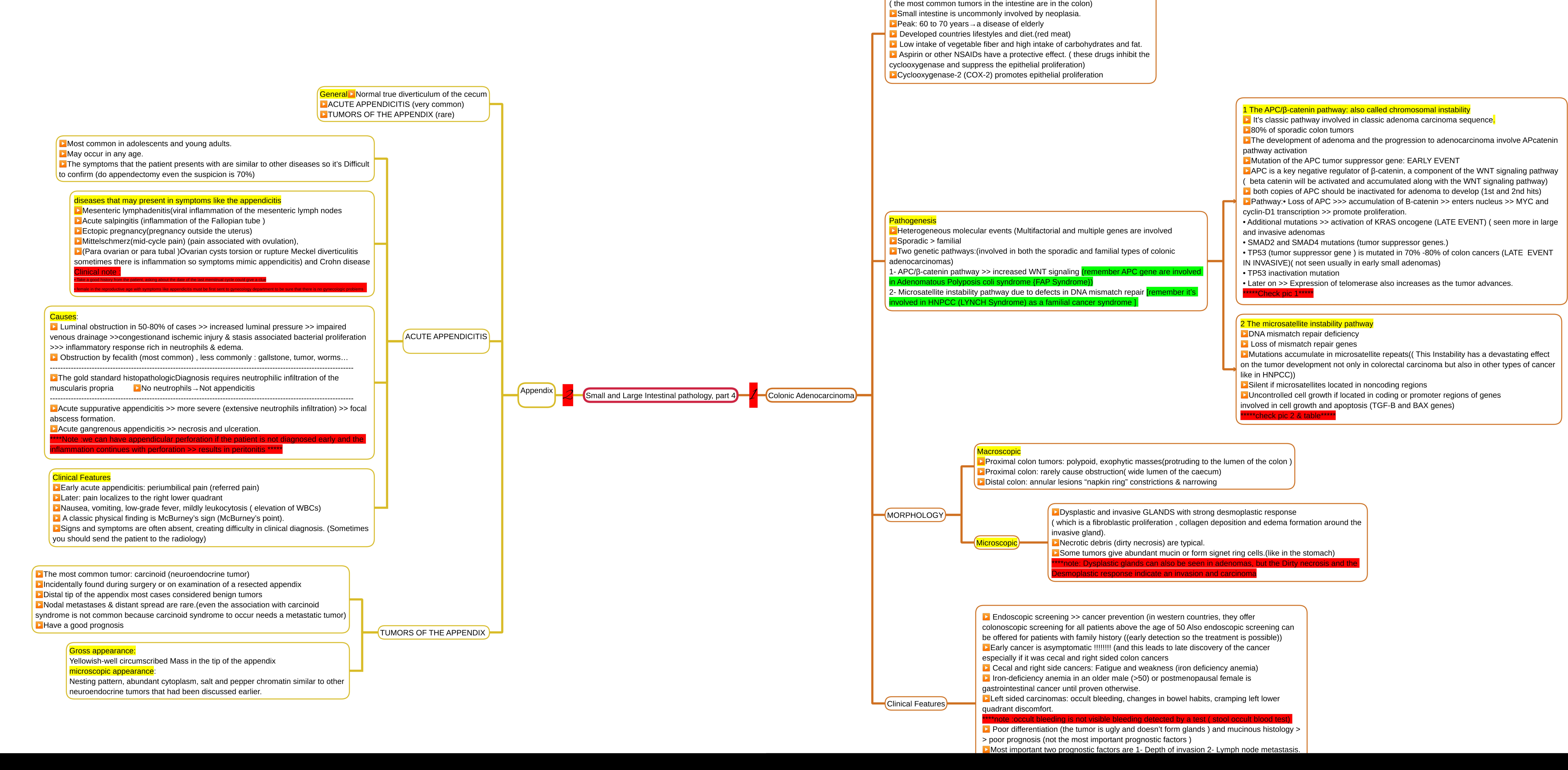
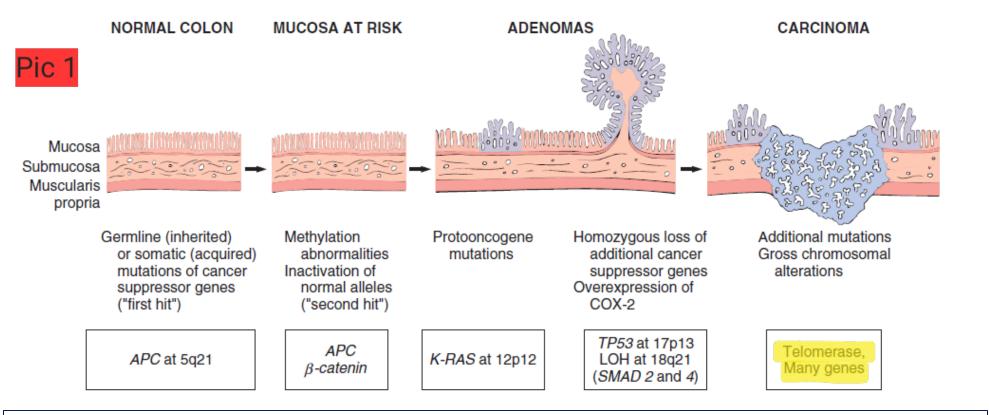
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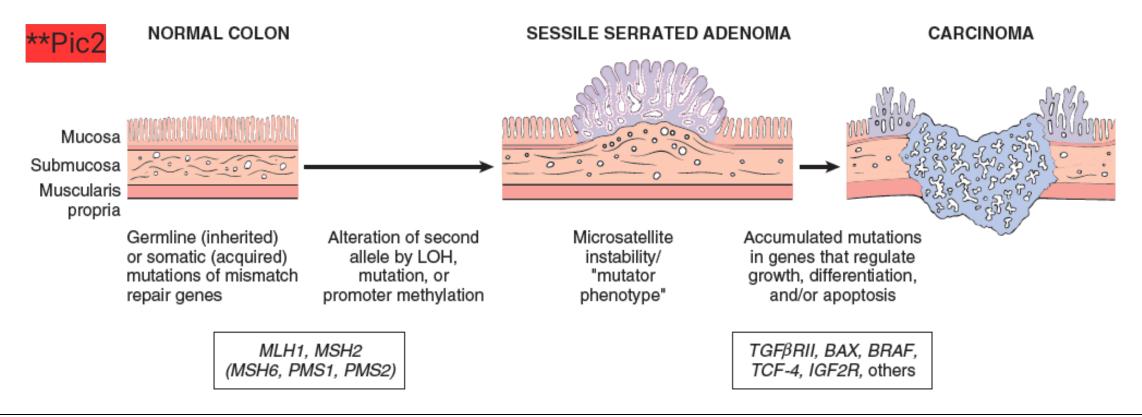


Most common malignancy of the gastrointestinal tract



- If a patient had the Germline mutation which can be inherited in familial cases like in (FAP) for ex, then he needs another copy of the gene to be mutated which can take place in a short time so here the tumor will appear at an earlier age since we already have the first hit inherited and the second mutation hit will appear later in life In sporadic cases we have the somatic mutation which is acquired later in life then we should have the second mutation to be acquired so this will take longer time and this is why familial cancers appear earlier than sporadic
- → 1st HIT mutation in APC gene, the 2nd HIT should take place resulting in beta catenin accumulation in cytoplasm so here the Mucosa is at risk (the mucosa starts its transformation) (note that we still don't have adenoma or dysplasia so these events occur very early even before the formation of a colonoscopically detected mass or lesion)

Then we will have later event which is K-RAS mutation which is Protooncogene so the development of adenoma will start as you can see in the 3rd picture in the figure, it starts very small then it will enlarge forming the large colonic adenoma. (As the adenoma enlarges as the risk for progression increases), then other mutations will accumulate like TP53, SMAD2&4 Later on, when the tumor becomes invasive (grossly visible cancer) \rightarrow more mutations – more chromosomal alterations will appear & the telomerase will. be activated



• As we said , this pathway starts with a mutation in DNA mismatch repair genes which are multiple genes including MLH1, MSH2, MSH6, PMS1 and PMS2 . So, the mutation in these genes could be a germline mutation (inherited) like in HNPCC or a somatic (acquired) mutation like in sporadic colorectal carcinoma, then the 2nd copy of the gene should be altered in order for subsequent steps to take place like the first and second hits in APC beta catenin pathway so both copies should be inactivated , leading to microsatellite instability which results in forming a peculiar special type of adenoma and it's different from the adenoma formed in the 1st pathway "APC/Beta Catenin" →The adenoma here is known as sessile serrated adenoma (it's not ordinary adenoma) which also has risk of transformation to an invasive adenocarcinoma by acquisition of more mutations in different genes involving apoptotic and cell proliferation genes



2	Etiology	Molecular Defect	Target Gene(s)	Transmission	Predominant Site(s)	Histology
	Familial adenomatous polyposis (70% of FAP)	APC/WNT pathway	APC	Autosomal dominant	None	Tubular, villous; typical adenocarcinoma
	Hereditary nonpolyposis colorectal cancer	DNA mismatch repair	MSH2, MLH I	Autosomal dominant	Right side	Sessile serrated adenoma; mucinous adenocarcinoma
	Sporadic colon cancer (80%)	APC/WNT pathway	APC	None	Left side	Tubular, villous; typical adenocarcinoma
	Sporadic colon cancer (10%–15%)	DNA mismatch repair	MSH2, MLH I	None	Right side	Sessile serrated adenoma; mucinous adenocarcinoma

- This table summarizes almost all what we have said about the mechanism &the pathogenesis of colorectal cancer >> memorize it well
- Remember that we have sporadic cancer and familial cancer (FAP and HNPCC syndrome)
- colonic adenocarcinoma is sporadic most of the time (80% involve the APC /WNT pathway and 10 to 15 % involve the DNA mismatch repair pathway which also seen in lynch syndrome)
- The sessile serrated adenoma is more commonly seen in the DNA mismatch repair genes pathway
- More classical adenomas (left side involvement) are typical for sporadic colon cancer

