



High-Risk Features Are Prognostic in dMMR/MSI-H Stage II Colon Cancer

Amr Mohamed ^{1*}, Renjian Jiang², Philip A. Philip³, Maria Diab⁴, Madhusmita Behera⁴, Christina Wu⁴, Olatunji Alese⁴, Walid L. Shaib⁴, Tyra M. Gaines⁴, Glen G. Balch⁵, Bassel F. El-Rayes⁴ and Mehmet Akce⁴

¹ Case Comprehensive Cancer Center, Case Western Reserve University, Cleveland, OH, United States, ² Winship Research Informatics Shared Resource, Winship Cancer Institute, Emory University, Atlanta, GA, United States, ³ Karmanos Cancer Institute, Wayne State University, Detroit, MI, United States, ⁴ Department of Hematology and Medical Oncology, Winship Cancer Institute, Emory University School of Medicine, Atlanta, GA, United States, ⁵ Division of Colorectal Surgery, Department of Surgery, Emory University School of Medicine, Atlanta, GA, United States

OPEN ACCESS

Edited by:

Zhanlong Shen, Peking University People's Hospital, China

Reviewed by:

Stefan Urbanski, University of Calgary, Canada Jun Li, The University of Sydney, Australia

*Correspondence: Amr Mohamed amr.mohamed@uhhospitals.org

Specialty section:

This article was submitted to Gastrointestinal Cancers: Colorectal Cancer, a section of the journal Frontiers in Oncology

Received: 08 August 2021 Accepted: 21 September 2021 Published: 25 October 2021

Citation:

Mohamed A, Jiang R, Philip PA, Diab M, Behera M, Wu C, Alese O, Shaib WL, Gaines TM, Balch GG, El-Rayes BF and Akce M (2021) High-Risk Features Are Prognostic in dMMR/MSI-H Stage II Colon Cancer. Front. Oncol. 11:755113. doi: 10.3389/fonc.2021.755113 **Background:** High-risk features, such as T4 disease, bowel obstruction, poorly/ undifferentiated histology, lymphovascular, perineural invasion, and <12 lymph nodes sampled, indicate poor prognosis and define high-risk stage II disease in proficient mismatch repair stage II colon cancer (CC). The prognostic role of high-risk features in dMMR/MSI-H stage II CC is unknown. Similarly, the role of adjuvant therapy in high-risk stage II CC with dMMR/MSI-H (≥1 high-risk feature) has not been studied in prospective trials. The aim of this analysis of the National Cancer Database is to evaluate the prognostic value of high-risk features in stage II dMMR/MSI-H CC.

Methods: Univariate (UVA) and multivariate (MVA) Cox proportional hazards (Cox-PH) models were built to assess the association between clinical and demographic characteristics and overall survival. Kaplan–Meier survival curves were generated with log-rank tests to evaluate the association between adjuvant chemotherapy in high-risk and low-risk cohorts separately.

Results: A total of 2,293 stage II CC patients have dMMR/MSI-H; of those, 29.5% (n = 676) had high-risk features. The high-risk dMMR/MSI-H patients had worse overall survival [5-year survival and 95%CI, 73.2% (67.3–78.1%) vs. 80.3% (76.7–83.5%), p = 0.0001]. In patients with stage II dMMR/MSI-H CC, the high-risk features were associated with shorter overall survival (OS) along with male sex, positive carcinoembryonic antigen, Charlson–Deyo score >1, and older age. Adjuvant chemotherapy administration was associated with better OS, regardless of the high-risk features in dMMR/MSI-H (log-rank test, p = 0.001) or not (p = 0.0006). When stratified by age, the benefit of chemotherapy was evident only in patients age ≥65 with high-risk features.

Conclusion: High-risk features are prognostic in the setting of dMMR/MSI-H stage II CC. Adjuvant chemotherapy may improve survival specifically in patients \geq 65 years and with high-risk features.

Keywords: high risk, stage II, adjuvant chemotherapy, colon, cancer

INTRODUCTION

Colorectal cancer is the third most common cancer and the second leading cause of cancer-related mortality in the United States. It is estimated that 104,270 new cases of colon cancer (CC) will be diagnosed in 2021 in the US (1). Approximately 28% of patients with CC have stage II disease at presentation (2). The risk stratification of patients with stage II CC is dependent on molecular and clinicopathologic features. A prognostic role of high-risk features, such as T4 disease, bowel obstruction, poorly/ undifferentiated histology, lymphovascular, perineural invasion, and <12 lymph nodes sampled, is well established, and high-risk features increase the risk of cancer recurrence and the benefit from adjuvant therapy in patients with microsatellite stable disease stage II CC (3-5). A subgroup of high-risk patients with stage II CC with T4 disease may have a statistically inferior survival compared to those with stage IIIa tumors (6). Adjuvant chemotherapy improves progression-free survival (PFS) and overall survival (OS) in stages II and III CC (7, 8). The benefit of adjuvant therapy in stage II CC is relatively small, and as such, it is not routinely administered (9). Patient preferences, treatment-related toxicities, and the risk characteristics of the tumor are considered in treatment decisions regarding adjuvant therapy.

Approximately 15% of colorectal cancers (CRCs) are dMMR/ MSI-H, and patients with dMMR/MSI-H colon cancer are more likely to have a stage II disease (10). Mismatch repair (MMR) proteins are nuclear enzymes that bind to areas of abnormal DNA and repair base-base mismatch during cellular proliferation and division (11). Defects in DNA mismatch repair genes (MLH1, MSH2, MSH6, and PMS2) can lead to insertion or deletion of repeating nucleotide sequences in a process known as microsatellite instability (MSI). One third of these dMMR/MSI-H CC cases are inherited, known as Lynch syndrome carriers, and the rest are sporadic. MLH1 is considered the most commonly affected in the sporadic cases, which is more common in older patients and associated with BRAF V600E mutation (12). The microsatellite instability status of a tumor impacts the prognosis and benefit to adjuvant chemotherapy in patients with stage II CC (10-12). Multiple retrospective studies have shown that stage II patients with dMMR/MSI-H CC have a reduced metastatic potential and a more favorable prognosis compared to those with proficient mismatch repair (pMMR) tumors (13-16). In addition, previous retrospective studies of stages II and III colon cancer patients, analyzing data from randomized adjuvant therapy clinical trials, showed that stage II colon cancer patients with dMMR/MSI-H status did not benefit from adjuvant 5-FU-based chemotherapy (3, 10, 12, 17). Furthermore, Sargent et al. showed a decrease in overall survival (hazard ratio, 2.95; 95%CI, 1.02 to 8.54; p = 0.04) in dMMR/MSI-H stage II patients who were treated with singleagent 5FU compared to surgery alone (10). Whether high-risk features are prognostic in patients with dMMR/MSI-H stage II CC is not well established. The role of adjuvant chemotherapy in

patients with high-risk dMMR/MSI-H stage II CC is not well defined. This study aimed to evaluate the prognostic value of high-risk features in dMMR/MSI-H stage II CC and their impact on adjuvant chemotherapy in high-risk stage II CC with dMMR/MSI-H.

MATERIALS AND METHODS

Patient Selection

The National Cancer Database (NCDB) is a large cancer directory that represents approximately 70% of all newly diagnosed cancers in the US. The inclusion criteria for this study included the following: International Classification of Diseases for Oncology, third edition, morphological codes (8020, 8140, 8144, 8210, 8211, 8480, 8481, and 8490) and topography codes (C18.0-9), in participant user data files between the years 2010 and 2013. MSI status information was not available for patients diagnosed before 2010. The primary outcome was the overall survival of patients with dMMR/MSI-H stage II with high-risk features.

Eligibility Criteria

Patient information was independently reviewed by two of the authors for the eligibility criteria. The patients were deemed eligible if they have dMMR/MSI-H stage II CC. Patients with mixed adeno-squamous histology, rectosigmoid location, and rectal cancer were excluded. Patients who received radiation therapy before or after surgery were excluded. Patients who received chemotherapy prior to surgery were excluded, as this may impact the pathologic stage at resection. High-risk features were defined as the following: <12 lymph nodes examined, lymphovascular invasion (LVI), positive surgical margin, pT4 tumor. No data were available for obstruction or perforation at diagnosis. Poor or undifferentiated histology was not included as a high-risk feature, as it is a good prognostic factor in dMMR/ MSI-H stage II CC (18). High-risk stage II CC was defined as having at least one high-risk feature. Institutional approval and informed consent were not required for this study since the patient information in the database is completely de-identified, and the database is legally accessible to the public.

Data Extraction and Statistical Analysis

The patient-specific covariates included were date of diagnosis, date of death, age, gender, race, tumor site, histology, insurance status, stage, presence of metastatic disease, co-morbid medical conditions, location of treatment, and treatment regimen (single or multi-agent chemotherapy). The treatment and clinical outcomes included overall survival rate. All data were checked for internal consistency.

Analysis

All patients in the analysis had dMMR/MSI-H stage II colon cancer. Univariate and multivariate analyses were conducted to identify the factors associated with patient outcome (OS). The clinical and demographic characteristics of the patients were

Abbreviations: dMMR, deficient mismatch repair; pMMR, proficient mismatch repair; MSI-H, microsatellite high; MSS, microsatellite stable.

summarized using descriptive statistics as appropriate for variable type and distribution (chi-square test for categorical variables and ANOVA for numerical variables). Univariate and multivariate Cox-PH models were built to assess the association between patient characteristics and survival. Backward selection with an alpha level of removal of.05 was used in the multivariate analysis. The Kaplan–Meier survival curves were generated with log-rank tests to evaluate the association between adjuvant chemotherapy in high-risk and low-risk cohorts separately. All analyses were performed with a significance level of 0.05 (twosided) with SAS Statistical Package, v9.4 (SAS institute, Inc., Cary, North Carolina).

RESULTS

Of the 249,571 patients with stage II colorectal cancer diagnosed between 2010 and 2013 in the NCDB database, 6,426 patients were determined to have dMMR/MSI-H status, and 2,293 met the inclusion criteria of the study (**Figure 1**). Females accounted for 58.2% of patients; 87.4% were Caucasian. The median age was 69 years (range, 21–90 years old). The most common tumor location was the ascending colon (32.5%), followed by the cecum (27.1%) and the transverse colon (13.1%). The sigmoid and the descending colon accounted for 9.7 and 5.0% of cases, respectively (**Table 1**).

In the entire cohort, 29.5% (n = 676) of patients were deemed to have a high-risk stage II CC. Positive margins, LVI, and less than 12 lymph nodes examined were observed as 3.0, 14.3, and 3.9%, respectively. pT4 was present in 14.0% of patients (**Table 1**). Of the high-risk patients, 36.1% (n = 244) received



 TABLE 1 | Demographics and clinical characteristics of patients.

Total 2.293 Gender 958 (41.8) Male 958 (41.8) Female 1.335 (58.2) Age 14.78 Median 69 (21-90) STD 14.78 Race 2004 (87.4) African American 188 (8.2) Others 101 (4.4) Year of diagnosis 2010 2011 566 (24.7) 2012 633 (27.6) 2013 681 (29.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 8 Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surger at primary site 100.4) Partial cole	Variable	N (%)
Gender 958 (41.8) Female 958 (41.8) Pemale 958 (41.8) Mage 69 (21-90) STD 14.78 Race 2004 (87.4) African American 188 (8.2) Others 101 (4.4) Year of diagnosis 101 (4.4) 2010 413 (18.2) 2011 566 (24.7) 2012 633 (27.6) 2013 681 (29.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 57 (23.4) 2+ 216 (9.4) 2 216 (9.4) Site 1 Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 22 (9.7) Overlapping 36 (1.6) Not otherwise specified 22 (9.7) Subtotal colectomy 42 (1.8) Total p	Total	2,293
Male 968 (41.8) Female 1,335 (58.2) Age 14.78 Meclian 69 (21-90) STD 14.78 Race 2004 (87.4) Mite 2,004 (87.4) African American 188 (8.2) Others 100 (413 (18.0) 2010 413 (18.0) 2011 563 (27.6) 2013 681 (29.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 2 Ascending colon 746 (32.5) Hepatic flexure 143 (62) Transverse colon 301 (13.1) Splenic flexure 143 (62) Transverse colon 301 (13.1) Splenic flexure 143 (62) Transverse colon 301 (13.1) Splenic flexure 155 (5.0) Stot 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Subtotal colectomy/hemicolectomy 1,52 (75.1) Total colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,540 (67.2) Poorly differentiated/moderately differentiated 1,540 (67.2)	Gender	,
Female 1,335 (58.2) Age 1 Median (69 (21–90) STD 14.78 Race 2,004 (87.4) Mite 2,004 (87.4) African American 188 (8.2) Others 101 (4.4) Year of diagnosis 2010 2010 413 (18.0) 2011 566 (24.7) 2012 63 (27.6) 2013 681 (29.7) Charison-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 24 216 (9.4) Site 216 (9.4) Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 143 (6.2) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80.3) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 21 (0.9) Partial colectomy <td>Male</td> <td>958 (41.8)</td>	Male	958 (41.8)
Age 69 (21–90) STD 14.78 Race 2,004 (87.4) African American 188 (8.2) Others 101 (4.4) Year of diagnosis 101 (4.1) 2010 413 (18.0) 2011 566 (24.7) 2012 633 (27.6) 2013 668 (29.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (6.2) Ascending colon 746 (32.5) Hepatic filexure 143 (6.2) Transverse colon 301 (13.1) Splenic filexure 143 (6.2) Transverse colon 301 (13.1) Splenic filexure 143 (6.2) Verlapping 36 (1.6) Not othenwise specified 21 (0.9) Sugmoid colon 22 (29.7) Overlapping 36 (1.6) Not othenwise specified 21 (0.9) Sugmoid colon 22 (1.2) Subtotal colectomy (487 (21.2) Subtotal colectomy	Female	1,335 (58.2)
Median 69 (21-90) STD 14.78 Race 2004 (87.4) African American 188 (8.2) Others 101 (4.4) Year of diagnosis 2010 2010 413 (18.0) 2011 566 (24.7) 2012 633 (27.6) 2013 681 (29.7) Charison-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 21 Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (2) Sutiat colectomy 487 (21.2) Sutiat colectomy 487 (21.2) Sutiat colectomy/themicolectomy 10 (0.4) Unknown 32 (1.4) Grade 10 (0.4)	Age	
STD 14.78 Race 2.004 (87.4) Atrican American 188 (8.2) Others 101 (4.4) Year of diagnosis 2010 2010 413 (18.0) 2011 566 (24.7) 2012 633 (27.6) 2013 681 (28.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 2 Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Sigmoid colon 125 (5.0) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site Partial colectomy Partial colectomy 42 (1.8) Total colectomy 10 (0.4) Unknown 32 (1.4) Gate 10 Vell differentiated/undifferentiated 753 (32.8) Pattia colectomy (167.2) 321 (14.0) <td>Median</td> <td>69 (21–90)</td>	Median	69 (21–90)
Race 2,004 (87.4) White 2,004 (87.4) African American 188 (8.2) Others 101 (4.4) Year of diagnosis 2010 2010 413 (18.0) 2011 566 (24.7) 2012 633 (27.6) 2013 681 (28.7) Charlson-Deyo score 1 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 80.4) Ascending colon 746 (32.5) Hepatic flexure 143 (62.2) Transverse colon 301 (13.1) Splenic flexure 143 (62.2) Transverse colon 301 (14.1) Splenic flexure 143 (62.2) Transverse colon 301 (15.1)	STD	14.78
White 2,004 (87.4) Atrican Armerican 188 (8.2) Others 101 (4.4) Year of diagnosis 2010 2011 566 (24.7) 2013 681 (29.7) Charison-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 2 Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 2 Partial colectomy 1,222 (75.1) Total colectomy 1,221 (75.1) Total proctocolectomy 10 (0.4) Uhknown 32 (1.4) Grade 2 Well differentit	Race	
African American 188 (8.2) Others 101 (4.4) Year of diagnosis 2010 2010 413 (18.0) 2011 566 (24.7) 2012 633 (27.6) 2013 681 (29.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 212 (9.7) Verato differentiated 216 (9.2) Veratory at primary site 80 (3.5) Descending colon 21 (0.9) Suggry at primary site 21 (0.9) Partial colectomy 42 (1.8) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Otal proctocolectomy 10 (2.9) Poorly differentiated/undifferentiated 753 (32.8) Pathological T stage 1151 (0.0)	White	2,004 (87.4)
Others 101 (4.4) Year of diagnosis 2010 2010 413 (18.0) 2011 566 (24.7) 2012 633 (27.6) 2013 681 (29.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 2 Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenci flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (9.2) Surgery at primary site 22 Partial colectomy 42 (1.8) Sutbotal colectomy/hemicolectomy 17,22 (75.1) Total proctocolectomy 10 (0.4) Uhnown 32 (1.4) Grade 2 Well differentiated/moderately differentiated 753 (32.8) Pathological	African American	188 (8.2)
Year of diagnosis 413 (18.0) 2010 413 (18.0) 2011 566 (24.7) 2013 681 (29.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 1 Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 10 (0.4) Unknown 32 (1.4) Grade 1 Vell differentiated/moderately differentiated 753 (32.8) Pathological T stage 1 T3 1,957 (85.3) T4 321 (14.0)	Others	101 (4.4)
2010 413 (18.0) 2011 566 (24.7) 2012 633 (27.6) 2013 681 (29.7) Charlson-Deyo score 0 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 21 Partial colectomy 487 (21.2) Subtal colectomy/hemicolectomy 10 (0.4) Unknown 32 (1.4) Grade 753 (32.8) Pathological T stage 753 (32.8) Pathological T stage 753 (32.9) Pigh-risk features ^a 767 (25.5) Not high risk 1,677 (75.5) Vigh risk features ^a 71 (14.0)	Year of diagnosis	
2011 566 (24.7) 2013 681 (29.7) Charlson-Deyo score 537 (23.4) 2+ 216 (9.4) 2t+ 216 (9.4) Site 621 (27.1) Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 22 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site Partial colectomy/hemicolectomy Partial colectomy/hemicolectomy 1,722 (75.1) Total colectomy/hemicolectomy 10 (0.4) Unknown 32 (1.4) Grade 1 Well differentiated/moderately differentiated 1,540 (67.2) Porty differentiated/undifferentiated 753 (32.8) Pathological T stage 13 T3 1,957 (85.3) T4 321 (14.0) LV 322 (14.3) P	2010	413 (18.0)
2012 633 (27.6) 2013 631 (29.7) Charlson-Deyo score	2011	566 (24.7)
2013 681 (29.7) Charlson-Deyo score 1 0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 21 Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total proctocolectomy 10 (0.4) Uhknown 32 (1.4) Grade 21 Well differentiated/undifferentiated 1,540 (67.2) Poorly differentiated/undifferentiated 1,540 (67.2) Poorly differentiated/undifferentiated 1,640 (67.2) Poorly differentiated/undifferentiated 1,540 (67.2) Poorly diffe	2012	633 (27.6)
Charlson-Deyo score 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 8 (0.4) Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 9 Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Grade 1 Well differentiated/moderately differentiated 1,540 (67.2) Poorly differentiated/moderately differentiated 1,540 (67.2) Poorly differentiated/moderately differentiated 1,617 (70.5) High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 1 T4 321 (14.0) LV 329 (14.3) Systemic therapy after surgery	2013	681 (29.7)
0 1,540 (67.2) 1 537 (23.4) 2+ 216 (9.4) Site 8 (0.4) Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 211 (0.9) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 100.4 Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 10 (0.4) Unknown 32 (1.4) Grade 100.04 Well differentiated/moderately differentiated 753 (32.8) Pathological T stage 73 T3 1,957 (85.3) T4 321 (14.0) Risk group per study 14(1.0) High risk 676 (29.5) Not high risk 1,617 (70.5) High risk 676 (29.5) Not high risk 1,617 (70.5) </td <td>Charlson–Deyo score</td> <td></td>	Charlson–Deyo score	
1 537 (23.4) 2+ 216 (9.4) Site Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site Partial colectomy Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total proctocolectomy 42 (1.8) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Grade 100.4) Well differentiated/moderately differentiated 753 (32.8) Pathological T stage 1 T3 1,957 (85.3) T4 321 (14.0) Kigh risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 1 T4 321 (14.0) LV 329 (14.3)	0	1,540 (67.2)
2+ 216 (9.4) Site 8 (0.4) Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not therwise specified 21 (0.9) Surgery at primary site Partial colectomy Partial colectomy/hemicolectomy 1,722 (75.1) Total colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 10 (0.4) Unknown 32 (1.4) Grade 753 (32.8) Pathological T stage 73 T3 1,957 (85.3) T4 321 (14.0) Kisk group per study 167 (20.5) High risk 676 (29.5) Not high risk 1,617 (70.5) High risk 676 (29.5) Not high risk 1,617 (70.5) High risk 676 (29.5) Not high risk	1	537 (23.4)
Site 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site Partial colectomy Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total protocolectomy 10 (0.4) Unknown 32 (1.4) Grade 10 (0.4) Well differentiated/moderately differentiated 1,5540 (67.2) Poorly differentiated/moderately differentiated 753 (32.8) Pathological T stage 13 T4 321 (14.0) Risk group per study 1,617 (70.5) High risk 676 (29.5) Not high risk 1,617 (70.5) High risk 676 (29.5) Not high risk 1,617 (70.5)	2+	216 (9.4)
Appendix 8 (0.4) Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site Partial colectomy Partial colectomy fremicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total protocolectomy 10 (0.4) Unknown 32 (1.4) Grade Vell differentiated/moderately differentiated Vell differentiated/undifferentiated 753 (32.8) Pathological T stage 1 T3 1,957 (85.3) T4 321 (14.0) Risk group per study 49 (3.9) High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 1 T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	Site	
Cecum 621 (27.1) Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site Partial colectomy Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Grade Vell differentiated/moderately differentiated Vell differentiated/undifferentiated 753 (32.8) Pathological T stage 1 T3 1,957 (85.3) T4 321 (14.0) Risk group per study 1 High risk 1,617 (70.5) High-risk features ^a 1 T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	Appendix	8 (0.4)
Ascending colon 746 (32.5) Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site Partial colectomy Partial colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Grade 1,540 (67.2) Poorly differentiated/moderately differentiated 753 (32.8) Pathological T stage 1 T3 1,957 (85.3) T4 321 (14.0) LVI 329 (14.3) <116 (17.05)	Cecum	621 (27.1)
Hepatic flexure 143 (6.2) Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 21 (0.9) Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total colectomy 10 (0.4) Unknown 32 (1.4) Grade 753 (32.8) Pathological T stage 753 (32.8) Pathological T stage 753 (32.8) Pathological T stage 71 T4 321 (14.0) LV 329 (14.3) T4 321 (14.0) LV 329 (14.3) <12 lymph node removed	Ascending colon	746 (32.5)
Transverse colon 301 (13.1) Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 21 Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Grade 21 (0.9) Well differentiated/moderately differentiated 753 (32.8) Pathological T stage 73 (32.8) T4 321 (14.0) High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 21 (14.0) LV 329 (14.3) <12 lymph node removed	Hepatic flexure	143 (6.2)
Splenic flexure 80 (3.5) Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 2 Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total proctocolectomy 10 (0.4) Uhknown 32 (1.4) Grade 753 (32.8) Well differentiated/moderately differentiated 753 (32.8) Pathological T stage 73 T3 1,957 (85.3) T4 321 (14.0) Risk group per study 1 High risk 676 (29.5) Not high risk 1,617 (70.5) High risk features ^a 7 T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	Transverse colon	301 (13.1)
Descending colon 115 (5.0) Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 2 Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Grade 753 (32.8) Pathological T stage 753 (32.8) Pathological T stage 753 (32.8) Pathological T stage 73 T3 1,957 (85.3) T4 321 (14.0) Risk group per study 1676 (29.5) Not high risk 1,617 (70.5) High risk 1,617 (70.5) VI 329 (14.3) <12 lymph node removed	Splenic flexure	80 (3.5)
Sigmoid colon 222 (9.7) Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 21 (0.9) Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total colectomy 10 (0.4) Unknown 32 (1.4) Grade 753 (32.8) Pathological T stage 753 (32.8) Pathological T stage 753 (32.8) Pathological T stage 753 (32.8) Pigh risk 1,617 (70.5) High risk 1,617 (70.5) High risk 1,617 (70.5) Not high risk 1,617 (70.5) VI 329 (14.3) <12 lymph node removed	Descending colon	115 (5.0)
Overlapping 36 (1.6) Not otherwise specified 21 (0.9) Surgery at primary site 21 (0.9) Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Grade 10 (0.4) Well differentiated/moderately differentiated 753 (32.8) Pathological T stage 73 T3 1,957 (85.3) T4 321 (14.0) Risk group per study 11 High risk 676 (29.5) Not high risk teatures ^a 74 T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	Sigmoid colon	222 (9.7)
Not otherwise specified21 (0.9)Surgery at primary sitePartial colectomy487 (21.2)Subtotal colectomy/hemicolectomy1,722 (75.1)Total colectomy42 (1.8)Total proctocolectomy10 (0.4)Uhknown32 (1.4)Grade1,540 (67.2)Well differentiated/moderately differentiated753 (32.8)Pathological T stage73T31,957 (85.3)T4321 (14.0)Risk group per study11High risk676 (29.5)Not high risk1,617 (70.5)High risk features ^a 21 (14.0)LVI329 (14.3)<12 lymph node removed	Overlapping	36 (1.6)
Surgery at primary site 487 (21.2) Partial colectomy 1,722 (75.1) Total colectomy 42 (1.8) Total colectomy 10 (0.4) Uhknown 32 (1.4) Grade 753 (32.8) Pathological T stage 76 (29.5) Not high risk 1,617 (70.5) High risk 1,617 (70.5) High risk 1,617 (70.5) Systemic features ^a 76 (29.5) Not high risk 1,617 (70.5) Systemic/surgery sequence 89 (3.9) <td>Not otherwise specified</td> <td>21 (0.9)</td>	Not otherwise specified	21 (0.9)
Partial colectomy 487 (21.2) Subtotal colectomy/hemicolectomy 1,722 (75.1) Total colectomy 42 (1.8) Total proctocolectomy 10 (0.4) Unknown 32 (1.4) Grade 32 (1.4) Well differentiated/moderately differentiated 7,540 (67.2) Poorly differentiated/undifferentiated 7,53 (32.8) Pathological T stage 73 T3 1,957 (85.3) T4 321 (14.0) Risk group per study 1,617 (70.5) High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 74 12 lymph node removed 89 (3.9) Positive margin 69 (3.0) Systemic therapy 1,862 (81.2) Systemic therapy after surgery 431 (18.8) Chemotherapy 108 (4.7) No chemotherapy 108 (4.7) Multiagent chemotherapy 292 (12.7) Chemotherapy tope and number of agents not documented 31 (1.4)	Surgery at primary site	
Subtotal colectomy/hemicolectomy1,722 (75.1)Total colectomy42 (1.8)Total proctocolectomy10 (0.4)Unknown32 (1.4)Grade32 (1.4)Well differentiated/moderately differentiated1,540 (67.2)Poorly differentiated/undifferentiated753 (32.8)Pathological T stage73T31,957 (85.3)T4321 (14.0)Risk group per study1,617 (70.5)High risk676 (29.5)Not high risk1,617 (70.5)High-risk features ^a 74T4321 (14.0)LVI329 (14.3)<12 lymph node removed	Partial colectomy	487 (21.2)
I otal colectomy42 (1.8)Total proctocolectomy10 (0.4)Unknown32 (1.4)Grade32 (1.4)Well differentiated/moderately differentiated1,540 (67.2)Poorly differentiated/undifferentiated753 (32.8)Pathological T stage1,957 (85.3)T4321 (14.0)Risk group per study1,617 (70.5)High risk676 (29.5)Not high risk1,617 (70.5)High-risk features ^a 221 (14.0)LVI329 (14.3)<12 lymph node removed	Subtotal colectomy/hemicolectomy	1,722 (75.1)
I otal proctocolectomy10 (0.4)Unknown32 (1.4)Grade	lotal colectomy	42 (1.8)
Unknown 32 (1.4) Grade	lotal proctocolectomy	10 (0.4)
GradeWell differentiated/moderately differentiated1,540 (67.2)Poorly differentiated/undifferentiated753 (32.8)Pathological T stage73T31,957 (85.3)T4321 (14.0) Risk group per study 1,617 (70.5)High risk676 (29.5)Not high risk1,617 (70.5)High-risk features ^a 229 (14.3)<12 lymph node removed	Unknown	32 (1.4)
Vvei differentiated/moderately differentiated 1,540 (67.2) Poorly differentiated/undifferentiated 753 (32.8) Pathological T stage 73 T3 1,957 (85.3) T4 321 (14.0) High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 21 (14.0) LVI 329 (14.3) <12 lymph node removed		1 5 40 (07 0)
Pothological T stage 753 (32.8) Pathological T stage 73 T3 1,957 (85.3) T4 321 (14.0) Risk group per study 1,617 (70.5) High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 221 (14.0) LVI 329 (14.3) <12 lymph node removed	Vell differentiated/moderately differentiated	1,540 (67.2)
T3 1,957 (85.3) T4 321 (14.0) Risk group per study High risk High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 1 T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	Poony dillerentiated/undillerentiated	103 (32.8)
T4 321 (14.0) Risk group per study High risk High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 1 T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	ratiological i stage	1 057 (95 2)
14 321 (14.0) Risk group per study 1 High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 21 (14.0) T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	13	321 (14 0)
High risk 676 (29.5) Not high risk 1,617 (70.5) High-risk features ^a 1 T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	Pisk group per study	521 (14.0)
Not high risk 1,617 (70.5) High-risk features ^a 1,617 (70.5) T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	High risk	676 (20 5)
High-risk features ^a 1,617 (10.0) High-risk features ^a 321 (14.0) LVI 329 (14.3) <12 lymph node removed	Not high risk	1 617 (70 5)
T4 321 (14.0) LVI 329 (14.3) <12 lymph node removed	High-risk features ^a	1,017 (70.0)
LVI 329 (14.3) <12 lymph node removed	T4	321 (14 0)
<12 lymph node removed		329 (14.3)
Positive margin 69 (3.0) Systemic/surgery sequence 1,862 (81.2) No systemic therapy after surgery 431 (18.8) Chemotherapy 1,862 (81.2) Single-agent chemotherapy 108 (4.7) Multiagent chemotherapy 292 (12.7) Chemotherapy type and number of agents not documented 31 (1.4)	<12 lymph node removed	89 (3.9)
Systemic/surgery sequence No systemic therapy Systemic therapy after surgery Chemotherapy No chemotherapy No chemotherapy Single-agent chemotherapy Multiagent chemotherapy 292 (12.7) Chemotherapy type and number of agents not documented 31 (1.4)	Positive margin	69 (3.0)
No systemic therapy 1,862 (81.2) Systemic therapy after surgery 431 (18.8) Chemotherapy 1,862 (81.2) Single-agent chemotherapy 1,862 (81.2) Single-agent chemotherapy 108 (4.7) Multiagent chemotherapy 292 (12.7) Chemotherapy type and number of agents not documented 31 (1.4)	Systemic/surgery sequence	50 (0.0)
Systemic therapy after surgery 431 (18.8) Chemotherapy 1,862 (81.2) Single-agent chemotherapy 108 (4.7) Multiagent chemotherapy 292 (12.7) Chemotherapy type and number of agents not documented 31 (1.4)	No systemic therapy	1,862 (81 2)
Chemotherapy 1,862 (81.2) No chemotherapy 1,862 (81.2) Single-agent chemotherapy 108 (4.7) Multiagent chemotherapy 292 (12.7) Chemotherapy type and number of agents not documented 31 (1.4)	Systemic therapy after surgery	431 (18.8)
No chemotherapy 1,862 (81.2) Single-agent chemotherapy 108 (4.7) Multiagent chemotherapy 292 (12.7) Chemotherapy type and number of agents not documented 31 (1.4)	Chemotherapy	.51 (10.0)
Single-agent chemotherapy 108 (4.7) Multiagent chemotherapy 292 (12.7) Chemotherapy type and number of agents not documented 31 (1.4)	No chemotherapy	1,862 (81 2)
Multiagent chemotherapy Chemotherapy type and number of agents not documented 31 (1.4)	Single-agent chemotherapy	108 (4.7)
Chemotherapy type and number of agents not documented 31 (1.4)	Multiagent chemotherapy	292 (12,7)
	Chemotherapy type and number of agents not documented	31 (1.4)

^aHigh-risk features not mutually exclusive.

adjuvant chemotherapy, of whom 72.1% (n = 176) received multiagent therapy and 23.4% (n = 57) received single-agent chemotherapy, and 4.5% (n = 11) received an unknown number of agents.

On univariate analysis, high-risk status, pT4A/B tumor (pT3 as reference), pathological stage IIB/C (pathological stage IIA as reference), <12 lymph nodes (\geq 12 lymph nodes as reference), positive margins (negative margins as reference), Charlson–Deyo score >1 (0 as reference), and elder age (continuous scale) at diagnosis were associated with worse overall survival (**Table 2**). On multivariate analysis, male sex, positive surgical margin (negative margin as reference), Charlson–Deyo score >1 (0 as reference), high-risk disease, and older age at diagnosis were associated with worse OS (**Table 2**).

High-risk dMMR/MSI-H patients had worse OS compared to non-high-risk dMMR/MSI-H patients in the entire cohort when not stratified by status of adjuvant chemotherapy administration [5-year survival and 95%CI: 73.2% (67.3–78.1%) *vs.* 80.3% (76.7– 83.5%), p = 0.0001, **Figure 2A**]. Median survival is not reachable in our cohort since none of the cohorts had more than 50% of patients who died at the end of follow-up; hence, 5-year survival was provided. In patients who received no adjuvant chemotherapy, the high-risk dMMR/MSI-H patients had worse OS [5-year survival and 95%CI: 69.8% (62.6–75.9%) *vs.* 78.4% (74.3–81.9%), p < 0.0001, **Figure 2B**].

High-risk dMMR/MSI-H patients who received adjuvant chemotherapy had better OS compared to those who had no chemotherapy [5-year survival and 95%CI: 78.0% (66.4–86.0%) vs. 69.8% (62.6–75.9%), p = 0.0011, **Figure 2C**]. In patients with no high-risk features, patients who received adjuvant chemotherapy had better OS [5-year survival and 95%CI: 94.3% (87.6–99.4%) vs. 78.4% (74.3–81.9%), p = 0.0006, **Figure 2D**]. In the patient groups by single/multi-agent adjuvant chemotherapy, single-agent and multi-agent chemotherapy patients demonstrated similar OS, which were both better than those with no chemotherapy. This finding is consistent in high-risk-feature (p = 0.01, **Figure 2E**) and nohigh-risk feature patients (p = 0.0024, **Figure 2F**). When patients with no high-risk features were stratified by age, chemotherapy was no longer associated with better OS. The overall survival in patients with no high-risk features and aged <65 was not different with chemotherapy *versus* no chemotherapy [5-year OS and 95%CI: 96.7% (89.4–99.0%) *vs.* 90.0%: (83.9–93.9%), p = 0.1068, **Figure 3A**]. The overall survival in the same cohort aged \geq 65 years was not different with chemotherapy *versus* with no chemotherapy [5-year survival and 95%CI: 96.1% (62.5–94.2%) and 70.9% (65.1–75.9%), p = 0.1070, **Figure 3B**].

The overall survival in patients with high-risk features was stratified by age. OS in patients aged <65 who received chemotherapy was not different than those of patients who did not receive chemotherapy [5-year OS and 95%CI: 81.7% (67.0–90.3%) vs. 81.8% (69.4–89.5%), p = 0.4209, **Figure 3C**]. The overall survival in the same cohort aged ≥65 years was superior with chemotherapy *versus* with no chemotherapy [5-year OS and 95%CI: 74.5% (56.2–86.1%) vs. 64.6% (55.5–72.3%), p = 0.0490, **Figure 3D**].

DISCUSSION

Prior reports and guidelines established the prognostic value of high-risk features in molecularly unspecified stage II CRC (9, 19–21). However, the therapeutic and prognostic implications of dMMR/MSI-H with high-risk clinicopathologic features have not been adequately studied. This study demonstrates that high-risk features are also prognostic in patients with stage II dMMR/MSI-H. This is the largest published study to establish the prognostic impact of high-risk features in dMMR/MSI-H stage II CRC.

The prognostic role of high-risk features in stage II dMMR/ MSI-H setting raises the question regarding the role of adjuvant therapy in this group of patients. In this study, we observed a significant overall survival benefit in patients with high-risk dMMR/MSI-H stage II CC who received adjuvant chemotherapy compared with those who received surgery only.

TABLE 2 | Overall survival (OS) by mismatch repair and treatment status in univariate and multivariate analysis.

	Univariate OS		p-values	Multivariate OS		p-values
	HR	95%CI		HR	95%CI	
Age	1.05 (1.04–1.06)		<0.001	1.04 (1.03–1.06)		<.001
Male gender	1.02 (0	.82–1.28)	0.843	1.36 (1.07-1.71)		0.004
High risk	1.56 (1	.24–1.97)	<0.001	1.71 (1.35–2.17)		< 0.001
Adjuvant chemotherapy	0.52 (0	.36–0.73)	<0.001	0.66 (0.44-0.98)		0.040
Pathological T4A ^a	1.72 (1	.19–2.49)	0.004			
Pathological T4B ^a	2.19 (1	.53-3.14)	< 0.001			
Pathologic stage IIB ^a	1.63 (1	.13–2.35)	0.009			
Pathologic stage IIC ^a	2.27 (1	.58–3.25)	<.001			
LVI ^a	1.34 (1	.00–1.81)	0.248			
<12 LNS ^a	2.06 (1	.35–3.15)	<.001			
Positive surgical margin ^a	2.79 (1	.79–4.34)	<.001			
Charlson-Deyo score 1	1.71 (1	.31–2.22)	<.001	1.45 (1.11–1.90)		<.001
Charlson–Deyo score 2+	3.89 (2	.93–5.17)	<.001	2.72 (2.03–3.65)		<.001

^aRemoved from the multivariate model to avoid collinearity with high risk.



FIGURE 2 | (A) Survival in high-risk (n = 676) versus no-high-risk (n = 1,617) patients in the entire cohort (n = 2,293) when not stratified by chemotherapy status. (B) Survival in patients who received no adjuvant chemotherapy (n = 1,862) in high-risk (n = 432) and no-high-risk (n = 1,430) patients. (C) Survival in high-risk patients (n = 676) who received adjuvant chemotherapy (n = 244) versus no adjuvant chemotherapy (n = 432). (D) Survival in patients with no high-risk features (n = 1,617) who received adjuvant chemotherapy (n = 187) versus no adjuvant chemotherapy (n = 1,430). (E) Survival in patients with high risk (n = 676) who received no adjuvant chemotherapy (n = 57), and multiagent chemotherapy (n = 176). (F) Survival in patients with no high-risk features who received no chemotherapy (n = 1,430), single-agent chemotherapy (n = 51), and multiagent chemotherapy (n = 116).

The benefit was associated with both multiagent and single-agent adjuvant chemotherapy. This is a novel finding that has a potential impact on clinical practice in the absence of data from clinical trials. The survival benefit persisted in patients >65 years when stratified by age. Sporadic dMMR/MSI-H is known to be associated with older age at diagnosis compared to germline dMMR/MSI-H (17, 22). In this study, patients with no high-risk features and dMMR/MSI-H stage II colon cancer also benefited from chemotherapy; however, age was an important confounding factor. Survival benefit was not evident in the low-risk group when stratified by age. We speculate that patients in the older age group of high-risk dMMR/MSI-H stage II CC of this study may have had a sporadic dMMR/MSI-H disease and may have derived more benefit from adjuvant chemotherapy compared to the younger population, which possibly may have had a higher rate of germline dMMR/MSI-H. It should be noted that no data were available in NCDB for BRAF status, MLH1 methylation status, and germline testing of family history.

Several prior studies, including retrospective analysis and observational reports, have attempted to address the role of adjuvant chemotherapy in stage II CC with high-risk features; however, these studies did not include the dMMR//MSI-H status (3, 23). Data about the role of adjuvant chemotherapy in highrisk dMMR/MSI-H stage II CC is limited. Tougeron et al.



(n = 83) versus no chemotherapy (n = 303).

reported the clinical outcomes of stage II and III dMMR/MSI-H CC patients treated between 2000 and 2011 in a multicenter retrospective French study (24). Sixty percent (n = 149) of the patients were deemed to have high-risk factors, and 22% (n = 33) of the high-risk patients were treated with adjuvant chemotherapy. The high-risk features included pT4, VELIPI criteria (vascular emboli, lymphatic invasion, or perineural invasion), poor/undifferentiated histology, less than eight lymph nodes examined, tumor perforation, and initial bowel obstruction. Patients who were treated with adjuvant FOLFOX and not the single agent 5-FU showed a trend for longer disease-free survival compared to surgery alone. Although it included specific chemotherapy regimen information, the number of patients in that study was much smaller than in the present study.

Control for variables that could have influenced the results, such as histopathologic features, tumor grade, age, and performance status, was performed. Significant limitations still exist in this analysis, and these include retrospective design, lack of randomization, and no individualized patient data regarding the specifics of chemotherapy or follow-up. The reasons why adjuvant chemotherapy was not administered is unknown. The precise chemotherapy agents administered were not available. The sporadic *versus* germline mutational status is unknown, and the prevalence of other genomic alterations is similarly unknown. The overall survival is not cancer specific in this study, as NCDB includes only all-cause overall survival. In addition, the high-risk features did not include obstruction or perforation, as there was no data available in the database. Tumor perforation was shown to be associated with interperitoneal tumor dissemination (25), which raises the concern of whether these patients actually have stage II disease. The reported incidence of perforation and obstruction in stage II CC is less than 10% (26–28), and the impact of tumor cell spillage on recurrence and survival increases the risks substantially and may contribute to an increase in the benefit of adjuvant therapy.

CONCLUSION

The prognostic value of high-risk features in dMMR/MSI-H stage II CC is confirmed. The prognostic value of high-risk features should be considered in adjuvant therapy discussions. Adjuvant chemotherapy may be associated with better OS in high-risk dMMR/MSI-H stage II CC patients, but significant limitations exist, including the retrospective nature of the data set. In the absence of randomized trials, the benefits and the risks of adjuvant therapy should be discussed with the patients with high-risk dMMR/MSI-H stage II CC. Further research needs to

be done in the low-risk dMMR/MSI-H stage II CC to confirm the lack of benefit from adjuvant chemotherapy.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/supplementary material. Further inquiries can be directed to the corresponding author.

AUTHOR CONTRIBUTIONS

AM and MA participated in collecting data, writing the manuscript, and editing the manuscript. MD participated in collecting data, writing the manuscript, and editing the manuscript. MB participated in the analysis of data and editing the manuscript. RJ, PP, CW, OA, WS, TG, and GB participated in editing the manuscript. BE-R participated in mentoring the

REFERENCES

- Siegel RL, Miller KD, Fuchs HE, Jemal A. Cancer Statistics, 2021. CA: A Cancer J Clin (2021) 71(1):7–33. doi: 10.3322/caac.21654
- Jemal A, Siegel R, Ward E, Hao Y, Xu J, Thun MJ. Cancer Statistics, 2009. CA Cancer J Clin (2009) 59(4):225–49. doi: 10.3322/caac.20006
- Verhoeff SR, van Erning FN, Lemmens VE, de Wilt JH, Pruijt JF. Adjuvant Chemotherapy Is Not Associated With Improved Survival for All High-Risk Factors in Stage II Colon Cancer. *Int J Cancer* (2016) 139(1):187–93. doi: 10.1002/ijc.30053
- Casadaban L, Rauscher G, Aklilu M, Villenes D, Freels S, Maker AJ. Adjuvant Chemotherapy is Associated With Improved Survival in Patients With Stage II Colon Cancer. *Cancer* (2016) 122(21):3277–87. doi: 10.1002/cncr.30181
- Andre T, de Gramont A, Vernerey D, Chibaudel B, Bonnetain F, Tijeras-Raballand A, et al. Adjuvant Fluorouracil, Leucovorin, and Oxaliplatin in Stage II to III Colon Cancer: Updated 10-Year Survival and Outcomes According to BRAF Mutation and Mismatch Repair Status of the MOSAIC Study. J Clin Oncol (2015) 33(35):4176–87. doi: 10.1200/JCO.2015.63.4238
- Bastos DA, Ribeiro SC, de Freitas D, Hoff PM. Combination Therapy in High-Risk Stage II or Stage III Colon Cancer: Current Practice and Future Prospects. *Ther Adv Med Oncol* (2010) 2(4):261–72. doi: 10.1177/ 1758834010367905
- Wolmark N, Rockette H, Mamounas E, Jones J, Wieand S, Wickerham DL, et al. Clinical Trial to Assess the Relative Efficacy of Fluorouracil and Leucovorin, Fluorouracil and Levamisole, and Fluorouracil, Leucovorin, and Levamisole in Patients With Dukes' B and C Carcinoma of the Colon: Results From National Surgical Adjuvant Breast and Bowel Project C-04. J Clin Oncol (1999) 17(11):3553–9. doi: 10.1200/JCO.1999.17.11.3553
- Wolmark N, Rockette H, Fisher B, Wickerham DL, Redmond C, Fisher ER, et al. The Benefit of Leucovorin-Modulated Fluorouracil as Postoperative Adjuvant Therapy for Primary Colon Cancer: Results From National Surgical Adjuvant Breast and Bowel Project Protocol C-03. *J Clin Oncol* (1993) 11 (10):1879–87. doi: 10.1200/JCO.1993.11.10.1879
- 9. NCCN. Clinical Practice Guidelines in Oncology. Colon Cancer (2019).
- Sargent DJ, Marsoni S, Monges G, Thibodeau SN, Labianca R, Hamilton SR, et al. Defective Mismatch Repair as a Predictive Marker for Lack of Efficacy of Fluorouracil-Based Adjuvant Therapy in Colon Cancer. J Clin Oncol (2010) 28(20):3219–26. doi: 10.1200/JCO.2009.27.1825

whole project and editing the manuscript. All authors read and approved the final manuscript.

FUNDING

The research reported in this publication was supported in part by the Winship Research Informatics Shared Resource of Winship Cancer Institute of Emory University and National Institutes of Health/ National Cancer Institute under award number P30CA138292.

ACKNOWLEDGMENTS

Part of the data presented in this study was presented at the 2018 ASCO Annual Meeting in Chicago, IL, United States. The data used in the study are derived from a de-identified National Cancer Database (NCDB) file. The NCDB is a joint project of the Commission on Cancer of the American College of Surgeons and the American Cancer Society.

- Kheirelseid EA, Miller N, Chang KH, Curran C, Hennessey E, Sheehan M, et al. Mismatch Repair Protein Expression in Colorectal Cancer. J Gastrointest Oncol (2013) 4(4):397–408. doi: 10.3978/j.issn.2078-6891.2013.021
- Kawakami H, Zaanan A, Sinicrope FA. Implications of Mismatch Repair-Deficient Status on Management of Early Stage Colorectal Cancer. *J Gastrointest Oncol* (2015) 6(6):676–84. doi: 10.3978/j.issn.2078-6891.2015.065
- Sinicrope FA, Yang ZJ. Prognostic and Predictive Impact of DNA Mismatch Repair in the Management of Colorectal Cancer. *Future Oncol* (2011) 7 (3):467–74. doi: 10.2217/fon.11.5
- Zhang CM, Lv JF, Gong L, Yu LY, Chen XP, Zhou HH, et al. Role of Deficient Mismatch Repair in the Personalized Management of Colorectal Cancer. Int J Environ Res Public Health (2016) 13(9). doi: 10.3390/ijerph13090892
- Sinicrope FA. DNA Mismatch Repair and Adjuvant Chemotherapy in Sporadic Colon Cancer. Nat Rev Clin Oncol (2010) 7(3):174–7. doi: 10.1038/nrclinonc.2009.235
- Popat S, Hubner R, Houlston RS. Systematic Review of Microsatellite Instability and Colorectal Cancer Prognosis. J Clin Oncol (2005) 23(3):609– 18. doi: 10.1200/JCO.2005.01.086
- Sinicrope FA, Foster NR, Thibodeau SN, Marsoni S, Monges G, Labianca R, et al. DNA Mismatch Repair Status and Colon Cancer Recurrence and Survival in Clinical Trials of 5-Fluorouracil-Based Adjuvant Therapy. J Natl Cancer Inst (2011) 103(11):863–75. doi: 10.1093/jnci/djr153
- Henson DE, Albores-Saavedra J, Corle D. Carcinoma of the Gallbladder. Histologic Types, Stage of Disease, Grade, and Survival Rates. *Cancer* (1992) 70(6):1493-7. doi: 10.1002/1097-0142(19920915)70:6<1493::aidcncr2820700608>3.0.co;2-u
- Quah HM, Chou JF, Gonen M, Shia J, Schrag D, Landmann RG, et al. Identification of Patients With High-Risk Stage II Colon Cancer for Adjuvant Therapy. *Dis Colon Rectum* (2008) 51(5):503–7. doi: 10.1007/s10350-008-9246-z
- Benson AB 3rd, Schrag D, Somerfield MR, Cohen AM, Figueredo AT, Flynn PJ, et al. American Society of Clinical Oncology Recommendations on Adjuvant Chemotherapy for Stage II Colon Cancer. J Clin Oncol (2004) 22 (16):3408–19. doi: 10.1200/JCO.2004.05.063
- Schmoll HJ, Van Cutsem E, Stein A, Valentini V, Glimelius B, Haustermans K, et al. ESMO Consensus Guidelines for Management of Patients With Colon and Rectal Cancer. A Personalized Approach to Clinical Decision Making. *Ann Oncol* (2012) 23(10):2479–516. doi: 10.1093/annonc/mds236

- 22. Cohen R, Buhard O, Cervera P, Hain E, Dumont S, Bardier A, et al. Clinical and Molecular Characterisation of Hereditary and Sporadic Metastatic Colorectal Cancers Harbouring Microsatellite Instability/DNA Mismatch Repair Deficiency. *Eur J Cancer* (2017) 86:266-74. doi: 10.1016/ j.ejca.2017.09.022
- O'Connor ES, Greenblatt DY, LoConte NK, Gangnon RE, Liou J-I, Heise CP, et al. Adjuvant Chemotherapy for Stage II Colon Cancer With Poor Prognostic Features. J Clin Oncol (2011) 29(25):3381–8. doi: 10.1200/ JCO.2010.34.3426
- Tougeron D, Mouillet G, Trouilloud I, Lecomte T, Coriat R, Aparicio T, et al. Efficacy of Adjuvant Chemotherapy in Colon Cancer With Microsatellite Instability: A Large Multicenter AGEO Study. J Natl Cancer Inst (2016) 108 (7). doi: 10.1093/jnci/djv438
- Koppe MJ, Boerman OC, Oyen WJG, Bleichrodt RP. Peritoneal Carcinomatosis of Colorectal Origin: Incidence and Current Treatment Strategies. Ann Surg (2006) 243(2):212-22. doi: 10.1097/ 01.sla.0000197702.46394.16
- Badia JM, Sitges-Serra A, Pla J, Martí Ragué J, Roqueta F, Sitges-Creus A. Perforation of Colonic Neoplasms. A Review of 36 Cases. *Int J Colorectal Dis* (1987) 2(4):187–9. doi: 10.1007/BF01649502
- 27. Mandava N, Kumar S, Pizzi WF, Aprile IJ. Perforated Colorectal Carcinomas. Am J Surg (1996) 172(3):236–8. doi: 10.1016/S0002-9610(96)00164-X
- 28. Lee IK, Sung NY, Lee YS, Lee SC, Kang WK, Cho HM, et al. The Survival Rate and Prognostic Factors in 26 Perforated Colorectal

Cancer Patients. Int J Colorectal Dis (2007) 22(5):467-73. doi: 10.1007/s00384-006-0184-8

Author Disclaimer: The American College of Surgeons and the Commission on Cancer have not verified and are not responsible for the analytic or statistical methodology employed or the conclusions drawn from these data by the investigator.

Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's Note: All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Copyright © 2021 Mohamed, Jiang, Philip, Diab, Behera, Wu, Alese, Shaib, Gaines, Balch, El-Rayes and Akce. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.