Utility of serum neuropilin-1 and angiopoietin-2 as markers of hepatocellular carcinoma

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ABSTRACT
This study aimed to assess the diagnostic value of two serum angiogenic markers neuropilin-1 (NRP-1) and angiopoietin-2 (ANG-2) in patients with hepatocellular carcinoma (HCC) and their relation to tumor characteristics. 149 subjects were recruited and divided into five groups: 50 patients with recently diagnosed HCC, 49 patients with cirrhosis on top of hepatitis C virus infection, and 50 healthy subjects. Serum NRP-1 and ANG-2 were estimated by ELISA. Alpha-fetoprotein (AFP) levels were measured using fluorescence immunoassay. Serum NRP-1 and ANG-2 levels were significantly higher in patients with HCC (2221.8±1056.8 pg/mL and 3018.5±841.4 pg/mL) than healthy subjects (219.3±61.8 pg/mL and 2007.7±904.8 pg/mL) and patients with cirrhosis (1108.9±526.6 pg/mL and 2179.1±599.2 pg/mL), respectively. In multivariate logistic regression analysis, NRP-1 and ANG-2 were the only independent factors of HCC development and correlated positively with each other (r=0.781, p<0.001). Receiver operating characteristic curve analysis showed that the area under the curve (AUC) of NRP-1 was higher than that of ANG-2 in discriminating HCC from patients with cirrhosis (0.801 vs 0.748, p=0.250) and healthy subjects (0.992 vs 0.809, p<0.001). The AUC of NRP-1 was detected to be increased (0.994) when combined estimation with AFP was performed. Elevated serum NRP-1 and ANG-2 levels were detected in patients with HCC with tumor numbers ≥3, tumor size ≥5 cm, tumor stages B/C according to the Barcelona Clinic Liver Cancer staging system, vascular invasion, and distant metastasis. In conclusion, NRP-1 is a potential serological marker for HCC diagnosis and is better than ANG-2. It is feasible to be estimated in combination with AFP to enhance its diagnostic power. High serum NRP-1 and ANG-2 levels are associated with advanced HCC tumor characteristics.

INTRODUCTION
Hepatocellular carcinoma (HCC) is the sixth most common malignancy and the fourth leading cause of cancer-related mortality worldwide in 2018.1 Most cases of HCC had developed on top of chronic hepatitis infections caused by either hepatitis B virus (HBV) or hepatitis C virus (HCV). In Egypt, due to the high prevalence of HBV and HCV chronic infections, HCC came on top of solid malignancies.2

Due to the absence of symptoms of early HCC and the lack of its effective screening strategies, most patients with HCC are diagnosed with late-stage disease after metastasis has occurred, resulting in a very poor prognosis and a low overall 5-year survival rate of <16%.3 Therefore, early detection of HCC is essential to improve patients’ survival.4 The American Association for the Study of Liver Diseases (AASLD) once recommended serum alpha-fetoprotein (AFP) as...
a screening test for HCC surveillance in patients with hepatic cirrhosis. However, one major disadvantage is that AFP levels can be falsely elevated in patients who have active hepatitis with no evidence of HCC. Also, about one-third of HCC cases have normal serum AFP especially when tumor size is less than 2 cm. The role of biomarkers in various malignancies has emerged, and thus searching for novel serological biomarkers is needed for screening HCC to reduce its high mortality.

Hepatic tumorigenesis is a multistep complicated process characterized by uncontrolled cellular proliferation and increased angiogenesis to enhance tumor growth. There is evidence that angiogenesis plays a vital role in the development and progression of HCC. Neuropilin-1 (NRP-1) is a transmembrane glycoprotein that is primarily found to play a role in neuronal axon guidance and embryonic angiogenesis. It was described as a coreceptor for vascular endothelial growth factors (VEGFs) and for secreted semaphorins in association with VEGF receptors or plexins, respectively. However, it was shown that NRP-1 also interacts with epidermal growth factor receptor (EGFR) on the cell surface, enhancing ligand-induced EGFR clustering and intracellular signaling. Additional growth factor-induced signaling cascades have been associated with NRP-1, although the implicated molecular mechanisms remain largely unclear. Thus, NRP-1 seems to act as a receptor hub on the cell surface, promoting multiple signaling cascades. NRP-1 can encourage different aspects of tumorigenesis, such as angiogenesis, cell survival, migration, invasion, and chemoresistance. NRP-1 is reported to be upregulated in several tumors including astrocytomas, breast, prostate, lung, pancreas, bile duct, gastric, and colon cancers. Overexpression of NRP-1 is closely correlated with the infiltration and migration of tumors. The contribution of NRP-1 and its ligands to tumor growth and metastasis has spurred a strong interest in NRP-1 antagonists used in combination with anti-VEGF chemotherapy as novel antiangiogenesis therapies. Previous studies have reported that NRP-1 is overexpressed in HCC tissue compared with normal tissue, but the use of serum NRP-1 as a diagnostic marker for HCC continues to be a subject of interest especially when compared with AFP.

Angiopoietins have been identified as ligands for vascular endothelial-specific Tie2 receptor tyrosine kinase and may be important growth factors in the generation of new blood vessels. Angiopoietin-2 (ANG-2), a member of the angiopoietins, is thought to have a remarkable role in angiogenesis. A previous study showed that increased ANG-2 can result in persistent disruption of the cellular crosstalk between endothelial cells and pericytes. Furthermore, high concentrations of ANG-2 are considered to function as antiapoptotic factors in endothelial cells by activating the phosphatidylinositol 3′-kinase/Akt signaling pathway. Levels of ANG-2 mRNA expressions have been reported to be significantly increased in HCC, non-small cell lung cancer, and gastric cancer compared with adjacent non-cancerous components. This increased expression may play a critical role in promoting tumor angiogenesis and progression.

The purpose of the current study was to measure serum levels of NRP-1 and ANG-2 to determine its potential diagnostic utility in patients with HCC and to determine their relation to the tumor characteristics in these patients.

PATIENTS AND METHODS
Patients and samples
This is a retrospective study carried out in the internal medicine, tropical medicine, clinical oncology and nuclear medicine, physiology and clinical pathology departments of Tanta University Hospitals, Egypt. A total of 149 subjects were included in our study in a convenience series, and they were divided into three groups: group I: 50 patients recently diagnosed with HCC; group II: 49 patients suffering from liver cirrhosis on top of HCV infection; and group III: 50 apparently healthy individuals matched in age and sex, which served as the control group. The diagnosis of HCC was made on histopathological confirmation of hepatic focal lesions, or if not available clinical and radiological HCC diagnosis based on the guidelines of the AASLD. The staging was made according to the Barcelona Clinic Liver Cancer (BCLC) staging system. Patients suffering from any malignant diseases other than HCC or those with HCC who received treatment were excluded from the study. Patients’ flow throughout this study is demonstrated in online supplemental figure S1.

Informed consent was obtained from both the patients and the control group.

All patients were subjected to the following: full history taking, thorough clinical examination, and radiological investigations including abdominal ultrasonography and CT. Lower limb edema was examined clinically, whereas ascites was assessed by physical examination and abdominal ultrasonography and graded according to the European Association for the Study of the Liver as follows: grade 1 was defined as mild ascites only detectable by ultrasound, grade 2 was defined as moderate ascites evident by moderate symmetrical distension of the abdomen, and grade 3 was defined as large or gross ascites with marked abdominal distension. The Child-Pugh-Turcotte class was determined and the Model for End-Stage Liver Disease (MELD) score was calculated for patients with HCC and cirrhosis according to published guidelines. Blood collection for patients and control groups
Whole blood was collected by standard venipuncture in VACUETTE Blood Collection Tubes (Greiner Bio-One, Krefeld, Germany), and a tube containing clot activator for complete blood picture (complete blood count) on fully automated cell counter (Erma, PCE 210N, Tokyo, Japan), and a tube containing clot activator for automated chemistry analyzer (Konelab Prime 60i, Vantaa, Finland) and AFP on automated fluorescence immunosassay.
analysts (Tosoh AIA 1800 ST, Tokyo, Japan). The results were interpreted separately by independent investigators, who were blinded to other laboratory results as well as clinical, radiological, and diagnostic findings of the study subjects.

**Serum NRP-1 determination**

Serum NRP-1 level was measured using a commercially available, quantitative sandwich enzyme immunoassay technique (Human Neuropilin-1, Abcam ELISA Kit, catalog number ab227901, Minneapolis, USA) in accordance with the manufacturer’s instructions. The sensitivity is 7 pg/mL. The intra- and interassay coefficient of variation (CV%) were 1.9% and 3%, respectively.

**Serum ANG-2 determination**

Serum ANG-2 level was measured using a commercially available, quantitative sandwich enzyme immunoassay technique (Human Angiopoietin-2, Quantikine ELISA Kit, catalog number DANG20, R&D Systems, Minneapolis, USA) in accordance with the manufacturer’s instructions. The sensitivity is 8.29 pg/mL. The intra- and interassay CV% were 4.2%–6.9% and 7.4%–10.4%, respectively.

**Statistical analysis**

The analyses and graphs of the current study results were performed using IBM SPSS V.24, MedCalc V.18, and GraphPad Prism. Shapiro-Wilk test was conducted to check for normal distribution of dependent variables. Normally distributed numerical results were compared between the studied groups by Student’s t-test. One-way analysis of variance with post-hoc test was used to compare the results among more than two groups. Non-parametric numerical results were compared by Mann-Whitney U test. Univariate and multivariate logistic regression analyses were used to determine the related covariables with HCC development. Pearson correlation was used for correlating serum AFP with NRP-1 and ANG-2 levels. These markers were combined using binary logistic regressions, and their diagnostic characteristics were determined by receiver operating characteristic (ROC) curve analyses. The optimal cut-off value for each marker was assessed via Youden’s index. P values were considered significant if they are less than 0.05.

**Sample size calculation**

We conducted a power analysis (G Power V.3.1 statistical software, Franz Faul, Universität Kiel Germany) based on a previous study. The findings indicated a minimum sample size of 66 samples (22 samples for each group) based on an α of 5% and a power of 95%. Considering a possible loss of about 10% of patients, we should use a minimum of 75 samples (25 samples for each group).

**RESULTS**

**Clinicopathological characteristics of study subjects**

A total of 149 subjects were recruited to this study as follows: 50 patients recently diagnosed with HCC, 49 patients with cirrhosis on top of HCV infection, and 50 healthy subjects as the control group. There was no significant difference in age and sex among the three groups (p=0.422, p=0.720), respectively. Regarding the laboratory data, there was a statistically significant difference in LFT and international normalized ratio (INR) among the three groups. Serum AFP, NRP-1, and ANG-2 levels were significantly higher in the HCC group than in the control and cirrhotic groups (p<0.001) (table 1). Patients with cirrhosis have a mean MELD score of 18.4±5.4. There were six cases with Child A, 28 with Child B, and 15 with Child C. Patients with HCC have a mean MELD score of 16.9±6.0, with no significant difference.

### Table 1  Demographic, clinical and laboratory characteristics of the studied groups

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Group I (HCC) (n=50)</th>
<th>Group II (cirrhosis) (n=49)</th>
<th>Group III (healthy control) (n=50)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age (years)</td>
<td>59.2±6.7</td>
<td>58.8±6.3</td>
<td>57.5±7.1</td>
<td>0.422</td>
</tr>
<tr>
<td>Gender, n (%)</td>
<td>27 (54.0)</td>
<td>23 (46.9)</td>
<td>27 (54.0)</td>
<td>0.720</td>
</tr>
<tr>
<td>Male</td>
<td>23 (46.0)</td>
<td>26 (53.1)</td>
<td>23 (46.0)</td>
<td></td>
</tr>
<tr>
<td>Female</td>
<td>26 (52.0)</td>
<td>33 (67.3)</td>
<td>–</td>
<td>0.120</td>
</tr>
<tr>
<td>Ascites, n (%)</td>
<td>34 (68.0)</td>
<td>39 (79.6)</td>
<td>–</td>
<td>0.190</td>
</tr>
<tr>
<td>Ascites, n (%)</td>
<td>2.2±1.2†</td>
<td>2.4±0.9†</td>
<td>0.6±0.2</td>
<td>&lt;0.001*</td>
</tr>
<tr>
<td>ALT (U/L)</td>
<td>76±32†</td>
<td>68±19†</td>
<td>20±6</td>
<td>&lt;0.001*</td>
</tr>
<tr>
<td>AST (U/L)</td>
<td>95±43†</td>
<td>79±25†</td>
<td>24±7</td>
<td>&lt;0.001*</td>
</tr>
<tr>
<td>ALK (U/L)</td>
<td>96±29†</td>
<td>106±27†</td>
<td>64±19</td>
<td>&lt;0.001*</td>
</tr>
<tr>
<td>Albumin (g/dL)</td>
<td>2.7±0.8†</td>
<td>2.5±0.4†</td>
<td>4.0±0.4</td>
<td>&lt;0.001*</td>
</tr>
<tr>
<td>INR</td>
<td>1.9±0.7†</td>
<td>2.1±0.7†</td>
<td>1.0±0.1</td>
<td>&lt;0.001*</td>
</tr>
<tr>
<td>AFP (ng/mL)</td>
<td>356.1±257.3†</td>
<td>155.2±65.6†</td>
<td>4.9±2.0</td>
<td>&lt;0.001*</td>
</tr>
<tr>
<td>NRP-1 (pg/mL)</td>
<td>2221.8±1056.6†</td>
<td>1108.9±526.8†</td>
<td>219.3±61.8</td>
<td>&lt;0.001*</td>
</tr>
<tr>
<td>ANG-2 (pg/mL)</td>
<td>3018.5±841.4†</td>
<td>2179.1±599.2</td>
<td>2007.7±904.8</td>
<td>&lt;0.001*</td>
</tr>
</tbody>
</table>

Data are expressed as mean±SD.

*P<0.05 significant.

†Significant with group III.

‡Significant with group II.

AFP, alpha-fetoprotein; ALK, alkaline phosphatase; ALT, alanine transaminase; ANG-2, angiopoietin-2; AST, aspartate transaminase; HCC, hepatocellular carcinoma; INR, international normalized ratio; NRP-1, neuropilin-1.
had vascular invasion, and distant metastasis. Thirty-three patients (66%) had tumor numbers >3, 19 (38%) had tumor size ≥5 cm, and 5 (10%) had distant metastasis. Moreover, 4 patients (8.0%) were presented in stage 0, 13 patients (26%) in stage A, 12 patients (24%) in stage B, and 21 patients (42%) in stage C according to the BCLC staging system.

Demographic and biochemical covariables in relation to HCC

Logistic regression analyses were performed to detect the related covariables for HCC development. In univariate analysis, total bilirubin, aspartate transaminase, albumin, INR, AFP, NRP-1, and ANG-2 were significantly associated with HCC development, while age and sex were non-significant variables. In multivariate analysis, AFP and NRP-1 remained the only significant independent factors with HCC (table 2).

Diagnostic value of NRP-1 and ANG-2

As serum NRP-1 and ANG-2 levels were significantly higher in patients with HCC than in healthy subjects and patients with cirrhosis, we used ROC curve analysis to evaluate the diagnostic efficacy of both markers in discriminating HCC from healthy subjects or patients with cirrhosis compared with AFP.

The area under the ROC curve (AUC) indicated that NRP-1 had the highest diagnostic efficacy in differentiating HCC from healthy subjects (AUC 0.992), which is slightly higher than that of AFP (AUC 0.983) with no significant AUC difference (0.009, p=0.431) and that of ANG-2 (AUC 0.809) with significant AUC difference (0.183, p<0.001) (table 3, figure 1). Moreover, the diagnostic efficacy of NRP-1 (AUC 0.801) in discriminating HCC from patients with cirrhosis was higher than that of AFP (AUC 0.762) with no significant AUC difference (0.039, p=0.465), and also higher than that of ANG-2 (AUC 0.748) with no significant AUC difference (0.053, p=0.250) (table 3, figure 1).

Using Pearson correlation analysis, serum NRP-1 and ANG-2 correlated positively with each other and with AFP (p<0.001) (figure 2). Linear regression analysis revealed that NRP-1 is independently related to AFP (p<0.001), whereas no significant relation was detected between ANG-2 and AFP (p=0.225). Therefore, we suggested that NRP-1 and AFP be considered as the best combination. Based on these results, the combined analysis of NRP-1 and AFP revealed that the AUC of combined AFP and NRP-1 (AUC 0.825) was slightly higher than that of NRP-1, AFP, and ANG-2 in differentiating HCC from patients with cirrhosis, with no significant AUC differences. Also, the AUC of combined AFP and NRP-1 (AUC 0.994) was slightly higher than that of NRP-1, AFP, and ANG-2 in differentiating HCC from healthy controls, with no significant AUC differences except with ANG-2 (0.184, p<0.001) (table 3, figure 1).

Biomarkers correlation with clinicopathological features in HCC

In patients with HCC, serum NRP-1 and ANG-2 levels were significantly higher in BCLC stages B and C than stages 0 and A. Moreover, serum NRP-1 and ANG-2 levels were significantly higher in patients with HCC with tumor numbers >3 (p<0.001, p<0.001), tumor size ≥5 cm (p=0.001, p=0.032), as well as the presence of vascular invasion (p=0.001, p<0.001) and distant metastasis (p=0.008, p=0.033), respectively (figure 3).

DISCUSSION

In the present study, we studied two serum angiogenetic markers, NRP-1 and ANG-2, in patients with newly diagnosed HCC, searching for their role in the early diagnosis of HCC and their relationship with tumor characteristics. Accordingly, 50 untreated patients with HCC were enrolled and their serum levels of NRP-1 and ANG-2 were estimated and compared with that of 49 patients with liver cirrhosis on top of HCV infection.
Table 3  ROC curve analysis of serum NRP-1 and ANG-2 performance characteristics among the studied groups

<table>
<thead>
<tr>
<th>Serum NRP-1</th>
<th>AUC</th>
<th>95% CI</th>
<th>P value</th>
<th>Cut-off</th>
<th>Sensitivity</th>
<th>Specificity</th>
</tr>
</thead>
<tbody>
<tr>
<td>Patients with HCC (group I) vs patients with liver cirrhosis (group II)</td>
<td>0.801</td>
<td>0.709 to 0.875</td>
<td>&lt;0.001*</td>
<td>&gt;1418</td>
<td>72.0</td>
<td>87.8</td>
</tr>
<tr>
<td>Patients with HCC (group I) vs healthy controls (group III)</td>
<td>0.992</td>
<td>0.949 to 1.000</td>
<td>&lt;0.001*</td>
<td>&gt;367</td>
<td>96.0</td>
<td>98.0</td>
</tr>
<tr>
<td>Patients with HCC (group I) vs patients with liver cirrhosis (group II)</td>
<td>0.748</td>
<td>0.651 to 0.830</td>
<td>&lt;0.001*</td>
<td>&gt;2780</td>
<td>60.0</td>
<td>73.5</td>
</tr>
<tr>
<td>Patients with HCC (group I) vs healthy controls (group III)</td>
<td>0.809</td>
<td>0.719 to 0.881</td>
<td>&lt;0.001*</td>
<td>&gt;2679</td>
<td>66.0</td>
<td>84.0</td>
</tr>
</tbody>
</table>

**Serum AFP**

| Patients with HCC (group I) vs patients with liver cirrhosis (group II) | 0.762  | 0.666 to 0.842 | <0.001* | >188.0  | 74.0        | 75.5        |
| Patients with HCC (group I) vs healthy controls (group III) | 0.983  | 0.934 to 0.998 | <0.001* | >7.9    | 96.0        | 90.0        |

**Combined analysis (serum NRP-1+AFP)**

| Patients with HCC (group I) vs patients with liver cirrhosis (group II) | 0.825  | 0.736 to 0.894 | <0.001* | >64.0   | 100.0       | 100.0       |
| Patients with HCC (group I) vs healthy controls (group III) | 0.994  | 0.952 to 1.000 | <0.001* | –       | 96.0        | 100.0       |

*P<0.05 significant.

**Abbreviations**: AFP, alpha-fetoprotein; ANG-2, angiopoietin-2; AUC, area under the curve; HCC, hepatocellular carcinoma; NRP-1, neuropilin-1; ROC, receiver operating characteristic.

and 50 apparently normal healthy individuals comparable in age and sex as the control group.

This study showed that serum NRP-1 levels were significantly higher in patients with HCC compared with the healthy subjects and patients with cirrhosis. Our results are in agreement with Lin et al.'s study,34 and Abugabal et al.,35 who reported that serum NRP-1 levels in patients with HCC were significantly higher than that of patients with cirrhosis and healthy control subjects. Moreover, our study revealed that patients with cirrhosis have significantly increased serum NRP-1 levels compared with healthy subjects. The same findings were reported by Lin et al.’s study.34

In addition, patients with HCC showed significantly higher serum ANG-2 levels than patients with cirrhosis and healthy subjects in the present study. In agreement with our findings, Chen et al.,33 Scholz et al.,36 and Sharma et al.37 reported that serum ANG-2 levels in patients with HCC were significantly higher than that in patients with cirrhosis and healthy subjects. However, Pestana et al.38 reported no difference in serum ANG-2 levels between HCC and cirrhosis. Moreover, no significant differences in serum ANG-2 levels between patients with cirrhosis and healthy controls were observed in this study, as also reported by Sharma et al.’s study.37 Contradictory to our results, Chen et al.,33 Scholz et al.,36 and Pestana et al.38 reported statistically significant differences in serum ANG-2 levels between patients with cirrhosis and healthy subjects.

Among the demographic and biochemical variables, NRP-1, ANG-2, and AFP were significantly associated with HCC development in univariate analysis. However, NRP-1 and AFP were the only independent factors for HCC development in the multivariate logistic regression analysis and correlated positively with each other. These results highlight the value of NRP-1 compared with ANG-2 as an independent factor of HCC. In the same context, Lin et al.34 reported a significant positive correlation between serum levels of NRP-1 and AFP. Pestana et al.38 and Chen et al.33 reported that serum ANG-2 levels were positively correlated with AFP.

In this study, we found that elevated serum NRP-1 and ANG-2 levels were detected in patients with HCC with increased tumor numbers >3, tumor size ≥5 cm, advanced tumor BCLC stages (B and C), and the presence of vascular invasion and distant metastasis. Similar findings were reported by many previous studies. For instance, Abugabal et al.35 detected that higher plasma levels of NRP-1 were significantly associated with advanced BCLC stages and the presence of vascular invasion. Lin et al.34 also reported that high serum NRP-1 levels were significantly associated with advanced HCC stages. Moreover, Zhang et al.30 reported that high NRP-1 expression was significantly associated with portal vein invasion, and Chen et al.31 reported that TNM (tumor, node, metastasis) staging and vascular...
and portal vein invasion are significantly associated with NRP-1 expression. Functionally, Bergé and his colleagues\(^3\) reported that NRP-1 expression in the liver of transgenic HCC mice is increased with disease progression, in both vascular and tumor compartments. On the other hand, Pestana \textit{et al}\(^{38}\) reported that higher ANG-2 levels were significantly associated with vascular invasion and high TNM stages. Also, Diaz-Sanchez \textit{et al}\(^{40}\) found that ANG-2 levels were correlated with vascular invasion and advanced BCLC stages. Moreover, Kuboki \textit{et al}\(^{41}\) demonstrated that high ANG-2 levels in the hepatic vein significantly were correlated with portal vein invasion. Furthermore, Llovet and his colleagues\(^{42}\) reported that baseline concentration of ANG-2 positively correlated with variables associated with poor outcomes in HCC, such as vascular invasion and distant metastasis. On the contrary, Sharma \textit{et al}\(^{37}\) reported that serum levels of ANG-2 had no association with portal vein invasion, tumor number, or tumor localization.

Unfortunately, the majority of the aforementioned studies have assessed NRP-1 expression in liver tissues that require

\[\text{Figure 2}\] Pearson correlation analyses between the studied serological markers (NRP-1, ANG-2, and AFP). (A) Positive significant correlation between serum AFP and NRP-1 (r=0.781, p<0.001); (B) positive significant correlation between serum AFP and ANG-2 (r=0.519, p<0.001); and (C) positive significant correlation between serum NRP-1 and ANG-2 (r=0.606, p<0.001). AFP, alpha-fetoprotein; ANG-2, angiopoietin-2; NRP-1, neuropilin-1.

\[\text{Figure 3}\] Distribution levels of serum NRP-1 and ANG-2 in patients with HCC showed significantly higher levels of serum NRP-1 and ANG-2 levels in (A, F) patients with HCC with BCLC stages B and C than those with BCLC stages 0 and A; (B, G) patients with HCC with tumor numbers >3 compared with those with tumor numbers ≤3 (p<0.001, p<0.001); (C, H) patients with HCC with tumor size ≥5 compared with those with tumor size <5 (p=0.001, p<0.032); (D, I) patients with HCC with vascular invasion compared with those with no vascular invasion (p=0.001, p<0.001); and (E, J) patients with HCC with distant metastasis compared with those with no distant metastasis (p=0.008, p<0.033); *p<0.05, **p<0.01, ***p<0.001. ANG-2, angiopoietin-2; BCLC, Barcelona Clinic Liver Cancer; HCC, hepatocellular carcinoma; NRP-1, neuropilin-1.
a liver biopsy, which is not applicable in the majority of patients with HCC. Hence, by estimating these markers in the serum, our study presents these markers as applicable non-invasive screening facility for HCC. However, the results of this study should be interpreted with caution and lead time bias should be considered when applying these biomarkers for screening purposes in HCC. The HCC cases included in this study were diagnosed from those seeking medical advice due to clinical manifestations of HCC; however, there is a period of time before the onset of clinical manifestations in which HCC has developed with no clinical manifestations, called the detectable preclinical phase of the disease (DPCD). The efficacy of these biomarkers may be changed whenever estimated during this period. Thus, further studies are required to investigate the efficacy of these biomarkers during DPCD.

In this study, the diagnostic efficacy of both angiogenic markers has been assessed using the ROC curve analysis. The AUC-ROC indicated that the diagnostic efficacy of NRP-1 is higher than AFP in differentiating HCC from patients with cirrhosis and healthy individuals. Similar findings were reported by Lin et al., who detected that the AUC of NRP-1 was 0.971 and higher than that of AFP (AUC=0.862). On the contrary, ANG-2 had a lower diagnostic efficacy than AFP in discriminating HCC from patients with cirrhosis and healthy subjects. However, Chen et al. revealed some contradictions in this regard and reported that the AUC for ANG-2 was 0.924 and better than that of AFP (AUC=0.902). For the first time, the diagnostic accuracy of both markers was compared in this study and the results revealed a higher diagnostic efficacy for NRP-1 over ANG-2 in discriminating HCC from patients with cirrhosis (0.801 vs 0.748, p=0.250) and healthy subjects (0.922 vs 0.809, P<0.001). Moreover, combined analysis of both NRP-1 and AFP was performed and revealed that its diagnostic power is slightly better than NRP-1 or AFP alone in differentiating HCC from healthy subjects, with much more significant difference detected with ANG-2.

There are some limitations to this study. First, it included a relatively small number of subjects; however, the G power analysis indicated the adequacy of our sample. Second, the measurement of NRP-1 and ANG-2 was performed by ELISA, which is a non-standardized method; however, the precision and sensitivity of the kits were acceptable. Third, lead time bias should be considered whenever applying these results for screening purposes in HCC. Fourth, the subjects were selected from a single center and certain criteria were applied, such as HCV infection, which may confer ascertainment bias. Therefore, it is recommended to extend this research to a large cohort and investigate the efficacy of these biomarkers during DPCD. In conclusion, the results of this study suggested that NRP-1 is a better marker for the diagnosis of HCC than AFP and ANG-2, and it is feasible to be estimated in combination with AFP to enhance its diagnostic power. Also, higher levels of NRP-1 and ANG-2 were associated with advanced HCC tumor characteristics.

Contributors MTAG, RAE: study design, methodology, laboratory investigations, original draft writing. RAE: methodology, resources, patient selection and clinical evaluation, original draft writing. MHE, MMM: methodology, clinical evaluation. SAD: methodology, resources, patient selection and clinical evaluation. RLY: data analysis. All authors revised and approved the final manuscript version.

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