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Contents

Thrice Monthly Volume 10 Number 26 September 16, 2022

REVIEW

- 9180** Assisting individuals with diabetes in the COVID-19 pandemic period: Examining the role of religious factors and faith communities

Eseadi C, Ossai OV, Onyishi CN, Ilechukwu LC

- 9192** Role of octreotide in small bowel bleeding

Khedr A, Mahmoud EE, Attallah N, Mir M, Boike S, Rauf I, Jama AB, Mushtaq H, Surani S, Khan SA

MINIREVIEWS

- 9207** Internet of things-based health monitoring system for early detection of cardiovascular events during COVID-19 pandemic

Dami S

- 9219** Convergence mechanism of mindfulness intervention in treating attention deficit hyperactivity disorder: Clues from current evidence

Xu XP, Wang W, Wan S, Xiao CF

- 9228** Clinical presentation, management, screening and surveillance for colorectal cancer during the COVID-19 pandemic

Akbulut S, Hargura AS, Garzali IU, Aloun A, Colak C

- 9241** Early diagnostic value of liver stiffness measurement in hepatic sinusoidal obstruction syndrome induced by hematopoietic stem cell transplantation

Tan YW, Shi YC

ORIGINAL ARTICLE

Case Control Study

- 9254** Local inflammatory response to gastroesophageal reflux: Association of gene expression of inflammatory cytokines with esophageal multichannel intraluminal impedance-pH data

Morozov S, Sentsova T

Retrospective Study

- 9264** Evaluation of high-risk factors and the diagnostic value of alpha-fetoprotein in the stratification of primary liver cancer

Jiao HB, Wang W, Guo MN, Su YL, Pang DQ, Wang BL, Shi J, Wu JH

- 9276** One-half layer pancreaticojejunostomy with the rear wall of the pancreas reinforced: A valuable anastomosis technique

Wei JP, Tai S, Su ZL

- 9285** Development and validation of an epithelial-mesenchymal transition-related gene signature for predicting prognosis

Zhou DH, Du QC, Fu Z, Wang XY, Zhou L, Wang J, Hu CK, Liu S, Li JM, Ma ML, Yu H

Observational Study

- 9303** Incidence and risk factor analysis for swelling after apical microsurgery

Bi C, Xia SQ, Zhu YC, Lian XZ, Hu LJ, Rao CX, Jin HB, Shang XD, Jin FF, Li JY, Zheng P, Wang SH

CASE REPORT

- 9310** Acute carotid stent thrombosis: A case report and literature review

Zhang JB, Fan XQ, Chen J, Liu P, Ye ZD

- 9318** Congenital ovarian anomaly manifesting as extra tissue connection between the two ovaries: A case report

Choi MG, Kim JW, Kim YH, Kim AM, Kim TY, Ryu HK

- 9323** Cefoperazone-sulbactam and ornidazole for *Gardnerella vaginalis* bloodstream infection after cesarean section: A case report

Mu Y, Li JJ, Wu X, Zhou XF, Tang L, Zhou Q

- 9332** Early-onset ophthalmoplegia, cervical dyskinesia, and lower extremity weakness due to partial deletion of chromosome 16: A case report

Xu M, Jiang J, He Y, Gu WY, Jin B

- 9340** Posterior mediastinal extralobar pulmonary sequestration misdiagnosed as a neurogenic tumor: A case report

Jin HJ, Yu Y, He W, Han Y

- 9348** Unexpected difficult airway due to severe upper tracheal distortion: A case report

Zhou JW, Wang CG, Chen G, Zhou YF, Ding JF, Zhang JW

- 9354** Special epithelioid trophoblastic tumor: A case report

Wang YN, Dong Y, Wang L, Chen YH, Hu HY, Guo J, Sun L

- 9361** Intrahepatic multicystic biliary hamartoma: A case report

Wang CY, Shi FY, Huang WF, Tang Y, Li T, He GL

- 9368** ST-segment elevation myocardial infarction in Kawasaki disease: A case report and review of literature

Lee J, Seo J, Shin YH, Jang AY, Suh SY

- 9378** Bilateral hypocalcaemic cataracts due to idiopathic parathyroid insufficiency: A case report

Li Y

- 9384** Single organ hepatic artery vasculitis as an unusual cause of epigastric pain: A case report

Kaviani R, Farrell J, Dehghan N, Moosavi S

- 9390** Congenital lipoid adrenal hyperplasia with Graves' disease: A case report

Wang YJ, Liu C, Xing C, Zhang L, Xu WF, Wang HY, Wang FT

- 9398** Cytokine release syndrome complicated with rhabdomyolysis after chimeric antigen receptor T-cell therapy: A case report
Zhang L, Chen W, Wang XM, Zhang SQ
- 9404** Antiphospholipid syndrome with renal and splenic infarction after blunt trauma: A case report
Lee NA, Jeong ES, Jang HS, Park YC, Kang JH, Kim JC, Jo YG
- 9411** Uncontrolled high blood pressure under total intravenous anesthesia with propofol and remifentanyl: A case report
Jang MJ, Kim JH, Jeong HJ
- 9417** Noncirrhotic portal hypertension due to peripheral T-cell lymphoma, not otherwise specified: A case report
Wu MM, Fu WJ, Wu J, Zhu LL, Niu T, Yang R, Yao J, Lu Q, Liao XY
- 9428** Resumption of school after lockdown in COVID-19 pandemic: Three case reports
Wang KJ, Cao Y, Gao CY, Song ZQ, Zeng M, Gong HL, Wen J, Xiao S
- 9434** Complete recovery from segmental zoster paresis confirmed by magnetic resonance imaging: A case report
Park J, Lee W, Lim Y
- 9440** Imaging findings of immunoglobulin G4-related hypophysitis: A case report
Lv K, Cao X, Geng DY, Zhang J
- 9447** Systemic lupus erythematosus presenting with progressive massive ascites and CA-125 elevation indicating Tjasma syndrome? A case report
Wang JD, Yang YF, Zhang XF, Huang J
- 9454** Locally advanced cervical rhabdomyosarcoma in adults: A case report
Xu LJ, Cai J, Huang BX, Dong WH
- 9462** Rapid progressive vaccine-induced immune thrombotic thrombocytopenia with cerebral venous thrombosis after ChAdOx1 nCoV-19 (AZD1222) vaccination: A case report
Jiang SK, Chen WL, Chien C, Pan CS, Tsai ST
- 9470** Burkitt-like lymphoma with 11q aberration confirmed by needle biopsy of the liver: A case report
Yang HJ, Wang ZM
- 9478** Common carotid artery thrombosis and malignant middle cerebral artery infarction following ovarian hyperstimulation syndrome: A case report
Xu YT, Yin QQ, Guo ZR
- 9484** Postoperative radiotherapy for thymus salivary gland carcinoma: A case report
Deng R, Li NJ, Bai LL, Nie SH, Sun XW, Wang YS
- 9493** Follicular carcinoma of the thyroid with a single metastatic lesion in the lumbar spine: A case report
Chen YK, Chen YC, Lin WX, Zheng JH, Liu YY, Zou J, Cai JH, Ji ZQ, Chen LZ, Li ZY, Chen YX

- 9502** Guillain-Barré syndrome and hemophagocytic syndrome heralding the diagnosis of diffuse large B cell lymphoma: A case report
Zhou QL, Li ZK, Xu F, Liang XG, Wang XB, Su J, Tang YF
- 9510** Intravitreal injection of conbercept for bullous retinal detachment: A case report
Xiang XL, Cao YH, Jiang TW, Huang ZR
- 9518** Supratentorial hemangioblastoma at the anterior skull base: A case report
Xu ST, Cao X, Yin XY, Zhang JY, Nan J, Zhang J

META-ANALYSIS

- 9524** Certain sulfonylurea drugs increase serum free fatty acid in diabetic patients: A systematic review and meta-analysis
Yu M, Feng XY, Yao S, Wang C, Yang P

LETTER TO THE EDITOR

- 9536** Glucose substrate in the hydrogen breath test for gut microbiota determination: A recommended noninvasive test
Xie QQ, Wang JF, Zhang YF, Xu DH, Zhou B, Li TH, Li ZP
- 9539** A rare cause of acute abdomen after a Good Friday
Pante L, Brito LG, Franciscatto M, Brambilla E, Soldera J
- 9542** Obesity is associated with colitis in women but not necessarily causal relationship
Shen W, He LP, Zhou LL
- 9545** Risk stratification of primary liver cancer
Tan YW

ABOUT COVER

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Risk stratification of primary liver cancer

You-Wen Tan

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Abstract

The risk stratification of primary liver cancer (PLC) discussed in a review of viral hepatitis and PLC could lead to misunderstandings by readers. For example, a single study or a small number of studies cannot comprehensively summarize the risk factors of PLC, is not included in the family history of liver cancer, and chronic hepatitis D is listed as a medium risk factor for the development of PLC. Currently, PLC prediction models with good clinical validation values have been applied clinically, such as the Toronto hepatocellular carcinoma risk index, REACH-B model, and PAGE-B model. Therefore, the Chinese, together with several research societies, have formulated the "Guideline for stratified screening and surveillance of primary liver cancer (2020 edition)." This guideline outlines PLC screening in at-risk populations, both in hospitals and communities. It is recommended to stratify the at-risk population into four risk levels: low-, intermediate-, high-, and extremely high-risk. This is highly recommended and applied in clinical practice.

Key Words: Risk factors; Model; Primary liver cancer; Hepatocellular carcinoma

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Core Tip: Primary liver cancer (PLC) prediction models with good clinical validation values have been applied clinically, such as the Toronto hepatocellular carcinoma risk index, REACH-B model, and PAGE-B model. Therefore, the Chinese, together with several research societies, have formulated the "Guideline for stratified screening and surveillance of primary liver cancer (2020 edition)." This guideline outlines PLC screening in at-risk populations, both in hospitals and communities. It is recommended to stratify the at-risk population into four risk levels: low-, intermediate-, high-, and extremely high-risk. This is highly recommended and applied in clinical practice.

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TO THE EDITOR

I have read an article with title: "Viral hepatitis and hepatocellular carcinoma: From molecular pathways to the role of clinical surveillance and antiviral treatment" with great interest[1]. This is a very good systematic review which fully discusses the etiology, risk factors, and pathogenesis of primary liver cancer (PLC). High-, medium-, and low-risk factors of PLC were analyzed. However, the term "risk factors for the development of PLC" in this article could lead to misunderstandings by readers.

First, a single study or a small number of studies cannot comprehensively summarize the risk factors of PLC. Perhaps a meta-analysis is more reasonable. A meta-analysis of chronic hepatitis B hepatocellular carcinoma (HCC) from China included 3165 cases and 10896 controls from 27 studies[2]. The results showed that, from the currently available evidence, Chinese people with a high viral load, not treated with antiviral treatment, with a family history of liver cancer, of male sex, and with chronic hepatitis B virus (HBV) infection are at risk of developing HCC. In another meta-analysis of risk factors for nonalcoholic fatty liver disease (NAFLD) and HCC, 18 studies involving 470,404 patients were included. In NAFLD patients before cirrhosis, the incidence of HCC was 0.03/100 person years. In patients with liver cirrhosis, the incidence rate was 3.78/100 person years[3]. However, it still needs to be noted that the risk factors of PLC that can be seen at present are for a specific disease, and there is still a lack of meta-analysis on the risk factors of all PLC. Many studies have analyzed PLC risk factors. Among them, the studies that have been highly recognized mentioned the Toronto HCC risk index (THRI) which takes into consideration five variables, including age, gender, platelet count, and the cause of liver cirrhosis (0, 0, 36, 54, and 97 for autoimmune liver disease, hepatitis C with sustained virological response [SVR], other liver diseases, fatty hepatitis, untreated chronic hepatitis C or chronic hepatitis B, respectively). A total of 366 points of the THRI model were constructed. The 5-year cumulative incidence rates of HCC in the low- (< 120 points), medium- (120-240 points), and high- (> 240 points) risk groups were 1.2% and 4.5%, 4%, respectively, and 15, and the 10-year cumulative incidence rates of HCC were 3%, 10%, and 32%, respectively[4].

The THRI model was also validated in China. For patients with cirrhosis classified into low-, medium-, and high-risk groups, the 5-year cumulative incidence rates of HCC were 0%, 13%, and 34%, respectively, and the area under the receiver operating characteristic curve (AUC) of the prediction model is 0.707[5]. The significance of the THRI model is that even in patients with liver cirrhosis who are at high risk of HCC, it is still necessary to conduct further risk stratification to distinguish the extremely high-risk population of HCC.

HBV infection is the main cause of HCC. Even without cirrhosis, there is only 6.5% chance of 2%-9.5% of HBV infected individuals to progress to HCC in their lifetime. Some scholars in China have built the REACH-B model with a total of 17 points, including sex, age, alanine aminotransferase (ALT) HBeAg status, and HBV DNA level, aiming at the risk stratification of HBV-infected individuals who have not received antiviral treatment; scores of 0-5, 6-11, and 12-17 are low, medium, and high risk, respectively [6]. The incidence of HCC in the lowest (0) and highest (17) scores within 3, 5, and 10 years is 0.0% and 23.6%, 0.0% and 47.4% and 0.0% and 81.6%, respectively.

HBV remains a risk factor for HCC after antiviral treatment. Based on the 5-year results of oral antiviral treatment with entecavir or tenofovir in 1,815 CHB patients in Europe, the PAGE-B model with three parameters, including age, sex, and platelet count, and a total score of 25 points was constructed to evaluate the risk of HCC in patients with HBV infection after antiviral treatment[7]. According to the score, the patients were divided into three groups: low-risk (0-9 points), medium-risk (10-17 points), and high-risk (18-25 points). The 5-year cumulative incidence of HCC in the low-, moderate-, and high-risk groups was 0%, 0%-4%, and 16%-17%, respectively. The Korean cohort verified that the PAGE-B model was equally effective in Asian populations, and the AUC for 5-year HCC predictive power was 0.77[8].

The PAGE-B team recently added two parameters, age and liver stiffness measurement; thus, the maximum total score is now 15 points and divided into low-risk (< 5 points), medium-risk (6-10 points), and high-risk (> 11 points) levels. The cumulative incidence of HCC in 12 years was 0.0%, 4.0%, 0.0%, and 13.8%. In conclusion, the 5-year cumulative incidence rate of HCC was still 5.5% in patients with chronic HBV infection treated with antiviral therapy, especially those classified in the high-risk group (7%-8.4%). However, the negative predictive rate of HCC in 5 and 12 years in the low-risk group is as high as 97.5%-100%[6,8]. In untreated individuals infected with HCV, serum HCV RNA, ALT, and HCV genotype 1b were independent predictors of HCC[9,10].

Risk prediction models have a long history in predicting HCC incidence rate in CHB patients. Currently, approved models are as follows: CU-HCC, GAG-HCC, Page-B, mPAGE-B, REACH-B and mREACH-B[11,12]. A meta-analysis used six models to perform AUC validation on 22 studies published between 2011 and 2020[13]. The AUC values of the six models ranged from 0.715 to 0.778. In

Table 1 Identification and stratification patients with high risk for liver cancer

Estimated annual incidence of HCC (%)	Distinguishing feature	Risk stratification model of liver cancer
Low risk of liver cancer (< 1)	(1) HBV infected patients in immune tolerance period; (2) HBV or HCV related chronic hepatitis with SVR acquired by antiviral therapy; and (3) ALT, normal PLCTelet, non viral liver disease	HBsAg positive, REACH-B score ≥ 5 (no antiviral treatment), or PAGE-B score ≤ 9
Moderate risk of liver cancer (1-3)	(1) HBV or HCV related chronic hepatitis of LLV without antiviral treatment or after antiviral treatment, aged < 40 yr; HBV or HCV related cirrhosis with SVR obtained by antiviral therapy; and (2) non viral cirrhosis with normal ALT or chronic non viral hepatitis with abnormal ALT	(1) HBsAg positive, REACH-B score 6-11 (no antiviral treatment) or PAGE-B score 10-17; and (2) THRI score of cirrhotic patients ≤ 240
High risk of liver cancer (> 3 and < 6)	(1) HBV or HCV related cirrhosis of LLV without or after antiviral treatment; (2) non-viral cirrhosis patients with diabetes or family history of liver cancer with first-degree relatives (B1); and (3) men, age > 40 yr old; women, age > 50 yr; No antiviral therapy for HBV/HCV related chronic hepatitis	(1) HBsAg positive, REACH-B score > 12 (no antiviral treatment) or PAGE-B score ≤ 18 ; and (2) THRI score of cirrhotic patients > 240
Extremely high risk for liver cancer (> 6)	(1) Hepatic nodule (1-2 cm) in abdominal US examination or LGDN and HGDN in pathology; (2) HBV and HCV related cirrhotic nodules (< 1 cm); and (3) synergistic risk factors such as no antiviral treatment, HBV or HCV related liver cirrhosis with diabetes or family history of liver cancer in first-degree relatives after treatment	–

ALT: Alanine aminotransferase; HCC: Hepatocellular carcinoma; SVR: Sustained virological response; THRI: Toronto hepatocellular carcinoma risk index.

the antiviral treatment subgroup, the AUC values of mREACH-B, GAG-HCC and mPAGE-B were 0.785, 0.760 and 0.778, respectively. In the subgroup of liver cirrhosis, the recognition performance of all models is very poor (AUC < 0.7). The clinical application of these models can improve patients' prognosis and aid them in making informed decisions about treatment. However, these models were derived from different cohorts with or without antiviral treatment and affected by many factors. There are no guidelines for publishing the same standardized guide to predict the risk of HCC among CHB patients[14-16].

Second, among the risk factors, there was no family history of PLC. In the HBV- or HCV-infected population, first-degree relatives with a family history of PLC significantly increased the risk of HCC [17-19]. Scholars in China have followed up 22472 residents and a total of 362268 people/year, and 374 cases of HCC have been detected. HBV patients with a family history of HCC were 2.5 times more likely to develop HCC than those without a family history[19]. A similar cohort of 7933 non-PLC patients and 201 patients with PLC from China also showed that the risk of HCC was 2.76 higher (95% CI: 1.88-4.05) in individuals without HBV infection but with a family history of PLC, but in the population with HBV positive and family history of primary HCC, the risk of HCC was 41.34 (95%CI: 23.69-72.12)[17]. Therefore, first-degree relatives infected with HBV and HCV with a family history of liver cancer had a significantly increased risk of HCC in at all stages, which requires special attention[18].

Third, untreated chronic hepatitis D virus infection was classified as a moderate risk factor in the analysis of HCC risk factors, with an OR of 3.9. HDV is a defective virus dependent on HBV infection. Chronic hepatitis D inevitably overlaps with chronic hepatitis B. As described in the article, chronic hepatitis D is considered the most serious form of chronic viral hepatitis, leading to rapid progression of liver cirrhosis and higher mortality. More than 10% of patients infected with chronic hepatitis D virus develop cirrhosis within 5 years of infection, and more than 80% of patients suffer from liver cirrhosis decompensation within 30 years[20]. Recent cohort studies have found that compared to patients with a single HBV infection, the risk of patients with HDV infection is nine times higher, although it is generally believed that HDV does not represent a major risk factor for the development of HCC[21,22]. In addition, persistent hepatitis D virus replication has been a risk factor for liver disease progression to cirrhosis and HCC[23]. A European study on the collaborative action of viral hepatitis showed that the incidence of HCC in patients with anti-hepatitis D virus-positive cirrhosis increased by 3.2 times compared with those with negative hepatitis[21]. Other recent studies have shown that patients with HDV/HBV co-infection have an increased risk of HCC compared to patients with HBV infection alone (adjusted HR = 9.30)[24]. A large study estimated that the risk of HCC in patients with chronic hepatitis D virus infection (RR = 3.9) was significantly higher than that in patients with a single HBV infection [25].

Therefore, the Chinese, together with several research societies have formulated the "Guideline for stratified screening and surveillance of primary liver cancer (2020 edition)"[26]. China has a high incidence of PLC, and chronic HBV infection is still the greatest risk factor for HCC. According to the high-risk factors, the guidelines for screening at-risk populations were divided into low-, intermediate-, high-, and extremely high-risk groups (Table 1) and propose monitoring and timing based on ultrasound findings and alpha fetoprotein level. Abdominal ultrasonography combined with serum

alpha-fetoprotein examination (routine surveillance) is recommended every 6 mo for patients at high risk of PLC. Routine surveillance every 3 mo and enhanced computer tomography/ magnetic resonance imaging examination every 6-12 mo are recommended for patients at an extremely high risk of PLC. The surveillance interval can be extended every 1 year or longer for those with low-risk or intermediate-risk PLC because their annual incidence of PLC is very low. However, the cost-effectiveness of these recommendations remains to be evaluated. We believe that the stratified analysis of risk factors and screening recommendations for PLC in China can be successfully implemented by doctors to aid the early diagnosis and treatment of PLC. Of course, large sample verification and observation are required.

FOOTNOTES

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