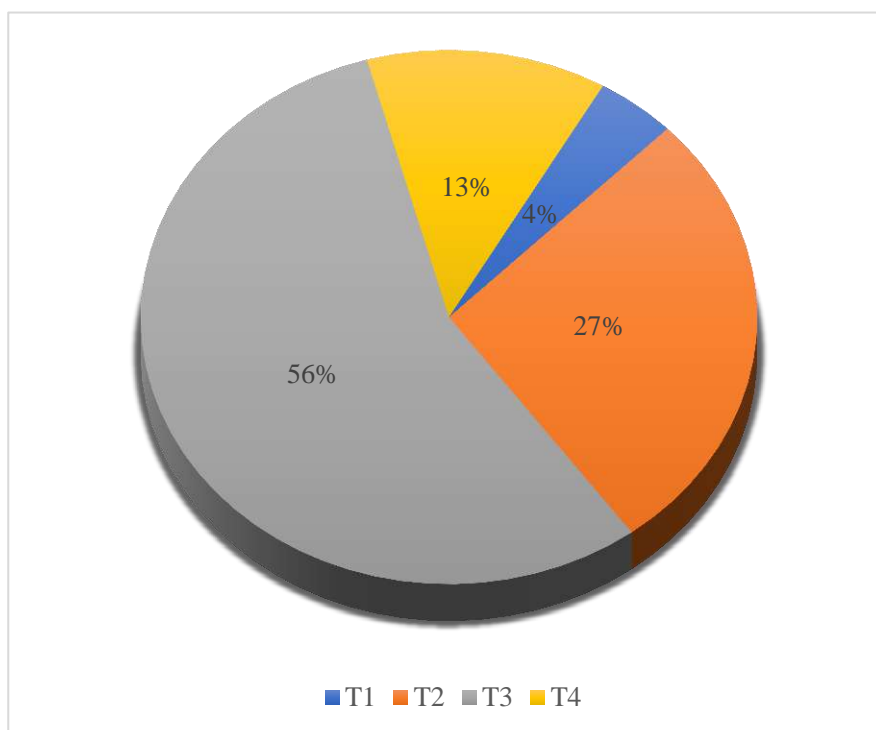


Chart 6: Categorical subjects distribution by tumor size

Categorical subjects distribution lymph node status (n=95):

| | Frequency | Percent |
|----------|-----------|---------|
| Negative | 55 | 57.9 |
| Positive | 40 | 42.1 |
| Total | 95 | 100.0 |

Table 10: Categorical subjects distribution lymph node status

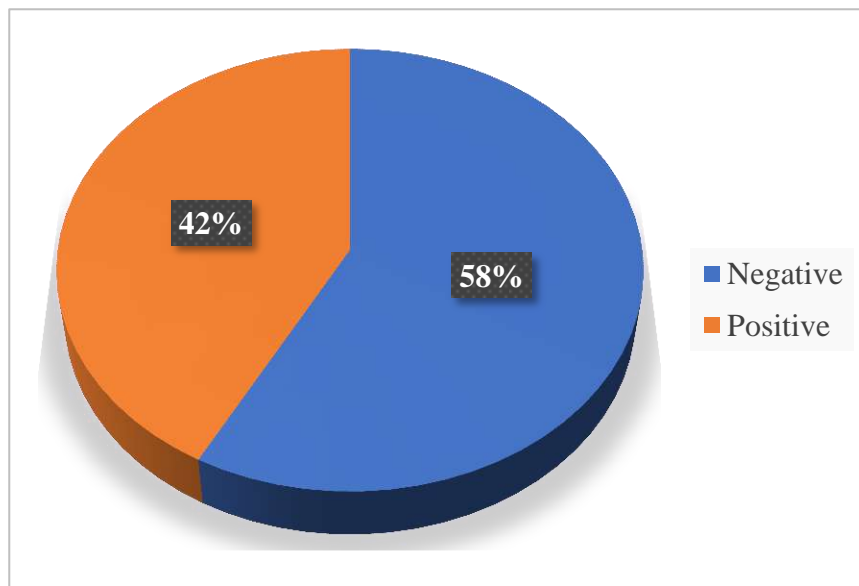


Chart 7: Categorical subjects distribution lymph node status

Distribution of subjects according to Vascular Invasion (n=95):

| | Frequency | Percent |
|---------|-----------|---------|
| Absent | 87 | 91.6 |
| Present | 8 | 8.4 |
| Total | 95 | 100.0 |

Table 11: Distribution of subjects according to Vascular Invasion

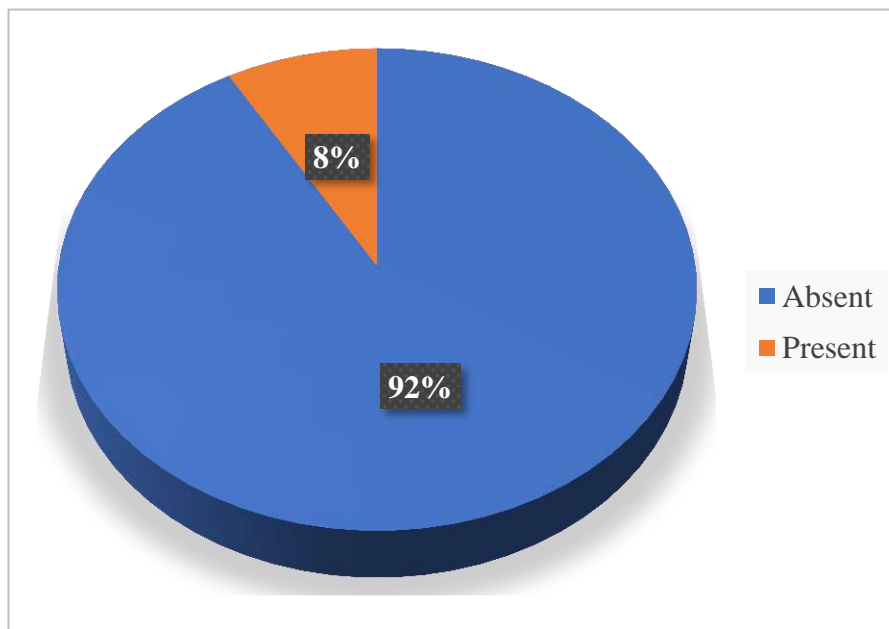


Chart 8: Distribution of subjects according to Vascular Invasion

Distribution of subjects according to Perineural invasion (n=95):

| | Frequency | Percent |
|---------|-----------|---------|
| Absent | 93 | 97.9 |
| Present | 2 | 2.1 |
| Total | 95 | 100.0 |

Table 12: Distribution of subjects according to Perineural invasion

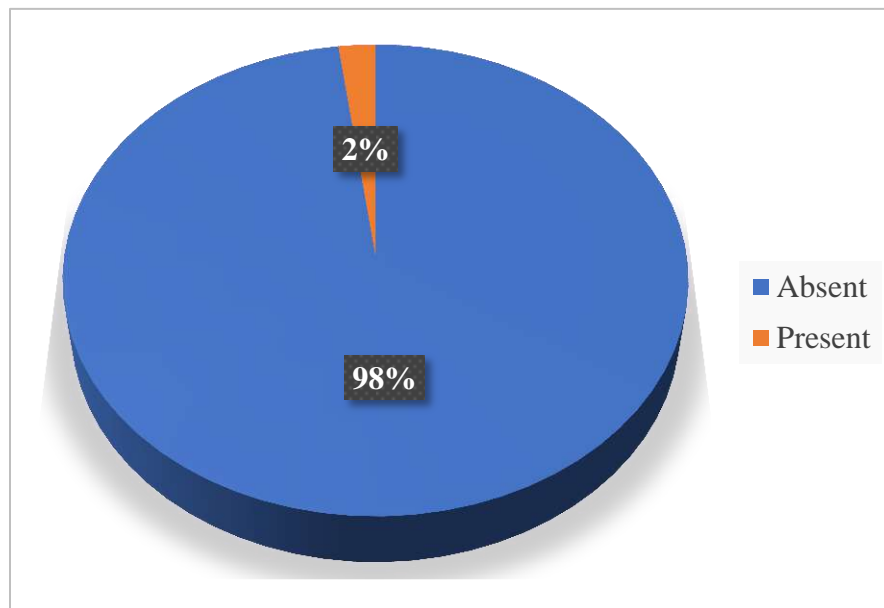


Chart 9: Distribution of subjects according to Perineural invasion

Categorical subjects distribution by Syndecan-1 expression (n=95):

| | Frequency | Percent |
|-------|-----------|---------|
| HIGH | 67 | 70.5 |
| LOW | 28 | 29.5 |
| Total | 95 | 100.0 |

Table 13: Categorical subjects distribution by Syndecan-1 expression

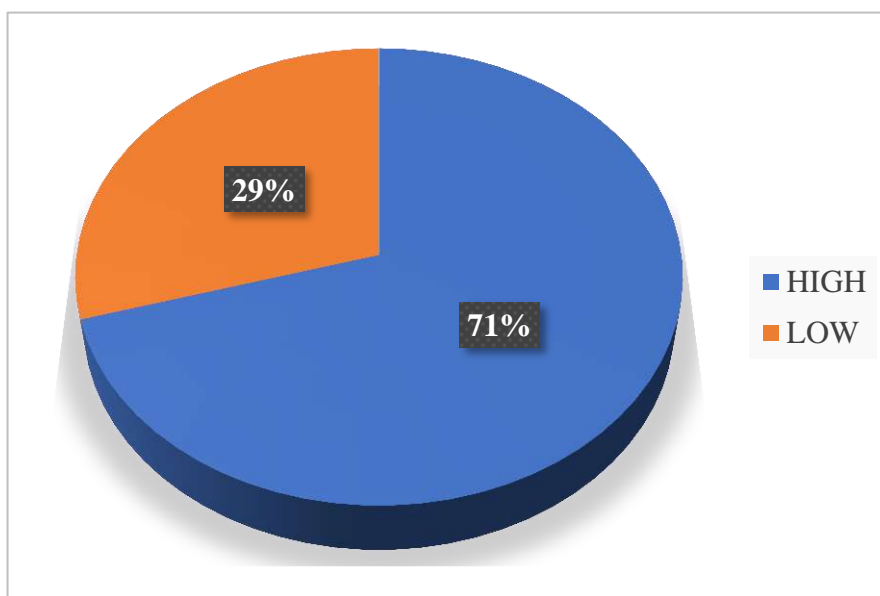


Chart 10: Categorical subjects distribution by Syndecan-1 expression

Distribution of subjects according to age group and expression

| | High expression | | Low expression | |
|----------|-----------------|-------|----------------|-------|
| | N | % | N | % |
| 20-29yrs | 1 | 33.3% | 2 | 66.7% |
| 30-39yrs | 4 | 80.0% | 1 | 20.0% |
| 40-49yrs | 10 | 66.7% | 5 | 33.3% |
| 50-59yrs | 15 | 75.0% | 5 | 25.0% |
| 60-69yrs | 24 | 72.7% | 9 | 27.3% |
| 70-79yrs | 11 | 73.3% | 4 | 26.7% |
| 80-89yrs | 2 | 50.0% | 2 | 50.0% |

Table 14: Distribution of subjects according to age group and expression

p-value 0.750, There was no discernible gap between high and low expression in this age bracket.

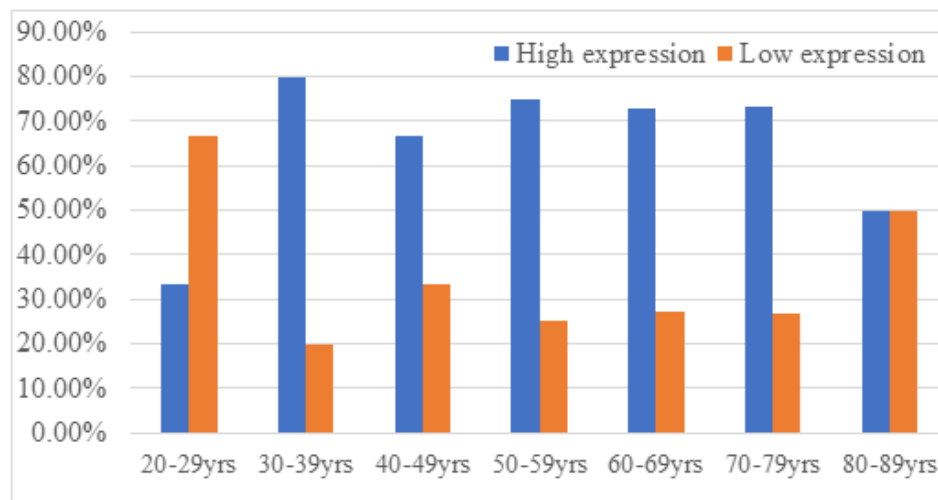


Chart 11: - Categorical subjects distribution by age group and Syndecan-1 expression.

Categorical subjects distribution by sex and Syndecan-1 expression

| | High expression | | Low expression | |
|--------|-----------------|-------|----------------|-------|
| | N | % | N | % |
| Female | 34 | 81.0% | 8 | 19.0% |
| Male | 33 | 62.3% | 20 | 37.7% |

Table 15: Categorical subjects distribution by sex and Syndecan-1 expression

p-value 0.069, statistically significant difference was not found between high and low expression for sex.

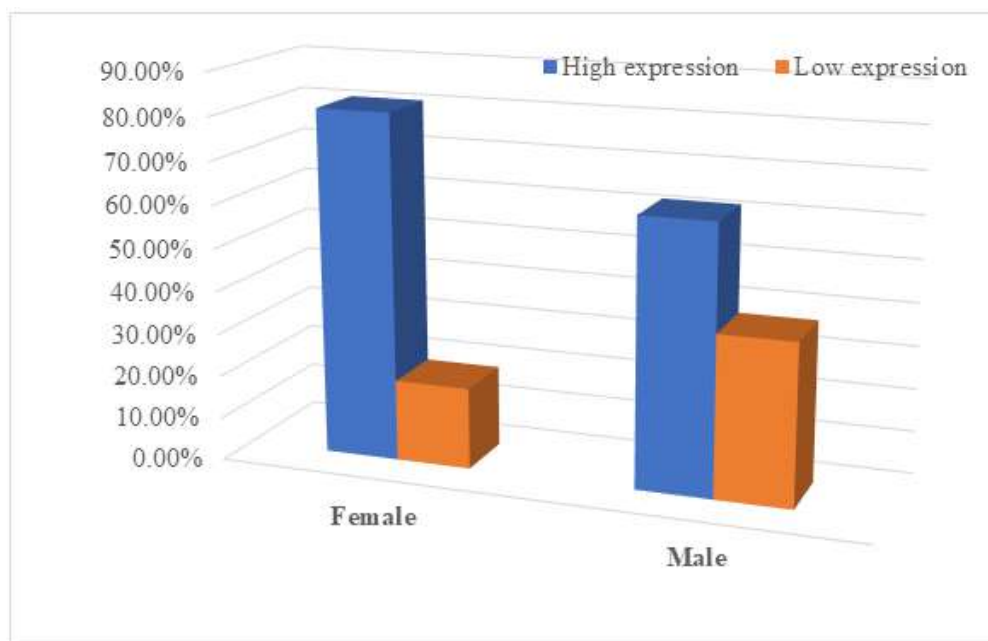


Chart 12: Categorical subjects distribution by sex and Syndecan-1 expression

Categorical subjects distribution by site and Syndecan-1 expression

| | High expression | | Low expression | |
|------------------|-----------------|--------|----------------|-------|
| | N | % | N | % |
| Ascending colon | 15 | 71.4% | 6 | 28.6% |
| Descending colon | 8 | 72.7% | 3 | 27.3% |
| Rectum | 27 | 65.9% | 14 | 34.1% |
| Sigmoid colon | 10 | 66.7% | 5 | 33.3% |
| Transverse colon | 7 | 100.0% | 0 | .0% |

Table 16: Categorical subjects distribution by site and Syndecan-1 expression

p-value 0.478, no statistically significant difference was found between high and low expression for the site.

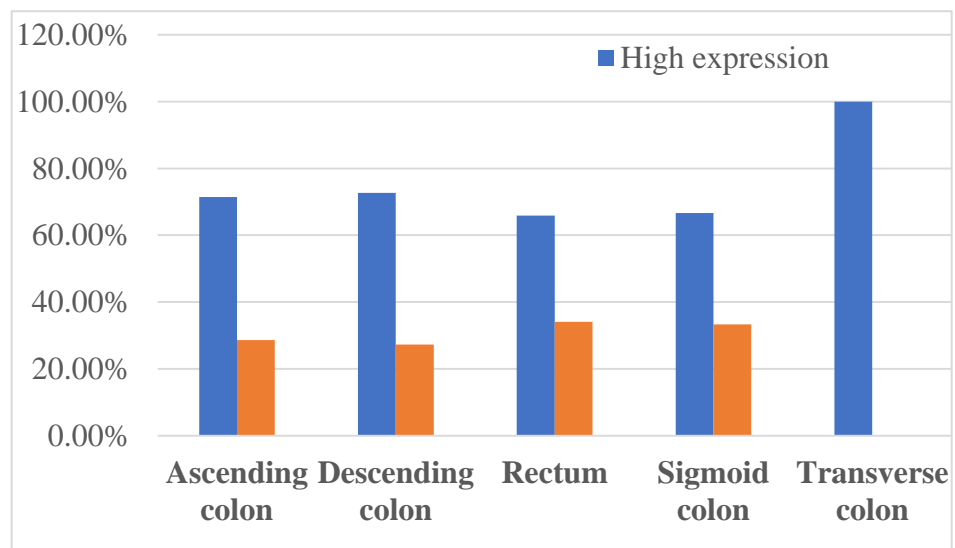


Chart 13: Categorical subjects distribution by site and Syndecan-1 expression

Categorical subjects distribution by malignancy grading and Syndecan-1 expression:

| | High expression | | Low expression | |
|---------------------------|-----------------|-------|----------------|-------|
| | N | % | N | % |
| Well-differentiated | 35 | 97.2% | 1 | 2.8% |
| Moderately differentiated | 30 | 75.0% | 10 | 25.0% |
| Poorly differentiated | 2 | 10.5% | 17 | 89.5% |

Table 17: Categorical subjects distribution by malignancy grading and Syndecan-1 expression

There was a statistically significant distinction between high and low expression with respect to malignancy grading was statistically significant ($p < 0.001$)

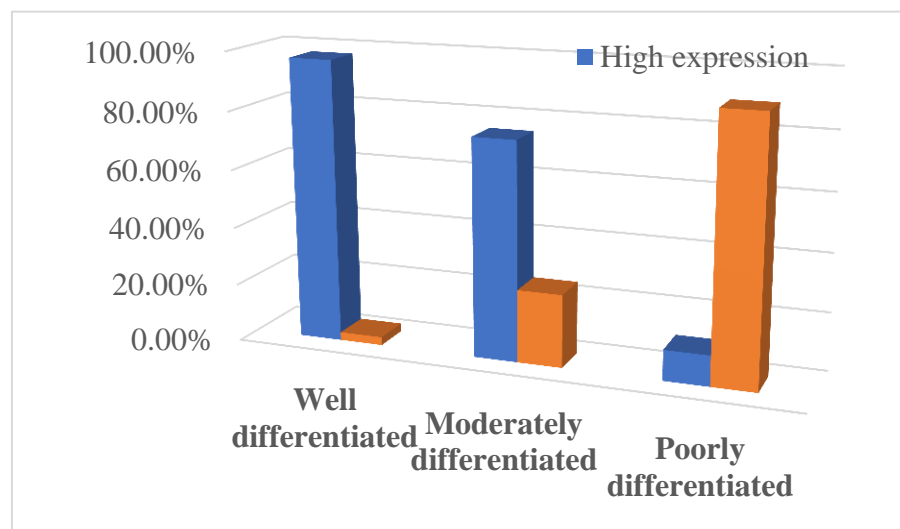


Chart 14: Categorical subjects distribution by malignancy grading and Syndecan-1 expression

Distribution of subjects according to staging and expression:

| | High expression | | Low expression | |
|-----|-----------------|-------|----------------|-------|
| | N | % | N | % |
| I | 20 | 80.0% | 5 | 20.0% |
| II | 18 | 62.1% | 11 | 37.9% |
| III | 29 | 70.7% | 12 | 29.3% |

Table 18: Distribution of subjects according to staging and expression

p-value 0.354, no statistically significant difference between high and low expression for staging was found.

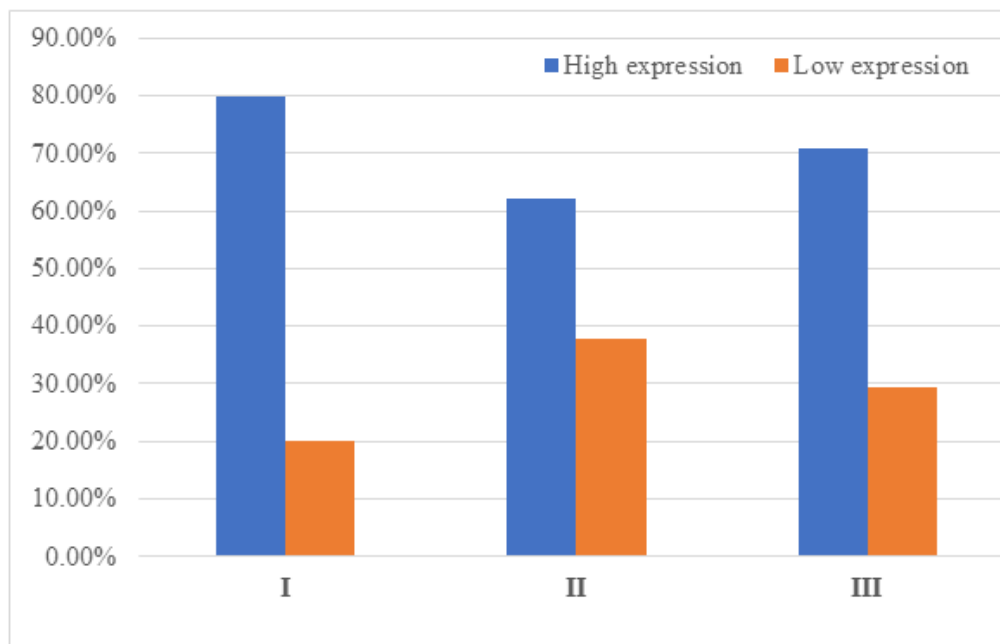


Chart 15: Distribution of subjects according to staging and expression

Categorical subjects distribution by tumor size and Syndecan-1 expression:

| | High expression | | Low expression | |
|----|-----------------|-------|----------------|-------|
| | N | % | N | % |
| T1 | 3 | 75.0% | 1 | 25.0% |
| T2 | 20 | 76.9% | 6 | 23.1% |
| T3 | 39 | 73.6% | 14 | 26.4% |
| T4 | 5 | 41.7% | 7 | 58.3% |

Table 19: Categorical subjects distribution by tumor size and Syndecan-1 expression

p-value 0.133, There was no discernible correlation between expression levels and tumour growth.

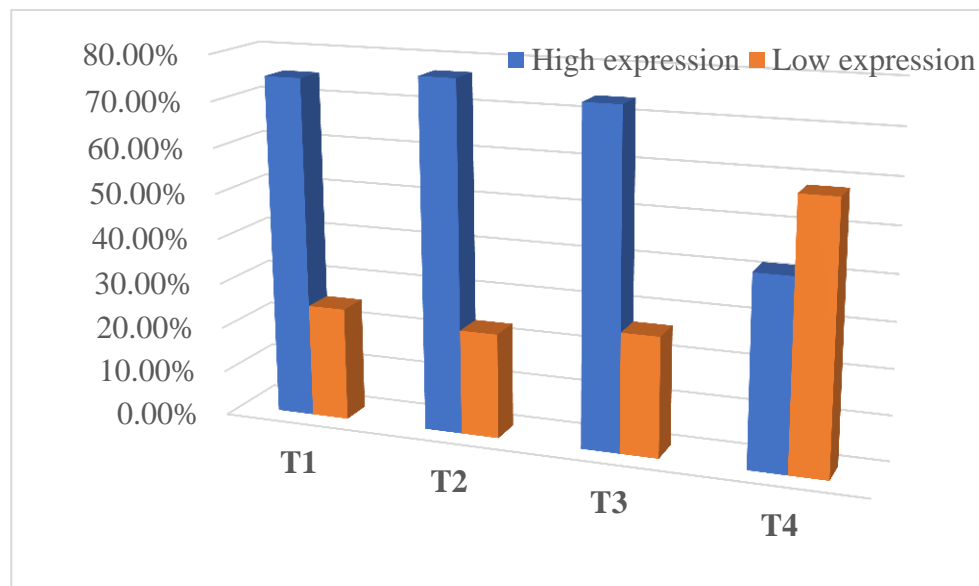


Chart 16: Categorical subjects distribution by tumor size and Syndecan-1 expression

Categorical subjects distribution by lymph node status and Syndecan-1 expression:

| | High expression | | Low expression | |
|----------|-----------------|-------|----------------|-------|
| | N | % | N | % |
| Negative | 39 | 70.9% | 16 | 29.1% |
| Positive | 28 | 70.0% | 12 | 30.0% |

Table 20: Categorical subjects distribution by lymph node status and Syndecan-1 expression

p-value 1.00, no statistically significant difference between high and low expression for the lymph node was found.

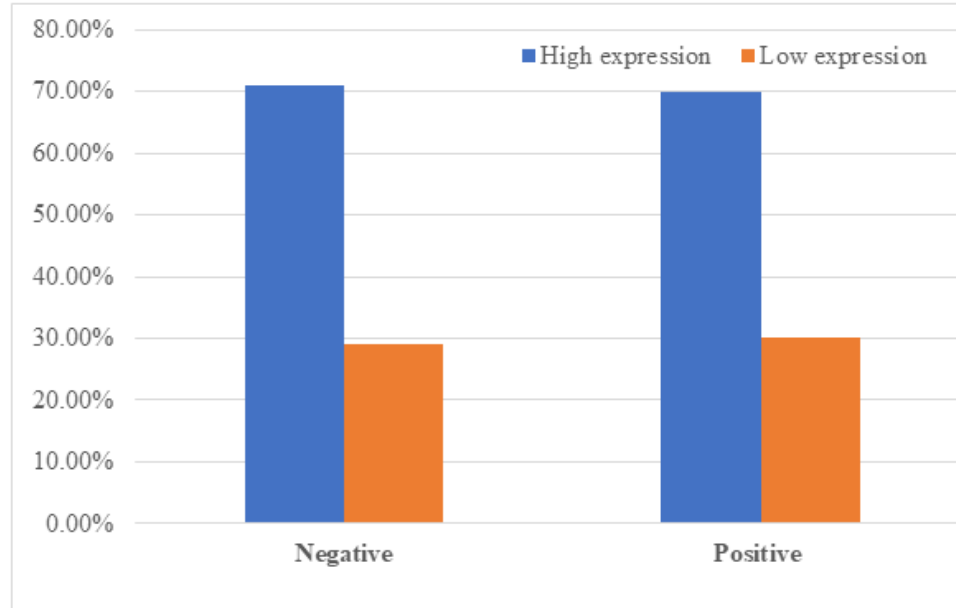


Chart 17: Categorical subjects distribution by lymph node status and Syndecan-1 expression

Categorical subjects distribution by Vascular Invasion and Syndecan-1 expression:

| | High expression | | Low expression | |
|---------|-----------------|-------|----------------|-------|
| | N | % | N | % |
| ABSENT | 60 | 69.0% | 27 | 31.0% |
| PRESENT | 7 | 87.5% | 1 | 12.5% |

Table 21: Categorical subjects distribution by Vascular Invasion and Syndecan-1 expression

p-value 0.429 showed no statistically significant difference between high and low expression for vascular invasion.

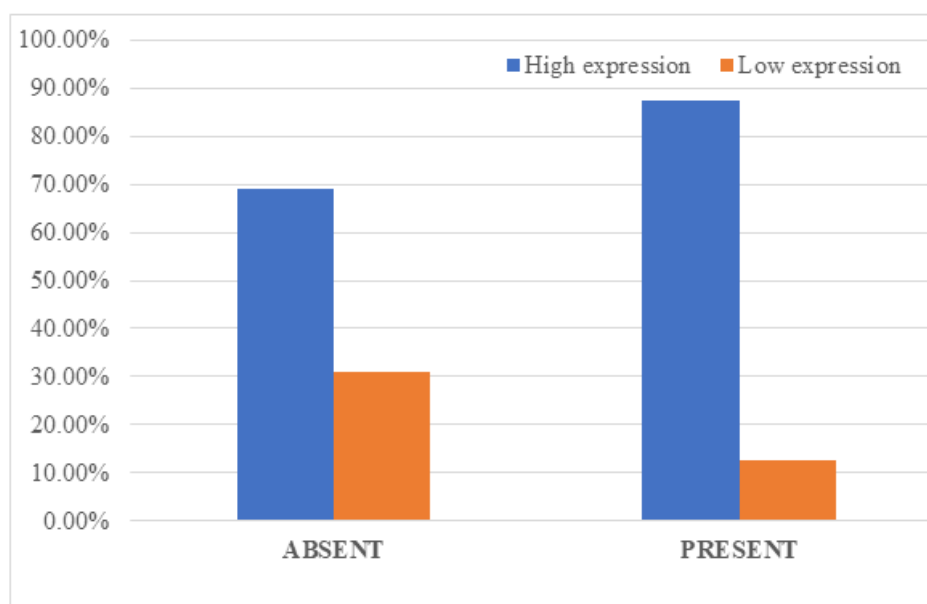


Chart 18: Categorical subjects distribution by Vascular Invasion and Syndecan-1 expression

Categorical subjects distribution by Perineural Invasion and Syndecan-1 expression:

| | High expression | | Low expression | |
|---------|-----------------|-------|----------------|-------|
| | N | % | N | % |
| ABSENT | 66 | 71.0% | 27 | 29.0% |
| PRESENT | 1 | 50.0% | 1 | 50.0% |

Table 22: Categorical subjects distribution by Perineural Invasion and Syndecan-1 expression

p-value 0.505, no statistically significant difference between high and low expression for Perineural invasion was found.

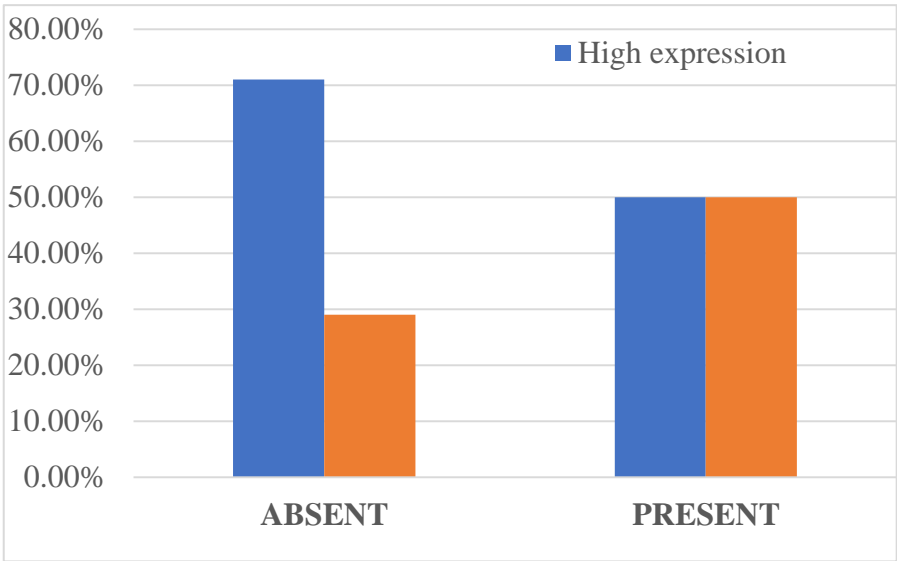


Chart 19: Categorical subjects distribution by Perineural Invasion and Syndecan-1 expression



Figure 5. Image showing proliferative growth in descending colon on cut section



Figure 6. Image of ulcero-proliferative growth with central necrosis and involving the serosa in sigmoid colon

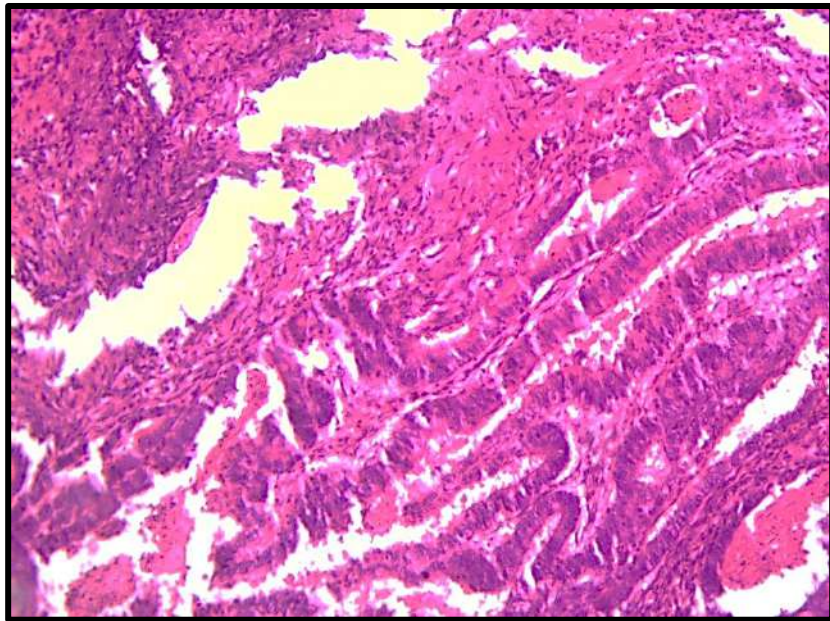


Figure 7: Microphotograph of H and E-stained section with 100x power showing Well Differentiated Adenocarcinoma

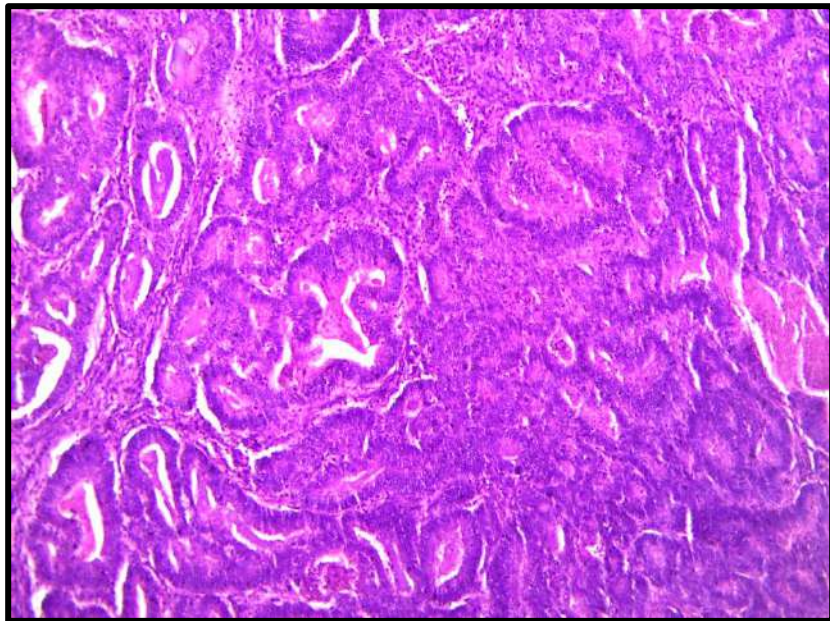


Figure 8: Microphotograph of H and E-stained section with 100x power showing Well Differentiated Adenocarcinoma

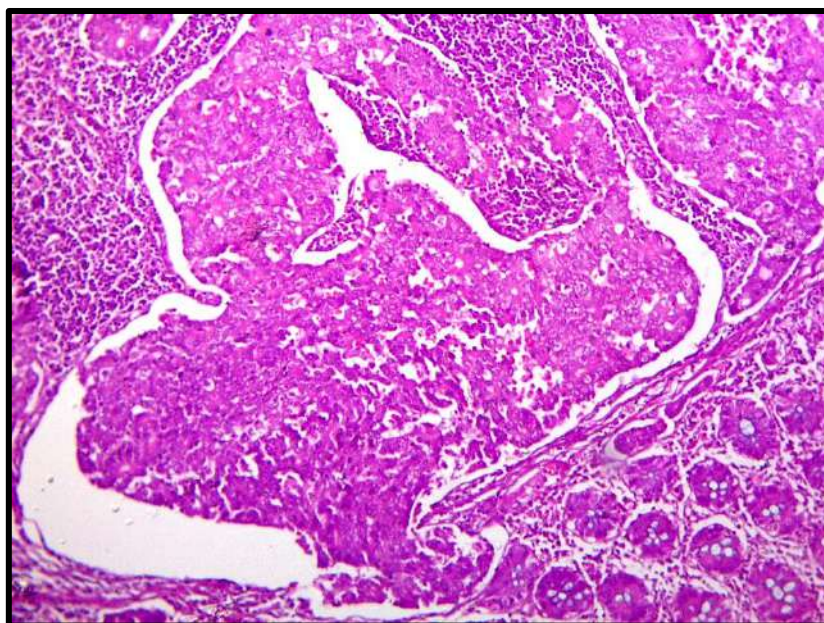


Figure 9: Microphotograph of H and E-stained section with 100x power showing Well Differentiated Adenocarcinoma

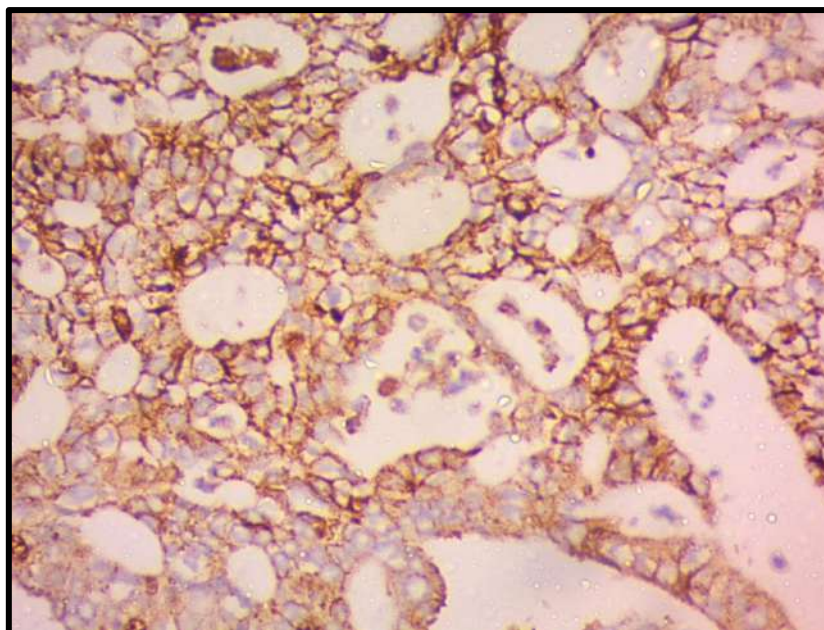


Figure 10. Microphotograph of Syndecan-1 IHC staining in 400X showing High expression in well differentiated colorectal carcinoma.

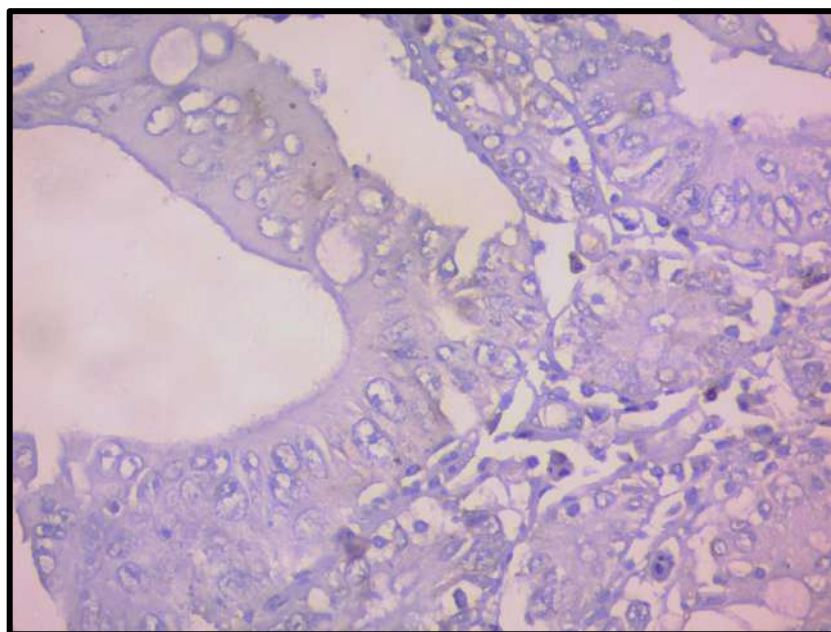


Figure 11. Microphotograph of Syndecan-1 IHC staining in 400X showing Low expression in poorly differentiated colorectal carcinoma.

DISCUSSION

DISCUSSION

CRCs are among the most frequently diagnosed cancers among men and women globally, according to the World Health Organization.⁴⁸

Australia and New Zealand have a high incidence of colorectal cancer, but Africa and south-central Asia have a lower incidence.⁴⁹

Age is one of the significant risk factors for the development of colorectal cancer. It is uncommon in people with age less than 40 years, and incidence increases with each successive decade.⁵⁰

Other factors that influence the outcome of colorectal carcinoma are sporadic CRCs or adenomatous polyps, inflammatory bowel disease, abdominopelvic radiation, cystic fibrosis, and hereditary CRC syndromes such as adenomatous polyposis syndromes and Lynch syndrome.²⁴

In addition to these factors CRC development is influenced by the use of red meat and the processed meat, use of tobacco in form of smoking or chewing, consumption of alcohol, and use of androgen deprivation therapy.⁴⁹

Eight prognostic factors are now judged to be clinically significant in colorectal carcinoma, and they are as follows, Serum CEA levels, tumor regression score, circumferential resection margin from the edge of tumor to nearest dissected margins, lymphovascular invasion, perineural invasion, microsatellite instability, KRAS, and NRAS mutation and BRAF mutation.⁴

With cellular interactions and also as adhesion molecule between cell and the matrix. In malignant transformation epithelial expression of Syndecan-1 is lost which is more evident in poorly differentiated CRCs. Syndecan-1 is also a target of therapy for

Indatuximab. So, in all cases of colorectal cancer, epithelial Syndecan-1 expression.

Comparison of Age Distribution with other studies:

| Study | Age |
|--------------------------------------|-----------------|
| Al-Maghrabi J. et al. (2021) (n=202) | <60 years 108 |
| | >60 years 94 |
| Li K et al. (2017) (n=477) | <60.5 years 211 |
| | >60.5 years 266 |
| Mitselou et al (2022) (n=69) | 40-81 years |
| The present study (n=95) | <60 years 54 |
| | >60 years 41 |

Table 23: Comparison of Age Distribution with other studies

In this study, out of 95 cases, 54 (56.84%) were in the age group ≤ 60 years, and 41 (43.15%) were in the age group >60 years.

Peak incidence was in the 60-69 age group (34.7%).

In other studies, such as Al-Maghrabi et al.⁴⁷ (n=202) ≤ 60 years were 108 and > 60 years were 94 subjects, Li K et al.⁵¹ (n=477) ≤ 60.5 years were 211 and > 60.5 years were 266 subjects, Mitselou et al.⁸ (n=69) age range was around 40-81 years.

Comparison of Sex Distribution with other studies:

| Study | Male % | Female % |
|--------------------------------------|--------|----------|
| Al-Maghrabi J. et al. (2021) (n=202) | 55 | 45 |
| Li K et al. (2017) (n=477) | 48 | 52 |
| Mitselou et al (2016) (n=69) | 61 | 39 |
| The present study (n=95) | 56 | 44 |

Table 24: Comparison of Sex Distribution with other studies

Though there is no sex predilection for colorectal carcinoma, most studies show a higher prevalence of colorectal carcinoma in males. In the present study, males 55% and females 45% were seen. Compared with other studies like Al-Maghrabi et al.⁴⁷ Mitselou et al.⁸ In a study by Li K et al.⁵¹ Females were 52%, and males were 48%.

Comparison of location of tumor distribution with other studies:

| Study | Site | |
|--------------------------------------|------------------|--------|
| Al-Maghrabi J. et al. (2021) (n=202) | Right colon | 25.7 % |
| | Left colon | 62.9 % |
| | Rectum | 11.4 % |
| Li K et al. (2019) (n=477) | Right colon | 74 % |
| | Left colon | 36 % |
| Theodoro T R et al (2022) (n=24) | Colon | 37.5 % |
| | Rectum | 62.5 % |
| The present study (n=95) | Ascending colon | 22.1 % |
| | Descending colon | 11.6 % |
| | Rectum | 43.2 % |
| | Sigmoid colon | 15.8 % |
| | Transverse colon | 7.4 % |

Table 24: Comparison of location of tumor distribution with other studies

In the present study, most of the cases were in the rectum (43.2%), followed by the sigmoid colon (15.8%), ascending colon (22.1%), descending colon (11.6%), and transverse colon (7.4%). Our study was similar to the survey done by Theodoro T R et al.⁵² where most of the cases were in the rectum (62.5%) followed by the colon (37.5%), and also with the study done by Li K et al.⁵¹ where colorectal carcinoma was primarily seen in right colon 74% than in left colon 36%.

Our study was contrary to the survey done by Al-Maghrabi et al.⁴⁷ where most of the cases seen in the left colon 62.9% followed by the right colon 25.7% and the rectum being 11.4%.

Comparison of Tumor Size with other studies:

| | | Al-Maghrabi J. et al. (2021) (n=202) | Li K et al. (2017) (n=477) | Theodoro T R et al (2022) (n=24) | Present study (n=95) |
|------------|--------|---|----------------------------------|--|----------------------------|
| Tumor size | <50 mm | 45% | 64% | 58.3% | 47.3% |
| | >50 mm | 55% | 36% | 41.7% | 54.7% |

Table 25: Comparison of Tumor Size with other studies

The present study had a tumor size ≥ 50 mm in 54.7% of cases. The present study was similar to the survey done by AL-Maghrabi et al. (55%).⁴⁷ But our study was contrary to other studies by Li K et al.⁵¹ (36%) and Theodoro T R et al.⁵² where they showed tumor size <50 mm being more cases.

Comparison of Histological Grading with other studies:

| | | Al-Maghrabi J. et al. (2021) (n=202) | Li K et al. (2017) (n=477) | Mitselou et al (2016) (n=69) | Theodoro T R et al (2022) (n=24) | Present study (n=95) |
|-----------------------|------------------------------|---|----------------------------------|---------------------------------|---|----------------------------|
| Malignancy Grading | Well Differentiated | 21.3% | 47.3 % | 4.35% | 8.3% | 37.9% |
| | Moderately Differentiated | 65.8% | 52.7% | 85.51% | 91.7% | 42.1% |
| | Poorly Differentiated | 12.9% | | 10.14% | | 20% |
| | | | | | | |

Table 26: Comparison of Histological Grading with other studies

In this study, most of the tumors 42.1% were graded as moderately differentiated, followed by 37.9 % of the well-differentiated tumor and 20% of the poorly differentiated tumor. This is similar to studies were done by Al-Maghrabi et al.⁴⁷ Li K et al.⁵¹ Mitselou et al., and Theodoro T R et al.⁵² where predominant tumors were graded as moderately differentiated.

Comparison of Pathological T Staging Distribution with other studies:

| | | Al-Maghrabi J. et al. (2021) (n=202) | Li K et al. (2017) (n=477) | Theodoro T R et al (2022) (n=24) | Present study (n=95) |
|-----------|----|--------------------------------------|----------------------------|----------------------------------|----------------------|
| T Staging | T1 | 12.9 % | 14.46 % (T1 and T2) | | 4.21 % |
| | T2 | 1.5 % | | 25 % | 27.36 % |
| | T3 | 15.8 % | 85.53 % (T3 and T4) | 75% | 55.78 % |
| | T4 | 73.8 % | | | 12.63 % |

Table 27: Comparison of Pathological T Staging Distribution with other studies

The current study had more cases in T2 and T3 stages (68.41%), which was similar to studies done by Al-Maghrabi et al.⁴⁷ (89.6%) Li K et al.⁵¹ (86.53%) and Theodoro T R et al.⁵²(75%)

Comparison of Pathological N Staging Distribution with other studies:

| | | Al-Maghrabi J. et al. (2021) (n=202) | Li K et al. (2017) (n=477) | heodoro T R et al (2022) (n=24) | Present study (n=95) |
|-----------|----------------|--------------------------------------|----------------------------|---------------------------------|----------------------|
| N Staging | N 0 | 54.9 % | 60.79% | 65.22 % | 57.9 % |
| | N 1 and N 2 | 45.1 % | 39.2 % | 34.78 % | 42.1 % |

Table 28: Comparison of Pathological N Staging Distribution with other studies

In the present study, 57.9% of cases were in N0 Stage, followed by 42.1% in N1 and N2. This is similar to other studies such as Al-Maghrabi et al.⁴⁷, Li K et al.⁵¹ and Theodoro T R et al.⁵² where N0 stage was seen in 54.9 %, 60.79%, and 65.22%, respectively.

Comparison of Vascular Invasion with other studies:

| | | Al-Maghrabi J. Et al. (2021) (n=202) | Mitselou et al (2016) (n=69) | Teodoro T R et al (2022) (n=24) | Present study (n=95) |
|----------------------|---------|---|------------------------------------|---------------------------------------|----------------------------|
| Vascular Invasion | Absent | 84.7 % | 27.54 % | 33.3 % | 91.6 % |
| | Present | 15.3 % | 72.46 % | 66.6 % | 8.4 % |

Table 29: Comparison of Vascular Invasion with other studies

Vascular invasion is an important prognostic factor in colorectal carcinoma. In the present study, only 8.4% of cases showed vascular invasion. The present study is similar to a study done by Al-Maghrabi et al.⁴⁷ which showed 15.3% of patients with vascular invasion. In contrast to other studies by Mitselou et al. 8 and Theodoro et al. 53, more vascular invasions were seen in 72.46% and 66.6% of cases.

Comparison of perineural Invasion with other studies:

| | | Theodoro T R et al (2022) (n=24) | The present study(n=95) |
|---------------------|---------|--|----------------------------|
| Perineural Invasion | Absent | 79.2 % | 97.9% |
| | Present | 20.8 % | 2.1% |

Table 30: Comparison of perineural Invasion with other studies

Perineural is a critical prognostic factor in colorectal carcinoma. In the present study, only 2.1% of cases had a perineural invasion, similar to a study done by Theodoro et al.⁵² which had 20.8% of patients with perineural invasion.

Comparison of expression of Syndecan-1 Scoring and Malignancy grading with other studies:

| | | High expression | Low expression |
|--|------------------------------|--------------------|----------------|
| | | % | % |
| The present study (n=95) | Well-differentiated | 97.2% | 2.8% |
| | Moderately differentiated | 75.0% | 25.0% |
| | Poorly differentiated | 10.5% | 89.5% |
| Al-Maghrabi J. Et al. (2021) (n=202) | Well-differentiated | 70.7% | 29.3% |
| | Moderately differentiated | 63.9% | 36.1% |
| | Poorly differentiated | 57.1% | 42.9% |

Table 31: Comparison of expression of Syndecan-1 Scoring and Malignancy grading with other studies

In the current investigation, well-differentiated adenocarcinomas displayed high expression of Syndecan-1 immunohistochemistry, whereas poorly differentiated adenocarcinomas displayed low expression of Syndecan. The current study is in contrast to a study done by Al-Maghrabi et al.⁴⁷ where high expression was in both well-differentiated adenocarcinomas and poorly differentiated adenocarcinomas.

| Parameter | | High expression | | Low expression | | Test | p value |
|---------------------|---------------------------|-----------------|---------|----------------|--------|-----------------|---------|
| | | N | % | N | % | | |
| Age | 20-29yrs | 1 | 33.30% | 2 | 66.70% | Chi-square test | 0.75 |
| | 30-39yrs | 4 | 80.00% | 1 | 20.00% | | |
| | 40-49yrs | 10 | 66.70% | 5 | 33.30% | | |
| | 50-59yrs | 15 | 75.00% | 5 | 25.00% | | |
| | 60-69yrs | 24 | 72.70% | 9 | 27.30% | | |
| | 70-79yrs | 11 | 73.30% | 4 | 26.70% | | |
| | 80-89yrs | 2 | 50.00% | 2 | 50.00% | | |
| Sex | Female | 34 | 81.00% | 8 | 19.00% | Chi-square test | 0.069 |
| | Male | 33 | 62.30% | 20 | 37.70% | | |
| Site | Ascending colon | 15 | 71.40% | 6 | 28.60% | Chi-square test | 0.478 |
| | Descending colon | 8 | 72.70% | 3 | 27.30% | | |
| | Rectum | 27 | 65.90% | 14 | 34.10% | | |
| | Sigmoid colon | 10 | 66.70% | 5 | 33.30% | | |
| | Transverse colon | 7 | 100.00% | 0 | 0.00% | | |
| Malignancy grade | Well differentiated | 35 | 97.20% | 1 | 2.80% | Chi-square test | <0.001 |
| | Moderately differentiated | 30 | 75.00% | 10 | 25.00% | | |
| | Poorly differentiated | 2 | 10.50% | 17 | 89.50% | | |
| Tumor size | T1 | 3 | 75.00% | 1 | 25.00% | Chi-square test | 0.133 |
| | T2 | 20 | 76.90% | 6 | 23.10% | | |
| | T3 | 39 | 73.60% | 14 | 26.40% | | |
| | T4 | 5 | 41.70% | 7 | 58.30% | | |
| Stage of the tumor | I | 20 | 80.00% | 5 | 20.00% | Chi-square test | 0.354 |
| | II | 18 | 62.10% | 11 | 37.90% | | |
| | III | 29 | 70.70% | 12 | 29.30% | | |
| Lymph node status | Negative | 39 | 70.90% | 16 | 29.10% | Chi-square test | 1 |
| | Positive | 28 | 70.00% | 12 | 30.00% | | |
| Vascular invasion | Absent | 60 | 69.00% | 27 | 31.00% | Chi-square test | 0.429 |
| | Present | 7 | 87.50% | 1 | 12.50% | | |
| Perineural invasion | Absent | 66 | 71.00% | 27 | 29.00% | Chi-square test | 0.505 |
| | Present | 1 | 50.00% | 1 | 50.00% | | |

Table 32: Syndecan-1 expression and its relation to clinicopathological parameters.

The relationship between Syndecan-1 expression and clinicopathological parameters:

In the current investigation, there was a statistically significant link between high Syndecan-1 expression in well-differentiated adenocarcinomas and low Syndecan-1 expression in poorly differentiated adenocarcinomas (p 0.001). In contrast to the current study, Al-Maghrabi et al.⁴⁷ found no statistically significant relationship between Syndecan-1 expression and malignancy grade (p=0.503).

In contrast to Al-Maghrabi et al.⁴⁷ and Li K et al.⁵¹ whose studies reported p 0.001 and p=1, respectively, respectively, our research revealed no statistically significant link between lymph node positivity and Syndecan-1 expression (p=1) and this could be because of low lymph node positivity in our study in comparison to other studies.

In contrast to the current study, where there was no statistical significance with the expression of Syndecan-1 and T stage of the tumour, Li K et al. had substantial statistical significance with the T stage of the tumour which had 85.53% of cases in T3 & T4 stage. Research has shown that SDC1 is coexpressed with EMT markers (E-cadherin and β -catenin) in CRCs and that this coexpression is regulated during epithelial-mesenchymal transition (EMT). The loss of SDC1 expression in carcinoma cells reduces cell adhesion to the extracellular matrix and enhances cell motility and invasion.⁵¹

Similar to studies by Al-Maghrabi et al. and Li K et al.^{47,51} there was no statistically significant relationship between the patient's age, sex, tumour location, tumour stage, vascular invasion, and perineural invasion in our investigation.

CONCLUSION

CONCLUSION

High Syndecan-1 expression is shown in well-differentiated adenocarcinomas, while low expression is seen in poorly differentiated adenocarcinomas of the colon and rectum and indirectly shows the outcomes of clinical treatments. IHC analysis of Syndecan 1 can be used to regularly assess prognostic relevance, which would aid in clinical outcome, as loss of Syndecan-1 expression is related with loss of differentiation.

SUMMARY

SUMMARY

- The present study was undertaken in the Department of Pathology, Sri Devaraj Urs Medical College, Tamaka, Kolar over a period of two years from October 2020 to November 2022.
- A total of 95 cases of Colorectal carcinoma who underwent surgical resection were studied. H & E Slides of all cases were reviewed and performed immunohistochemistry against Syndecan-1. Expression of Syndecan-1 was evaluated and correlated with clinicopathological data of cases such as age, sex, histological grading, lymph node status and staging.
- Peak incidence was in the 60-69 years age group (34.7%). Most common site of the tumor was rectum (43.2%). Majority of cases were in Stage III(43.15%).
- Expression of Syndecan-1 by immunohistochemistry was scored as low-expression group (scores 0-2) and a high-expression group (scores 3-6) from the total score based on membranous/cytoplasmic staining & percentage of positively stained cells.
- 70.5 % of cases demonstrated High expression & 29.5% of cases had Low expression of Syndecan-1
- High expression of Syndecan-1 in Well differentiated carcinomas and Low expression of Syndecan-1 in poorly differentiated carcinomas was significantly correlated with malignancy grade of the colorectal carcinomas.

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BIBLIOGRAPHY

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ANNEXURES

ANNEXURE - I INFORMED CONSENT FORM

INFORMED CONSENT FORM

STUDY TITLE: The immunohistochemical expression of syndecan 1(CD 138) and its correlation with staging and grading of colorectal carcinoma.

I, _____ have read or have been read to me the patient information sheet and understand the purpose of the study, the procedure that will be used, the risk and benefits associated with my involvement in the study and the nature of information will be collected and disclosed during the study.

I have had my opportunity to ask my questions regarding various aspects of the study and my questions are answered to my satisfaction.

I, the undersigned, agree to participate in this study and authorize the collection and disclosure of my personal information for the dissertation.

Name and signature / thumb impression

Date:

Place:

Name and signature / thumb impression

Date:

Place:

(Witness/Parent/ Guardian/ Husband)

ANNEXURE –II

PATIENT INFORMATION SHEET

PATIENT INFORMATION SHEET:

STUDY TITLE: The immunohistochemical expression of syndecan 1(CD 138) and its correlation with staging and grading of colorectal carcinoma.

PLACE OF STUDY: Department of Pathology, Sri Devaraj Urs Medical College, Kolar.

The main aim of the study is to find the role of syndecan 1 in colorectal carcinoma. The specimens will be collected from the department of pathology, SDUMC, Kolar. This study will be approved by the institutional ethical committee. The information collected will be used only for dissertation and publication. There is no compulsion to agree to participate. You are requested to sign / provide thumb impression only if you voluntarily agree to participate in the study. All information collected from you will be kept confidential and will not be disclosed to any outsider. Your identity will not be revealed. You will not receive any monetary benefits to participate in this research. This informed consent document is intended to give you a general background of study. Please read the following information carefully and discuss with your family members. You can ask your queries related to study at any time during the study. If you are willing to participate in the study you will be asked to sign an informed consent form by which you are acknowledging that you wish to participate in the study and entire procedure will be explained to you by the study doctor. You are free to withdraw your consent to participate in the study any time without explanation and this will not change your future care.

For any clarification you are free to contact the investigator.

PRINCIPAL INVESTIGATOR: Dr Sudarshan K

ANNEXURE III

STUDY PROFORMA

PATIENT PROFORMA

Name:

Age:

Hospital Number:

Anonymised Sample No:

Chief complaint:

History of presenting illness:

Past history:

Personal history:

Local examination:

Histopathological diagnosis:

Gross:

Microscopy:

Stage of disease:

KEY TO MASTER CHART

S. No = SERIAL NUMBER

UNIQUE HOSPITAL IDENTIFICATION NUMBER YEAR=YEAR OF BIOPSY

BIOPSY No= BIOPSY NUMBER

AGE= AGE IN YEARS

SEX: M= MALE F= FEMALE

SPECIMEN TYPE: APR= ABDOMINOPERINEAL

RESECTION TNM=TUMOUR NODE METASTASIS

STAGING LN=LYMPHNODE

LN + = LYMPH NODE POSITIVE

LVI= LYMPHOVASCULAR INVASION

PNI= PERINEURAL INVASION

TOTAL IHC SCORE = TOTAL IHC SCORE

| S. NO | HOSPITAL NO | YEAR | BIOPSY NO | A G E | S E X | SPECIMEN TYPE | SITE | HISTOPATHOLOGY DIAGNOSIS | MALIGNANCY GRADING | GROWTH | TNM | STA GING | TUMOR SIZE | LN + | LVI | PNI | Total IHC score |
|-------|-------------|------|-----------|-------|-------|--------------------|------------------|--------------------------|---------------------------|------------------------|------------|----------|------------|------|--------|--------|-----------------|
| 1 | 438421 | 2008 | 1002 | 35 | M | APR | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 3 |
| 2 | 432848 | 2008 | 736 | 31 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50MM | 0 | ABSENT | ABSENT | 6 |
| 3 | 413277 | 2008 | 406 | 48 | M | APR | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T4N0MX | II | <50MM | 0 | ABSENT | ABSENT | 0 |
| 4 | 458197 | 2008 | 1510 | 50 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T2N1MX | III | >50MM | 2 | ABSENT | ABSENT | 3 |
| 5 | 354632 | 2008 | 1264 | 70 | M | HEMICOLECTOMY | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T4N0MX | II | >50MM | 0 | ABSENT | ABSENT | 0 |
| 6 | 453464 | 2008 | 1344 | 68 | M | ANTERIOR RESECTION | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50MM | 0 | ABSENT | ABSENT | 6 |
| 7 | 458410 | 2008 | 1373 | 75 | M | HEMICOLECTOMY | TRANSVERSE COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T3N1MX | II | <50MM | 0 | ABSENT | ABSENT | 6 |
| 8 | 453520 | 2008 | 1743 | 40 | F | ANTERIOR RESECTION | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T1N0MX | I | <50MM | 0 | ABSENT | ABSENT | 6 |
| 9 | 470873 | 2008 | 1789 | 20 | M | HEMICOLECTOMY | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T1N0MX | I | >50MM | 0 | ABSENT | ABSENT | 6 |
| 10 | 470610 | 2008 | 1792 | 45 | M | HEMICOLECTOMY | TRANSVERSE COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N1MX | I | <50MM | 0 | ABSENT | ABSENT | 5 |
| 11 | 486531 | 2009 | 1510 | 60 | F | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N1MX | I | <50MM | 0 | ABSENT | ABSENT | 6 |
| 12 | 501634 | 2010 | 1388 | 48 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T4N1MX | III | >50MM | 1 | ABSENT | ABSENT | 6 |
| 13 | 525369 | 2010 | 1937 | 52 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N1MX | I | <50MM | 0 | ABSENT | ABSENT | 6 |
| 14 | 579655 | 2010 | 1504 | 47 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | PROLIFERATIVE | T4N1MX | III | <50MM | 2 | ABSENT | ABSENT | 5 |
| 15 | 630751 | 2011 | 782 | 25 | F | APR | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3aIBrnmix | II | >50MM | 0 | ABSENT | ABSENT | 3 |
| 16 | 638844 | 2011 | 1736 | 65 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N1MX | III | >50MM | 3 | ABSENT | ABSENT | 0 |
| 17 | 733193 | 2011 | 1804 | 28 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T3N1MX | II | >50MM | 0 | ABSENT | ABSENT | 0 |
| 18 | 743282 | 2011 | 1038 | 54 | F | APR | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3aIBrnmix | II | >50MM | 0 | ABSENT | ABSENT | 5 |
| 19 | 789154 | 2011 | 1461 | 46 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N1MX | III | >50MM | 2 | ABSENT | ABSENT | 3 |
| 20 | 791325 | 2011 | 1664 | 60 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T2N1MX | III | >50MM | 2 | ABSENT | ABSENT | 6 |
| 21 | 733105 | 2011 | 1994 | 52 | F | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N1MX | III | >50MM | 2 | ABSENT | ABSENT | 6 |
| 22 | 762580 | 2012 | 8 | 65 | F | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T3N1MX | III | >50MM | 2 | ABSENT | ABSENT | 4 |
| 23 | 770859 | 2012 | 146 | 75 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N1MX | III | >50MM | 2 | ABSENT | ABSENT | 5 |
| 24 | 816143 | 2012 | 1358 | 67 | M | APR | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T1N1MX | I | <50MM | 0 | ABSENT | ABSENT | 4 |
| 25 | 835745 | 2012 | 1866 | 51 | F | ANTERIOR RESECTION | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T3N1MX | III | >50MM | 1 | ABSENT | ABSENT | 4 |
| 26 | 840354 | 2012 | 1814 | 36 | M | ANTERIOR RESECTION | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 6 |
| 27 | 837910 | 2012 | 1790 | 73 | F | APR | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N1MX | III | >50MM | 2 | ABSENT | ABSENT | 6 |
| 28 | 841155 | 2012 | 2213 | 85 | F | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T4N1MX | III | >50MM | 1 | ABSENT | ABSENT | 2 |
| 29 | 836409 | 2012 | 2480 | 45 | F | APR | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCERATIVEINFILTRATIVE | T3N1MX | III | <50MM | 2 | ABSENT | ABSENT | 4 |
| 30 | 843857 | 2012 | 286 | 63 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 6 |

| | | | | | | | | | | | | | | | | | |
|----|--------|------|------|----|---|--------------------|------------------|----------------|---------------------------|-------------------------|----------|-----|-------|---|---------|---------|---|
| 31 | 86745 | 2012 | 2387 | 49 | F | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1MX | III | >50MM | 1 | PRESENT | ABSENT | 5 |
| 32 | 863324 | 2012 | 2480 | 74 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 6 |
| 33 | 819843 | 2012 | 349 | 80 | F | ANTERIOR RESECTION | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1MX | III | >50MM | 1 | ABSENT | ABSENT | 4 |
| 34 | 883254 | 2012 | 1167 | 62 | M | APR | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | PROLIFERATIVE | T2N1MX | I | <50MM | 0 | ABSENT | ABSENT | 6 |
| 35 | 899146 | 2012 | 1501 | 64 | M | APR | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1MX | III | <50MM | 2 | ABSENT | ABSENT | 6 |
| 36 | 882182 | 2013 | 298 | 55 | F | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N1MX | III | >50MM | 2 | ABSENT | ABSENT | 4 |
| 37 | 878863 | 2013 | 331 | 60 | M | ANTERIOR RESECTION | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1MX | III | >50MM | 1 | ABSENT | ABSENT | 0 |
| 38 | 883102 | 2013 | 427 | 67 | M | APR | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | PROLIFERATIVE | T2N1MX | I | <50MM | 0 | ABSENT | ABSENT | 5 |
| 39 | 879624 | 2013 | 443 | 46 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 5 |
| 40 | 903057 | 2013 | 851 | 60 | M | APR | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N1MX | III | >50MM | 1 | ABSENT | ABSENT | 3 |
| 41 | 958439 | 2013 | 2098 | 65 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N2MX | III | >50MM | 4 | ABSENT | ABSENT | 0 |
| 42 | 940966 | 2013 | 1844 | 65 | F | ANTERIOR RESECTION | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | PROLIFERATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 6 |
| 43 | 928495 | 2013 | 1652 | 65 | F | ANTERIOR RESECTION | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50MM | 0 | ABSENT | ABSENT | 5 |
| 44 | 854002 | 2013 | 1714 | 70 | F | APR | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50MM | 0 | ABSENT | ABSENT | 5 |
| 45 | 981042 | 2014 | 223 | 60 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N1MX | III | >50MM | 1 | ABSENT | ABSENT | 2 |
| 46 | 940314 | 2014 | 588 | 73 | F | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | POORLY DIFFERENTIATED | PROLIFERATIVE | T4N1N1MX | IIB | <50MM | 3 | ABSENT | ABSENT | 2 |
| 47 | 931225 | 2014 | 893 | 64 | M | APR | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T1N1MX | I | <50MM | 0 | ABSENT | ABSENT | 2 |
| 48 | 967345 | 2014 | 1305 | 56 | F | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 4 |
| 49 | 32776 | 2014 | 1989 | 50 | F | APR | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 5 |
| 50 | 155065 | 2015 | 1533 | 60 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1MX | III | >50MM | 1 | PRESENT | ABSENT | 4 |
| 51 | 195235 | 2015 | 783 | 46 | M | HEMICOLECTOMY | DESCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | >50MM | 0 | ABSENT | ABSENT | 2 |
| 52 | 213889 | 2015 | 1439 | 54 | M | APR | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50MM | 0 | ABSENT | ABSENT | 0 |
| 53 | 208706 | 2015 | 3187 | 70 | M | HEMICOLECTOMY | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T4N1N1MX | IIB | >50MM | 0 | ABSENT | ABSENT | 0 |
| 54 | 254597 | 2016 | 1885 | 45 | F | HEMICOLECTOMY | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N0MX | IIA | <50MM | 0 | ABSENT | ABSENT | 0 |
| 55 | 289663 | 2016 | 2936 | 60 | F | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | POORLY DIFFERENTIATED | PROLIFERATIVE | T4N1N1MX | IIB | <50MM | 3 | ABSENT | ABSENT | 3 |
| 56 | 305665 | 2016 | 2067 | 58 | F | HEMICOLECTOMY | DESCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | >50MM | 0 | ABSENT | ABSENT | 3 |
| 57 | 218304 | 2016 | 48 | 68 | M | APR | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50MM | 0 | ABSENT | ABSENT | 5 |
| 58 | 304816 | 2016 | 2001 | 76 | F | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50MM | 0 | ABSENT | ABSENT | 6 |
| 59 | 428218 | 2017 | 1207 | 45 | M | APR | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N2MX | IIC | <50MM | 6 | PRESENT | PRESENT | 0 |
| 60 | 402489 | 2017 | 474 | 45 | F | APR | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1MX | III | <50MM | 3 | PRESENT | PRESENT | 6 |
| 61 | 502643 | 2017 | 2504 | 84 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N0MX | II | <50MM | 0 | ABSENT | ABSENT | 2 |
| 62 | 541581 | 2018 | 382 | 63 | F | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | IIA | <50MM | 0 | ABSENT | ABSENT | 6 |
| 63 | 553372 | 2018 | 1515 | 30 | M | APR | RECTUM | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1MX | III | <50MM | 2 | ABSENT | ABSENT | 4 |

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|----|--------|------|------|----|---|--------------------|-----------------------|-------------------------|---------------------------|-------------------------|-----------|------|-------|----|---------|--------|---|
| 64 | 548316 | 2018 | 613 | 70 | M | ANTERIOR RESECTION | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | III | <50mm | 0 | ABSENT | ABSENT | 5 |
| 65 | 550703 | 2018 | 782 | 56 | M | ANTERIOR RESECTION | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50mm | 0 | ABSENT | ABSENT | 5 |
| 66 | 616361 | 2018 | 2030 | 57 | F | HEMICOLECTOMY | DESCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N0M0 | II | <50mm | 0 | ABSENT | ABSENT | 6 |
| 67 | 655446 | 2019 | 253 | 65 | M | HEMICOLECTOMY | DESCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1bMX | IB | <50mm | 3 | ABSENT | ABSENT | 5 |
| 68 | 728086 | 2019 | 1592 | 55 | F | HEMICOLECTOMY | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50mm | 0 | ABSENT | ABSENT | 6 |
| 69 | 700297 | 2019 | 728 | 72 | M | HEMICOLECTOMY | TRANSVERSE COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1MX | III | >50mm | 2 | ABSENT | ABSENT | 4 |
| 70 | 683154 | 2019 | 1467 | 65 | F | APR | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50mm | 0 | ABSENT | ABSENT | 5 |
| 71 | 841770 | 2020 | 707 | 60 | F | HEMICOLECTOMY | DESCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | <50mm | 0 | ABSENT | ABSENT | 5 |
| 72 | 844349 | 2020 | 838 | 75 | F | HEMICOLECTOMY | RECTOSIGMOID | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N0MX | I | <50mm | 0 | ABSENT | ABSENT | 6 |
| 73 | 84806 | 2020 | 991 | 40 | F | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T2N0MX | I | >50mm | 0 | ABSENT | ABSENT | 5 |
| 74 | 846813 | 2020 | 1285 | 45 | F | HEMICOLECTOMY | RECTUM | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T2N0MX | I | <50mm | 0 | ABSENT | ABSENT | 6 |
| 75 | 867528 | 2020 | 1466 | 62 | F | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N1aMX | III | >50mm | 1 | ABSENT | ABSENT | 6 |
| 76 | 866322 | 2020 | 1530 | 65 | F | HEMICOLECTOMY | TRANSVERSE COLON | ADENOCARCINOMA | WELL DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | IIA | >50mm | 0 | ABSENT | ABSENT | 6 |
| 77 | 875828 | 2020 | 1732 | 65 | M | HEMICOLECTOMY | ASCENDING COLON | ADENOCARCINOMA | POORLY DIFFERENTIATED | ULCEROPROLIFERATIVE | T4aNO0MX | IB | <50mm | 0 | ABSENT | ABSENT | 2 |
| 78 | 848200 | 2020 | 1209 | 35 | M | HEMICOLECTOMY | RECTUM | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCEROPROLIFERATIVE | T3N0MX | II | <50mm | 0 | ABSENT | ABSENT | 6 |
| 79 | 891338 | 2021 | 172 | 68 | M | HEMICOLECTOMY | SIGMOID COLON | ADENOCARCINOMA | MODERATELY DIFFERENTIATED | ULCERATIVE/INFILTRATIVE | T3N0MX | II | <50mm | 0 | ABSENT | ABSENT | 6 |
| 80 | 903275 | 2021 | 578 | 65 | M | Hemicolecotomy | Rectosigmoid junction | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T3N2bMx | IIIC | >50mm | 2 | Absent | Absent | 3 |
| 81 | 943507 | 2021 | 1550 | 68 | M | Hemicolecotomy | Descending colon | Adenocarcinoma | Well differentiated | Ulceroproliferative | T3N1bMx | IIIB | >50mm | 2 | Absent | Absent | 6 |
| 82 | 35236 | 2021 | 1830 | 60 | F | Hemicolecotomy | Ascending colon | Adenocarcinoma | Well differentiated | Ulceroproliferative | T3N0Mx | IIIA | >50mm | 0 | Absent | Absent | 5 |
| 83 | 40225 | 2021 | 1951 | 55 | F | APR | Rectum | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T3N0Mx | IIA | <50mm | 0 | Absent | Absent | 3 |
| 84 | 48038 | 2021 | 2174 | 70 | F | Hemicolecotomy | Descending colon | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T2N0Mx | I | <50mm | 0 | Absent | Absent | 2 |
| 85 | 46320 | 2021 | 2188 | 60 | F | Hemicolecotomy | Descending colon | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T2N1bMx | IIIA | >50mm | 1 | Absent | Absent | 6 |
| 86 | 64673 | 2022 | 463 | 50 | F | Hemicolecomy | Descending colon | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T4a1x2bMx | IIIC | >50mm | 11 | Present | Absent | 4 |
| 87 | 50567 | 2022 | 574 | 56 | M | Hemicolecomy | Descending colon | Adenocarcinoma | Well differentiated | Infiltrative | T4a1Vc1Mx | IIIB | <50mm | 3 | Present | Absent | 6 |
| 88 | 71096 | 2022 | 684 | 53 | M | Hemicolecomy | Rectum | Adenocarcinoma | Moderately differentiated | Proliferative | T3N0Mx | II | >50mm | 0 | Absent | Absent | 3 |
| 89 | 77098 | 2022 | 837 | 60 | F | APR | Rectum | Adenocarcinoma | Moderately differentiated | Infiltrative | T4bVc1Mx | IIIC | >50mm | 1 | Absent | Absent | 4 |
| 90 | 76836 | 2022 | 863 | 54 | M | Hemicolecomy | Rectosigmoid junction | Adenocarcinoma | Well differentiated | Infiltrative | T3N0Mx | IIA | >50mm | 0 | Absent | Absent | 5 |
| 91 | 63632 | 2022 | 884 | 76 | F | Hemicolecomy | Transverse colon | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T3N2aMx | IIIB | >50mm | 1 | Present | Absent | 4 |
| 92 | 76366 | 2022 | 972 | 50 | M | Hemicolecomy | Transverse colon | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T3N1Mx | IIIB | >50mm | 1 | Present | Absent | 5 |
| 93 | 84786 | 2022 | 1160 | 72 | F | Total colectomy | Caecum | Mucinous Adenocarcinoma | Well differentiated | Infiltrative | T2N2aMx | IIIB | >50mm | 5 | Absent | Absent | 4 |
| 94 | 112259 | 2022 | 1782 | 52 | M | Hemicolecomy | Transverse colon | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T2N0Mx | I | >50mm | 0 | Absent | Absent | 5 |
| 95 | 122895 | 2022 | 1815 | 85 | M | Hemicolecomy | Descending colon | Adenocarcinoma | Moderately differentiated | Ulceroproliferative | T3N1aMx | III | <50mm | 1 | Absent | Absent | 4 |