Use of the Informatics for Integrating Biology and the Bedside (i2b2) Population to Test Serum Bilirubin Levels and Risk for Inflammatory Bowl Diseases and the Involvement of Uridine Glucuronosyltransferase Genes

Carla J. Gallagher Lincoln University Lincoln University, PA, USA cgallagher@lincoln.edu

ABSTRACT

Chronic inflammation associated with inflammatory bowel disease (IBD) results in increased oxidative stress that damages the colonic microenvironment. A low level of serum bilirubin, an endogenous antioxidant, has been associated with increased risk for Crohn's disease (CD), but no study has tested another common IBD ulcerative colitis (UC). Bilirubin is metabolized in the liver by uridine glucuronosyltransferase 1A1 (UGT1A1) exclusively. Genetic variants cause functional changes in UGT1A1 which result in hyperbilirubinemia, which can be toxic to tissues if untreated and results in a characteristic jaundiced appearance. Approximately 10% of the Caucasian population is homozygous for the microsatellite polymorphism UGT1A1*28, which results in increased total serum bilirubin levels due to reduced transcriptional efficiency of UGT1A1 and an overall 70% reduction in UGT1A1 enzymatic activity. The aim of this study was to examine whether bilirubin levels are associated with the risk for ulcerative colitis (UC). Using the Informatics for Integrating Biology and the Bedside (i2b2), a large case-control population was identified from a single tertiary care center, Penn State Hershey Medical Center (PSU). Similarly, a validation cohort was identified at Virginia Commonwealth University Medical Center. Logistic regression analysis was performed to determine the risk of developing UC with lower concentrations of serum bilirubin.

Permission to make digital or hard copies of part or all of this work for personal or classroom use is granted without fee provided that copies are not made or distributed for profit or commercial advantage and that copies bear this notice and the full citation on the first page. Copyrights for third-party components of this work must be honored. For all other uses, contact the Owner/Author.

ACM-BCB'18, August 29-September 1, 2018, Washington, DC, USA. © 2018 Copyright is held by the owner/author(s). ACM ISBN 978-1-4503-5794-4/18/08. DOI: https://doi.org/10.1145/3233547.3233638 From the PSU cohort, a subset of terminal ileum tissue was obtained at the time of surgical resection to analyze UGT1A1 gene expression (which encodes the enzyme responsible for bilirubin metabolism). Similar to CD patients, UC patients also demonstrated reduced levels of total serum bilirubin. Upon segregating serum bilirubin levels into quartiles, risk of UC increased with reduced concentrations of serum bilirubin. These results were confirmed in our validation cohort. UGT1A1 gene expression was up-regulated in the terminal ileum of a subset of UC patients. Lower levels of the antioxidant bilirubin may reduce the capability of UC patients to remove reactive oxygen species leading to an increase in intestinal injury. One potential explanation for these lower bilirubin levels may be up-regulation of UGT1A1 gene expression, which encodes the only enzyme involved in conjugating bilirubin. Therapeutics that reduce oxidative stress may be beneficial for these patients.

CCS CONCEPTS

Medical Informatics

KEYWORDS

health informatics, bedside, IBD

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