



IBD PATHOGENESIS AND ITS RELEVENCE TO TREATMENT

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PGIMER

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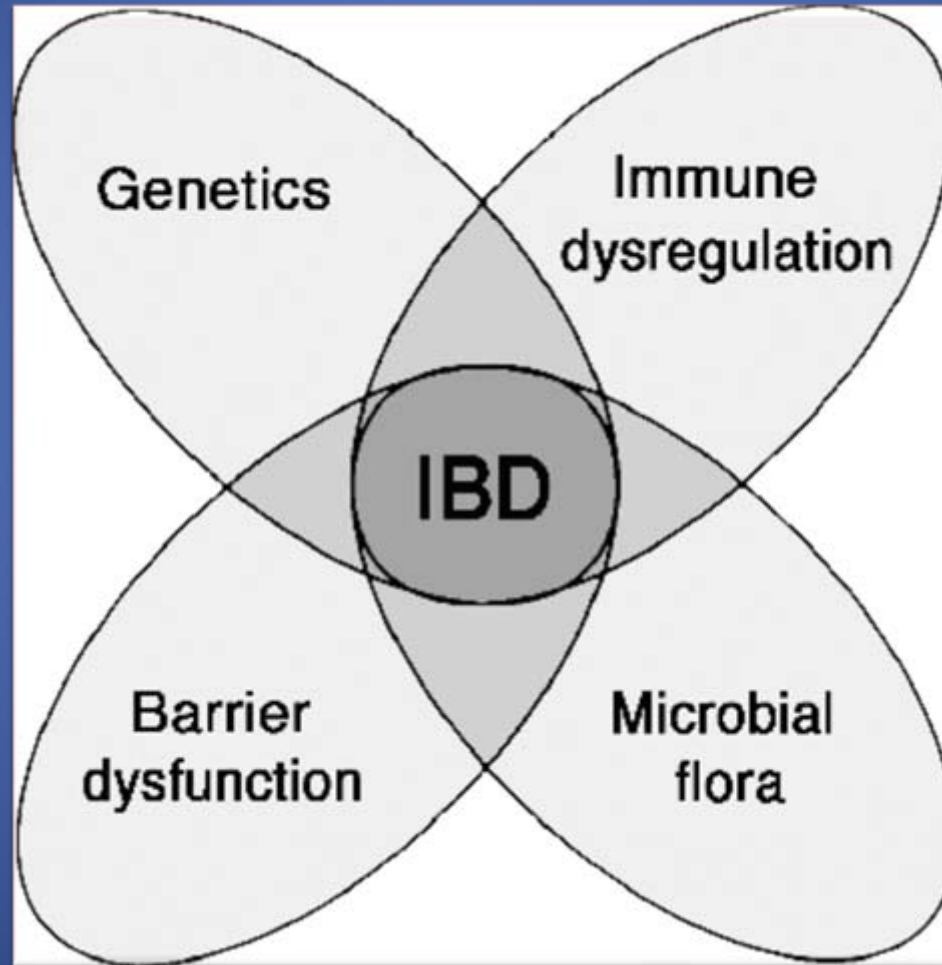
INTRODUCTION

- IBD - immune-mediated disease
- Complex pattern/interplay of host genetics and environmental influences
- Substantial progress in understanding the pathophysiology
- Translated into newer, more effective therapies
- Improved the quality of life of patients with IBD

OUTLINE

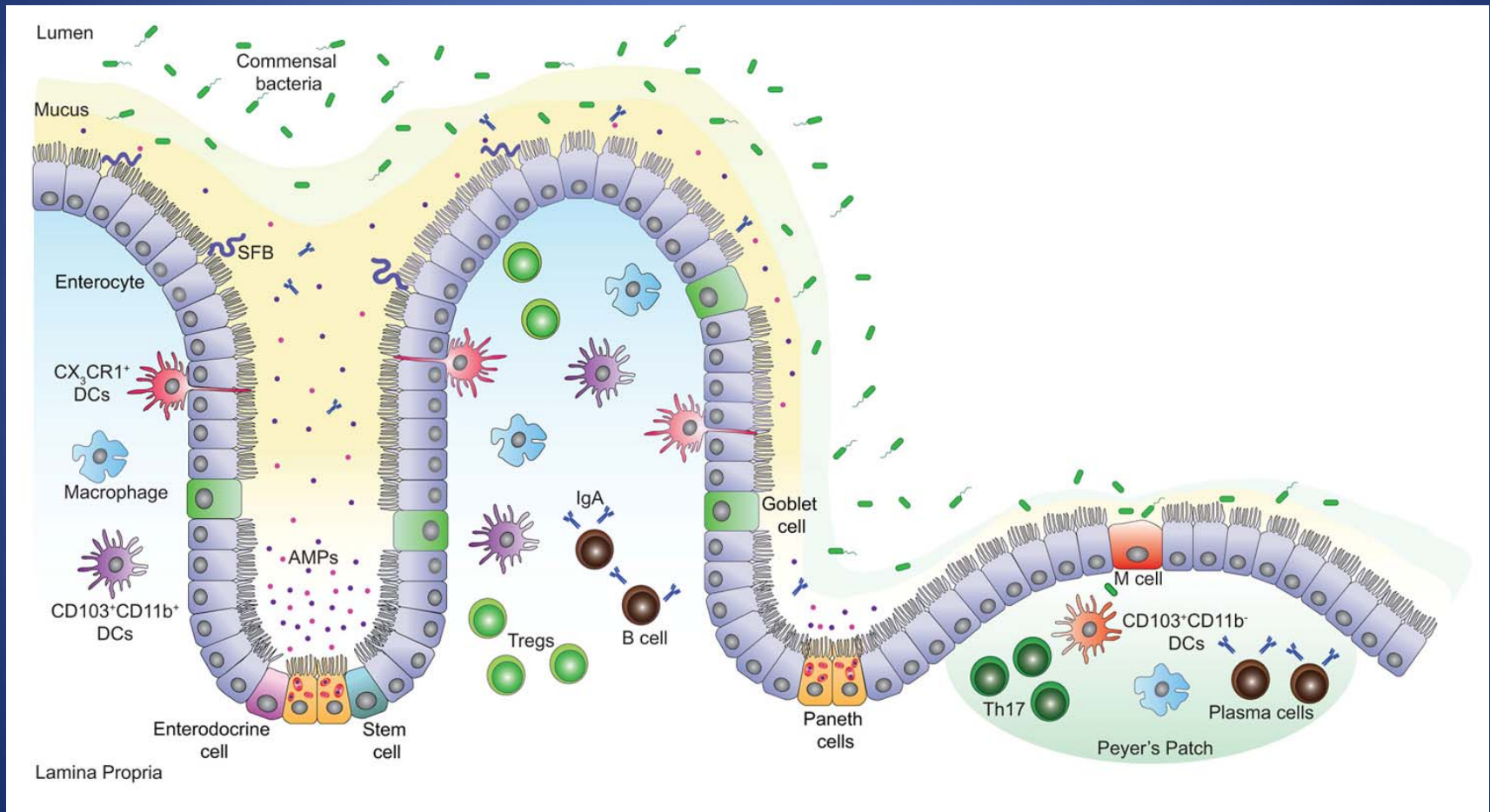
- Overview of gut immunity
 - Innate immunity
 - Intestinal barrier
 - Innate immune cells
 - Adaptive immunity
- Gut microbiome
- Evidence of dysfunction in pathogenesis
- Relevance for therapy

MULTIFACTORIAL PATHOGENESIS

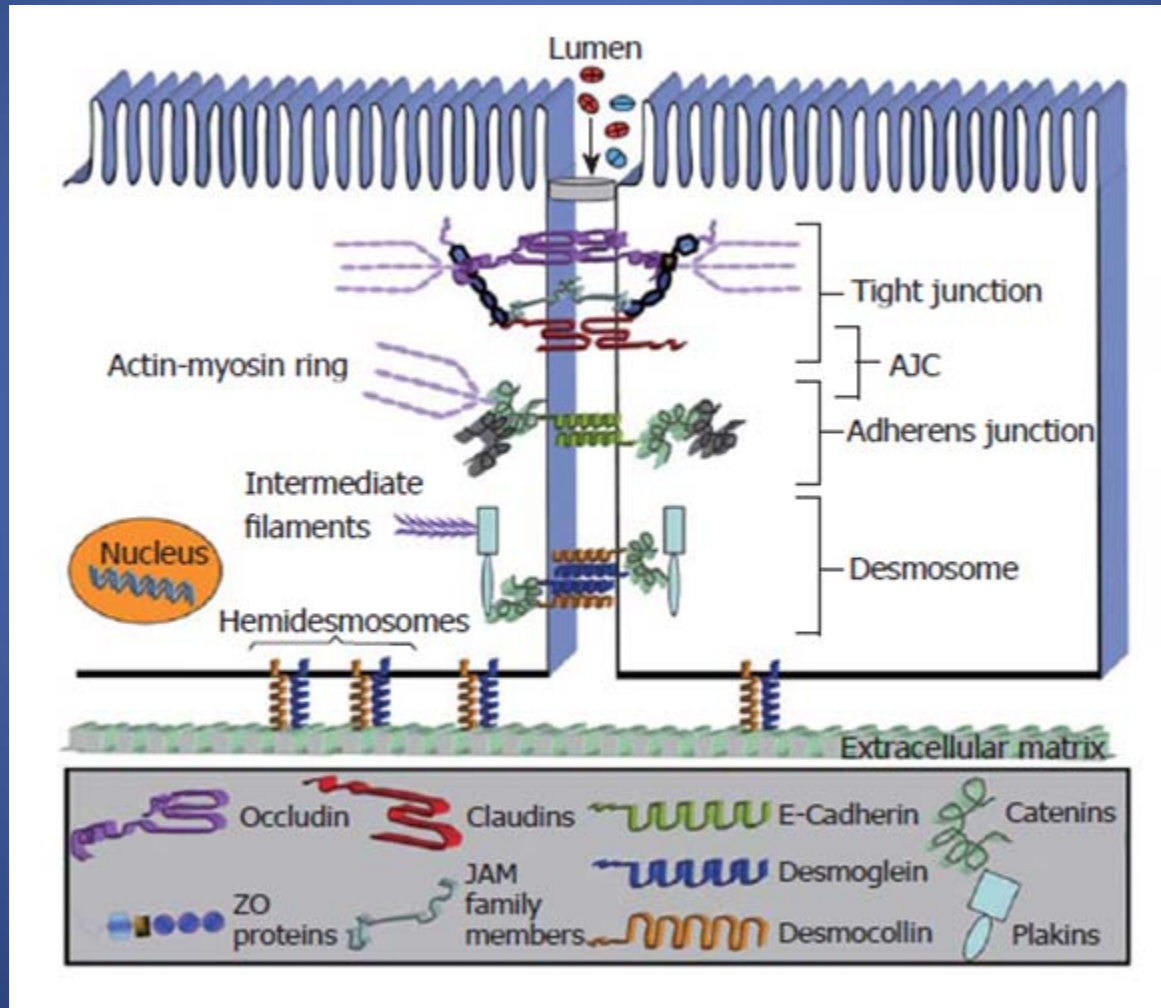


INNATE IMMUNITY

Epithelial barrier



EPITHELIAL BARRIER



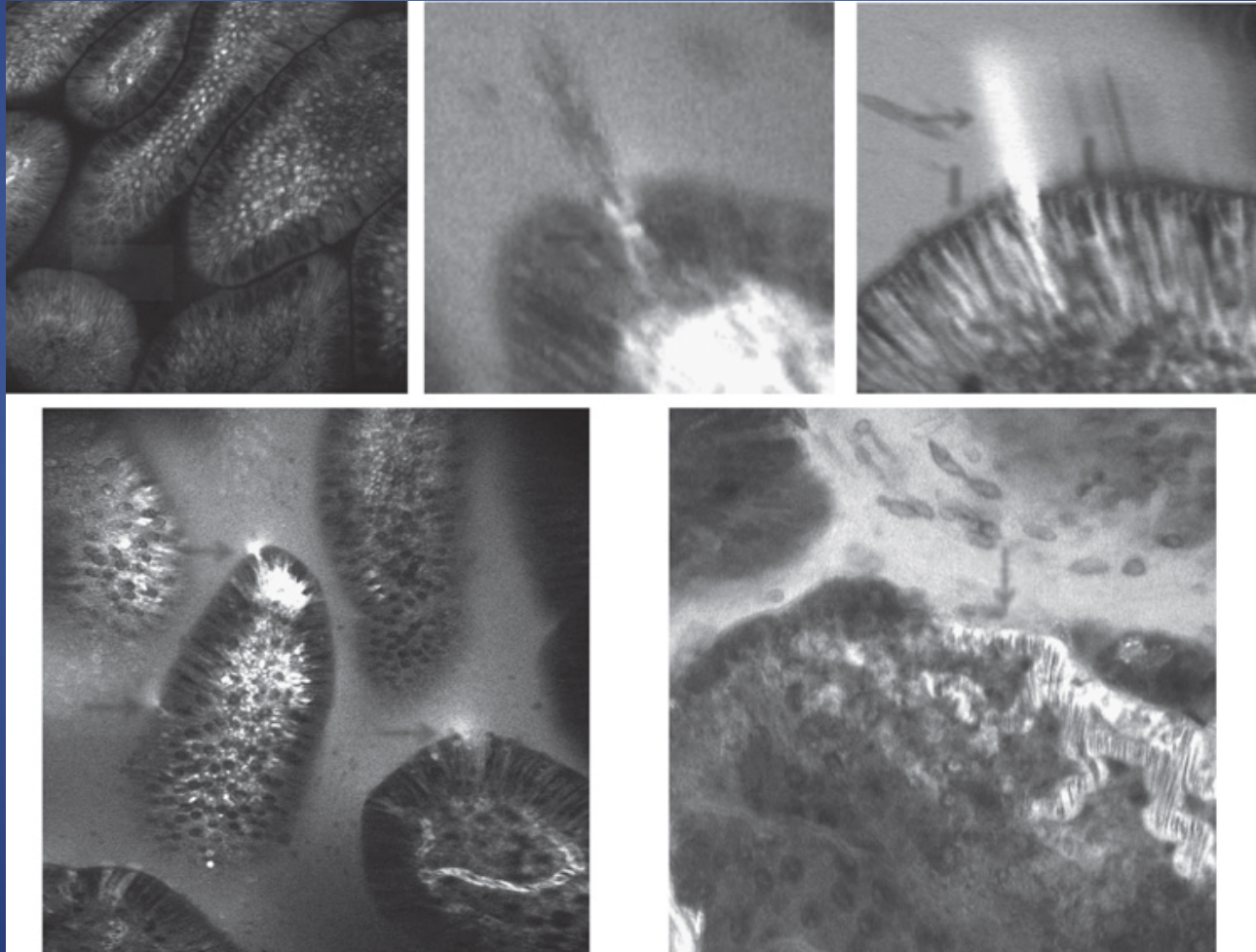
BARRIER DYSFUNCTION

- Abnormal permeability established in CD patients
- GWAS studies – CD risk locus MUC 19 – intestinal mucus layer
- Barrier dysfunction directly has been observed by confocal endomicroscopy
 - Predictive of IBD relapse
- Polymorphic variations in several IBD-associated genes primarily affect epithelial permeability
- Epithelial cell death (paneth cells) – caspase 8 deletion – CD in humans

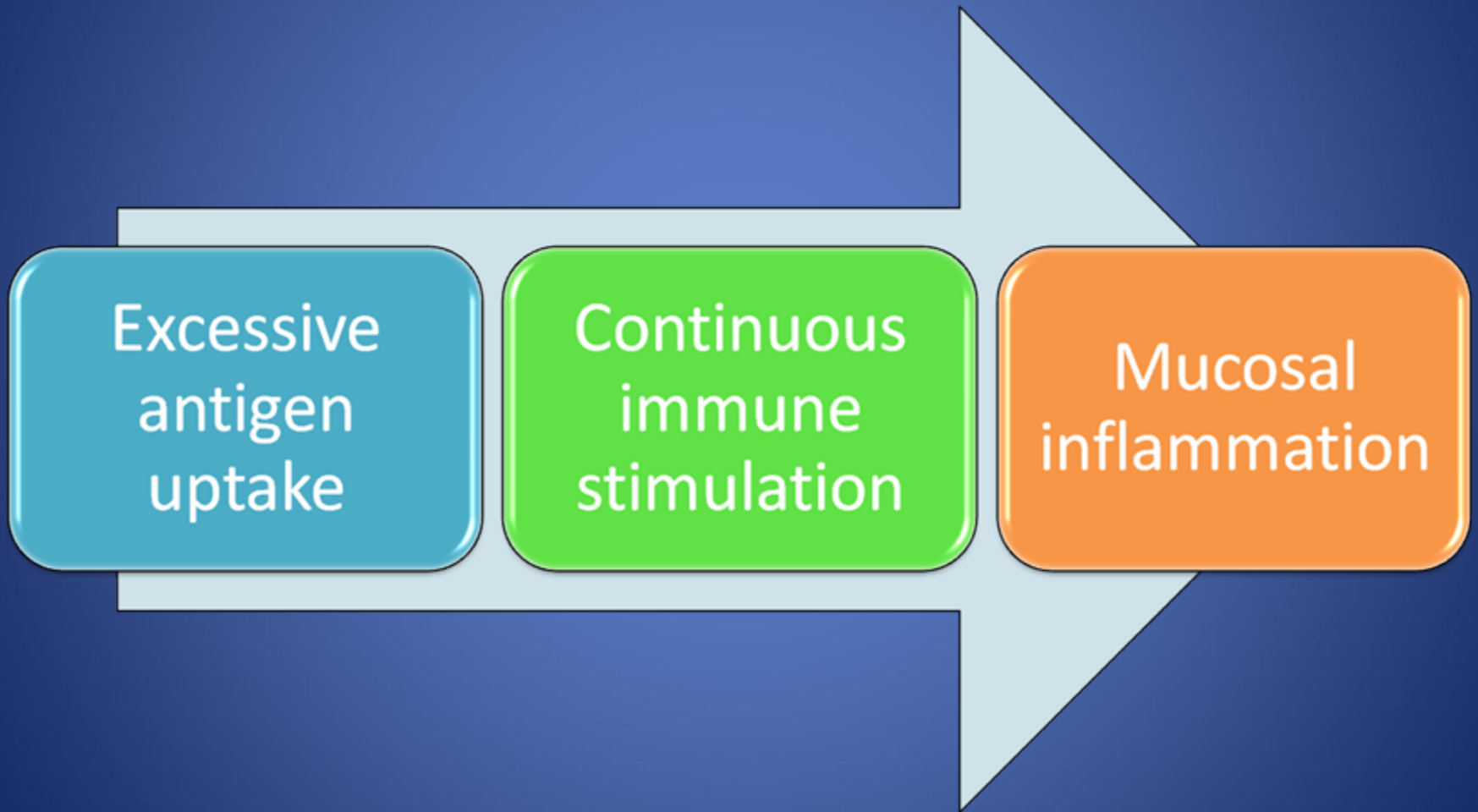
*Am J Gastroenterol 1997, Nature 2012, Gut 2012,
Int J Colorectal Dis 2012, Nature 2011*

LOSS OF BARRIER FUNCTION

Confocal endomicroscopy



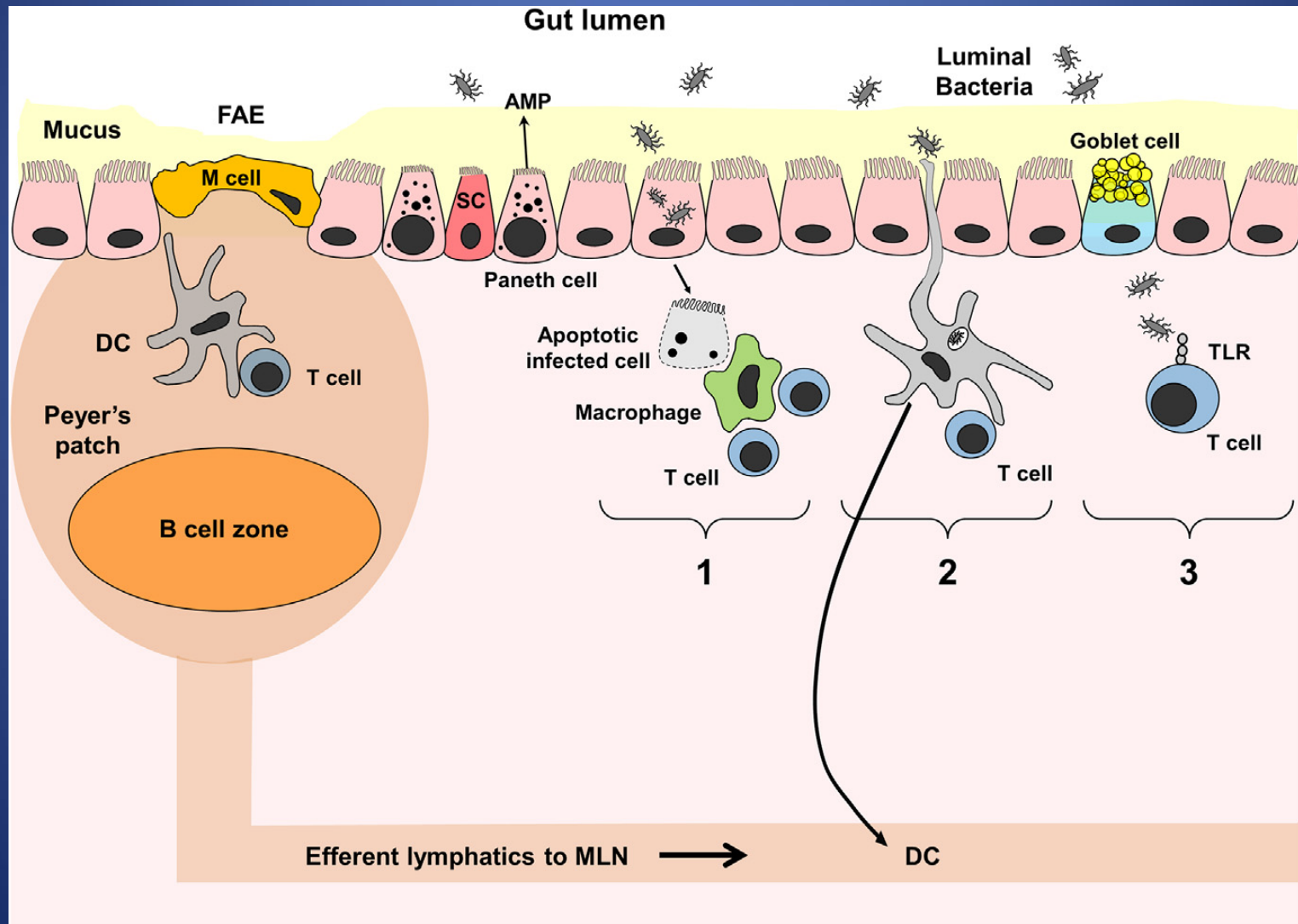
EPITHELIAL PERMIABILITY



INNATE IMMUNITY

- Rapid and less specific response to invading microorganisms or toxic macromolecules
- Innate immune cells
 - Macrophages
 - Dendritic cells
 - Atypical lymphocytes and NKT
- Pathogen recognition receptors (PRR) and pathogen-associated molecular patterns (PAMPs)
- Autophagy pathway

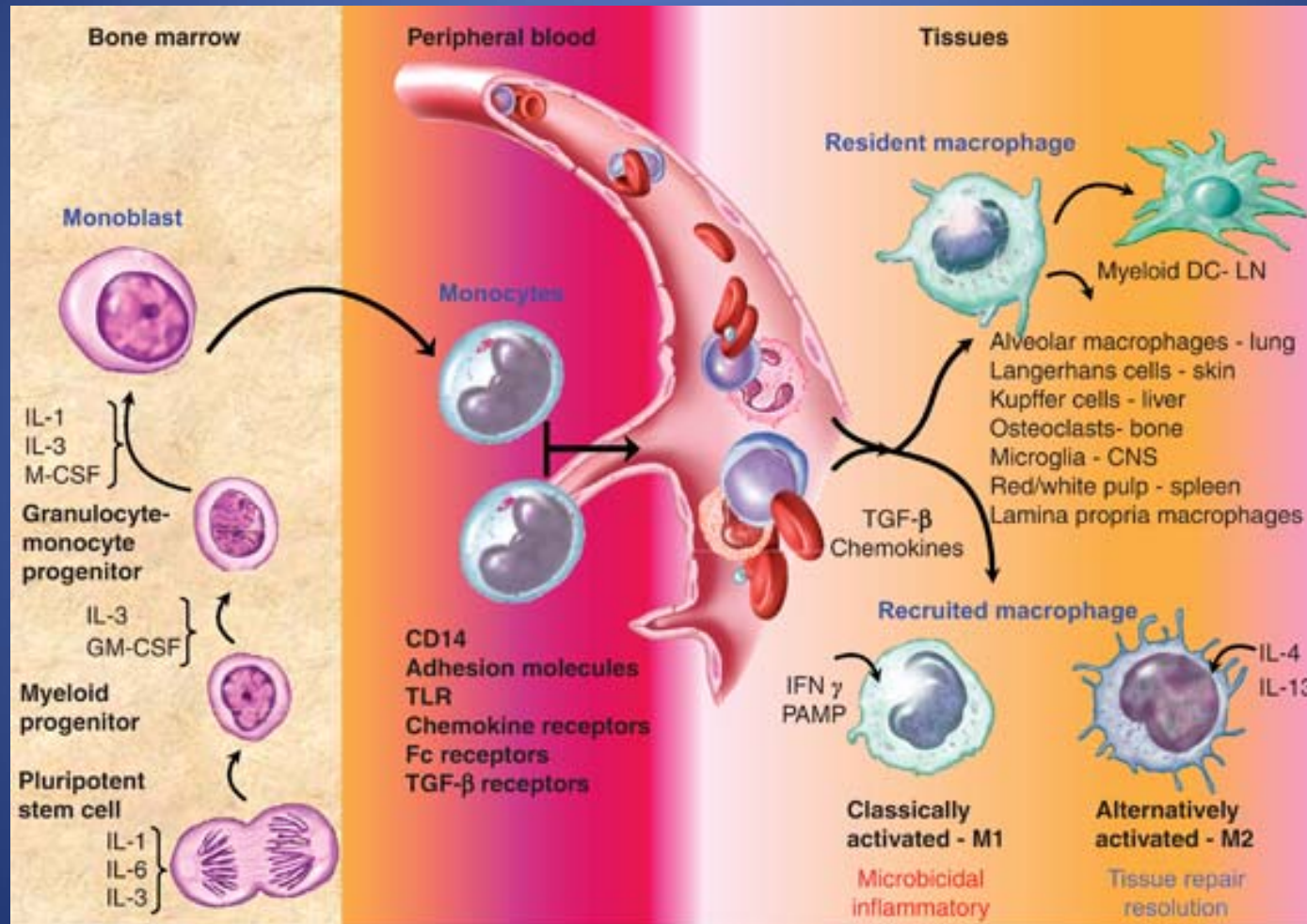
INTESTINAL MUCOSAL IMMUNE SYSTEM



PRR and PAMPs

- PRR
 - Membrane bound (ie, Toll-like receptors [TLRs], C-type lectin receptor) or
 - Cytoplasmic (ie, nucleotide-binding oligomerization domain family members [NODs], retinoic acid–inducible gene 1–like receptor)
- PAMPs (eg, lipopolysaccharide and peptidoglycan)
 - highly conserved molecules on microbes as they are central to survival

MACROPHAGES



MACROPHAGES

Setting of pathogen invasion/inflammation

Convert to a proinflammatory phenotype (ligation of their PRR)



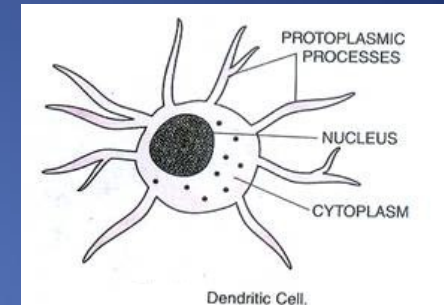
Phagocytosis + secretion of cytokines
IL-1, IL-6, IL-8, TNF- α and TGF- β and recruits cells



Critical link between innate and adaptive immunity

DENDITIC CELLS

- Phagocytic and APCs
- MHC class II molecules
- Normal physiologic state – low levels of costimulatory molecules –
 - T regulatory cells (Tregs)
 - Anti-inflammatory cytokines – IL-4, IL-10 and TGF- β
- Proinflammatory microenvironment -migrate to T cell areas of the GALT
 - induce effector responses
 - Induce mucosal homing receptor $\alpha 4\beta 7$ and chemokine receptor CCR9 on T cells



ATYPICAL LYMPHOCYTES

- $\gamma\delta$ TCR chains
- >10% of SI intraepithelial lymphocytes
- Do not depend on thymus for development
- Do not recognize antigen in association with MHC class I or II
- Effectors against pathogens and tumors and also act as APCs

Natural Killer T cells (NKT cells)

- Mature in the thymus and recognize lipid antigen (presumably bacterial)
- On activation, they secrete large quantities of proinflammatory cytokines and
- readily kill infected cells or tumor cells
- Produce large amounts of proinflammatory cytokine IL-13 have been found in the intestine of patients with UC

INNATE IMMUNE CELLULAR DYSFUNCTION

- The most recent metaanalysis GWAS in IBD
 - susceptibility genes involved in innate mucosal defense (NOD2, CARD9, REL,SLC1A) and
 - antigen presentation (ERAP2, LNPEP)
- Small bowel CD is the loss-of-function polymorphisms in the bacterial sensing gene CARD15/NOD2

INNATE IMMUNE CELLULAR DYSFUNCTION

- In CD -defective inflammatory response to injury and bacterial products
- Failure of clearance of bacteria and inflammatory debris
 - Disruption in the autophagy pathway
 - GWAS - ATG16L1 risk allele contributes to Paneth cell dysfunction
- Multiple ER stress-related genes like XBP1 have been associated with IBD

NOD 2 GENE AND CD

- 2-fold risk for CD heterozygotes and an approximately 20-fold risk FOR homozygotes or complex heterozygotes
- Mutation- reduced NFkB expression
- But NFkB elevated in CD
- Multiple theories to explain this

*Nature 2001, Biochem Biophys Res
Commun 2013*

TARGETS FOR THERAPY

- TNF α inhibitors – Infliximab etc
- IL-12/23 - Ustekinumab
- IL-13 –
 - QAX576 IL-13,
 - Anrukinzumab,
 - Tralokinumab
- TLR- DIMS0150 and BL-7040

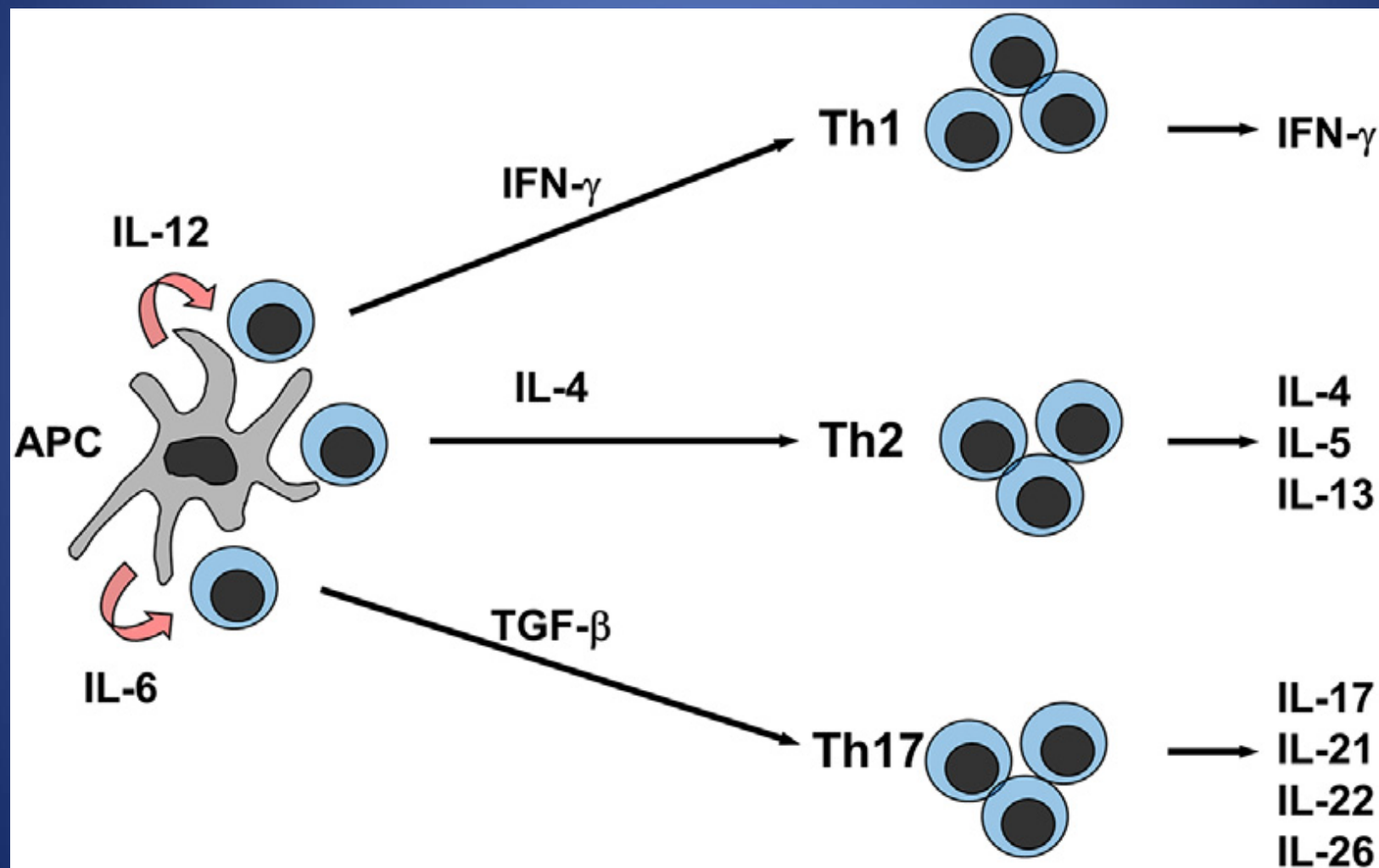
ADAPTIVE IMMUNITY

- Central role in human and experimental IBD models
- Adaptive immune cells
 - T cells
 - Tregs
 - B cells

T CELLS

- CD4+ AND CD8+ - equal proportions in LP
- CD4+ essential role in the pathogenesis
- Costimulation via CD28 (on T cells) and B7.1 or B7.2(on APC) -T-cell activation
- Costimulation mediated by cytotoxic T-lymphocyte antigen 4 (CTLA-4) (on T cells) by B7.1 or B7.2 results in T cell inhibition

T HELPER CELLS



ACTIVATED T CELLS

Cognate antigen+T cell – proinflammatory cytokines – TNF α

Upregulate adhesion molecules, alter the blood flow, endothelial cell shape, and vascular permeability to enhance the migration of inflammatory cells

Structural alterations in the inflamed tissue, including ulceration

Tregs

- Subpopulation of CD4+ T cells
- Restrain not only effector T cells but also innate inflammatory leukocytes
- suppress mucosal inflammation and murine colitis
- Evidence that effector T cells obtained from patients with IBD are refractory to suppression by Tregs

ADAPTIVE IMMUNE CELLULAR DYSFUNCTION

- T cell–driven animal models of colitis mimic human IBD
- IL-23 receptor variant associated with CD impairs the IL-23–induced Th17 effector function and is a protective genetic variant
- Established and emerging therapies – destruction of activated effector T cells or the blockade of T cell–derived proinflammatory cytokines

ADAPTIVE IMMUNE CELLULAR DYSFUNCTION

- Clonal populations of T cells (ie, expansion of T cells in response to persistent and specific antigens) – CD
- Animal model of IBD, similar T-cell clones
- Patients with IBD have antibodies directed against particular microbial antigens

TARGETS FOR THERAPY

| Biologic Target | Antibody/Drug | Mechanism of Action | CD, UC, or Both |
|-----------------------|---------------|--|-----------------|
| CCR9 | CCX282-B | Inhibition of CCR9 | CD |
| | CCX 025 | Inhibition of CCR9 | CD |
| IL-21 | PF 05230900 | IL-21 receptor antagonist | CD |
| IL-13 | QAX576 | IL-13 antagonist | CD |
| | Anrukinzumab | IL-13 antagonist | UC |
| | Tralokinumab | IL-13 antagonist | UC |
| IL-17 | Vidofludimus | Inhibitor of IL-17 A and IL-17F | Both |
| IL-12/23 | Ustekinumab | Blockade of IL-12/23 | CD |
| IL-18 | GSK1070806 | Blockade of soluble IL-18 | CD |
| IL-6 and IL-6R | Tocilizumab | Inhibitor of IL -6 | CD |
| | PF04236921 | Inhibitor of IL -6 | CD |
| IP-10 | MDX 1100 | Blockade of interferon- γ inducible protein (IP-10 or CXCL10) | UC |
| IRAK4/TRAF6/ MyD88 | RDP58 | Disrupts IRAK4/TRAF6/MyD88 signaling and reduces production of proinflammatory cytokines | Both |
| JAK3 | Tofacitinib | Inhibition of JAK3 | Both |

TARGETS FOR THERAPY

| | | | |
|-------------------------------|--|---|------|
| MAdCAM-1 | PF-547659 | Blocks MAdCAM-1 | Both |
| NF- κ B | HE3286 | Synthetic steroid that modulates NF- κ B activity | UC |
| NKG2D | NN8555 | Anti-NKG2D receptor monoclonal antibody | CD |
| PKC | AEB071/Sotrastaurin | PKC inhibitor | UC |
| T Cell | Laquinimod | Reduces IL-17 level and interferes with migration of T cells | CD |
| TLR | DIMS0150 | Blockade of Toll-like receptor | UC |
| | BL-7040 | Blockade of Toll-like receptor | UC |
| TNF- α | Infliximab | Neutralization of TNF- α | Both |
| | Adalimumab | Neutralization of TNF- α | Both |
| | Certolizumab pegol | Neutralization of TNF- α | CD |
| | Golimumab | Neutralization of TNF- α | UC |
| | Debiaerse | Vaccine against TNF- α consisting of a TNF- α derivative TNF- α kinoid | CD |
| Effector T cells, B cells | Antigen specific Type 1 regulatory cells (OvaSave) | Autologous ova expanded regulatory T cells injected | CD |
| α 4 integrin | AJM-300 | Blockade of α 4 integrin | CD |
| α 4 integrin | Natalizumab | Blockade of α 4 integrin | Both |
| α 4 β 7 integrin | Vedolizumab | Blockade of α 4 β 7 integrin | Both |
| β 7 integrin | Etrolizumab (aka rHuMab β 7) | Anti- β 7 integrin | UC |

GUT MICROBIOTA

- Pivotal role
- No germs: No IBD (No IBD in germ free rats)
- The flora in IBD:
 - More often has pathogens
 - More adherent bacteria
 - Reduced diversity : ecology disturbed
 - Increased Enterobacteriaceae; Decreased Firmicutes
 - ? Role of fungus (ASCA +itivity)

MICROBES AND IBD

- Change of flora effective in Rx
 - Antibiotics
 - Probiotics
 - Elemental diet
- Infection precipitates flares

PATHOGENIC OR ALTERED COMMENSAL BACTERIA

- Enhanced epithelial adherence
- Epithelial invasion
- Resistance to killing
- Acquisition of virulence factors
- Stimulation of innate and adaptive immunity
- production of butyrate causes poor epithelial integrity
- Increased production of toxic metabolites like H₂S

MICROBIAL PATHOGENS

- *Mycobacterium avium paratuberculosis (MAP)*
 - May infect genetically susceptible CD patients with intracellular bacterial killing defects due to ATG16L1, NOD2, or NCF4 polymorphism
- Adherent invasive *E.coli*
 - Identified from inflamed ileal mucosa of CD
 - E coli DNA in 80% of granulomas
 - High titres of Anti-Ecoli Ab in 55% of Crohns disease
- Enterotoxigenic *Bacteroides fragilis* in 19% of IBD
- *Enterococcus faecalis*

DYSBIOSIS

- Alteration of indigenous microbiota alters
 - Dominant antigens
 - Metabolic function of the gut
- Intestinal mucosal bacteria found at concentration greater than 10^9 /ml in
 - 95% of IBD patients
 - 65% of IBS patients
 - 35% of healthy controls
- CD occurs in intestinal segments with the highest bacterial concentrations

TARGETS FOR THERAPY

- Probiotics
- Antibiotics
- Limited success

TO SUMMARIZE

| TARGET | THERAPY |
|--------------------|---|
| Epithelial barrier | |
| Innate immunity | Biologicals |
| Adaptive immunity | Steroids, Immunosuppressive medications, Biologicals, Immunotherapy |
| Gut microbiota | Probiotics, Antibiotics |

CONCLUSION

- Studies of mucosal immunity have led to recent advances in therapy
- GWAS data and studies on microbiome – unraveled the complex interaction between host and environment
- Many aspects of mucosal immunity remain unclear
- Clinical phenotype doesn't correspond to the immunophenotype of patients