

FAP, MAP, and PPAP

Marco Novelli





FAP, MAP, NAP and PPAP

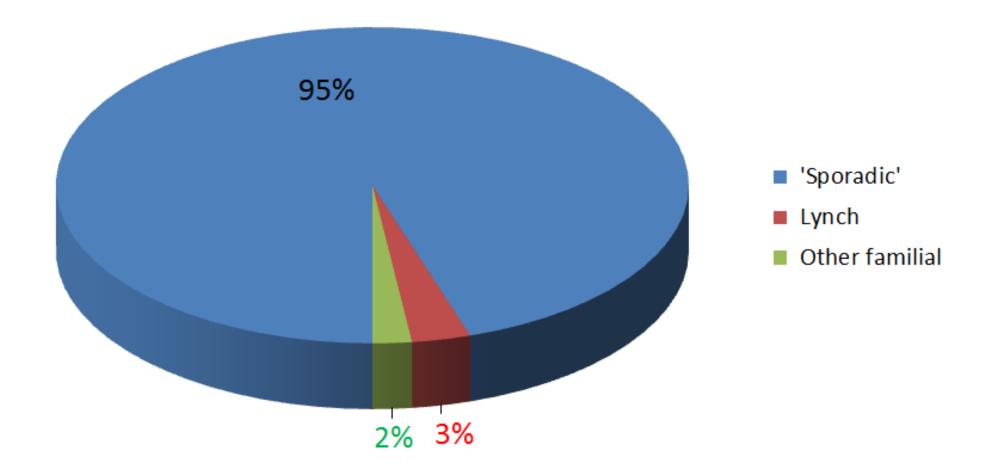
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Overview

- History of polyposis
- Familial adenomatous polyposis
- MYH-associated polyposis
- NTLH1-associated polyposis
- Other polyposis syndromes

Colorectal cancer



History of polyposis

- First description of colorectal polyposis:
 - Menzel 1721, Berlin
- first histologically verified case of adenomatous polyposis:
 - Sklifasowski 1881, Russia

X. POLYADENOMA TRACTUS INTESTINALIS.

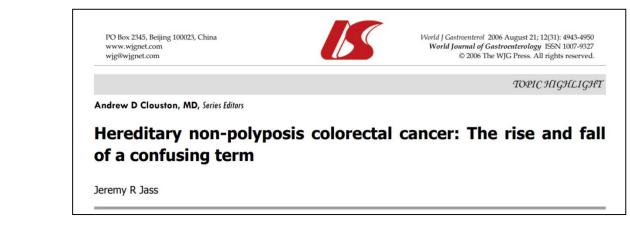
Проф. Н. В. Склифосовскаго.

Различные патологические процессы въ нижней части кишечника сопровождаются функціональными разстройствами, въ числѣ которыхъ самыя постоянныя—

 1882 Cripps termed "disseminated polypus of the rectum" – inherited predisposition.

Lynch syndrome

- "Cancer family syndrome" Lynch 1966.
 - Lynch syndrome 1
 - Lynch syndrome 2
- Lynch named the condition 'Hereditary Non-Polyposis Colorectal Carcinoma' (HNPCC), 1985.
- Jass 2006 terminology HNPCC should be phased out.



- Familial adenomatous polyposis 1881
- Peutz–Jeghers syndrome 1921
- Juvenile polyposis 1939
- Gardner's syndrome 1951
- Turcot syndrome 1959
- Cowden's syndrome 1963
- Lynch syndrome (HNPCC) -1966 (1985)
- Muir–Torre syndrome 1967-8
- Hyperplastic polyposis -1980

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Gardner's syndrome - 1951

Turcot syndrome 1959

- Cowden's syndrome 1963
- Lynch syndrome (HNPCC)-1966 (1985)

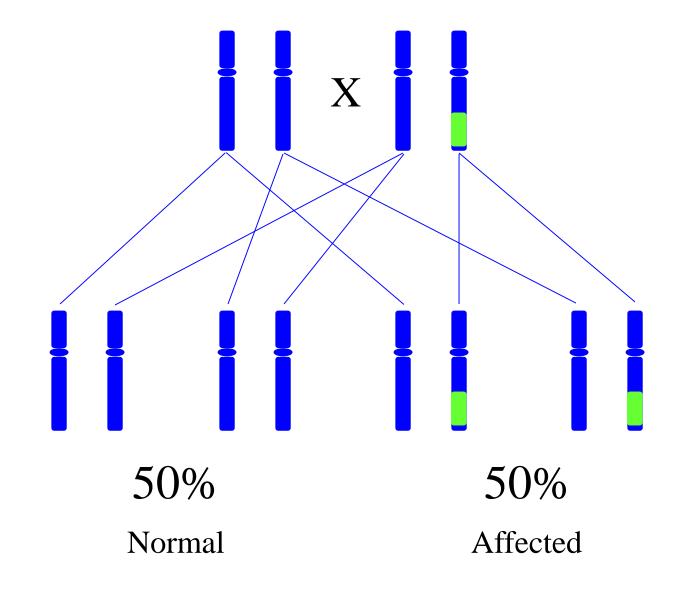
Muir_Torre syndrome - 1967-8

• Hyperplastic Serrated polyposis -1980

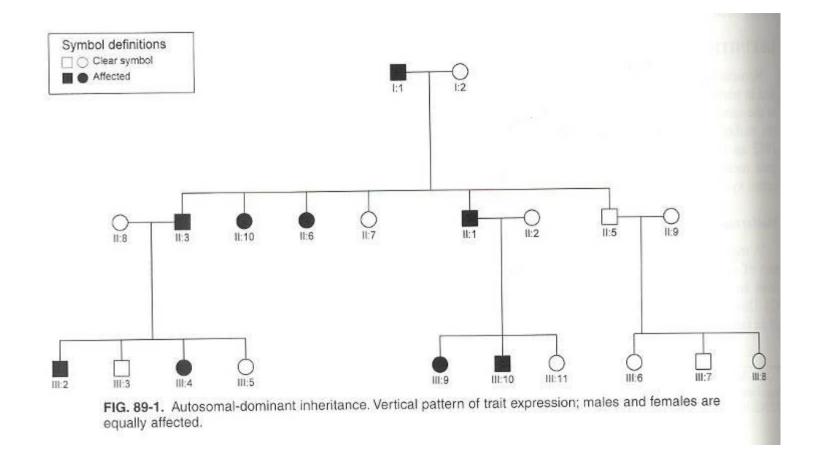
- Lynch syndrome
- Familial adenomatous polyposis
- MYH-associated polyposis
- NTHL1-associated polyposis
- Serrated polyposis
- Hereditary mixed polyposis
- Polymerase proof reading-associated polyposis
- Peutz-Jeghers syndrome
- Juvenile polyposis
- PTEN hamartoma syndromes (Cowden's)
- Constitutional mismatch repair deficiency

Mendelian inheritance

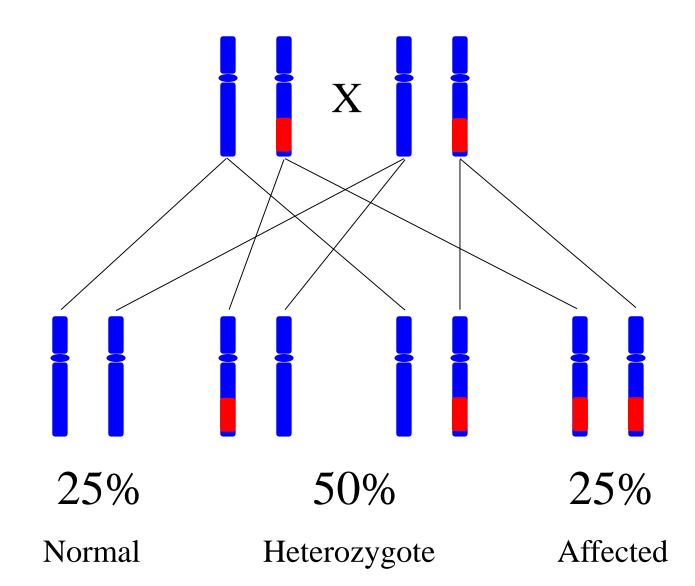
Mendelian Autosomal Dominant Inheritance



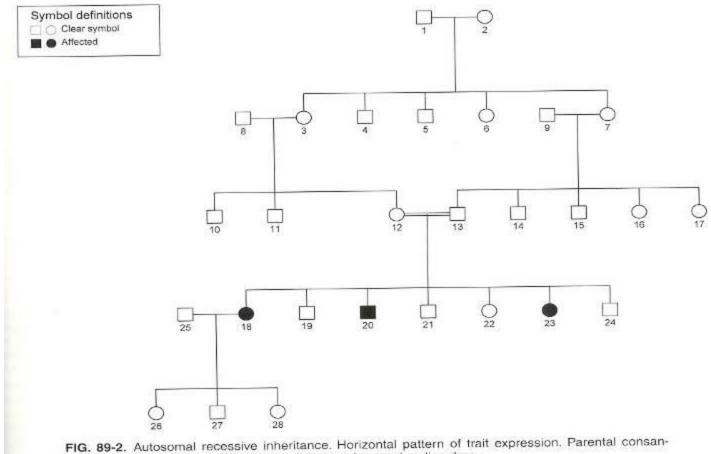
Autosomal Dominant Inheritance



Mendelian Autosomal Recessive Inheritance

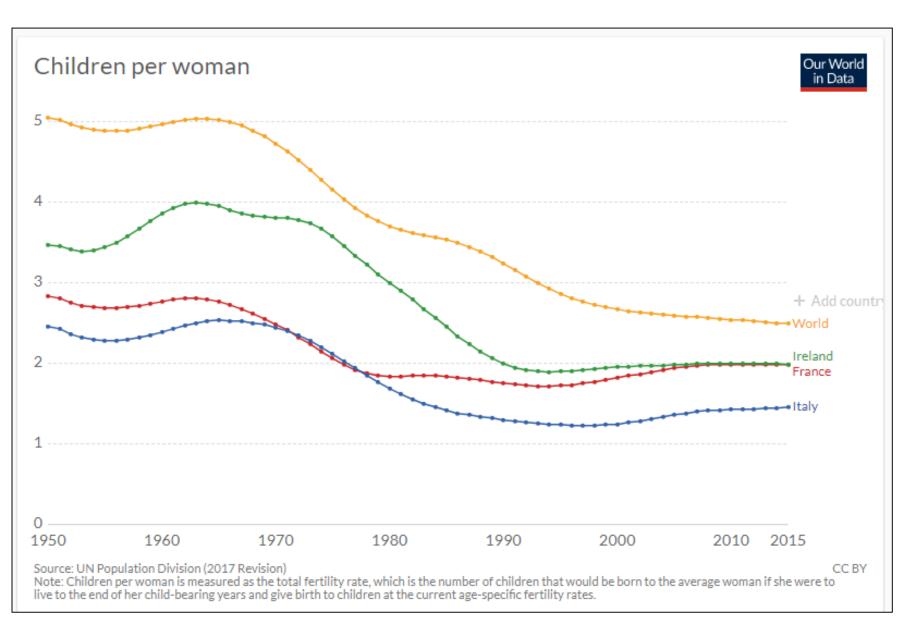


Autosomal Recessive Inheritance

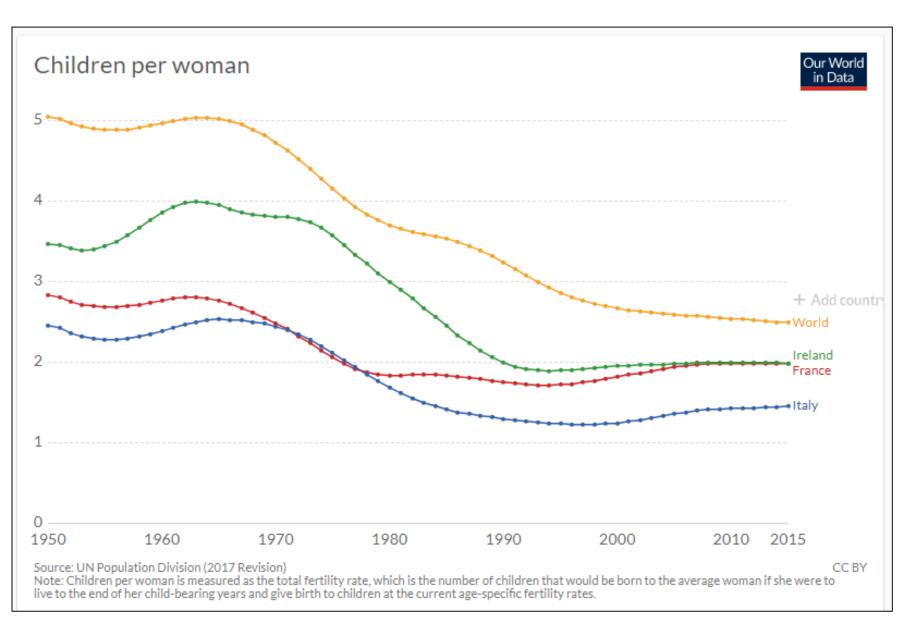


guinity is often present in families with autosomal recessive disorders.

Historical reduction in family size



Historical reduction in family size



→ Unlikely to be a family history in recessively inherited syndromes

- Lynch syndrome
- Familial adenomatous polyposis
- MYH-associated polyposis
- NTHL1-associated polyposis
- Serrated polyposis
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Dominant
Recessive
Uncertain

Familial adenomatous polyposis (FAP)



> 100 colorectal polyps





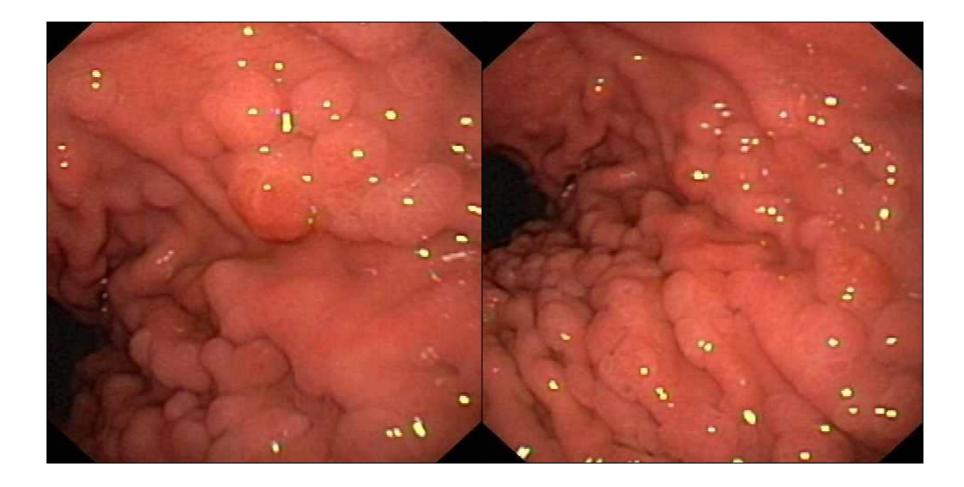
Familial adenomatous polyposis

- Due to a germline mutation in the APC gene (5q22)
- > 100 adenomatous polyps predominantly in large intestine (typically 1000s).
- Colorectal adenocarcinomas < 40 years old (10-15 years after appearance of polyps).
- Adenomas in small intestine and stomach.
- Extra-intestinal manifestations.
- Commonest cause death now duodenal carcinoma and desmoids.



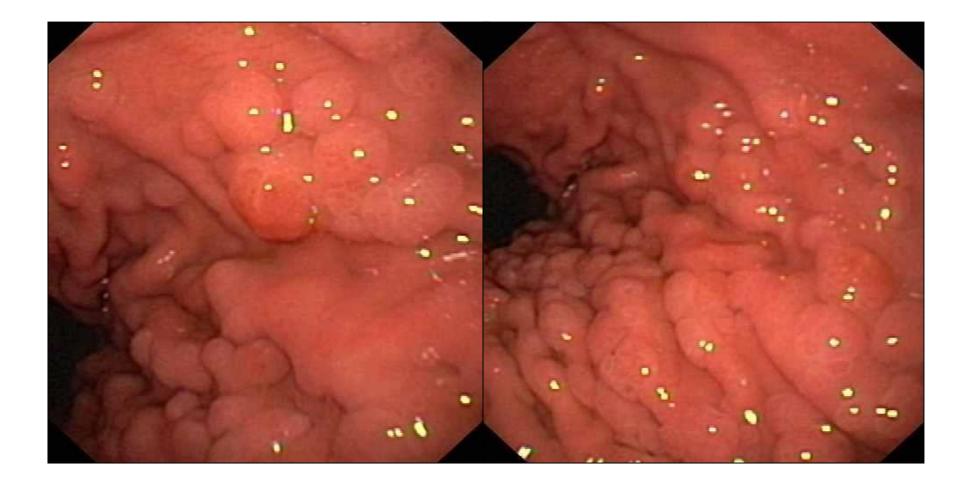
Extra-intestinal manifestations of FAP

- Gastric fundic gland polyps.
- CHRPEs.
- Sebaceous cysts.
- Osteomas (jaw).
- Desmoid tumours.
- Papillary carcinoma of thyroid (Cribriform-morular variant).
- Hepatoblastoma.
- Brain tumours (Glioblastoma M).



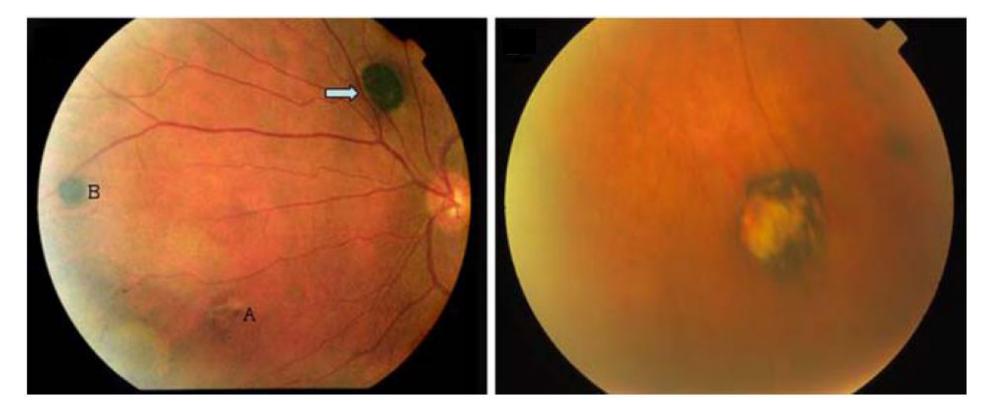
Fundic gland polyps :

- 0.8 2% normal population.
- 12.5 84% FAP patients



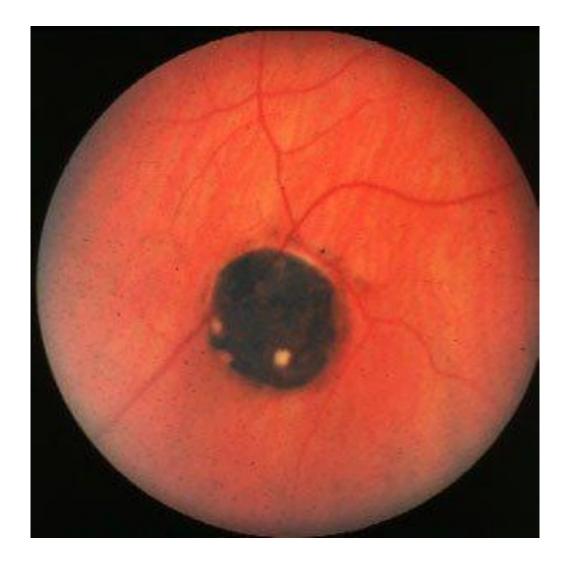
Fundic gland polyps :

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 - **Common with PPIs**
- 12.5 84% FAP patients

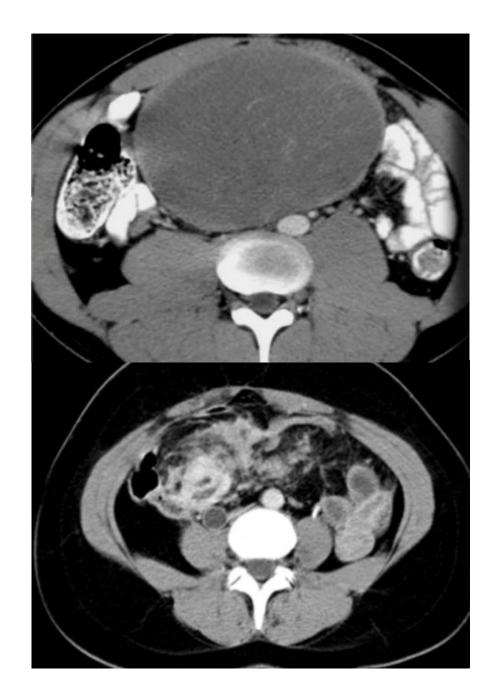


Congenital Hyperpigmentation of the Retinal Pigment Epithelium (CHRPE).

- Normal population 5% (0.3 40%)
- FAP 66-92%



CHRPE in FAP are typically bilateral with a de-pigmented halo.



Desmoid tumours

- Rare tumours 2-4/million per annum.
- 10 30% incidence in FAP.
- 7.5% of desmoids occur on a background of FAP.
- Sporadic tumours have a somatic mutation in CTNNB1 (β catenin).



Desmoid tumours

Frizzled

recepto

B-catenin

ZNRF3

B-catenin

LPR

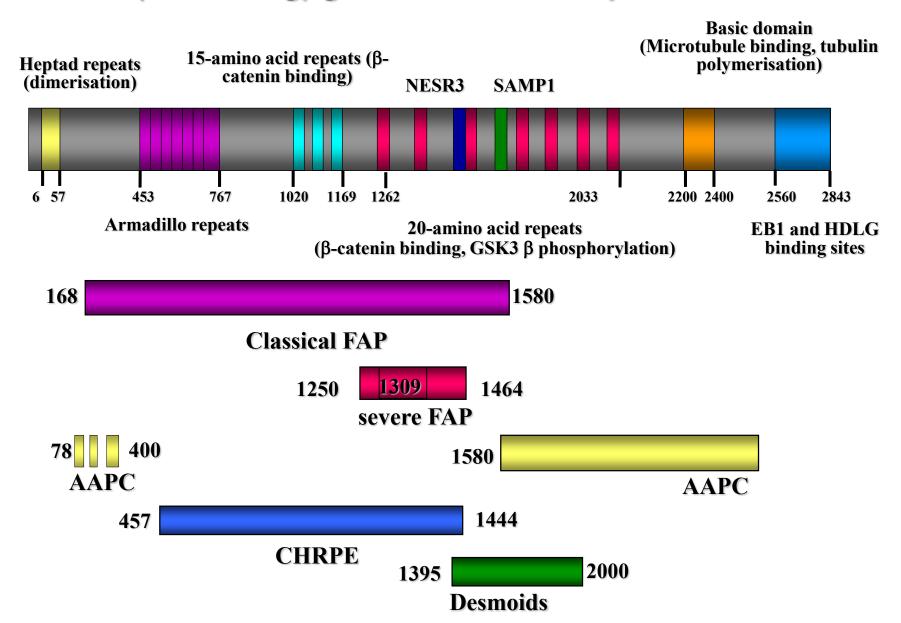
Inactive GSK-3f

B-catenin

accumulation

- Rare tumours 2-4/million per a
- 10 30% incidence in FAP.
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- Sporadic tumours have a somatic mutation in CTNNB1 (β catenin).

APC protein domains and FAP phenotype association with (truncating) germline mutation position



Attenuated FAP

- Typically less than 100 adenomas in colon.
- Mean 25 adenomas (0 470).
- Concentrated in proximal colon.
- Some extra-colonic manifestations.
- Colorectal carcinoma in 70% by 80 years of age.

MUTYH-Associated Polyposis

MYH-Associated Polyposis

MAP

Discovery of MYH-associated polyposis

- Family with multiple adenomas and CRC
- No known pathogenic APC or MMR germline mutation
- 3 affected sibs 11 tumours
 - 15/18 somatic APC mutations G:C>T:A
 - → Suggested mutations from oxidative DNA damage.

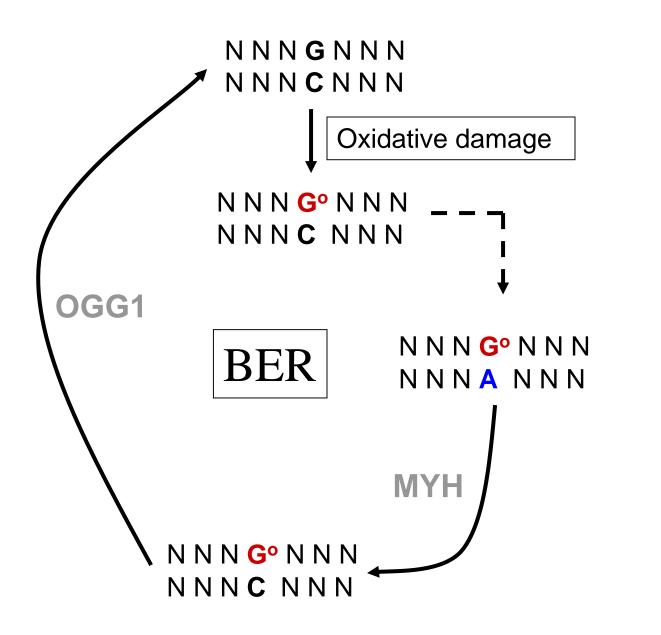
Al-Tassan *et al* Nat Genet 30:227-232, 2002

Oxidative damage of DNA

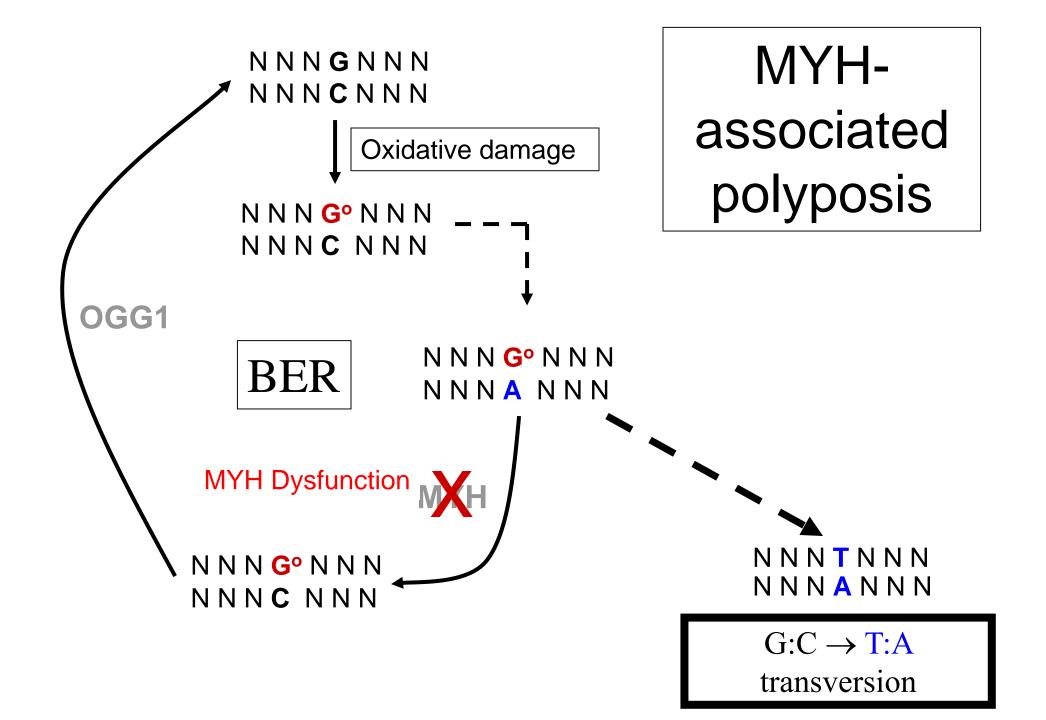
- Oxidative damage 10⁴ lesions/cell/day.
- 8-oxo-G most deleterious.
- 8-oxo-G mis-pairs with adenine residues

Normal G:C Damaged G^o:A G:C \rightarrow T:A mutation

• Role of MYH is to excise mispaired adenines (Base excision repair BER).







MYH-associated polyposis

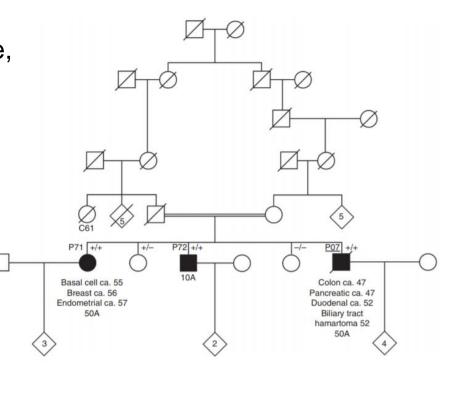
- Autosomal recessive inheritance.
- Penetrance approaching 100%.
- Clinically similar to attenuated FAP.
 - Typically <100 polyps(but may show classical FAP phenotype).
 - Some extra colonic manifestations.
 - May also have serrated polyps
- Extracolonic tumours
 - Duodenal, urinary tract, ovarian and ? breast carcinomas.
 - Sebaceous skin tumours,
 - Desmoid tumours (case report).

MYH-associated Polyposis

- Due to biallelic mutations in MYH gene.
- Thought that there is a gradual accumulation oxidative damage causing tumorigenesis
 - $APC \rightarrow$ classical adenomas
 - $KRAS \rightarrow$ serrated polyps
- Mean age cancer diagnosis 50 years old.
- Up to 30% of European patients with > 15 100 adenomas carry biallelic MYH mutations.

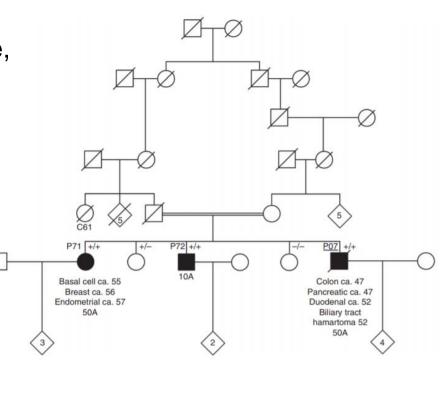
NTHL1-associated polyposis

- NTHL1 is a base excision repair gene (like MYH).
- Broader spectrum of mutations than MAP but more C:G>T:A (MAP C:G>A:T).
- Autosomal recessive inheritance, high penetrance.
- Probably 1/5 as common as MAP.
- Spectrum of tumour types:
 - colorectal carcinoma
 - endometrial carcinoma
 - duodenal carcinoma
 - skin (BCC)
 - others?

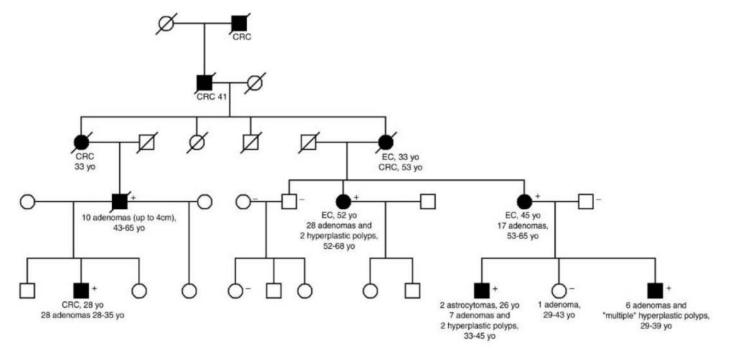


NTHL1-associated tumour syndrome

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- Autosomal recessive inheritance, high penetrance.
- Probably 1/5 as common as MAP.
- Spectrum of tumour types:
 - colorectal carcinoma
 - endometrial carcinoma
 - duodenal carcinoma
 - skin (BCC)
 - bladder, breast, others?



Polymerase Proofreading-associated polyposis (PPAP)



- Germline POLE and POLD1 mutations.
- Major catalytic and proof reading subunits of the Pole and Pol δ enzyme complexes
- Mutations impair proof reading \rightarrow hyper mutation

Polymerase Proofreading-associated polyposis (PPAP)

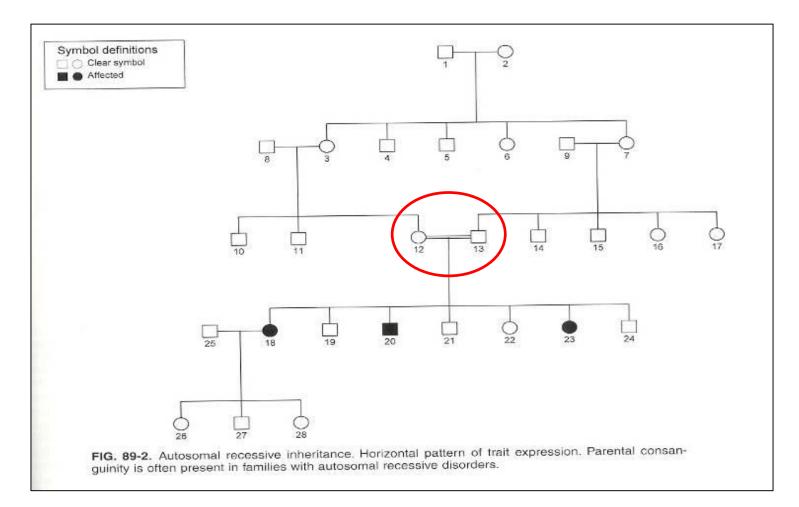
- Autosomal dominant with high penetrance.
- Multiple colorectal adenomas (3 10s).
- Early-onset CRC (mean age 36) may be synchronous, often lymphocyte rich.
- POLD1 Endometrial adenocarcinomas, breast carcinoma and brain tumours.
- POLE Endometrial, small bowel, ovarian, pancreatic and brain tumours.
- Tumours microsatellite stable but have multiple point mutations.
- POLE can also be mutated in sporadic CRC (<2%).

Briggs S, Tomlinson I. Germline and somatic polymerase ϵ and δ mutations define a new class of hypermutated colorectal and endometrial cancers. J Pathol. 2013 Jun;230(2):148-53.

Congenital Mismatch Repair Deficiency (Childhood cancer syndrome)

- "Autosomal recessive" Lynch syndrome.
- Biallelic mutations in PMS2, MSH2, MSH6 or MLH1 (compound or homozygous).
- Phenotype:
 - Café-au-lait spots (NF1 type features).
 - Adenomatous polyposis and colorectal cancers.
 - Brain tumours (Gliomas, PNETs, meduloblastomas).
 - Haematological malignancies (leukaemias and diffuse large B cell lymphoma).
- Accounts for some cases of Turcot syndome.

Congenital Mismatch Repair Deficiency



• Most commonly seen in asian and arabic populations where consanginous marriages are common



Congenital Mismatch Repair Deficiency

- Café-au-lait spots.
- Adenomatous polyposis and colorectal carcinomas.
- High grade brain tumours.
- Haematological malignancies.

Serrated polyposis

Table 2. World Health Organization diagnostic criteria for the diagnosis of serrated polyposis syndrome (SPS)

Any one of the following

- (1) At least five serrated polyps proximal to the sigmoid colon, two of which are >10 mm in diameter
- (2) Any number of serrated polyps proximal to the sigmoid colon in an individual who has a first-degree relative with SPS
- (3) Greater than 20 serrated polyps of any size distributed throughout the colon
 - Genetics, natural history and phenotype of this syndrome remain poorly characterised.
 - Family history CRC in 40-60%
 - ?50% chance of developing colorectal carcinoma.
 - 26% of carcinomas synchronous/metachronous.

Serrated polyposis

- Polyp spectrum
 - Hyperplastic polyps.
 - Sessile serrated lesions.
 - Traditional serrated adenomas.
 - Conventional adenomas.

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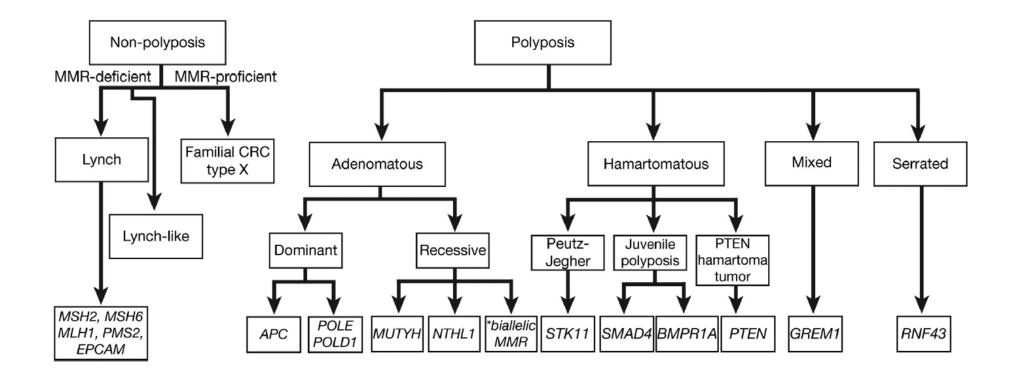
\rightarrow PHENOTYPIC OVERLAP WITH MAP

LEARNING POINTS

- Adenomatous polyposis syndromes rare (?1% CRC).
- Classical FAP autosomal dominant inheritance but several new syndromes autosomal recessive.

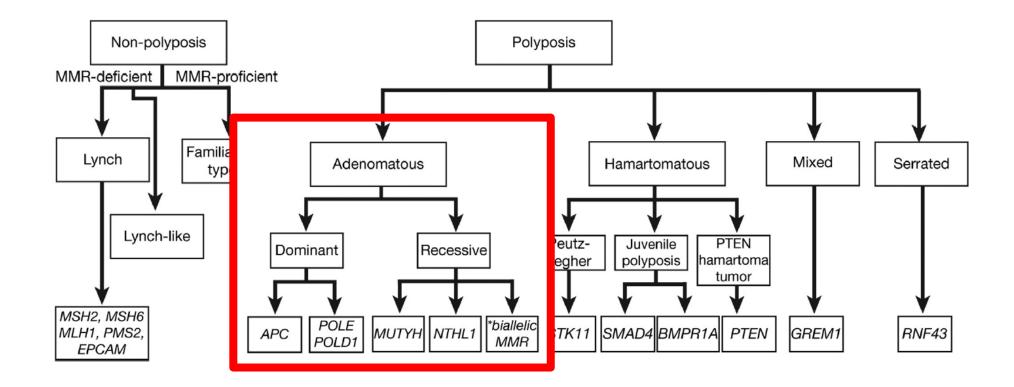
- MAP + NAP and Constitutional mismatch repair deficiency.

- Two pathways to colorectal carcinogenesis:
 - Genomic instability via p53
 - Hypermutation MMR, BER, DNA polymerase malfunction.
- Every increasing spectrum of extra-intestinal malignancies.
- MAP and Serrated polyposis may show phenotypic overlap
 Serrated polyps and adenomas



Recent Discoveries in the Genetics of Familial Colorectal Cancer and Polyposis

Laura Valle Clinical Gastroenterology and Hepatology 2017;15:809–819



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Thank you for your attention

