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<https://doi.org/10.1016/j.humpath.2019.04.015>

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“Histology of colorectal adenocarcinoma with double somatic mismatch-repair mutations is indistinguishable from those caused by Lynch syndrome”—reply



Dear Editor,

Thank you to Drs Demetra Georgiou, Vicki Kiesel, Angela F. Brady, and Kevin Monahan for sharing their experiences with Lynch-like syndrome in the United Kingdom in response to our article [1]. We appreciate the difficulty of knowing how to manage patients with unexplained mismatch repair deficiency (dMMR) appropriately and found the authors' data very interesting. One detail we would like to note is the difference between management recommendations in the United Kingdom and the United States. The authors say that “in the UK, [patients with unexplained dMMR] and their

first degree relatives are eligible for 2 yearly colonoscopy as per LS guidelines [2].” In the United States, current management guidelines from the National Comprehensive Cancer Network's Genetic/Familial High-Risk Assessment: Colorectal version 1.2018 state that “if double somatic [MMR] mutations are identified [via tumor sequencing] or if [tumor sequencing] does not help clarify the result, *it is recommended that these patients and their close relatives be managed based on their family history and NOT as if they have Lynch syndrome.* Regardless of the results of tumor sequencing, these patients and their close relatives should be managed based on their family history and NOT as if they have Lynch syndrome unless their family history warrants it” [3]. One of the biggest factors contributing to the shift from managing dMMR patients as if they had Lynch syndrome without an identified germline mutation to managing them based on their family history was the discovery that 45%-69% of patients with unexplained dMMR actually have double somatic mutations, not a missed germline mutation [4-7].

We would also like to update that although we found no histological differences between colorectal tumors due to Lynch syndrome or double somatic MMR mutations, we have found clinical differences between the 2 patient groups. We recently published the characteristics that differentiate colorectal cancer patients with Lynch syndrome from those with double somatic MMR mutations or unexplained dMMR [8]. We found that patients with double somatic MMR mutations were more likely to have loss of MLH1/PMS2 proteins in their tumor, and patients with Lynch syndrome were more likely to have a family history of Lynch-associated cancers and/or a personal history of multiple primary cancers [8]. We hope that our new publication will help to provide further insight into a complex clinical situation.

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<https://doi.org/10.1016/j.humpath.2019.04.017>

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