FOLFIRI Combined with Bevacizumab as First-Line Treatment for Metastatic Colorectal Cancer Patients with Hyperbilirubinemia after UGT1A1 Genotyping

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Introduction

The current standard first-line treatment for metastatic colorectal cancer (mCRC) includes FOLFOX (oxaliplatin plus infusional 5-fluorouracil and leucovorin) or FOLFIRI (5-fluorouracil, leucovorin, and irinotecan) plus biological agents. Irinotecan is a camptothecin derivative that exerts cytotoxic effects on cancer cells by inhibiting topoisomerase I, and FOLFIRI has been proven to be effective for the first-line treatment of patients with mCRC [1, 2]. However, to date, there is no standard protocol for mCRC patients with bilirubin >2 mg/dl. Relevant studies of the clinical benefits of chemotherapy for patients with severe hyperbilirubinemia and jaundice are lacking [3].

UGT1A1*28 is considered the main predictor of toxicity in irinotecan-treated mCRC. UGT1A1 genotyping, established as a predictive marker for irinotecan-induced severe neutropenia/diarrhea, has been approved by the Food and Drug Administration of the USA [4]. However, there is no predictive marker for irinotecan-based che-
motherapy for the treatment of mCRC with severe liver dysfunction and prolonged hyperbilirubinemia. Hence, we report a case of severe hyperbilirubinemia with extensive liver metastasis from mCRC with FOLFIRI plus bevacizumab as first-line treatment.

Case Report

A 46-year-old male was referred to our hospital and was hospitalized with an initial presentation of a 3-month history of constipation, abdominal distension, poor appetite, body weight loss, and jaundice. Colonoscopy revealed a nearly 4/5 circumferential obstructing ulcerative tumor in the rectosigmoid colon at approximately 12 cm from the anal verge, and pathological examination of a colonoscopic biopsy showed adenocarcinoma. No specific family history or past history was identified. Physical findings revealed icteric sclera and palpable hepatomegaly from the epigastrium to the right upper abdomen. Computed tomography (CT) of the abdomen and pelvis showed a huge rectosigmoid colon tumor approximately 6 cm in maximal diameter with more than 20 hepatic metastases (>70% liver replacement). The patient was finally diagnosed with rectosigmoid colon cancer with multiple liver metastases, UICC stage IV, based on the above imaging study. Abnormal laboratory findings included a total bilirubin level of 5.94 mg/dl, an alanine aminotransferase level of 276 IU/l, an aspartate aminotransferase level of 125 IU/l, and a carcinoembryonic antigen level of 12,587 ng/ml. Genomic DNA, extracted from the peripheral blood, was subjected to PCR sequencing for genotyping of the promoter region of the $UGT1A1$ gene. The (TA)$_n$ repeat of this gene showed 6/6 repeats. Due to poor liver function, jaundice, and the results of genotyping, the recommended dose of irinotecan ($180 \text{ mg/m}^2$) was underdosed in this mCRC patient with $UGT1A1$ $6/6$ or $UGT1A1$ $6/7$ gene polymorphism. The patient was treated with a regimen of FOLFIRI at an initial dosage of 120 mg/m$^2$ administered over 2 h on day 1 followed by 5-fluorouracil (400 mg/m$^2$ as an i.v. bolus infusion followed by 2,400 mg/m$^2$ i.v. infusion over a 46-hour period plus bevacizumab 5 mg/kg, repeated every 2 weeks). After 4 cycles of therapy, with the irinotecan dose being escalated from 120 to 180 mg/m$^2$, the bilirubin level of the patient significantly decreased to 0.78 mg/dl, the alanine aminotransferase level decreased to 27 IU/l, the aspartate aminotransferase level decreased to 27 IU/l, the hepatic metastases were noted on the CT scan. The arrows indicate hepatic metastases in a+b; colon tumor in c+d.
noscopy showed that the primary tumor had prominently shrunk (fig. 2). The patient received 12 cycles of FOLFIRI plus bevacizumab with clinical benefit. He survived 9 months from the time of first-line treatment with FOLFIRI combined with bevacizumab.

Discussion

This patient with rectosigmoid cancer and prolonged grade 4 hyperbilirubinemia was successfully treated with a regimen of FOLFIRI plus bevacizumab. After 4 cycles of this treatment, the therapy was well tolerated, with normalization of liver functions and predominant tumor shrinkage. We believe that, after excluding the possibility of obstructive jaundice, the use of novel genomic DNA analysis of the blood as in our current study can result in more favorable outcomes for such patients. mCRC patients with severe hyperbilirubinemia and liver dysfunction in which pharmacokinetic clearance is impaired often have a limited performance status and are at increased risk for chemotherapy-related complications [5, 6]. In mCRC with liver metastasis and hyperbilirubinemia, there could be tumor burden-related (nonobstructive) or obstructive jaundice, and we should differentiate between them using laboratory examinations and imaging studies. The severe hyperbilirubinemia noted in these patients is more reflective of the underlying pathogenesis. Bilirubin data are currently used to exclude certain patients from receiving therapy (i.e. those with a serum bilirubin concentration of >2 mg/dl should not be treated with irinotecan) [5, 7].

Irinotecan is converted into 7-ethyl-10-hydroxycamptothecin (SN-38) by a carboxylesterase and finally metabolized through the uridine diphosphate glucuronosyl transferase enzyme (UGT), predominantly by the UGT1A1 isoenzyme. Recently, Marcuello et al. [8] conducted a genotype-directed dose-finding study of irinotecan in combination with fluorouracil/leucovorin as first-line treatment for advanced CRC. Patients with homozygous polymorphism of the UGT1A1 promoter (UGT1A1*28) were more frequently associated with severe toxicity following irinotecan treatment. The genotype of the promoter polymorphism of the UGT1A1 gene in our patient was either UGT1A1 6/6 or UGT1A1 6/7. Individuals who are homozygous for the UGT1A1*28 allele are at increased risk for toxicity following the initiation of FOLFIRI treatment, and a reduced initial dose should be considered for patients known to be homozygous for the UGT1A1*28 allele (7 repeats). Heterozygous patients (carriers of one variant allele and one wild-type allele which results in intermediate UGT1A1 activity) may be at increased risk for toxicity; however, clinical results have been variable and such patients have been shown to tolerate normal starting doses [9]. Furthermore, the possible role of bevacizumab as a protective agent against the toxicity of irinotecan in these patients might need to be investigated. The addition of bevacizumab has been previously reported to improve the pathological response and protect against hepatic injury in patients treated with oxaliplatin-based chemotherapy for colorectal liver metastases [10].

Conclusion

The FOLFIRI regimen was successfully used in our patient with liver function abnormalities caused by mCRC without concerns regarding toxicity.
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