



Article title: Review of Hereditary non-polyposis colorectal cancer

Authors: Maneesh Singh [1]

Affiliations: kansas city university [1]

Orcid ids: 0000-0003-4859-2546[1]

Contact e-mail: meshusingh123@gmail.com

License information: This work has been published open access under Creative Commons Attribution License <http://creativecommons.org/licenses/by/4.0/>, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited. Conditions, terms of use and publishing policy can be found at <https://www.scienceopen.com/>.

Preprint statement: This article is a preprint and has not been peer-reviewed, under consideration and submitted to ScienceOpen Preprints for open peer review.

DOI: 10.14293/PR2199.001096.v1

Preprint first posted online: 25 September 2024

Introduction.

Hereditary non-polyposis colorectal cancer is also known as Lynch syndrome. It is autosomal dominant and is one of the most common causes of inherited colon cancer (1,2). A family history of colorectal cancer in first-degree relatives in two generations, and one case before the age of 50 is seen with Lynch syndrome (3).

Abstract

The purpose of this paper is to provide a review of hereditary non-polyposis colorectal cancer. This paper covers the causes, epidemiology, pathophysiology, histology, diagnosis, and treatment for the disease.

Body.

Cause

The cause of Lynch syndrome is a germline mutation in DNA mismatch repair. This leads to defective repair of DNA causing abnormal cell growth and tumor formation (4).

Epidemiology

The epidemiology of Lynch syndrome is as follows: it has a prevalence of about 1% of 6% in the white race population. Colorectal cancer is usually found about 10 to 15 years earlier than the general population and there is no variation in diagnosis between men and women (1).

Pathophysiology

The following is the pathophysiology of Lynch syndrome. In Lynch Syndrome, there is defective DNA mismatch repair. This can result in a base substitution from an insertion or deletion in a newly formed strand of DNA. Normal DNA mismatch repair identifies the mismatch and repairs it. However, mutations with the mismatch repair genes can lead to inadequate DNA repair causing malignant transformation. Mutations in the following genes lead to defective mismatch repair:

- hMLH1 on chromosome 3p22
- hMSH2 and hMSH6 on chromosome 2p16
- hPMS1 on chromosome 2q32 and hPMS2 on chromosome 7p22
- hMSH3 on chromosome 5q14.1
- EXO1 on chromosome 1q43.

Approximately 75 to 80% of Lynch syndrome is caused by mutations in hMLH1, hMSH2, and hMSH6 genes (1,5)

Histology

The histology of Lynch syndrome is characterized by mucus poorly differentiated signet ring cells. There is a lymphocytic infiltrate with a germinal center (6).

Add images

Diagnosis

Mutations in any one of hMLH1, hMSH2, and hMSH6 genes establishes the diagnosis of lynch syndrome.

The Amsterdam II criteria is used to diagnose patients with Lynch syndrome. The criteria are listed below:

1. Three or more relatives with histology showing Lynch syndrome cancers such as colorectal cancer, endometrial cancers, small bowel cancers or kidney cancers. One individual must be a first degree relative and the other two must have familial adenomatous polyposis ruled out.
2. At least two generations with Lynch syndrome cancers

3. At least one cancer diagnosed before age 50 (1,7).

The Bethesda criteria guidelines are used to identify individuals who requires tumor testing for microsatellite instability testing. Patient with the following should undergo genetic testing for microsatellite instability:

1. Colorectal cancer in individuals less than 50 years old
2. Lynch syndrome associated tumors
3. Colorectal cancer in individuals less than 60 years old, along with microsatellite instability histology.
4. At least one first degree relative with the Lynch syndrome, tumor such as colorectal, endometrial, stomach, small bowel, ovarian or Biliary.
5. Colorectal cancer in two or more first-degree relatives with Lynch syndrome tumors (1,7).

Clinical features of Lynch syndrome include the following: constipation, diarrhea, abdominal pain, bloating, blood in the stool, iron deficiency anemia, unexplained weight loss or generalized fatigue (1).

Treatment

Treatment for colorectal cancer is usually done with a total abdominal colectomy along with an ileorectal anastomosis. This is followed by annual endoscopies for rectal surveillance (8).

A segmental colectomy can be done for individuals who are not candidates for total colectomy. A prophylactic hysterectomy with bilateral salpingo-oophorectomy can be done for women undergoing colectomy (8).

Immune therapy targeting programmed death receptor – 1 has shown to be helpful with patients with metastatic colorectal cancer (1).

Cancer prevention strategies include prophylactic hysterectomy and bilateral salpingo-oophorectomy surgery, and oral contraceptives (1,9).

Conclusion.

Hereditary non-polyposis colorectal cancer, also known as Lynch syndrome, is caused by mutations in the genes for DNA mismatch repair. This is due to microsatellite instability. Diagnosis is made with genetic testing revealing mutations in MLH1, MSH2, MSH6, and PMS2) or EPCAM genes. Amsterdam II criteria can be used for diagnosis. Treatment is with total abdominal colectomy (1).

References

1. Ojha SK, Laslett N. Hereditary Nonpolyposis Colon Cancer. [Updated 2023 Jul 16]. In: StatPearls [Internet]. Treasure Island (FL): StatPearls Publishing; 2024 Jan-. Available from: <https://www.ncbi.nlm.nih.gov/books/NBK564511/>
2. Giráldez MD, Castellví-Bel S, Balaguer F, Gonzalo V, Ocaña T, Castells A. Lynch syndrome in colorectal cancer patients. *Expert Rev Anticancer Ther.* 2008 Apr;8(4):573-83. doi: 10.1586/14737140.8.4.573. PMID: 18402524.
3. Coffin E, Dhooge M, Abou Ali E, Dermine S, Lavole J, Palmieri LJ, Chaussade S, Coriat R. Syndrome de LYNCH : identification et prise en charge [Identification and management of patients with Lynch syndrome]. *Presse Med.* 2019 Sep;48(9):904-914. French. doi: 10.1016/j.lpm.2019.07.011. Epub 2019 Sep 24. PMID: 31561847.
4. Singh AK, Talseth-Palmer B, McPhillips M, Lavik LAS, Xavier A, Drabløs F, Sjursen W. Targeted sequencing of genes associated with the mismatch repair pathway in patients with endometrial cancer. *PLoS One.* 2020 Jul 7;15(7):e0235613. doi: 10.1371/journal.pone.0235613. PMID: 32634176; PMCID: PMC7340288.
5. Cerretelli G, Ager A, Arends MJ, Frayling IM. Molecular pathology of Lynch syndrome. *J Pathol.* 2020 Apr;250(5):518-531. doi: 10.1002/path.5422. PMID: 32141610.
6. Hemminger JA, Pearlman R, Haraldsdottir S, Knight D, Jonasson JG, Pritchard CC, Hampel H,

Frankel WL. Histology of colorectal adenocarcinoma with double somatic mismatch-repair mutations is indistinguishable from those caused by Lynch syndrome. *Hum Pathol*. 2018 Aug;78:125-130. doi: 10.1016/j.humpath.2018.04.017. Epub 2018 May 1. PMID: 29723603; PMCID: PMC6296362.

7. Cohen SA, Pritchard CC, Jarvik GP. Lynch Syndrome: From Screening to Diagnosis to Treatment in the Era of Modern Molecular Oncology. *Annu Rev Genomics Hum Genet*. 2019 Aug 31;20:293-307. doi: 10.1146/annurev-genom-083118-015406. Epub 2019 Mar 8. PMID: 30848956.

8. Syngal S, Brand RE, Church JM, Giardiello FM, Hampel HL, Burt RW; American College of Gastroenterology. ACG clinical guideline: Genetic testing and management of hereditary gastrointestinal cancer syndromes. *Am J Gastroenterol*. 2015 Feb;110(2):223-62; quiz 263. doi: 10.1038/ajg.2014.435. Epub 2015 Feb 3. PMID: 25645574; PMCID: PMC4695986.

9. Lu KH, Loose DS, Yates MS, Nogueras-Gonzalez GM, Munsell MF, Chen LM, Lynch H, Cornelison T, Boyd-Rogers S, Rubin M, Daniels MS, Conrad P, Milbourne A, Gershenson DM, Broaddus RR. Prospective multicenter randomized intermediate biomarker study of oral contraceptive versus depo-provera for prevention of endometrial cancer in women with Lynch syndrome. *Cancer Prev Res (Phila)*. 2013 Aug;6(8):774-81. doi: 10.1158/1940-6207.CAPR-13-0020. Epub 2013 May 2. PMID: 23639481; PMCID: PMC3737517.