

# Gastrointestinal Cancer

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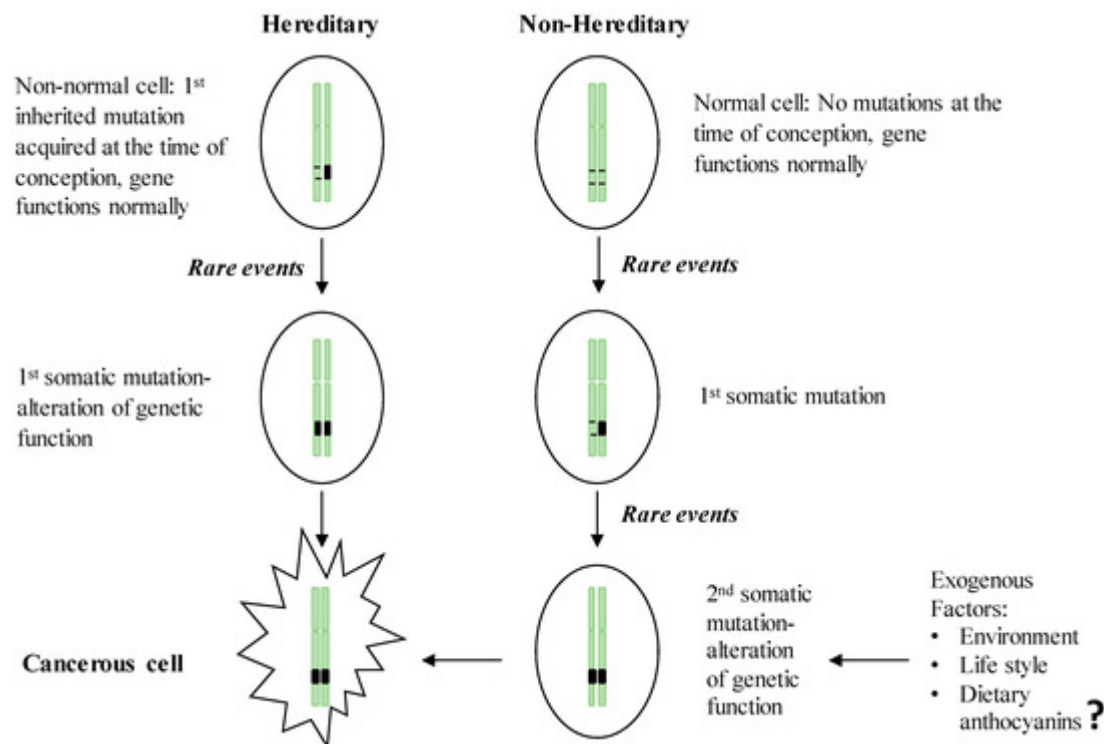
Gastrointestinal (GI) cancer is a heterogeneous cancer that tends to occur in the more common sporadic forms rather than the rare inherited forms. The process of initiation and formation of neoplastic cells in the GI tract can be classified into four main mechanisms: (i) inherited transmission of mutations; (ii) exposure to different carcinogens; (iii) chronic inflammatory conditions/microbial dysbiosis; and (iv) sporadic mutations and epigenetic changes.

cancer, digestive system,

carcinogenesis

## 1. Hereditary GI Cancers

Hereditary GI cancers represent a phenotypically diverse group of diseases involving malignant tumors of the digestive tract, extra-GI cancers, and benign abnormalities characterized by inherited genetic mutations transmitted from parent to child. However, no more than 3%–5% of GI cancers have shown a clear hereditary basis [\[1\]](#). The esophagus, stomach, colon, small intestine, and pancreas have been identified as the organs most likely to inherit germline mutations [\[2\]](#). The best known inherited malignant tumors are associated with the GI tract, representing monogenic hereditary diseases that result from mutations in a single gene [\[3\]](#). Despite the specific differences in the genes involved, inherited GI cancers share a common set of characteristics: (i) the majority of GI cancers are detectable in the early stages of life; (ii) these cancers follow an autosomal dominant inheritance mechanism in which the neoplasm occurs in 1st degree relatives; and (iii) the formation of multiple tumors [\[3\]\[4\]](#). In the hereditary form of GI cancers, the first genetic mutation in one of the alleles of a predisposition gene is acquired at the time of conception, and the somatic mutation of the second allele is then acquired via environmental insult, lifestyle practices or other exogenous factors ([Figure 1](#)). Once the two alleles of a specific predisposition gene are mutated, gene function is completely inactivated, leading to carcinogenesis. Compared to the sporadic form of GI cancer, which requires two somatic events during the inactivation of the predisposition gene, hereditary cancers present a higher risk because they need only one somatic mutation event, which explains the early onset of hereditary cancers [\[5\]](#). In parallel with the advancement of DNA technologies, the genetic mutations responsible for hereditary GI cancers have been widely documented ([Table 1](#)). These hereditary GI cancers include Cowden syndrome, MUTYH-associated polyposis, hereditary pancreatic cancer, Lynch syndrome, Peutz-Jeghers syndrome, familial adenomatous polyposis (FAP), attenuated FAP, serrated polyposis syndrome, and hereditary gastric cancer. Cancer-causing mutations can be initiated in three main classes of predisposition genes, oncogenes, tumor suppressor genes, and DNA repair genes, which are involved in establishing genetic stability.



**Figure 1.** Two-hit theory of the initiation of hereditary and non-hereditary cancer. People with a hereditary susceptibility to GI cancers harbor an inherited genetic mutation on one of the chromosomes at the time of conception and receive the 1<sup>st</sup> somatic mutation due to the endogenous (e.g., chronic inflammation) or exogenous (e.g., exposure to carcinogens) rare events which in turn inactivate the full function of the respective gene and initiate neoplastic transformation. Non-inherited forms of GI cancer occur by acquiring two somatic mutations in later life, resulting in the inactivation of a gene leading to the initiation of malignancy.

**Table 1.** Hereditary basis of GI cancers.

Type of the Cancer	Syndrome	Associated Germline Mutations	Reference
Esophageal	Familial Barrett's esophagus, Familial esophageal adenocarcinoma	MSR1, ASCC1 and CTHRC1	[6]
	Tylosis with esophageal cancer-squamous cell carcinoma	RHBDF2	[7]
Gastric	Diffuse hereditary gastric cancer-adenocarcinoma	CDH1 (E-cadherin)	[8]
Pancreatic	Hereditary pancreatitis	PRSS1, CFTR, SPINK1, CTSC	[9]
	Hereditary breast and ovarian cancer	BRCA1/2	
	Peutz-Jeghers syndrome	STK11/LKB1	

Type of the Cancer	Syndrome	Associated Germline Mutations	Reference
	Familial atypical multiple mole melanoma syndrome	CDKN2A/p16	
	Familial adenomatous polyposis	APC	
	Familial adenomatous polyposis	APC	
	Lynch syndrome	EPCAM, MLH1, MSH2, MSH6, PMS2	
Colorectal	MYH associated polyposis	MUTYH	<a href="#">[10]</a> <a href="#">[11]</a>
	Peutz-Jeghers syndrome	STK11	
	Juvenile polyposis syndrome	SMAD4, BMPR1A	
	Attenuated Familial adenomatous polyposis	APC	
	Familial adenomatous polyposis	APC	
	Lynch syndrome	Mutations in mismatch repair genes	<a href="#">[12]</a>
Small intestine			
	Juvenile polyposis syndrome	SMAD4	
	Peutz-Jeghers syndrome	STK11	
	$\alpha$ -1 antitrypsin deficiency	SERPINA1	
	Hereditary hemochromatosis	HFE	
Liver	Hereditary tyrosinemia type 1	FAH	<a href="#">[13]</a> <a href="#">[14]</a> <a href="#">[15]</a> <a href="#">[16]</a>
	Glycogen storage disease type 1	G6PC, SLC37A4	
	Wilson's disease	ATP7B	<a href="#">[17]</a>
	Niemann-park disease	SMPD1 AND NPC1 OR NPC2	
Biliary	Bile salt export pump deficiency	ABCB11	<a href="#">[18]</a>

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1. Accardi, F. et al. Accumulation of sporadic mutations can occur due to factors such as exposure to carcinogens [2], a westernized diet [19][20], diets rich in salt [21][22], obesity [23][24], chronic alcohol consumption [25][26], and chronic inflammation [27].
2. Rahner, N.; Steinke, V. Hereditary cancer syndromes. *Dtsch. Arztebl.* 2008, 105, 706–714.
3. The relationships between carcinogens, diet, inflammation, and GI cancers are multiple and complex. Exposure to carcinogens can initiate cancer development via somatic mutations that include point mutations, deletions, additions, and modified methylation of DNA [28]. There are several cellular mechanisms to protect DNA from carcinogen-induced mutations and to identify and correct these mutations before they give rise to malignancy. In spite of these protective mechanisms, the GI tract is continuously exposed to chemical and biological carcinogens, often due to diets that act as carriers of preformed carcinogens [2]. Among known carcinogens, tobacco smoke hydrocarbons are one of the most potent, being comprised of more than 60 mutagens and cancer-causing chemicals directly linked to esophageal [29], pancreatic [29], and gastric cancers [30][31]. In addition, exposure to airborne occupational carcinogens such as cement dust, quartz dust, and diesel exhaust fumes increases the risk of gastric cancer [32].
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