

UPDATE ON INFLAMMATORY BOWEL DISEASES ETHIOPATOGENICS

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ABSTRACT

THE INFLAMMATORY BOWEL DISEASES (IBD) CROHN'S DISEASE (CD) AND ULCERATIVE COLITIS (UC) ARE CHRONIC IMMUNOLOGIC DISEASES THAT CAN LEAD TO DECREASED QUALITY OF LIFE AND INVALIDITY; IBD ARE CHARACTERIZED BY ABERRANT IMMUNE RESPONSES TO INTESTINAL MICROBIOTA IN GENETICALLY SUSCEPTIBLE HOSTS THAT LEAD TO CHRONIC EXCESSIVE INFLAMMATION AND IN TIME TO PROGRESSIVE GUT WALL DESTRUCTION. IBD HAVE AN EUROPEAN INCIDENCE OF 12 TO 25 PER 100,000 PERSON-YEARS, HIGHER FOR UC THAN CD, WITH A PREVALENCE OF 0.5-1%.

ENVIRONMENTAL FACTORS LIKE CIGARETTE SMOKING, POLLUTION, CHANGES IN DIET AND EXCESSIVE USE OF ANTIBIOTICS ARE INCRIMINATED IN IBD PATOGENESIS.

THERE IS THOUGHT TO EXIST AN IMBALANCE IN THE INNATE AND ACQUIRED IMMUNE RESPONSES TO NORMAL INTESTINAL MICROBIOTA THAT ACTIVATES PRO-INFLAMMATORY CYTOKINES AND LEADS TO A CHRONIC INFLAMMATORY PROCESS AROUND THE GUT BARRIER. THIS IN TIME LEADS TO INTESTINAL BARRIER DESTRUCTION AND EXPOSURE TO EVEN MORE BACTERIAL PARTICLES THAT FURTHER AGGRAVATE THE INFLAMMATORY PROCESS.

GENETIC STUDIES HAVE PROVIDED MANY SUSCEPTIBLE LOCI FOR IBD. IT IS KNOWN FROM THE LITERATURE THAT GENETIC FACTORS ACCOUNT FOR 13.6% OF CD AND 7.5% FOR UC VARIANCE. IN THIS LITERATURE REVIEW WE HIGHLIGHT SOME ETHIOPATHOGENIC LEADS THAT ARE RECENT TARGETS OF CLINICAL AND FUNDAMENTAL IBD RESEARCH.

KEY WORDS: IBD GENETICS, IMMUNOLOGY, THERAPEUTIC TARGETS

INTRODUCTION

Crohn's disease (CD) and ulcerative colitis (UC) are chronic inflammatory immune mediated diseases that affect genetic predisposed individuals and can interfere with quality of life and working ability. Inflammatory bowel disease's (IBD) incidence in Europe is 12.7 (CD) and 24.3 (UC) per 100,000 person-years while prevalence is around 0.5-1% [1].

In genetically susceptible individuals the relationship between aberrant immune responses and intestinal microbiota, that can be changed by diet habits [2], leads to gut barrier

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destruction and higher permeability for pathogenic bacteria. This attracts in time chronic inflammation, edema, erosion and ulcerations that can manifest with fatigue, abdominal pain, weight loss and chronic bloody diarrhea. Due to a high variability and multiple pathogenic mechanisms there is still no clear etiopathogenic mechanism of IBD.

MAIN TEXT

1. Environmental factors have an important role in IBD pathogenesis, for example cigarette smoking is a known risk factor for CD because it interferes with the function of NOD2 gene [3]. There is also a north-south, west-east prevalence gradient in IBD that is probably due to environmental factors and diet. Current theories support the fact that an active hygiene, secondary to the high standard of life in western society, leads to a lower exposure to bacterial antigens in childhood and a higher incidence of IBD [4]. Changing nutrition habits in infants with decreasing breast feeding (known to be a protective factor), excess of antibiotic regimes, proton pump inhibitors and western diet rich in processed food, fat, animal proteins and polyunsaturated fatty acids and low in vegetables, fiber and fruits play an important role in changing intestinal microbiota [5] and increasing IBD prevalence.

2. Unlike general population, IBD patients have an aberrant immunologic response to external antigenic stimuli that is increased in duration and severity [6]. Alterations in gut wall start by loss of tight junctions between epithelial cells and destruction of mucosal layer which increases intestinal permeability to luminal bacterial antigens [7].

This alterations are further amplified by loss of inhibitory immune mechanisms, macrophages and numerous pro-inflammatory cytokines activation. Until recently we considered that immune reaction to antigens was different in CD as opposed to UC. In CD there was a lymphocyte T helper 1 (Th1) mediated reaction that differed to the Th2 activation seen in UC patients. A new pathogenic theory common for both CD and UC shows that the hyperactivity of cellular immune response is mediated by the presence of Th17 lymphocytes that are implicated in the secretion of numerous cytokines: interleukin (IL) 17, IL 22, IL 23, IL 26 [8][9][10].

There is another theory that the chronic inflammation in IBD is secondary to an energy imbalance. This imbalance is due to alterations in the oxidative metabolism from the intestinal epithelial cells [11][12] and to cell-specific endoplasmic reticulum (ER) stress which induces dysfunctions in Paneth cells, goblet cells that produce mucus and that alters the enterocyte barrier [13][14]. The ER stress contributes to pathogenic mechanisms of insulin-resistance in type 2 diabetes and this proves that there is a common pathogenic way in both IBD and metabolic diseases [15].

Mitochondria are essential cellular organelles for maintaining energetic equilibrium by adenosine triphosphate (ATP) mediated energy intake, apoptosis regulation and oxygen free radicals production. The paper published by Bertran and his colleagues proves that excessive levels of oxidized molecules, inhibited mitochondria and hypersensitivity to oxidative stress are implicated in IBD pathogenesis [16].

3. Genetic susceptibility in IBD patients was demonstrated in numerous genome-wide association studies. Nevertheless, it has been proven that there is no simple model, Mendelian type, that could explain IBD transmission. Mono-zygotic twins have up to 67% concordance for developing CD [17] and 15-20% for UC. By genotyping, in the last years there have been found numerous susceptibility loci associated with IBD. There are more than 40 confirmed loci and over 160 susceptibility loci [18]. Most of the confirmed loci are common for CD and UC like the genes for lymphocyte Th 17 mediated immune response (IL23R, IL12B, JAK 2, STAT 3) [19][20]. The genes that differ between the two IBD's are a good premise for future research and new therapeutic molecules, specific for each inflammatory bowel disease in

part. Crohn's disease is associated with autophagocytosis genes: ATG16L1, IRGM and NOD2 while UC genetic specificity is in relation to the genes that codify the gastro-intestinal epithelial barrier: HNF4a, E-caderin, LAMB 1 and IL 10 [21]. It's interesting that some genes like NOD 2 and PTPN22 predispose to CD but are a protective genetic element against UC. By genotyping rises the possibility to discover new molecules that are specific for each IBD pathogenesis in part and also the possibility to identify genetic markers that are able to predict disease outcome and therapeutic response. A recent study identified mutations that are part of NFkB inflammatory path; this particular mutations can predict the rate of response to anti tumor necrosis factor alfa (anti-TNF α) molecules - a very potent immunosuppressant class of drugs [22]. Bank S. et al paper also identified new possible therapeutic targets (IL 1 β , IL-6, IFN- γ) that can become very useful in the future.

4. IBD patients have a particular intestinal microbiota: a decrease in non-pathogenic species like Bifidobacteria, Lactobacili and Firmicutes and an increase in pathogenic species like Bacteroides and Escherichia coli [23]. In healthy subjects autophagy mechanisms are efficient enough to eliminate intracellular pathogens from digestive tract but Hang Thi Thu Nguyen and his colleagues have recently demonstrated that in CD patients Adherent Invasive Escherichia coli (AIEC) suppresses autophagy gene expression (ATG 5, ATG16L1) and that leads to the inability of clearing this bacteria from the intracellular space [24]. AIEC persistence maintains a chronic inflammatory intestinal process with excessive pro-inflammatory cytokines. This is just one of the papers that highlights the importance of gut microbiota in IBD etiopathogenic mechanisms and the utility of prebiotics and probiotics to obtain and maintain disease remission. Also, microbiota pathological diversity in CD and UC is reason to justify increasingly more often fecal transplant as a new way to treat selected IBD patients.

CONCLUSIONS

Even though all this etiopathogenic leads are valuable therapeutic targets for fundamental and clinical gastroenterology, immunology and genetics research we are still far away from the moment when finding "a cause and one etiologic treatment" for all IBD forms and patients.

General diversity of etiopathogenic mechanisms as well as global heterogeneity of IBD makes mandatory permanent literature update for every IBD specialist and also underlines the need to better understand local geographical particularities and trends in Crohn's disease and ulcerative colitis.

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