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### Accuracy of Cell-Free DNA (cfDNA) in Colorectal Cancer Diagnosis

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ABSTRACT

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#### 1. Introduction

Colorectal cancer is the third most common type of cancer globally after lung cancer and breast cancer, and is the second leading cause of cancer death in the world.<sup>1</sup> In Indonesia, GLOBOCAN 2020 data states that the incidence of new cases of colorectal cancer is in third place, namely 8.6%. In men, colorectal cancer is the second most common cancer (11.9%) and in women it is the fourth most common cancer (5.8%).<sup>1</sup> Technological and diagnostic developments in the last two decades have led to an increase in average life expectancy in colorectal cancer patients, although

## mortality rates are still high and are generally associated with late diagnosis.<sup>2</sup> There is limited ability to screen, so that patients come in an advanced stage or general condition that has greatly decreased due to delays in diagnosis, causing treatment to be more complex and risky. As a result, morbidity and mortality will be greater. Therefore, early diagnosis is very important for the survival of patients, because patients detected in stages I-II have a survival rate of up to 90%.<sup>3</sup> This number dropped drastically in stage IV, which was only about 14%. Detection of cancer-

Background: The requirement for malaria elimination to be recognized is to prove

Background. Colorectal cancer is the third most common type of cancer following

lung and breast cancer with the second most common cause of cancer-related death

globally. Delayed diagnosis due to limited source and modality to perform early diagnosis lead to advanced-stage condition which contributes to higher morbidities

and mortalities. Recent diagnosis of colorectal cancer depends on biopsy of suspected tissues, either obtained surgically or per colonoscopy. Colorectal cancer detection through cell-free DNA measurement allowing small-size cancer being detected even

in early stage. cfDNA originated from derivates of increased and abnormality

apoptosis-necrosis pathway from cancer lesion, therefore can be managed as specific tumor marker. **Methods:** Diagnostic test was performed at the Gastroentero Hepatology Outpatient Unit and Internal Medicine Inpatient Unit of Dr. Mohammad Hoesin General Hospital Palembang from March to June 2021. Data processing and

analysis using SPSS version 26.0 for Windows. **Results:** Among 50 subjects included, 39 subjects (78%) are diagnosed with colorectal cancer, whereas 11 subjects (22%) as control. The median of cfDNA result is 59,71 ng/mL with 92,3%

sensitivity, 90,9% specificity, 95,2% positive predictive value, 82,3% negative predictive value, and 92,4% accuracy rate. Combination the measurement of cfDNA,

Conclusion: The study reveals that cell-free DNA (cfDNA) demonstrated a very

CEA, and CA19-9 appears to have better AUC instead of single measurement.

promising accuracy rate in diagnosing colorectal cancer.

associated DNA abnormalities through peripheral blood allows small colorectal cancers to be detected at an early stage compared with currently available tumor markers, fecal occult blood tests, and currently available imaging modalities.<sup>4</sup>

In patients with cases of malignancy, identification of DNA in serum or plasma is referred to as circulating free DNA (cfDNA). cfDNA is derived from increased and abnormal apoptotic-necrosis pathways in cancerous lesions. This abnormal DNA degradation and secretion causes an increase in the levels of fragmented DNA of various sizes, and can be identified by biomolecular examination techniques in the blood. Tumor cells that release cfDNA are able to provide an overview of the genetic and epigenetic changes of the tumor origin that make cfDNA has an important role in the theory of genometastases. Several cfDNA diagnostic tests in colorectal cancer have shown conflicting results. Current studies detect cfDNA with a variety of different primary lesions. The study by Wang et al compared several primers in diagnosing colorectal cancer and found that the Alu and GADPHgenes (glyceraldehyde-3-phosphate dehydrogenase) are sensitive markers in diagnosing colorectal cancer. Alu is a short strand of DNA derived from the action of restriction endonucleases by Arthrobacter luteus (Alu). The Alu sequence is element transposable the most abundant, containing more than one million copies scattered throughout the human genome. Umetani et al found two primers new, ALU247 and ALU115, which were used in quantitative PCR detection of long and short chain DNA fragments. This study found a correlation between tumor progression and the ratio of long-shortchain DNA fragments in the blood of colorectal cancer patients using quantitative PCR of Alu sequences, the most frequently repeated sequence in the human genome. Detection of cfDNA with Alu marker can be a potential biomarker in screening and early detection of colorectal cancer. Meanwhile, other studies such as El-Gayar's study showed that cfDNA levels did not provide a good enough accuracy test result in diagnosing colorectal cancer. This difference may occur due to differences in methods and racial differences in the patient sample studied.3

#### 2. Methods

This type of research is a diagnostic test. The research was conducted at the Gastroentero Hepatology Polyclinic and Inpatient Department of Internal Medicine, Dr. Mohammad Hoesin Palembang which was carried out from March 2021 - June 2021. The research sample was taken by non- probability consecutive sampling. All research samples that meet the inclusion criteria from December 2020 - March 2021, as many as 50 samples, were included in the study sequentially. The samples were then divided into two groups according to the histopathological findings found, namely the colorectal cancer group or the group without colorectal cancer. The inclusion criteria: All colorectal cancer suspects who came to the Gastroentero Hepatology Polyclinic and Inpatient Department of Internal Medicine, Dr. Mohammad Hoesin Palembang, all colorectal cancer patients who have never undergone chemotherapy and/or radiotherapy, are willing to participate in the study by signing anform informed consent.

Geographical basic data retrieval in the form of age, gender, education level and occupation. The collection of clinical baseline data which included a family history of cancer, a history of comorbid diseases, a history of habits related to colorectal cancer (smoking, alcoholism, and diet), clinical symptoms related to colorectal cancer and body mass index. Blood specimen collection for examination at the Eureka Research Laboratory, Palembang, Indonesia, which includes complete routine blood (hemoglobin, leukocytes, platelets, blood chemistry (kidney function, liver function), colorectal cancer biomarkers (CEA, CA19-9), and cfDNA, in accordance with the protocols and procedures of the Eureka Research Laboratory.

The data that has been collected is processed using the SPSS version 26.0 program which will be displayed in the form of narration, tables and graphs. Numerical data is expressed as the appropriate median, minimum and maximum. Qualitative data is expressed as frequency and percentage for quantitative data that are not normally distributed, the comparison between the two groups was carried out using the Mann-Whitney test (non-parametric t test). The comparison between the 3 groups was carried out using the Kruskal-Wallis test and the post-Hoc "Schefe test" was used for pairwise comparisons. Spearman-rho is used to test the correlation between numerical variablesCurve Receiver Operating Characteristic (ROC) plotted for evaluation of the diagnostic test. All tests have two sides. The p value < 0.05 was considered significant.

#### 3. Results

Research variables included in the general characteristics of this study consisted of age, gender, body mass index (BMI) group, history of low fiber diet, smoking history, and alcohol history. Of the 50 research subjects, the average age of the patients was  $53.14 \pm 10.18$  years. The older group without cancer had an older mean age, although the mean difference was not significant. The group of patients with cancer had a mean age of  $48.91\pm13$  years and the group of

patients without a history of KKSV had a mean age of 54.33+9.09. As many as 25 (50%) patients were male and males were more commonly found in the cancer group, while females were more commonly found in the non-cancer group. As many as 40% of patients underweight and cancer group tend to be more underweight. There was no difference in gender and nutritional status in the two groups of research subjects. Based on the history of risk factors, it is known that most of the study participants have a lowfiber diet, do not smoke, and do not consume alcohol. The risk factors for consuming a low-fiber diet were found to be significantly more in the adenocarcinoma group. Meanwhile, there was no significant difference between smoking history and alcohol history in the two groups.

	Subjects (n = 50,%)	cancer (n = 39,%)	without cancer (n = 11%)	р
Age (years) *	53.14 ± 10.18	48.91 ± 13	54.33 <u>+</u> 9.09	<b>0.12</b> <sup>1</sup>
Gender				
Male Female	25 (50%) 25 (50%)	20 (51.3%) 19 (48.7%)	5 (45.5%) 6 ( 54.5%)	0.732
Nutritional Status				
Underweight Normoweight	20 (40%) 30 (60%)	18 (46.2%) 21 (53.8%)	2 (18.2%) 9 (81.8% )	0.16 <sup>3</sup>
Low Fiber Diet				
Yes None	46 (92%) 4 (8%)	38 (97.4%) 1 (2.6%)	8 (72.7%) 3 (27.3%)	0 ,03 <sup>3#</sup>
Smoking				
Yes None	23 (46%) 27 (54%)	19 (48.7%) 20 (51.3%)	4 (36.4%) 7 (63.6%)	0.472
Alcohol Consumption				
Yes No	7 (14%) 43 (86%)	5 (12.8%) 34 (87.2%)	2 (18.2%) 9 (81.8%)	0.64 <sup>3</sup>

Values are enclosed in the mean + SD or value (% percentage).

\*Kolmogorov Smirnov test p>0.05: data normally distributed, values attached in mean+SD

<sup>1</sup>Independent T-Test, <sup>2</sup>Chi Square tests, <sup>3</sup>Fisher Exact Test Cancer vs. No Cancer Group. #Significant p<0.05

The results of the examination of cfDNA, CEA, and CA19-9 are presented in table 2. From 50 research subjects, it was found that cfDNA in research subjects had a median value of 59.71 ng/mL, a minimum value of 8.43 ng/mL, and a maximum value of 157.86

ng/mL. cfDNA values were found to be significantly higher in the group of patients with cancer. The group with cancer also had higher CEA and CA19-9 values than the group without cancer.

Table 2. Results of examination of tumor markers in both groups					
Colorectal Tumor Markers	Subjects (n=50)	Colorectal Cancer (n=39)	Without Cancer (n=11)	р	
cfDNA (ng/mL)	59.71 (8.43-157, 86)	66.57 (25.86-157.86)	44 (8.43-56.43)	< 0.001	
CEA (ng/mL) CA19-9 (U/mL)	106 (1.10-984)	126 (1.10-984)	2, 20 (1.10-42.50) 3.50 (1.20-12)	< 0.001	
	49 (1.2-652)	88.60 (2-652)	5.50 (1.20-12)	< 0.001	

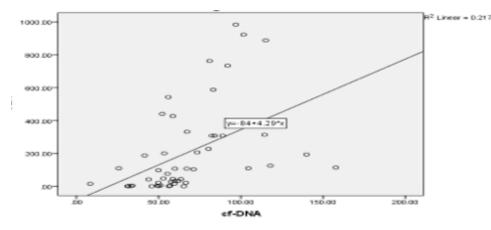
Distributed data abnormal with the Kolmogorov-Smirnov Test. The median difference test between cancer and no cancer groups with the test *Mann-Whitney*.

The results of the Spearman's rho test showed that there was a significant relationship between the levels of cfDNA and CEA (p=<0.001) with a strong positive correlation strength (r=0.635). Meanwhile, cfDNA and CA19-9 had a moderate correlation value (r=0.572) which was statistically significant (p=<0.001). The presentation of data regarding the correlation of cfDNA levels with CEA and CA19-9 in suspected colorectal cancer patients at Dr. Mohammad Hoesin Hospital Palembang can be seen in Table 3 and Figure 1.

Table 3. Correlation between cfDNA levels with CEA and CA19-9 levels in research subjects

	Correlation relationship between	R	р	
 cfDNA	CEA	0.635	< 0.001	
	CA19-9	0.572	< 0.001	
 ala mha taat	the n velue means if n<0.05. Correlation as	trangeth is norme mostly if no0 0	recall if r=0.01	

Spearman's rho test, the p value means if p<0.05. Correlation strength is very weak if r<0.2, weak if r=0.21-0.4, medium if r=0.41-0.6, strong if r=0.61-0.8 and very strong if >0.8



#### Correlation of cfDNA levels with serum CEA levels of patients suspected of colorectal cancer

#### Correlation of cfDNA levels with serum CA19-9 levels of patients suspected of colorectal cancer

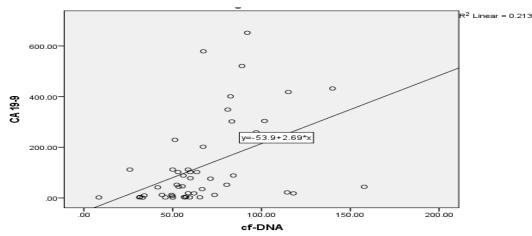


Figure 1. cfDNA correlation graph with CEA and CA19-9

Figure 2 shows that for cfDNA levels, the optimal cut-off point between sensitivity and specificity is at point 13 which when viewed in the analysis table is 50.64 ng/dL with sensitivity 92.3%, specificity 90.9%, positive predictive value 95.2%, negativevalue 82.3%, area under the curve predictive0.945 and accuracy

92.4%. The group of patients based on this level was the group with low serum cfDNA levels, namely <50.64ng/mL and the group with high serum cfDNA levels, 50.64ng/dL.

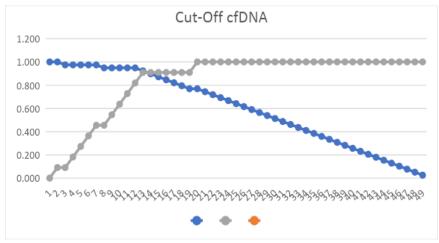


Figure 2. Cutoff point of serum cfDNA

Table 4. Diagnostic bivariate analysis of cfDNA compared to histopathological examination results in colorectal cancer diagnosis

cfDNA	Colorectal Cancer			
(cutoff 50.64 ng/mL)	Cancer	Without Cancer		
High cfDNA	37 (81.8%)	2 (5, 2%)		
low cfDNA	2 (18.2%)	9 (94.8%)		

Table 5. Diagnostic significance of cfDNA versus histopathological examination results in colorectal cancer diagnosis

Diagnostic Significance	Value (95%CI)		
AUC	0.945 (0.842-0.990)		
Sensitivity	92 ,3% (79.1%-98.4%)		
Specificity	90.9%(58.7%-99.8%)Prediction		
PositiveValue	95.2%(81.4%-99.7%) Prediction		
Value Negative	82.3%(48.7%-97.1%)		
Accuracy	92.4%(80.1%-97.4%)		

To determine the diagnostic significance of CEA, CA19-9, compared with the diagnostic value of cfDNA was analyzed bivariate between each diagnostic marker examined. In addition, analysis with combination markers was also carried out to determine the diagnostic value compared to the histopathological examination results. The calculation results are shown in table 5.

Table 6. Diagnostic significance between combination markers in colorectal cancer diagnosis

Markers (cutoff)	AUC	Sensitivity	Specificity	Positive predictive value (PPV)	Negative predictive value (NPV)	Diagnostic accuracy
cfDNA (50.64ng/mL)	0.945	92 ,3%	90.9%	95.2%	82.3%	92.4%
CEA (25ng/ml)	0.942	87.2%	90.9%	97.1%	66.8%	87.9%
CA19-9 ( 15U/ml)	0.935	87.1%	100%	100%	69.3%	90.1%
cfDNA and CEA	0.977	94.9%	81.8%	94.9%	91.8%	92.3%
cfDNA and CA19-9	0.988	87.2%	100%	100%	64.2%	90.1%

CfDNA, CEA, and CA19-9 examinations provide excellent AUC results with AUC's 0.945, 0.942, and 0.935, respectively. A combination examination of

#### 4. Discussion

Findings El-Gayar et al who analyzed the accuracy of cfDNA in Egypt using the nanodrop method against ALU markers as in this study showed lower cfDNA levels with a median level of 4.6ng/mL (range 1.1-48ng/mL).<sup>31</sup> This difference in the distribution of values may be influenced by differences in the course of the patient's disease, differences in race, and differences in laboratory analysis methodologies. To date, there is no universally used cfDNA assay standard.

The cfDNA value was found to be significantly higher in the group of patients with colorectal cancer

cfDNA with CEA and cfDNA with CA19-9 provides better AUC examination results than cfDNA, CEA, or CA19-9 examinations themselves.

(p<0.001). The results are similar to those of El-Gayar et al., Mead et al., Hao et al., and Zaher et al. who found higher cfDNA in the colorectal cancer group than in the noncancerous lesion group and in healthy controls.<sup>31,40,51-52</sup> Almost all previous reports using the ALU marker showed consistent results of significantly higher cfDNA levels in the colon cancer group than in the group without colon cancer. In healthy individuals, cfDNA is mostly derived from leukocytes. While in the case of cancer, cfDNA is likely derived from apoptosis and necrosis of cancer cells or surrounding cells and possibly through active release by cancer cells.<sup>53</sup>

CEA and CA 19-9 are the two major tumor markers

for colorectal cancer. CEA is atumor marker broad spectrum and is one of the many markers used in colorectal cancer. This study showed that the group with colorectal cancer had higher CEA and CA19-9 values than the group without cancer. These results are consistent with the studies of Flamini et al, Qi et al, Hao et al, and El-Gayar et al, which also showed CEA and CA19-9 values in the colorectal cancer group were higher than in the non-cancer group.<sup>31,49-50,53</sup> Various studies have shown that an increase in serum CEA in colorectal cancer patients indicates a worse prognosis than in patients with normal CEA levels. CA19-9 was also found to be elevated in colorectal cancer. Shin Nishiumi et al explained that the expression of CA19-9 in colorectal cancer patients plays a role in the diagnosis of colorectal cancer.55 László Herszényi et al also concluded in their study that in some colorectal cancer patients, there was a continuous increase in serum levels of CEA and CA19-9 about 2-5 months before they were declared relapse or metastases.<sup>56</sup> The results of this study are consistent with various studies.

The results of the Spearman's rho correlation test showed that there was a significant relationship between cfDNA levels and CEA with a strong positive correlation strength. Meanwhile, cfDNA and CA19-9 had a statistically significant moderate correlation value. This result contradicts the findings of Qi et al in 2013 who reported a weak and insignificant correlation between cfDNA and CEA (r=0.232, p=0.210) and between cfDNA and CA19-9 (r=-0.158, p=0.397). Qi et al suggested that cfDNA may be an independent predictor of colonic carcinoma progression which was not correlated with CEA or CA19-9.

Salem et al's study in Egypt showed results that were in line with this study.<sup>67</sup> The study showed a significant positive correlation between the DNA integrity index cfDNA markers of ALU and CEA, although the correlations proved tended to be weak (r=0.39, p=0.03). The El-Gayar study also showed results in line with moderate correlations of cfDNA with CA19-9 (r=0.4, p<0.001) and CEA (r=0.45, P<0.001).<sup>31</sup> This study has consistently shown that cfDNA correlates with CA19-9 and CEA so that cfDNA can be used as a tumor marker in the diagnosis of colorectal cancer.

#### 5. Conclusion

In this study, cfDNA has excellent accuracy in diagnosing colorectal cancer.

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