Laboratory Examination in Hepatocellular Carcinoma

Anggraini D

1 Departement of Clinical Pathology, Faculty of Medicine, Universitas Baiturrahmah, Padang, Indonesia
Email : debieanggraini@fk.unbrah.ac.id

Abstract

Hepatocellular carcinoma (HCC) is a malignant tumor originating from hepatocyte cells including primary malignant tumors of the epithelial liver. Hepatocellular carcinoma is the fifth most malignant in the world and the third largest cause of death of all malignancies. Diagnosis of HCC is obtained from the patient's history, examination of the patient, by imaging (ultrasonography, MRI or CT scan) and an increase in serum tumor biomarkers of alpha feto protein (AFP) (> 400 ng/mL). Early diagnosis is important to increase the patient's life expectancy and only 30-40% of patients are diagnosed at an early stage. Biomarkers in HCC not only help in diagnosing but also predict prognosis or recurrence and in choosing therapy. The stability of miRNA in blood circulation makes miRNA an ideal candidate for use in the diagnosis and treatment of cancer. Objective: to briefly review all current information about HCC. Conclusion: HCC diagnosis is obtained from the patient's history, physical examination, imaging (ultrasonic, MRI or CT scan that shows the presence of a typical liver mass for HCC) and an increase in AFP serum (> 400 ng/mL). Biomarker examination and evaluation is still very limited, one of which is the examination of MicroRNAs

Keywords — Biomarkers, hepatocellular carcinoma, microRNAs.

Katakunci — Biomarkers, karsinoma hepatoseluler, mikroRNAs.

Abstract

Hepatocellular carcinoma (HCC) adalah tumor ganas yang berasal dari sel hepatosit termasuk tumor ganas primer hati epitel. Karsinoma hepatoseluler adalah yang paling ganas ke lima di dunia dan penyebab kematian terbesar ketiga dari semua keganasan. Diagnosis HCC diperoleh dari riwayat pasien, pemeriksaan pasien, dengan pencitraan (ultrasonografi, MRI atau CT scan) dan peningkatan biomarker tumor serum protein alpha feto protein (AFP) (> 400 ng/mL). Diagnosis dini penting untuk meningkatkan harapan hidup pasien dan hanya 30-40% pasien didiagnosis pada tahap awal. Biomarker di HCC tidak hanya membantu dalam mendiagnosis tetapi juga memprediksi prognosis atau rekurensi dan dalam memilih terapi. Stabilitas miRNA dalam sirkulasi darah menjadi kandidat miRNA yang ideal untuk digunakan dalam diagnosis dan pengobatan kanker. Tujuan: Tujuan: untuk meninjau secara singkat semua informasi terkini tentang HCC. Kesimpulan: Diagnosis kanker hati didapat dari riwayat pasien, pemeriksaan fisik, pencitraan (ultrasonik, MRI atau CT scan) yang menunjukkan adanya massa hati yang khas untuk kanker hati) dan peningkatan serum AFP (> 400 ng/mL). Pemeriksaan dan evaluasi biomarker masih sangat terbatas, salah satunya adalah pemeriksaan MicroRNA

Keywords— Biomarker, Karsinoma Hepatoselular, mikroRNAs
I. INTRODUCTION

Hepatocellular carcinoma (HCC) is a malignant tumor originating from hepatocyte cells, including primary malignant tumors of the liver epithelial. Hepatocellular carcinoma accounts for 90% of primary liver cancer, which is the fifth most malignant in the world, and the third leading cause of death in all malignancies. The annual incidence of HCC continues to increase in each country and it is estimated that there are 500,000 to one million cases each year, including Indonesia.

II. RISK FACTORS OF HEPATOCELULAR CARCINOMA

Some risk factors of HCC include hepatitis B and C virus infections (HBV and HCV), aflatoxin B1 exposure (AFB1), chronic alcohol abuse and metabolic syndrome. About 54.4% of HCC cases are associated with HBV infection, so it is very interesting to examine the underlying mechanisms.

Three factors that influence the development of HCC in chronic HBV-infected individuals include the integration of viral DNA into the host cell genome, chronic inflammation due to the host's immune response to HBV infection and HBx protein which modulates the cellular signal transduction pathway. Hepatitis B virus X protein (HBx) is a major regulator protein encoded by the X-region HBV genome with a length of 154 amino acids, 17 kDa, which has an important role in the trans-activation of viral gene expression at the transcriptional level. HBx protein can induce genetic changes in tumor suppressor genes and oncogenes and epigenetic aberrations in HCC. HBx proteins regulate non-coding RNA (ncRNAs), consisting of microRNAs (miRNAs) and long non-coding RNA (lncRNA), which play a role in HBV-related hepatocarcinogenesis, modulate cell proliferation, cell cycle, anti-apoptosis, invasion and metastasis.

III. MICRORNAS AND HEPATOCELULAR CARCINOMA

MicroRNAs are small RNA molecules measuring 21-25 nucleotides, part of a non-coding RNA that acts as a regulator of posttranscriptional gene expression at the level of mRNA, which regulates liver development and homeostasis. MiRNA dysfunction is strongly associated with the tumor process because HBx protein regulates ncRNAs including the MiRNA to act as an oncogen or tumor suppressor gene that can develop or inhibit cancer. As a multifunctional protein, even though HBx protein does not have a DNA-binding domain, HBx protein can modulate ncRNAs indirectly, which interacts with various transcription factors such as p53, c-myc and nuclear factors kappa B and activates cellular signal transduction pathways, which change expression host genes and leads to the development of HCC.

Tumor gene protein 53 (p53) is a transcription factor that is a tumor suppressor, has a multifunction in controlling DNA replication and repair, maintaining genomic stability, cell cycle progression and programmed cell death. The presence of mutations and polymorphisms in p53, which cause tumor suppressors to become inactive, are often associated with the development of HCC in humans, where 30% of cases of HCC are caused by mutations that cause inactivation of the p53 gene. In addition, it was found that dysregulation of the p53 miRNA pathway in HCC would affect the expression of target mRNA involved in cell cycle regulation, apoptosis, angiogenesis, DNA repair and metastasis.

Disregulation of the p53 miRNA pathway can have an oncogenic effect or eliminate tumor suppressor function in HCC. Increased dysregulation of miRNA in HCC includes miR-221, miR-222 which causes tumor suppressor proteins p21, p27 and p57 to be suppressed, this will affect cyclins.
dependent kinase (CDK2) and cyclin E so that cell cycle development occurs in HCC 14. In addition, anti-apoptotic molecules such as Bcl-2, Bcl-w, Bcl-xl and Mcl-1, modulated by miRNAs such as miR-122, miR-34a, miR 125b and miR-let7, on HCC there is decreased dysregulation of these miRNAs resulting in failure to modulate the effects of anti-apoptotic molecules 14.

The stability of miRNA in blood circulation makes miRNA an ideal candidate for use in the diagnosis and treatment of cancer 17. The role of miR-221 in the pathogenesis of HCC has been widely reported, but in studies in China, differences in serum miR-221 levels in HCC patients and normal controls were not significant, so further research is needed to determine the diagnostic value of this miR-221 18.

Luo’s research, et al. Stated that the sensitivity and specificity of miR-122 in HCC in China were 70.6% and 67.1% respectively, if miR-122 was combined with alpha-fetoprotein (AFP) serum, the sensitivity and specificity increased respectively, each being 87.1% and 98.8% 18. Hung’s study, et al., The sensitivity and specificity of miR-let7 in HCC patients with chronic hepatitis B were 84.8% and 50%, respectively 19.

Bharali’s study, et al., The sensitivity and specificity of miR-34a in HCC patients in India were 81.8% and 82.7% 20, respectively. Xu’s research, et al. 2018 in China the sensitivity and specificity of miR-125b in HCC patients were 83% and 96%, respectively 21.

IV. DIAGNOSIS OF HEPATOCELLULAR CARCINOMA

Sensitive and specific initial diagnostic markers are needed to diagnose HCC, but at present the diagnosis is still dependent on serum alpha-fetoprotein (AFP) levels and ultrasonography (USG) 18,22. High morbidity in HCC is associated with a lack of markers of early detection of the disease and very poor prognosis, thus limiting choices in chemotherapy, adjuvant therapy, or surgical procedures. Besides that, examination of AFP levels and ultrasound screening in HCC-suspected patients sometimes underdiagnosis and false positives 18. Therefore, exploration in predictors of noninvasive early diagnosis of HCC 17,18,23 because early detection of HCC is very important for successful treatment and better patient survival 18.

V. CONCLUSION

HCC diagnosis is obtained from the patient's history, physical examination, imaging (ultrasonic, MRI or CT scan that shows the presence of a typical liver mass for HCC) and an increase in AFP serum (> 400 ng / mL). Biomarker examination and evaluation is still very limited, one of which is the examination of MicroRNAs

DAFTAR PUSTAKA


[6] Liu S, Koh SSY, Lee CGL. Hepatitis B Virus X Protein and Hepatocarcinogenesis. (Figure 1):1–14.


[23] Lou J, Zhang L, Lv S. Biomarkers for Hepatocellular Carcinoma. 2017;