# The gut has a complex microbial community (microbiome)

Dominant gut phyla:

Bacteroidetes, Firmicutes, Actinobacteria, Proteobacteria, Verrucomicrobia

Predominant families in the:



Donaldson, Lee and Mazmanian, Nature Rev Microbiol. 14: 20, 2016

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Mice from Taconic but not mice from Jackson have Segmented Filamentous Bacterium (SFB), which attaches itself to the epithelium in the small intestines and grows as a chain. Ivanov et al, Cell 139: 485, 2009.

#### Immune control of gut microbes



Hooper and MacPherson, Nature Rev Immunol. 10: 159, 2010

#### Immune control of gut microbes



Donaldson, Lee and Mazmanian, Nature Rev Microbiol. 14: 20, 2016

### IgA Background

- IgA is secreted in mucosal tissue and is transported across mucosal epithelial barriers by the poly-Ig receptor
- Poly-Ig receptor is cleaved; the part that stays bound to IgA is "secretory component"



#### IgA in small intestines



### IgA Background

- Some IgA is T cell-dependent; some IgA is T cellindependent
- Some IgA is produced in germinal center responses and exhibits somatic mutation
- Deep sequencing of IgA reveals a diverse response that includes a few specificities produced abundantly
- Gut bacteria that have IgA on them can be revealed by staining with a fluorescent anti-IgA and using flow cytometry and cell sorting followed by sequencing of 16S rRNA

### Sorting IgA-coated bacteria



Palm et al. Cell 158: 1000-10, 2014

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**Fotal** gA+ Bacteroidales (S24-7 gen.) \* UC Clostridiales Oscillospira Clostridiales gen. UC Lachnospiračeae Bacteroides Rikenellaceae gen. Lachnospiraceae gen. Coprococcus Ruminococcu's (Lachno) Ruminococcus Turicibacter Lactobacillus \* Anaerostipes Mucispirillum Adlercreutzia UC Ruminococcaceae Roseburia Unclassified Bacteria Erysipelotrichaceae gen. Bilophila Dehalobacterium Anaeroplasma Ruminococcaceae gen. Dorea Mogibacteriaceae gen. Clostridium Sutterella Coprobacillus UC Erysipelotrichaceae \* Prevotellaceae gen.

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#### ICI: relative abundance IgA+/ relative abundance IgA-

Mice: 4 taxons enriched in IgA+ coated; 22 taxons enriched in IgA-

Palm et al. Cell 158: 1000-10, 2014

Conclusion: bugs differ in what proportion is coated with IgA or not coated

#### Formulate two or three hypotheses and

come up with **one** experimental approach to address one or more of these hypotheses

Tools to consider:

-Flow cytometry to identify bugs that are coated with IgA or not coated (and relative proportions)

-Introduction of individual species or sets of species into germ free mice

-Mice or people with diseases, immunodeficiencies, and/or different diets

Hypothesis one: amount of antigen available to the immune system.

#### Most IgA plasma cells in gut are dependent on gut bacteria (B6 mice)



Hapfelmeier et al. Science 328: 1705-9, 2010

HA107 gavaged 6 times over 14 days; IgA plasma cells (green) evaluated after 4 weeks total (blue: DAPI staining of nuclei) (ASF: altered Schaedler flora, a mixture of 8 mouse gut bacteria that creates a relatively stable community)

#### Threshold for IgA response to E coli K-12





Live bugs induce IgA >10X better than heat-killed bugs

live E. coli are 10x better stimulus than heat killed E. coli

Hypothesis two: location of bugs relative to gut epithelium is important (bug must penetrate mucus layer and/or invade tissue to induce IgA response)

## Consortia of IgA+ vs. IgA- microbes: different degrees of IgA induction



Palm et al. Cell 158: 1000-10, 2014

## Consortia of IgA+ vs. IgA- microbes: invasion of the mucus layer by the IgA+ bugs



Palm et al. Cell 158: 1000-10, 2014 16S rRNA universal probe DAPI

Hypothesis three: bugs found in lumen of colon AND not in small intestines do not induce much IgA



intestines induce specific IgA; most bugs in colon may not induce IgA

Bunker et al. Immunity 43: 541, 2015.

Duodenum

3

2

Log<sub>10</sub>(IgA<sup>+</sup>/IgA<sup>-</sup>)

3

Hypothesis four: more "pathogenic" bugs preferentially induce IgA

## Sorting IgA-coated bacteria from people with Inflammatory Bowel Disease



Palm et al. Cell 158: 1000-10, 2014

#### Consortia of IgA+ vs. IgA- microbes



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Palm et al. Cell 158: 1000-10, 2014

## Consortia of IgA+ vs. IgA- microbes: inflammatory action



## A commensal gut microbe uses IgA to help it colonize the mucus layer



Bacteriodes fragilis, an abundant commensal bacterium (which is naturally coated with IgA), requires IgA to colonize the mucus layer well (green arrow: bacteria; yellow arrow: epithelium; in the gut it induces expression of a capsular polysaccharide that induces strong IgA production

Donaldson et al. Science 360: 795-800, 2018

# Microbiota: an important variable in animal research

#### Accounting for reciprocal hostmicrobiome interactions in experimental science

Thaddeus S. Stappenbeck<sup>1</sup> & Herbert W. Virgin<sup>1</sup>

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### Sorting IgA-coated bacteria: role of location in IgA induction



Bunker et al. Immunity 43: 541-553, 2015