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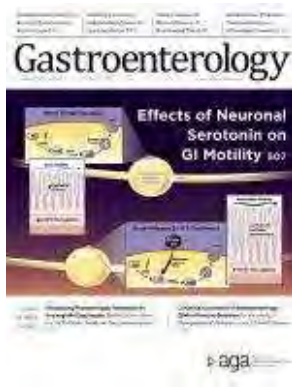
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AGA Abstracts

Su1660 – Colorectal Cancer Screening: Compliance with Multitarget Stool Dna Testing Among Medicare Beneficiaries

Rebecca Swartz, Emily Weiser, Philip Parks, Jack Van Thomme, Paul Limburg, Barry M. Berger

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Appendix SWARTZ02

performance. **Conclusions** A machine learning approach using cfDNA achieved high sensitivity and specificity in a large, predominantly early-stage, colorectal cancer cohort. The possibility of systematic technical and site-specific biases warrants similar confounder analyses in other retrospective, as well as prospective, studies. Prospective validation of the presented machine learning method and evaluation of a multi-analyte approach are underway.

Su1659

POSITIVE FAECAL OCCULT BLOOD TEST AFTER A PREVIOUS NEGATIVE COLONOSCOPY - TO SCOPE OR NOT?

Nivedita Rattan, Sulak Anandabaskaran, Diana Aston, George G. Ermerak, Watson Ng, Milan S. Bassan, David Abi-Hanna, Jenn H. Koo

Introduction Colorectal carcinoma (CRC) is the second-most commonly diagnosed cancer and second-highest cause of cancer mortality in Australia. Currently, there is no consensus to inform clinicians of the optimal approach to manage patients presenting with positive faecal occult blood test (FOBT) but recent negative colonoscopy. This study aims to determine the likelihood of detecting significant colorectal pathology in these patients and the clinical, pathological and colonoscopic factors associated with advanced pathology. **Methods:** This study recruited patients who underwent colonoscopy following positive FOBT from March 2015 – April 2018 at a tertiary hospital in Sydney, Australia. Data collected included patient demographics and clinical characteristics, previous and current colonoscopy findings. The primary outcome was to identify the prevalence of CRC and advanced adenoma (size >10mm, high-grade dysplasia or villous architecture) in a patient with positive FOBT but previous negative colonoscopy (defined as no polyp or cancer). The secondary outcomes include the timeframe to detection and prevalence of advanced lesions on current colonoscopy in patients with or without prior colonoscopy. Statistical analysis using chi-square tests and binary logistic regression models assessed independent predictors of advanced neoplasia. **Results:** Colonoscopies were performed on 740 patients (male 52.8%; median age 64 years old). There were 27 (3.6%) patients with CRC, 362 (48.9%) with at least one adenoma and 188 (25.4%) with an advanced adenoma. There were 219 patients (29.6%) who had a previous colonoscopy. Of these, 111 had a previous negative index colonoscopy and in this cohort, there was 1 CRC detected, 16 with advanced adenomas and 45 (40.5%) with adenomas. Among patients with a negative index colonoscopy within 4 years, none had CRC and 7 had an advanced adenoma. Of the 452 patients with no prior colonoscopy, there were 22 CRCs (4.9%) and 129 patients (28.5%) with an advanced adenoma on their current colonoscopy. Multivariate analysis identified 3 significant independent predictors for the detection of advanced adenomas/CRC - male sex (OR 2.07; 95%CI 1.44-2.98; $P < 0.001$), age (OR 1.04; 95%CI 1.02-1.07; $P < 0.001$) and no previous colonoscopy (OR 2.39 95%CI 1.59-3.62; $P < 0.001$). A personal history of polyps, family history of CRC, current abnormal symptoms and quality of bowel preparation of the previous colonoscopy were not significant predictors. **Conclusion** A previous negative colonoscopy done within 4 years conferred a 100% negative predictive value for detection of new CRC despite a positive FOBT. Any prior colonoscopy carried a better prognosis for detection of advanced pathology in a patient with current positive FOBT. These findings could potentially provide guidelines to managing patients with positive FOBT but previous negative colonoscopy.

Su1660

COLORECTAL CANCER SCREENING: COMPLIANCE WITH MULTI-TARGET STOOL DNA TESTING AMONG MEDICARE BENEFICIARIES

Rebecca Swartz, Emily Weiser, Philip Parks, Jack Van Thomme, Paul Limburg, Barry M. Berger

Introduction The multi-target stool DNA (mt-sDNA) test is a non-invasive colorectal cancer (CRC) screening assay included in the United States Preventive Services Task Force, American Cancer Society, and Multi-Society Task Force guidelines for CRC screening of average-risk adults ages 50 years and older. Mt-sDNA testing is performed by a single national reference laboratory (Exact Sciences Laboratories [ESL], Madison, WI) and includes a nationwide 24/7 patient navigation system (compliance program) to support test completion through patient education and periodic reminders. The majority of active patient navigation activity occurs during the 60 days following each mt-sDNA test order. We focus on Medicare beneficiaries, as Medicare covers mt-sDNA screening with no out-of-pocket expenses, removing the potential impact of financial barriers on compliance. The impact of this systematic approach to CRC screening for Medicare beneficiaries is reported here. **Methods** Medicare beneficiaries ages 65-85 years with a valid mt-sDNA order between September 1, 2016 and August 31, 2017 were identified from the ESL database. Compliance was defined as completion of the mt-sDNA test (positive or negative result) within 365 days of the order date. To assess factors associated with early versus late compliance, compliance rates within 30, 60, 90, 180, and 365 days of the order date were calculated. Compliance by age, sex, and Medicare coverage type were examined. Descriptive statistics were used to identify and compare the data. **Results** Mt-sDNA test compliance was 71.1% among a population of 368,494 Medicare beneficiaries ages 65-85 years that was 64% female. Compliance was similar among 5-year age cohorts though individuals ages 75-85 years had slightly higher compliance. The median time to compliance decreased with age, from 28 days for individuals ages 65-69 years to 24 days for individuals ages 80-85 years. Compliance was similar for traditional Medicare versus Medicare Advantage coverage and by patient sex (Table 1). Among individuals complying within 365 days of their mt-sDNA order, 61.6% complied within 30 days and 90.7% complied within 60 days, which correlates with the time when the mt-sDNA test's patient compliance program is most active (Table 2). **Conclusions** Data from this observational study show that a high percentage (71%) of Medicare beneficiaries who had an mt-sDNA test order completed the test, with 91% completing the test within 60 days. These findings suggest that mt-sDNA test features (e.g. non-invasive, high sensitivity, molecularly-based, three-year screening interval) coupled with built-in patient navigation can enhance CRC screening compliance in the Medicare population. Patient, provider, CRC screening program, and healthcare delivery system characteristics should be studied to understand why some patients do not comply or remain unscreened.

Table 1. mt-sDNA Orders, Completed Tests, Compliance, and Time to Compliance for the Study Population

Category Subgroup	Number of mt-sDNA Test Orders	Number of Tests Completed Within 365 Days of Order	Compliance Rate (% Tests Completed Within 365 Days of Order) [1]	Median Time to Compliance (Days)	IQR Time to Compliance (Days)
Overall Medicare population	368,494	262,084	71.1%	27	20, 37
Age					
65-69 years	141,191	99,764	70.7%	28	21, 39
70-74 years	121,566	86,220	70.9%	27	20, 37
75-79 years	70,502	50,998	72.3%	25	20, 35
80-85 years	35,235	25,102	71.2%	24	19, 34
Sex [2]					
Male	132,806	95,433	71.9%	26	20, 35
Female	235,617	166,601	70.7%	28	21, 38
Coverage type					
Traditional Medicare	250,800	178,922	71.3%	27	20, 37
Medicare Advantage Plan	117,694	83,162	70.7%	27	20, 37

1. Percentage of individuals with a valid mt-sDNA order who were successfully screened (positive or negative mt-sDNA result) within 365 days of their order date
2. Patient sex is available for orders only when included on the test order form
IQR = interquartile range; mt-sDNA = multi-target stool DNA test.

Mt-sDNA Orders, Completed Tests, Compliance, and Time to Compliance for the Study Population

Table 2. Cumulative Compliance with mt-sDNA Testing over 365 Days Following Order Date

Category	Within 30 Days	Within 60 Days	Within 90 Days	Within 180 Days	Within 365 Days
Compliance rate [1]	61.6%	90.7%	94.4%	97.8%	100.0%
Completed tests	161,449	237,603	247,495	256,432	262,084

1. Percentage of individuals with a valid mt-sDNA order who were successfully screened (positive or negative mt-sDNA test result) within 30, 60, 90, 180, or 365 days of their order date
Mt-sDNA = multi-target stool DNA test.

Cumulative Compliance with mt-sDNA Testing over 365 Days Following Order Date

Su1661

A FAVORABLE SURVIVAL OF THE SCREENING-DETECTED COLORECTAL CANCER IN YOUNG ADULTS (<50): IMPLICATIONS FOR COLORECTAL CANCER SCREENING FOR YOUNG ADULTS

Kazunori Takada, Kinichi Hotta, Kenichiro Imai, Sayo Ito, Yoshihiro Kishida, Masao Yoshida, Noboru Kawata, Naomi Kakushima, Kohei Takizawa, Hirotooshi Ishiwatari, Hiroyuki Matsubayashi, Hiroyuki Ono

Backgrounds and Aims: Recent increasing incidence and mortality of colorectal cancer (CRC) among young adults (<50 years) in the Western countries has been paid attentions. Screening for young adults is being considered but controversy due to limited evidence of its effectiveness. Most countries begin CRC screening at age of 50, whereas Japan begins at age of 40 mainly using fecal immunochemical test. Here, we aimed to clarify characteristics of CRC of young adults (<50) and the long-term outcomes of screening-detected CRC of young adults by comparisons with detection age and the detection mode using an institutional cancer registry database. **Methods:** This is a retrospective cohort study using a CRC registry of a Japanese tertiary cancer center. All consecutive patients who were diagnosed as CRC between January 2007 and December 2016 were included. CRCs were staged based on the 8th edition of the Union for International Control Criteria. When multiple CRCs were identified, a CRC at the most advanced stage only was considered. Patients were categorized into 3 groups according to age: Age <50 group, Age 50-75 group, Age >75 group. CRCs were categorized into screening-detected CRC (SD-CRC) and non-screening-detected CRC (non-SD-CRC). We calculated overall survival (OS) using the Kaplan-Meier method and adjusted hazard ratio (HR) using Cox regression models. **Results:** During median follow-up period of 4.6 year, 4345 patients with 4345 CRCs were identified. Age <50 group had 361 patients (8.3%). Age <50 group showed a significantly lower rate of SD-CRC (29.4 vs. 35.8% in Age 50-75 group). Higher rate of rectal cancer (50.4 vs. 43.3% in Age 50-75 group), increased proportion of family history of CRC (22.2 vs. 16.8% in Age 50-75 group), and decreased proportion of Stage I CRCs (15.2 vs. 30.3%) were noted. SD-CRC group of Age <50 showed increased proportion of Stage I CRCs (29.2 vs. 9.4% in non-SD-CRC group of Age <50) (Table 1). Adjusted HR for Age <50 group was 1.12 (95% CI, 0.90-1.41) (Figure 1A). SD-CRC group demonstrated a significantly better prognosis than non-SD-CRC group (HR, 0.51; 95% CI: 0.42-0.61) (Figure 1B). Notably, SD-CRC group in Age <50 group also demonstrated significantly more favorable OS than the non-SD-CRC group (HR, 0.50; 95% CI: 0.26-0.95), similar to in the Age 50-75 group (HR, 0.46; 95% CI: 0.36-0.95) and in the Age >75 group (HR, 0.59; 95% CI: 0.42-0.82). **Conclusion:** We found that young adult patients (<50) were more likely to present with advanced stage and had a lower rate of SD-CRC. Nevertheless, patients with SD-CRC at age <50 showed a significantly more favorable overall survival than non-SD-CRC patients. This implicates a potential benefit of screening for young adults (<50).

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Multi-Target Stool DNA Testing for Colorectal Cancer Screening: Emerging Learning on Real-world Performance

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Abstract

Purpose of Review

Multi-target stool DNA (MT-sDNA) was approved in 2014 for use in screening average-risk patients for colorectal cancer (CRC). Here, we highlight recent literature from post-market

studies to provide an update on clinical use and utility not possible from pre-approval studies.

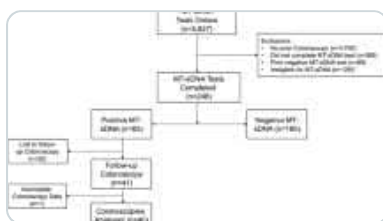
Recent Findings

MT-sDNA has been included in major society guidelines as an option for colorectal cancer screening, and has seen exponentially increasing use in clinical practice. MT-sDNA appears to be attracting new patients to CRC screening, and patient adherence to diagnostic colonoscopy after a positive MT-sDNA test is high. Approximately two-thirds of these patients are found to have colorectal neoplasia (CRN), 80% of whom have at least one right-sided lesion; 1 in 3 will have advanced CRN. High yield of CRN is due not only to post-screening increase in probability but also likely improved endoscopist attention. In those with a negative high-quality colonoscopy after positive MT-sDNA test (“false positive MT-sDNA”), further interventions do not appear to be necessary.

Summary

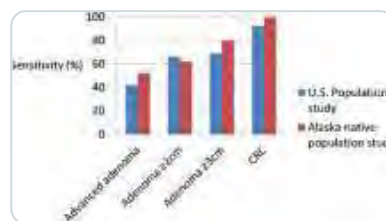
MT-sDNA is a promising tool to improve rates and quality of CRC screening. Further investigation should examine MT-sDNA performance in populations at increased risk for CRC, and as an interval test after colonoscopy to detect potentially missed lesions.

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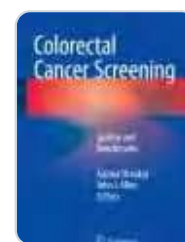
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Introduction

Colorectal cancer (CRC) is a major cause of cancer-related death in the USA [1], with approximately 150,000 new diagnoses and 50,000 deaths from CRC projected in 2019 alone [2]. With widespread screening initiatives to detect CRC and pre-cancerous polyps, CRC-related morbidity and mortality have decreased in recent years [2]. The US Preventive Services Task Force (USPSTF) recommends screening for CRC in average-risk adults between the ages of 50 and 75 [3], while more recent recommendations from the American Cancer Society (ACS) suggest initiating screening at age 45 in light of the increasing rate of CRC diagnosed in younger patients [4,5,6,7,8]. However, despite these guidelines and extensive public health initiatives, approximately one-third of screen-eligible adults in the USA remain unscreened [9, 10].

Several methods are recommended for CRC screening [3, 4, 11]. Colonoscopy is the most widely used modality in the USA [9] and permits simultaneous diagnostic and therapeutic intervention for colorectal neoplasia (CRN). However, screening colonoscopy has several notable drawbacks, including reduced sensitivity in the proximal colon [12,13,14,15,16,17,18] and inter-operator variability in colonoscopy quality [19,20,21,22]. Furthermore, its invasive nature; risk of complications; and the inconvenience of preparation, sedation, and time away from work are recognized barriers to patients considering screening [23]. Several non-invasive screening modalities are endorsed as an alternative to colonoscopy for use in average-risk patients. Guaiac-based fecal occult blood testing (gFOBT) has the strongest body of evidence demonstrating a reduction in CRC-related mortality [24,25,26,27]. However, dietary and medication interactions, poor performance in the proximal colon, and issues with adherence to the recommended yearly screening interval have led to decreased utilization of this test [28,29,30]. The introduction of fecal immunochemical testing (FIT) addressed some of these limitations, with FIT included as a first-line option in major society screening guidelines [3, 11]. A recent meta-analysis suggested moderate-to-high sensitivity and specificity for CRC and advanced adenoma (AA) in a single-use setting, but performance varied by assay and positivity cutoff [31]. Concerns remain regarding reduced sensitivity for right-sided and sessile serrated

precursor lesions [28, 29, 32] in addition to persistent poor adherence to required annual testing [30, 33, 34, 35, 36].

Multi-target stool DNA testing

Multi-target stool DNA (MT-sDNA) is a newer stool-based screening modality, with one commercial product available to patients in the USA (Cologuard™, Exact Sciences, Madison, WI). This test detects abnormal DNA markers (aberrantly methylated *BMP3* and *NDRG4*, mutant *KRAS*) shed from neoplastic cells in the gastrointestinal tract. Another marker, β -actin, serves as a control to ensure adequate quantity of DNA in the sample, while an immunochemical assay for hemoglobin detects evidence of gastrointestinal bleeding. These components are measured quantitatively and incorporated into a logistic regression algorithm containing mathematical interaction variables for each marker. These interaction terms result in numerous marker combinations that can trigger a pre-specified threshold score, resulting in a positive or negative test result that is more than the sum of its individual components [37].

MT-sDNA testing is performed by the patients in the comfort of their own home. Following a provider order, a manufacturer-based patient navigation office contacts the patient and arranges sample collection via a courier-delivered kit [38]. This system also provides patient education and test completion reminders which have been shown to result in adherence to testing of over 70% in Medicare beneficiaries [38]. Following delivery of the MT-sDNA kit, patients collect a whole stool sample, swab a portion to collect the FIT sample, add a preservative buffer to the whole stool, seal the container, and mail the samples in one box back to the laboratory for analysis.

MT-sDNA testing was approved for screening in average-risk adults by the US Food and Drug Administration (FDA) in 2014, following a large screen-setting trial by Imperiale et al. (DeeP-C trial), comparing MT-sDNA and FIT. Overall, the sensitivity of MT-sDNA for detecting CRC in this study was 92% (vs 74% for FIT, $p = 0.002$), and 42% for advanced pre-cancerous lesions (vs 24% for FIT, $p < 0.001$) [39]. A second study performed in an Alaska Native cohort confirmed these findings, and re-demonstrated increasing sensitivity by adenoma size, with 80% sensitivity for lesions > 3 cm, and 100% sensitivity for CRC [40]. Specificity was 87–94% across these two studies, and increased in younger patients (Table

1) [39, 40]. MT-sDNA appears to have similar performance for the detection of both conventional adenomas and sessile serrated polyps (SSPs). Large SSPs and hyperplastic polyps (≥ 1 cm) were included in the sensitivity estimate for advanced pre-cancerous lesions in the DeeP-C study [39]. For greater clarity, a post hoc analysis of DeeP-C data was performed on all lesions ≥ 1 cm with histology confirmed for SSP. The median size was 1.2 cm (range 1–3), and the majority (28/29 (93%)) were located proximal to the splenic flexure; only one SSP contained dysplasia. In this setting, MT-sDNA had a sensitivity of 55% (95% CI, 36–74%) and a specificity of 91% [41]. MT-sDNA performance characteristics for the individual endpoint category of large hyperplastic polyp have not been estimated and traditional serrated adenomas were not identified by screening colonoscopy in the DeeP-C study; these endpoints are deserving of future study.

Table 1 Multi-target stool DNA demonstrates higher specificity in younger patients with normal colonoscopy

Population modeling studies performed as part of a USPSTF review suggest that programmatic use of MT-sDNA would result in fewer lifetime colonoscopies than FIT and a stronger benefit-to-harms ratio than colonoscopy (Table 2) [3]. As a result of the above findings, MT-sDNA testing was subsequently endorsed by the USPSTF in their 2016 guidelines for CRC screening [3]. Utilization of MT-sDNA has been increasing exponentially since then, with nearly 3 million screens performed to date [42].

Table 2 Screening with multi-target stool DNA testing results in fewer lifetime colonoscopies than FIT, and has a superior benefit-to-harm ratio compared to colonoscopy

Outcomes in patients undergoing multi-target stool DNA testing

Since the introduction of MT-sDNA in 2014, data have begun to emerge regarding its “real-world” performance. MT-sDNA appears to be attracting new patients to screening. One recent study demonstrated high utilization of MT-sDNA in almost 400 previously non-adherent average-risk Medicare patients, with 51% harboring advanced neoplastic lesions; fewer than 20% had no neoplastic findings at diagnostic colonoscopy [[43](#)]. Furthermore, we recently presented data from a cohort of over 2000 MT-sDNA-positive patients showing that approximately 25% had never been screened for CRC and were > 10 years overdue [[44](#)]. While there may be many reasons for patients to go without screening, these findings suggest that MT-sDNA is attracting previously non-adherent patients, a critical outcome in the ongoing long-term fight to reduce CRC-attributable mortality.

Of all patients screened with MT-sDNA, 14–16% will have positive results [[39](#), [44](#), [45](#)]. Following a positive MT-sDNA test, it is imperative that patients undergo diagnostic colonoscopy to evaluate for CRN. While previous estimates of adherence to diagnostic colonoscopy ranged from 75 to 96% [[43](#), [45](#)], our recent analysis showed a diagnostic colonoscopy completion rate of ~ 90% in a large multi-practice setting [[44](#)]. Reasons for non-adherence to recommended diagnostic colonoscopy vary, but are commonly related to a wish to avoid colonoscopy, in addition to medical comorbidities that preclude sedation and procedural intervention [[45](#)]. These factors illustrate the importance of both provider and patient education prior to pursuing MT-sDNA testing in order to ensure selection of appropriate patients for screening with this test.

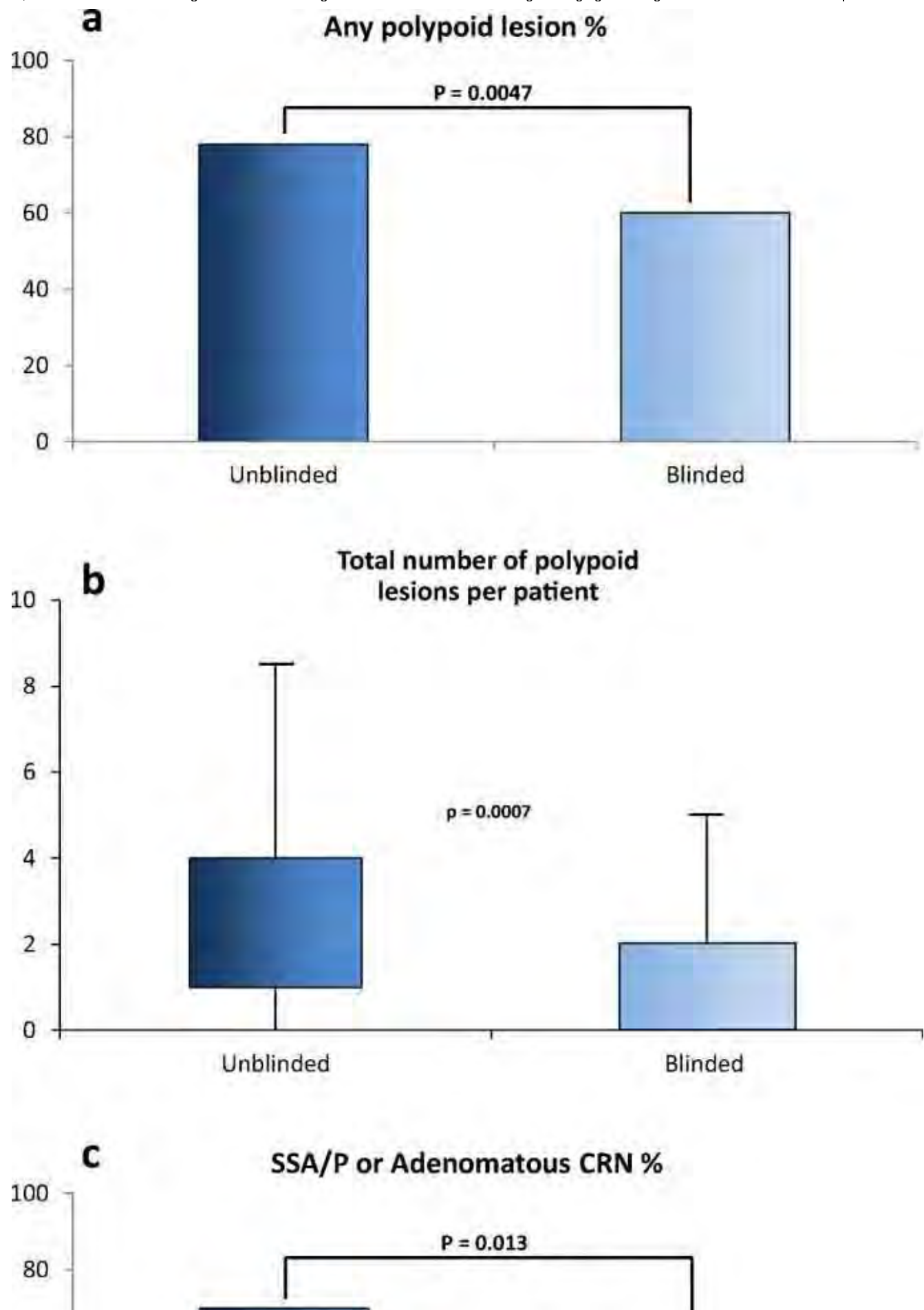
In patients with positive MT-sDNA testing who undergo diagnostic colonoscopy, our data show that approximately two-thirds will be found to have at least one neoplastic lesion [[44](#)]. Among patients with CRN, 40% will have at least one advanced lesion (defined as CRC, adenomas, or SSPs with at least one of the following characteristics: ≥ 1 cm in size, high-grade dysplasia, or having $\geq 25\%$ villous elements), and ~ 1% will have CRC. SSPs will be detected in about half of patients with positive tests.

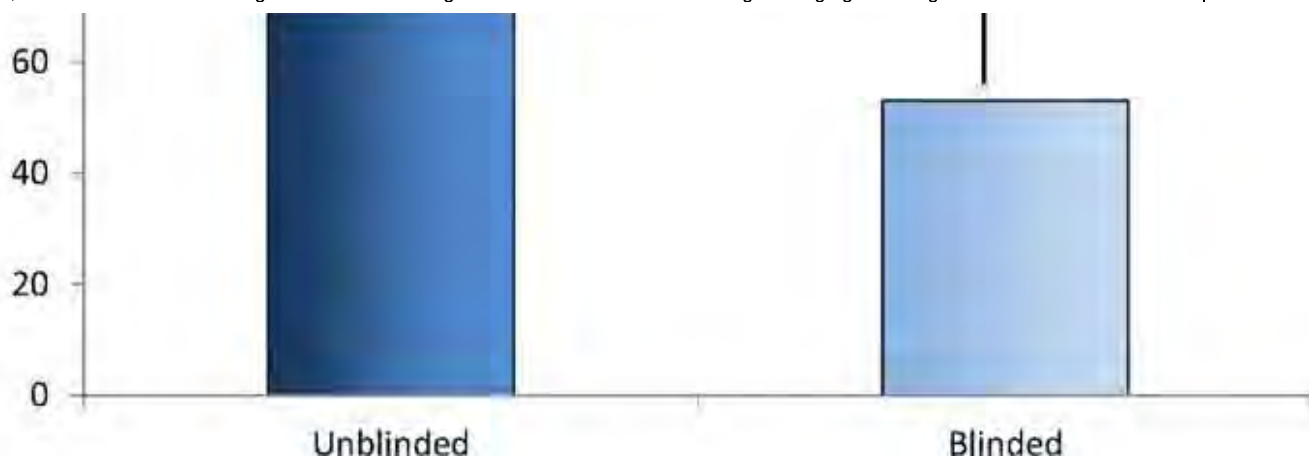
Importantly, MT-sDNA testing appears to perform well in detecting right-sided (proximal) neoplasia. In our multi-site cohort of patients with positive MT-sDNA tests, over 50% of patients overall (and 80% of those with neoplastic lesions) were found to have proximal CRN [[44](#)]. The reason for the high yield of proximal CRC is likely multifactorial, although

improved detection resulting from the multiple markers in MT-sDNA offers one potential explanation. Right-sided lesions tend to be flat and less likely to bleed [46], and exfoliation of DNA markers is more consistent throughout the colon [47]; therefore, MT-sDNA has an advantage over other non-invasive methods for the detection of right-sided neoplasia.

These point estimates for positive predictive value appear higher than reported in the pre-FDA approval trial, where advanced lesions were found in 24% of those with positive MT-sDNA tests [39]. This is likely in part due to underlying differences in the populations being studied, or in differences in those opting to use the test in clinical practice versus the clinical trial setting. However, there are several other factors that likely contribute to the high yield of MT-sDNA for CRN, both proximal and distal. First, by screening an average-risk population with MT-sDNA and performing further testing only in those with positive results, this creates an enriched population in whom the pre-test probability of CRN at diagnostic colonoscopy is increased. Furthermore, studies have shown that endoscopist knowledge of a positive MT-sDNA test significantly increases quality metrics such as ADR at diagnostic colonoscopy compared with blinded endoscopists, likely related to longer withdrawal time and increased attention to subtle lesions (Fig. 1) [48]. Moreover, recent data suggest that while improvement in quality and yield of diagnostic colonoscopy is observed in all endoscopists in a large academic practice, the greatest improvement in polyp detection is seen in gastroenterologists with lower baseline detection rates [49].

Fig. 1





The **a** proportion of patients with polyps, **b** number of polyps per patient, and **c** proportion of patients with sessile serrated and adenomatous polyps found at diagnostic colonoscopy after positive multi-target stool DNA test is higher in unblinded endoscopists compared with blinded endoscopists. Used with permission from Elsevier.

“False positive” MT-sDNA tests

A common question among physicians and other health care providers is what should be done in the setting of a positive MT-sDNA test followed by a negative colonoscopy (a “false positive” MT-sDNA test). However, before addressing this question, negative colonoscopy needs to be more clearly defined. The original studies investigating MT-sDNA testing labeled true positive tests as those in which only advanced neoplasia or CRC was detected [39]. However, we have shown that approximately 40% of lesions detected by MT-sDNA are non-advanced [44]; a positive MT-sDNA test in these situations is therefore unlikely to be biologically false, but rather to have detected exfoliated DNA from a neoplasm not meeting arbitrary size-based criteria. With this in mind, knowing that 14% of all MT-sDNA tests are positive and that 67% of these patients have CRN found at diagnostic colonoscopy, less than 5% of all patients screened with MT-sDNA would have no polyps found at colonoscopy ($14.1\% \times (100\% - 67\%) = 4.7\%$). Even when including only advanced CRN and CRC, 10% ($14.1\% \times (100\% - 28\%)$) would have false positive results.

In the clinical situation of a false positive MT-sDNA test, the next step is to ensure that a diagnostic colonoscopy of sufficient quality was performed. In clinical trials of MT-sDNA, only those with high-quality colonoscopy, defined as having good or excellent bowel preparation, photographic evidence of cecal intubation, and withdrawal time of ≥ 6 min, were included in primary study outcomes [21, 39, 40, 50]. Assuming a high-quality

colonoscopy, current guidelines suggest that no further testing be performed [11]. These recommendations are based on several studies that found no increased risk of CRC in this specific population. One such study prospectively evaluated 30 patients with initial false positive MT-sDNA testing (evaluated by both colonoscopy and esophagogastroduodenoscopy (EGD)), who underwent repeat MT-sDNA and colonoscopy/EGD 1 year later. In this cohort, only two advanced lesions were found on repeat testing; however, information on initial colonoscopy quality was not reported [51]. Another study retrospectively analyzed approximately 1000 patients with either false positive or true negative MT-sDNA tests, and found no increased rate of aerodigestive cancers in a per-protocol analysis at a median follow-up of 4 years [52]. More recently, a multi-site retrospective cohort study performed by Berger et al. evaluated approximately 1200 patients with negative colonoscopy over a median of 5 years, and stratified patients by concordant (negative) or discordant (positive) MT-sDNA results. They found no increased incidence of aerodigestive cancers in either group, with rates that were equivalent to the general population based on Surveillance, Epidemiology, and End Results (SEER) Program data [53]. Therefore, based on currently available evidence, clinicians should not recommend repeat colonoscopy, EGD, or other further testing in patients with positive MT-sDNA, a negative high-quality diagnostic colonoscopy, and no localizing signs or symptoms that would mandate evaluation.

Future directions

Even with increasing utilization of MT-sDNA and the emergence of post-approval performance data, several opportunities for further study remain. While the current recommended testing interval for MT-sDNA is 3 years, this is based on manufacturer's recommendations and computer modeling, and further investigation is needed into MT-sDNA performance with different testing frequencies. Additional study of patient adherence to both the testing schedule and diagnostic colonoscopy in other populations would also help to further inform test performance in the real world.

MT-sDNA testing may benefit additional patient populations. In response to the ACS recommendation to begin screening for CRC at age 45, studies conducted in archival tissue specimens demonstrated that MT-sDNA markers in primary CRC tumors among those 45–49 years old at diagnosis are similar to those seen in tumors of patients aged 50–64 [54]; methylated DNA markers were all substantially higher than observed in control tissues

across these age ranges. As a result of these data and those expected from a clinical trial utilizing MT-sDNA in patients age 45–49 ([ClinicalTrials.gov](https://clinicaltrials.gov/ct2/show/study/NCT03728348) Identifier: NCT03728348), the FDA recently expanded the label for MT-sDNA to include those age 45–84.

The FDA's current labeling excludes those at increased risk for CRC due to conditions such as inflammatory bowel disease (IBD), personal history of advanced CRN, family history of early CRC, and signs of blood loss, among others. Initial data from our group's evaluation of off-label use of MT-sDNA testing (~10% of tests) in patients with these risk factors show that test performance is similar to those at average risk [55]. Furthermore, in a research study of 192 patients with IBD undergoing MT-sDNA testing for surveillance, the test demonstrated similar sensitivity for both CRC and CRN when compared with the general population [56]. However, further study is needed before extending MT-sDNA use into these increased-risk populations.

Finally, MT-sDNA may have important utility as an interval test following apparently negative colonoscopy. In our group's analysis of data in patients undergoing MT-sDNA testing within 5 years of a negative colonoscopy, the yield of all CRN and most CRN subtypes was the same as in those undergoing testing 10 or more years after colonoscopy. When these patients were further stratified by whether their prior colonoscopy was complete or incomplete (e.g., poor prep, no cecal intubation), neoplastic yield remained similar between groups [57]. Advanced neoplasms were found in 16–17% of those screened early with MT-sDNA; these findings could not be explained by known elevations in baseline risk. Taken together, these findings suggest that lesions are likely being missed at screening colonoscopies. MT-sDNA should therefore be studied as a tool between colonoscopies to identify those in which polyps were likely to have been missed, and thereby reduce the risk of interval CRC.

Conclusion

MT-sDNA is a stool-based screening test with high sensitivity and specificity for both CRN and CRC. As its use has increased over the past 5 years after FDA approval, it has attracted a large population of previously non-adherent patients to screening. A majority of patients with a positive test are found to have CRN at diagnostic colonoscopy, with most harboring right-sided lesions. While further investigation is necessary into additional potential uses,

MT-sDNA has firmly cemented itself as a convenient, non-invasive, and effective tool for widespread screening in the ongoing fight against CRC.

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Ethics declarations

Conflict of Interest

Mayo Clinic and Exact Sciences (Madison, WI) own intellectual property under which Dr. Kisiel is an inventor and may receive royalties in accordance with Mayo Clinic policy. Jason D. Eckmann declares that he has no conflict of interest. Derek W. Ebner that he has no conflict of interest.

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Appendix ECKMANN02



Multi-Target Stool DNA Testing for Colorectal Cancer Screening: Emerging Learning on Real-world Performance

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Abstract

Purpose of Review Multi-target stool DNA (MT-sDNA) was approved in 2014 for use in screening average-risk patients for colorectal cancer (CRC). Here, we highlight recent literature from post-market studies to provide an update on clinical use and utility not possible from pre-approval studies.

Recent Findings MT-sDNA has been included in major society guidelines as an option for colorectal cancer screening, and has seen exponentially increasing use in clinical practice. MT-sDNA appears to be attracting new patients to CRC screening, and patient adherence to diagnostic colonoscopy after a positive MT-sDNA test is high. Approximately two-thirds of these patients are found to have colorectal neoplasia (CRN), 80% of whom have at least one right-sided lesion; 1 in 3 will have advanced CRN. High yield of CRN is due not only to post-screening increase in probability but also likely improved endoscopist attention. In those with a negative high-quality colonoscopy after positive MT-sDNA test (“false positive MT-sDNA”), further interventions do not appear to be necessary.

Summary MT-sDNA is a promising tool to improve rates and quality of CRC screening. Further investigation should examine MT-sDNA performance in populations at increased risk for CRC, and as an interval test after colonoscopy to detect potentially missed lesions.

Introduction

Colorectal cancer (CRC) is a major cause of cancer-related death in the USA [1], with approximately 150,000 new diagnoses and 50,000 deaths from CRC projected in 2019 alone [2]. With widespread screening initiatives to detect CRC and pre-cancerous polyps, CRC-related morbidity and mortality have decreased in recent years [2]. The US Preventive Services Task Force (USPSTF) recommends screening for CRC in average-risk adults between the ages of 50 and 75 [3], while more recent recommendations from the American Cancer Society (ACS) suggest initiating screening at age 45 in light of the increasing rate of CRC diagnosed in younger patients [4•,5,6,7,8]. However, despite these guidelines and extensive public health initiatives, approximately one-third of screen-eligible adults in the USA remain unscreened [9, 10•].

Several methods are recommended for CRC screening [3, 4•, 11•]. Colonoscopy is the most widely used modality in the USA [9] and permits simultaneous diagnostic and therapeutic intervention for colorectal neoplasia (CRN). However, screening colonoscopy has several notable drawbacks, including reduced sensitivity in the proximal colon [12–18•] and inter-operator variability in colonoscopy quality [19–22]. Furthermore,

its invasive nature; risk of complications; and the inconvenience of preparation, sedation, and time away from work are recognized barriers to patients considering screening [23]. Several non-invasive screening modalities are endorsed as an alternative to colonoscopy for use in average-risk patients. Guaiac-based fecal occult blood testing (gFOBT) has the strongest body of evidence demonstrating a reduction in CRC-related mortality [24–27]. However, dietary and medication interactions, poor performance in the proximal colon, and issues with adherence to the recommended yearly screening interval have led to decreased utilization of this test [28–30]. The introduction of fecal immunochemical testing (FIT) addressed some of these limitations, with FIT included as a first-line option in major society screening guidelines [3, 11•]. A recent meta-analysis suggested moderate-to-high sensitivity and specificity for CRC and advanced adenoma (AA) in a single-use setting, but performance varied by assay and positivity cutoff [31]. Concerns remain regarding reduced sensitivity for right-sided and sessile serrated precursor lesions [28, 29, 32•] in addition to persistent poor adherence to required annual testing [30, 33–36].

Multi-target stool DNA testing

Multi-target stool DNA (MT-sDNA) is a newer stool-based screening modality, with one commercial product available to patients in the USA (Cologuard™, Exact Sciences, Madison, WI). This test detects abnormal DNA markers (aberrantly methylated *BMP3* and *NDRG4*, mutant *KRAS*) shed from neoplastic cells in the gastrointestinal tract. Another marker, *β-actin*, serves as a control to ensure adequate quantity of DNA in the sample, while an immunochemical assay for hemoglobin detects evidence of gastrointestinal bleeding. These components are measured quantitatively and incorporated into a logistic regression algorithm containing mathematical interaction variables for each marker. These interaction terms result in numerous marker combinations that can trigger a pre-specified threshold score, resulting in a positive or negative test result that is more than the sum of its individual components [37].

MT-sDNA testing is performed by the patients in the comfort of their own home. Following a provider order, a manufacturer-based patient navigation office contacts the patient and arranges sample collection via a courier-delivered kit [38]. This system also provides patient education and test completion reminders which have been shown to result in adherence to testing of over 70% in Medicare beneficiaries [38]. Following delivery of the MT-sDNA kit, patients

collect a whole stool sample, swab a portion to collect the FIT sample, add a preservative buffer to the whole stool, seal the container, and mail the samples in one box back to the laboratory for analysis.

MT-sDNA testing was approved for screening in average-risk adults by the US Food and Drug Administration (FDA) in 2014, following a large screen-setting trial by Imperiale et al. (DeeP-C trial), comparing MT-sDNA and FIT. Overall, the sensitivity of MT-sDNA for detecting CRC in this study was 92% (vs 74% for FIT, $p = 0.002$), and 42% for advanced pre-cancerous lesions (vs 24% for FIT, $p < 0.001$) [39]. A second study performed in an Alaska Native cohort confirmed these findings, and re-demonstrated increasing sensitivity by adenoma size, with 80% sensitivity for lesions > 3 cm, and 100% sensitivity for CRC [40••]. Specificity was 87–94% across these two studies, and increased in younger patients (Table 1) [39, 40••]. MT-sDNA appears to have similar performance for the detection of both conventional adenomas and sessile serrated polyps (SSPs). Large SSPs and hyperplastic polyps (≥ 1 cm) were included in the sensitivity estimate for advanced pre-cancerous lesions in the DeeP-C study [39]. For greater clarity, a post hoc analysis of DeeP-C data was performed on all lesions ≥ 1 cm with histology confirmed for SSP. The median size was 1.2 cm (range 1–3), and the majority (28/29 (93%)) were located proximal to the splenic flexure; only one SSP contained dysplasia. In this setting, MT-sDNA had a sensitivity of 55% (95% CI, 36–74%) and a specificity of 91% [41]. MT-sDNA performance characteristics for the individual endpoint category of large hyperplastic polyp have not been estimated and traditional serrated adenomas were not identified by screening colonoscopy in the DeeP-C study; these endpoints are deserving of future study.

Population modeling studies performed as part of a USPSTF review suggest that programmatic use of MT-sDNA would result in fewer lifetime colonoscopies than FIT and a stronger benefit-to-harms ratio than colonoscopy (Table 2) [3]. As a result of the above findings, MT-sDNA testing was subsequently endorsed by the USPSTF in their 2016 guidelines for CRC screening [3]. Utilization of MT-sDNA has been increasing exponentially since then, with nearly 3 million screens performed to date [42].

Table 1. Multi-target stool DNA demonstrates higher specificity in younger patients with normal colonoscopy

Patient Population	Specificity
Imperiale et al [39]	
No CRC or advanced adenoma*, all ages	86.6%
Negative colonoscopy, all ages	90%
No CRC or advanced adenoma*, age 50–64	91.5%
Negative colonoscopy, age 50–64	94%
Redwood et al [40]	
No CRC or advanced adenoma*, all ages	91%
Negative colonoscopy, all ages	93%

*Patients with non-advanced lesions, hyperplastic polyps and other findings (e.g. diverticulitis) included in specificity calculation

Table 2. Screening with multi-target stool DNA testing results in fewer lifetime colonoscopies than FIT, and has a superior benefit-to-harm ratio compared to colonoscopy

Screening Modality	Life years gained*	CRC deaths averted*	Complications (GI/CV)*	Colonoscopies*	Life years gained/ complication	Life years gained/ colonoscopy	CRC deaths averted/ complication	CRC deaths averted colonoscopy
MT-sDNA	226	20	9	1714	25	0.13	2.2	0.01
FIT	244	22	10	1757	24	0.14	2.2	0.01
Colonoscopy	270	24	15	4049	18	0.07	1.6	0.006

Adapted from: Bibbens-Domingo, et al. [3]

*Per 1000 persons screened

Outcomes in patients undergoing multi-target stool DNA testing

Since the introduction of MT-sDNA in 2014, data have begun to emerge regarding its “real-world” performance. MT-sDNA appears to be attracting new patients to screening. One recent study demonstrated high utilization of MT-sDNA in almost 400 previously non-adherent average-risk Medicare patients, with 51% harboring advanced neoplastic lesions; fewer than 20% had no neoplastic findings at diagnostic colonoscopy [43•]. Furthermore, we recently presented data from a cohort of over 2000 MT-sDNA-positive patients showing that approximately 25% had never been screened for CRC and were > 10 years overdue [44]. While there may be many reasons for patients to go without screening, these findings suggest that MT-sDNA is attracting previously non-adherent patients, a critical outcome in the ongoing long-term fight to reduce CRC-attributable mortality.

Of all patients screened with MT-sDNA, 14–16% will have positive results [39, 44, 45]. Following a positive MT-sDNA test, it is imperative that patients undergo diagnostic colonoscopy to evaluate for CRN. While previous estimates of adherence to diagnostic colonoscopy ranged from 75 to 96% [43•, 45], our recent analysis showed a diagnostic colonoscopy completion rate of ~90% in a large multi-practice setting [44]. Reasons for non-adherence to recommended diagnostic colonoscopy vary, but are commonly related to a wish to avoid colonoscopy, in addition to medical comorbidities that preclude sedation and procedural intervention [45]. These factors illustrate the importance of both provider and patient education prior to pursuing MT-sDNA testing in order to ensure selection of appropriate patients for screening with this test.

In patients with positive MT-sDNA testing who undergo diagnostic colonoscopy, our data show that approximately two-thirds will be found to have at least one neoplastic lesion [44]. Among patients with CRN, 40% will have at least one advanced lesion (defined as CRC, adenomas, or SSPs with at least one of the following characteristics: ≥ 1 cm in size, high-grade dysplasia, or having ≥ 25% villous elements), and ~1% will have CRC. SSPs will be detected in about half of patients with positive tests.

Importantly, MT-sDNA testing appears to perform well in detecting right-sided (proximal) neoplasia. In our multi-site cohort of patients with positive MT-sDNA tests, over 50% of patients overall (and 80% of those with neoplastic lesions) were found to have proximal CRN [44]. The reason for the high yield of proximal CRC is likely multifactorial, although improved detection resulting from the multiple markers in MT-sDNA offers one potential explanation. Right-sided lesions tend to be flat and less likely to bleed [46], and exfoliation of DNA markers is more consistent throughout the colon [47]; therefore, MT-sDNA has an advantage over other non-invasive methods for the detection of right-sided neoplasia.

These point estimates for positive predictive value appear higher than reported in the pre-FDA approval trial, where advanced lesions were found in 24% of those with positive MT-sDNA tests [39]. This is likely in part due to underlying differences in the populations being studied, or in differences in those opting to use the test in clinical practice versus the clinical trial setting. However, there are several other factors that likely contribute to the high yield of MT-sDNA for CRN, both proximal and distal. First, by screening an average-risk population with MT-sDNA and performing further testing only in those with positive results, this creates an enriched population in whom the pre-test probability of CRN at diagnostic colonoscopy is increased. Furthermore, studies have shown that endoscopist knowledge of a positive MT-sDNA test significantly increases quality metrics such as ADR at diagnostic colonoscopy compared with blinded endoscopists, likely related to longer withdrawal time and increased attention to subtle lesions (Fig. 1) [48••]. Moreover, recent data suggest that while improvement in quality and yield of diagnostic colonoscopy is observed in all endoscopists in a large academic practice, the greatest improvement in polyp detection is seen in gastroenterologists with lower baseline detection rates [49].

“False positive” MT-sDNA tests

A common question among physicians and other health care providers is what should be done in the setting of a positive MT-sDNA test followed by a negative colonoscopy (a “false positive” MT-sDNA test). However, before addressing this question, negative colonoscopy needs to be more clearly defined. The original studies investigating MT-sDNA testing labeled true positive tests as those in which only advanced neoplasia or CRC was detected [39]. However, we have shown that approximately 40% of lesions detected by MT-sDNA are non-advanced [44]; a positive MT-sDNA test in these situations is therefore unlikely to be biologically false, but rather to have detected exfoliated DNA from a neoplasm not meeting arbitrary size-based criteria. With this in mind, knowing that 14% of all MT-sDNA tests are positive and that 67% of these patients have CRN found at diagnostic colonoscopy, less than 5% of all patients screened with MT-sDNA would have no polyps found at colonoscopy ($14.1\% \times (100\% - 67\%) = 4.7\%$). Even when including only advanced CRN and CRC, 10% ($14.1\% \times (100\% - 28\%)$) would have false positive results.

In the clinical situation of a false positive MT-sDNA test, the next step is to ensure that a diagnostic colonoscopy of sufficient quality was performed. In clinical trials of MT-sDNA, only those with high-quality colonoscopy, defined as having

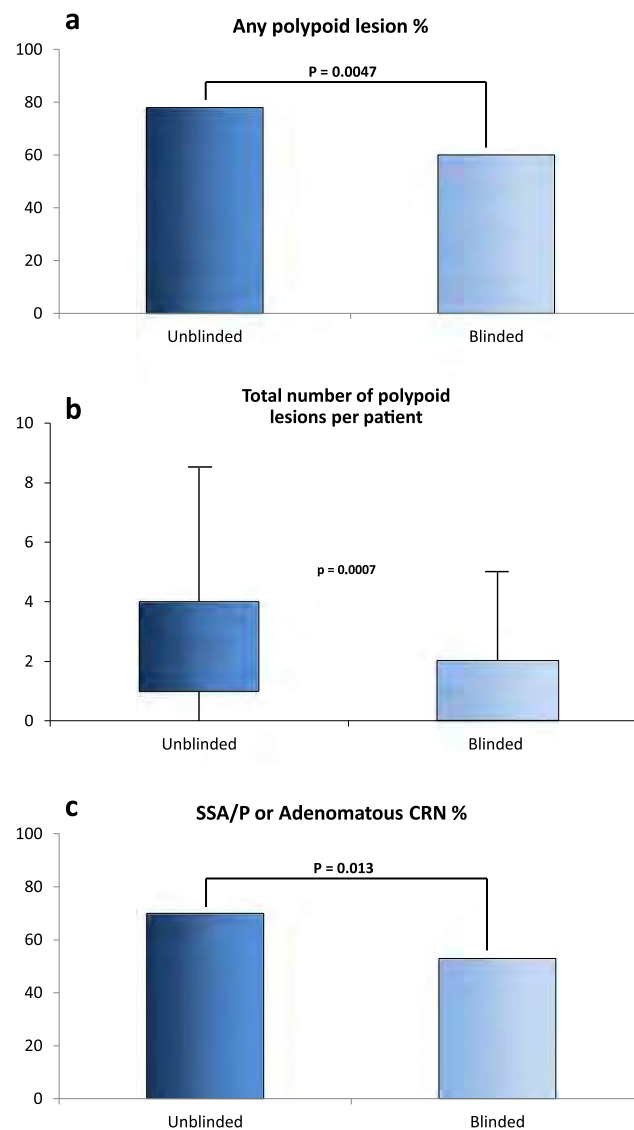


Fig. 1. The **a** proportion of patients with polyps, **b** number of polyps per patient, and **c** proportion of patients with sessile serrated and adenomatous polyps found at diagnostic colonoscopy after positive multi-target stool DNA test is higher in unblinded endoscopists compared with blinded endoscopists. Used with permission from Elsevier.

good or excellent bowel preparation, photographic evidence of cecal intubation, and withdrawal time of ≥ 6 min, were included in primary study outcomes [21, 39, 40••, 50]. Assuming a high-quality colonoscopy, current guidelines suggest that no further testing be performed [11•]. These recommendations are based on several studies that found no increased risk of CRC in this specific population. One such study prospectively evaluated 30 patients with initial false positive MT-sDNA testing (evaluated by both colonoscopy and esophagogastroduodenoscopy (EGD)), who underwent repeat MT-sDNA and colonoscopy/EGD 1 year later. In this cohort, only two advanced lesions were found on repeat testing; however, information on initial colonoscopy quality was not reported [51•]. Another study

retrospectively analyzed approximately 1000 patients with either false positive or true negative MT-sDNA tests, and found no increased rate of aerodigestive cancers in a per-protocol analysis at a median follow-up of 4 years [52•]. More recently, a multi-site retrospective cohort study performed by Berger et al. evaluated approximately 1200 patients with negative colonoscopy over a median of 5 years, and stratified patients by concordant (negative) or discordant (positive) MT-sDNA results. They found no increased incidence of aerodigestive cancers in either group, with rates that were equivalent to the general population based on Surveillance, Epidemiology, and End Results (SEER) Program data [53•]. Therefore, based on currently available evidence, clinicians should not recommend repeat colonoscopy, EGD, or other further testing in patients with positive MT-sDNA, a negative high-quality diagnostic colonoscopy, and no localizing signs or symptoms that would mandate evaluation.

Future directions

Even with increasing utilization of MT-sDNA and the emergence of post-approval performance data, several opportunities for further study remain. While the current recommended testing interval for MT-sDNA is 3 years, this is based on manufacturer's recommendations and computer modeling, and further investigation is needed into MT-sDNA performance with different testing frequencies. Additional study of patient adherence to both the testing schedule and diagnostic colonoscopy in other populations would also help to further inform test performance in the real world.

MT-sDNA testing may benefit additional patient populations. In response to the ACS recommendation to begin screening for CRC at age 45, studies conducted in archival tissue specimens demonstrated that MT-sDNA markers in primary CRC tumors among those 45–49 years old at diagnosis are similar to those seen in tumors of patients aged 50–64 [54]; methylated DNA markers were all substantially higher than observed in control tissues across these age ranges. As a result of these data and those expected from a clinical trial utilizing MT-sDNA in patients age 45–49 ([ClinicalTrials.gov](https://clinicaltrials.gov/ct2/show/study/NCT03728348) Identifier: NCT03728348), the FDA recently expanded the label for MT-sDNA to include those age 45–84.

The FDA's current labeling excludes those at increased risk for CRC due to conditions such as inflammatory bowel disease (IBD), personal history of advanced CRN, family history of early CRC, and signs of blood loss, among others. Initial data from our group's evaluation of off-label use of MT-sDNA testing (~10% of tests) in patients with these risk factors show that test performance is similar to those at average risk [55]. Furthermore, in a research study of 192 patients with IBD undergoing MT-sDNA testing for surveillance, the test demonstrated similar sensitivity for both CRC and CRN when compared with the general population [56•]. However, further study is needed before extending MT-sDNA use into these increased-risk populations.

Finally, MT-sDNA may have important utility as an interval test following apparently negative colonoscopy. In our group's analysis of data in patients undergoing MT-sDNA testing within 5 years of a negative colonoscopy, the yield of all CRN and most CRN subtypes was the same as in those undergoing testing 10 or more years after colonoscopy. When these patients were further stratified by whether their prior colonoscopy was complete or incomplete (e.g.,

poor prep, no cecal intubation), neoplastic yield remained similar between groups [57]. Advanced neoplasms were found in 16–17% of those screened early with MT-sDNA; these findings could not be explained by known elevations in baseline risk. Taken together, these findings suggest that lesions are likely being missed at screening colonoscopies. MT-sDNA should therefore be studied as a tool between colonoscopies to identify those in which polyps were likely to have been missed, and thereby reduce the risk of interval CRC.

Conclusion

MT-sDNA is a stool-based screening test with high sensitivity and specificity for both CRN and CRC. As its use has increased over the past 5 years after FDA approval, it has attracted a large population of previously non-adherent patients to screening. A majority of patients with a positive test are found to have CRN at diagnostic colonoscopy, with most harboring right-sided lesions. While further investigation is necessary into additional potential uses, MT-sDNA has firmly cemented itself as a convenient, non-invasive, and effective tool for widespread screening in the ongoing fight against CRC.

Compliance with Ethical Standards

Conflict of Interest

Mayo Clinic and Exact Sciences (Madison, WI) own intellectual property under which Dr. Kisiel is an inventor and may receive royalties in accordance with Mayo Clinic policy. Jason D. Eckmann declares that he has no conflict of interest. Derek W. Ebner that he has no conflict of interest.

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A study of 30 patients with positive MT-sDNA testing and negative follow-up colonoscopy, who were subsequently evaluated with repeat MT-sDNA and endoscopy. Only 2 patients had persistently positive MT-sDNA with advanced neoplasia on repeat testing; no colorectal cancer was found.

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This study of ~1000 patients compared patients with “false positive” MT-sDNA testing to those with “true negative” testing. Results showed no increase in likelihood of aerodigestive cancer in the false-positive group over 4 years, with incidence of aerodigestive cancers in the false positive group equivalent to SEER estimates in the general population.

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This retrospective study of 1200 patients undergoing MT-sDNA testing with either concordant (negative MT-sDNA and colonoscopy) or discordant (positive MT-sDNA and negative colonoscopy) results showed an increase in incidence of aerodigestive cancers in the discordant group compared with the concordant group. Rates of aerodigestive cancers were statistically similar in the discordant group compared with SEER data for the general population.

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