

Germline lynch syndrometesting: Are we there yet?

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INTRODUCTION

- Lynch syndrome is the most common of all hereditary cancer predisposition syndromes accounting for approximately 3-5% of all colorectal cancers (CRC) and endometrial cancers (EC).¹
- Identifying these patients helps predict prognosis, optimize treatment, plan future surveillance, and family screening.
- Currently, the most cost-effective and best screening strategy is screening patients with Mismatch repair (MMR) deficiency/Microsatellite instability (MSI) for further genetic testing.

OBJECTIVE

- To evaluate MMR/MSI across all tumors, along with uptake of germline testing for Lynch syndrome in India.

METHODS

- This is a single centre cross sectional study in which MMR and MSI status was evaluated across all tumours, along with uptake of germline testing for Lynch syndrome from 2021 to 2015.
- Mismatch repair status was assessed by testing the somatic tumour by either immunohistochemistry (IHC), for MLH1, PMS2, MSH2 & MSH6 (on Ventana platform), or MSI by fragment length analysis using Promega kit (SeqStudio/Applied biosystems).
- The germline testing was performed on peripheral blood using custom DNA panel for the four MMR genes (*MLH1*, *PMS2*, *MSH2*, *MSH6*), interrogating single nucleotide variants and indels by Next generation sequencing (Ion Torrent).

RESULTS

- 288 (19.9%) of 1447 cases tested for MMR/MSI, revealed MMR deficiency (dMMR)/MSI.
- 198 of the 288 dMMR/MSI cases were CRCs (68.8%), followed by Endometrial cancers (21.9%).
- Germline lynch testing was done in 77 of 288 cases (26.7%).
- 51 of 77 (66.2%) cases tested, revealed a germline mutation, with the highest frequency in *MLH1* gene (60.8%).
- 60 of 288 cases (20.8%), did not undergo germline testing, had insurance coverage (p=0.0005).
- 131 of the 198 (66.1%) dMMR/MSI CRCs were right sided.
- 42/61 CRCs tested (68.9%), revealed a germline mutation, most commonly implicating *MLH1* gene (66.7%), (p<0.0001).
- Most germline CRCs were right sided (64.3%), (p=0.2561).
- Germline CRCs showed a higher frequency of node negative stage (86.1%), (p=0.0195).
- 25 of 26 germline CRCs with *MLH1/PMS2* mutation, also revealed *MLH1/PMS2* loss on IHC testing (p<0.0001).
- 45 of 63 (71.4%) dMMR/MSI ECs, were low grade, while 44 (69.8%) had a node negative disease.
- 8 of 14 (57.1%) dMMR/MSI ECs tested, were germline mutant.

Fig 1: Distribution of dMMR/MSI cases according to cancer type

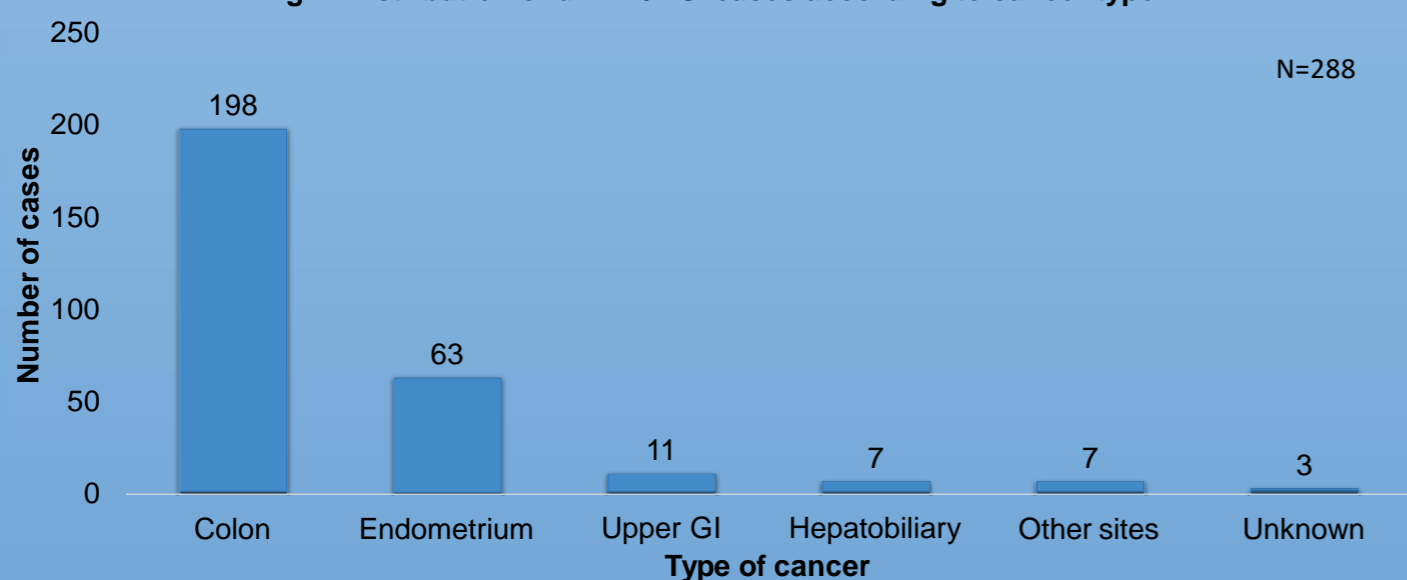


Fig 2: Extent of germline testing

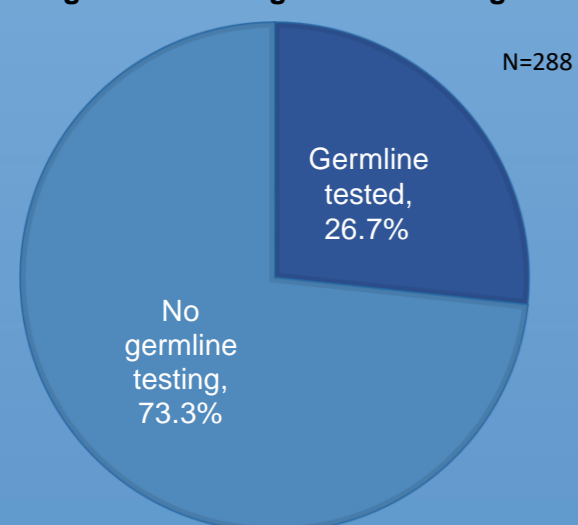


Fig 3: Extent of germline mutation

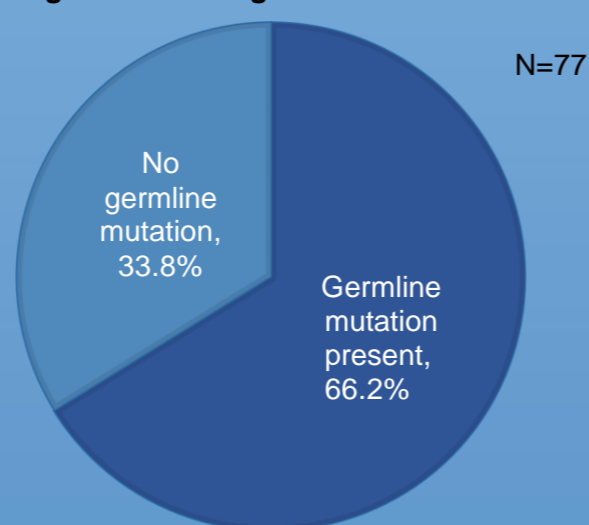


Fig 4: Distribution of genetic mutations

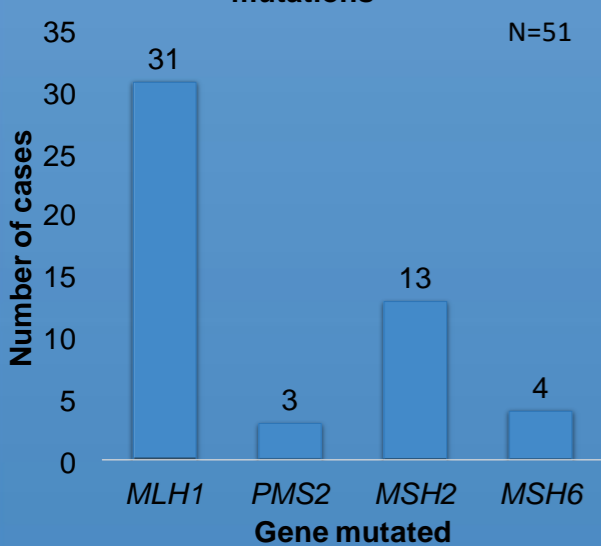


Fig 5: Nodal status in dMMR/MSI cancers

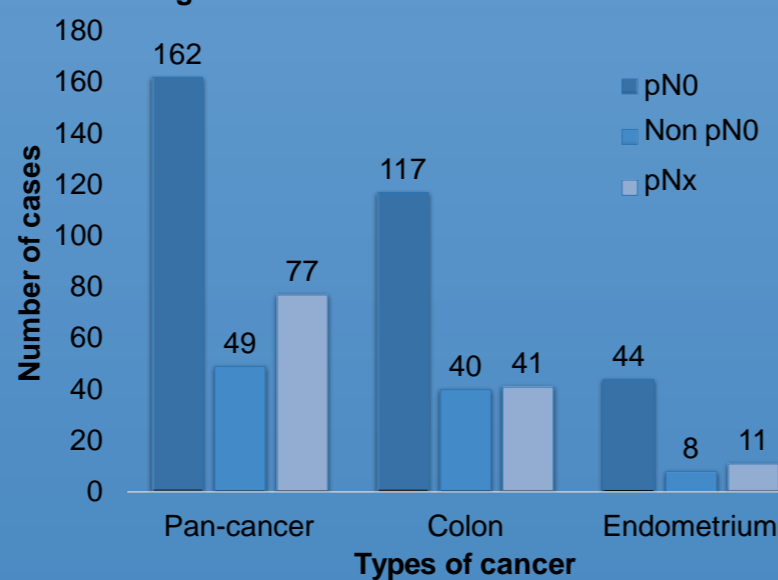


Fig 6: Nodal status in germline cancers

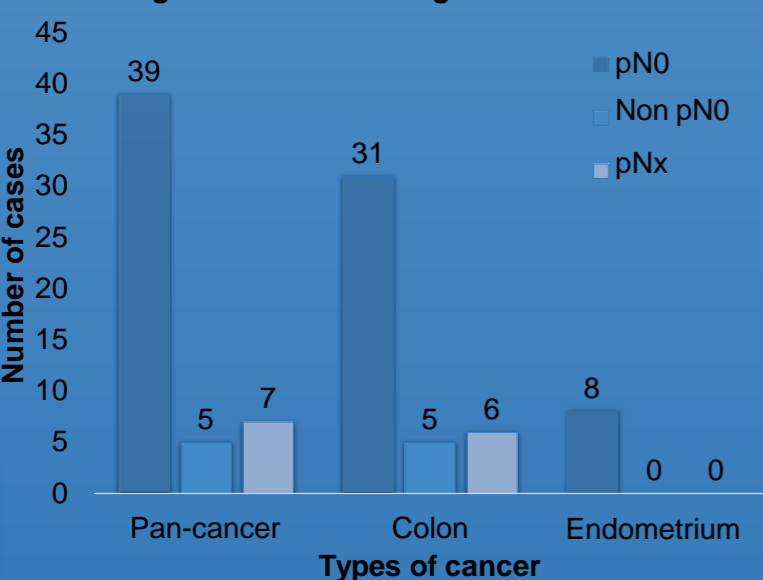
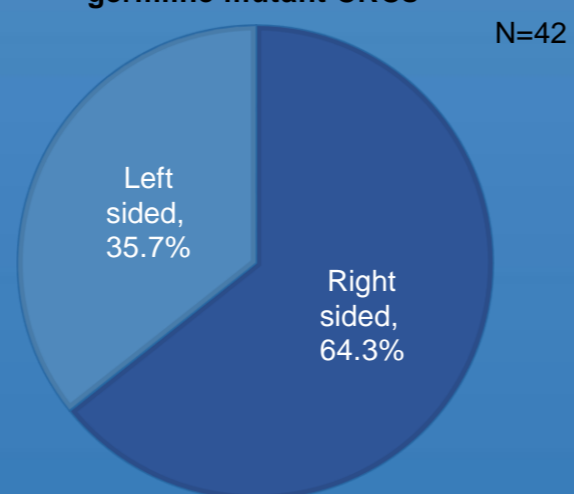


Fig 7: Anatomical distribution of germline mutant CRCs



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DISCUSSION

- Guidelines recommend that all CRCs, and lately all ECs, be assessed for dMMR/MSI, followed by germline testing, in order to increase identification of patients with Lynch syndrome.
- In the present study, the referral rate of genetic testing for Lynch syndrome was 26.7%, compared to an Australian study that revealed a referral rate of 11% in dMMR CRCs.²
- Meanwhile, Muller et al, reported that the highest referral rate was seen in Non-Hispanic whites (21.2%), while it was as low as 10.9% in Hispanic patients, in the United states.³
- In contrast, a study from Iceland revealed that, genetic testing was done in almost 80% cases.⁴

CONCLUSION

- Germline testing helps in early identification of recurrence & in screening of kindred. In our experience, the uptake of germline testing is low in India, owing to socioeconomic restraints.
- This unmet need can be met by clinicians and molecular pathologists with adequate counselling, helping eradicate the associated social stigmata.