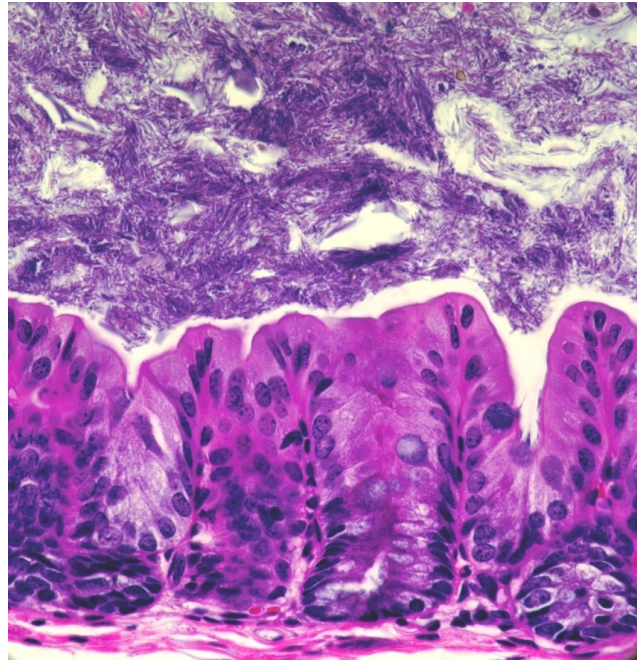


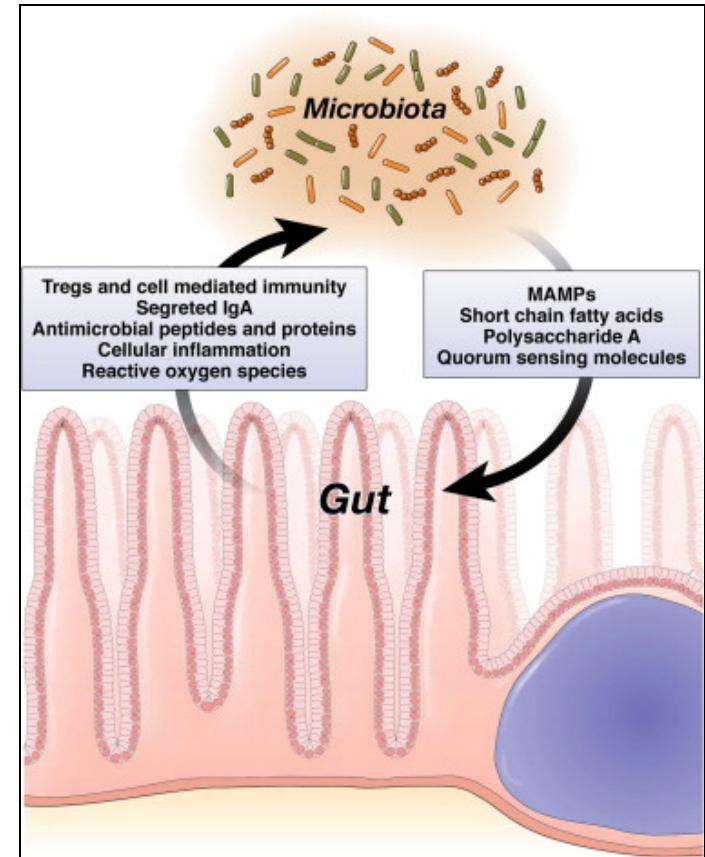
Control of Epithelial Homeostasis by the Microbiota



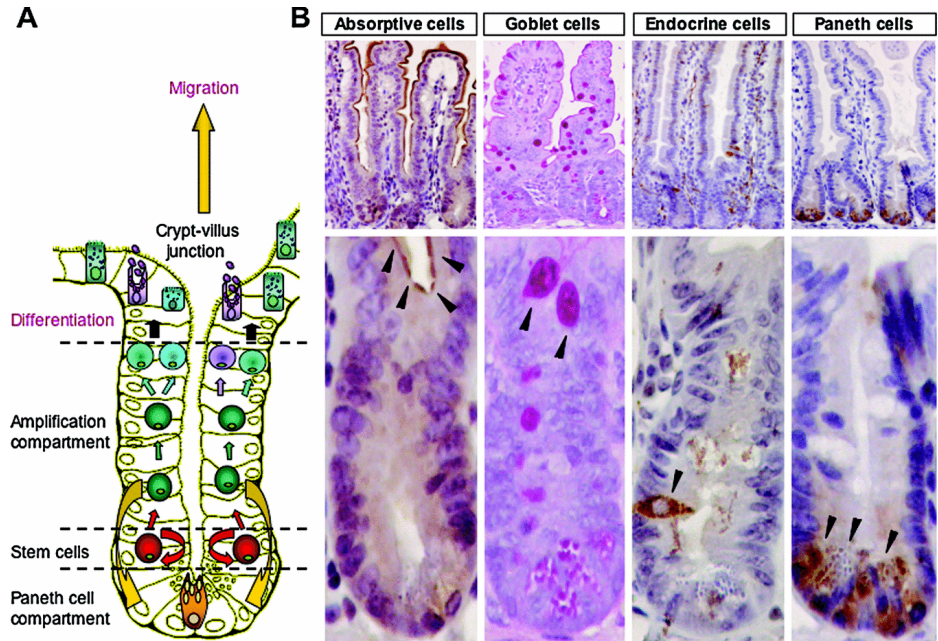
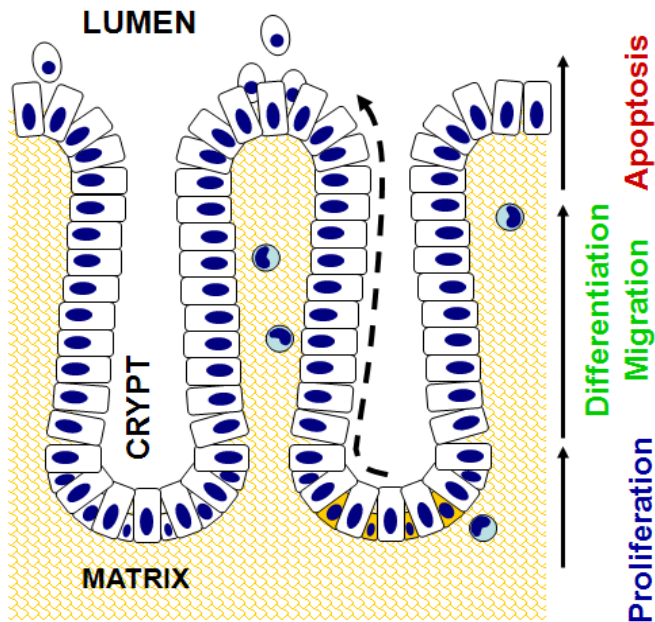
Andrew S. Neish M.D.
Department of Pathology
Emory University School of Medicine
Atlanta GA, USA

Effects of the microbiota on the gut

- Competitive exclusion of pathogens
- Metabolic/nutritional/energy utilization
 - Vitamin synthesis
 - SCFA as energy source- role in obesity
- Adaptive Immune Regulation,
 - Induction of Immunosuppressive T cells (Tregs)
- Innate Immune Regulation
 - Dampening of inflammatory responses
- Epithelial development and survival
 - Cytoprotective effects of PRR signaling
 - Stimulation of barrier function, IEC restitution, proliferation



The Intestinal Epithelia (motility, proliferation, differentiation)



Germ-free studies

- Small intestinal crypts exhibit a slower turnover of the epithelial cells, with crypt to villus transit time doubling
- Markedly attenuated regenerative responses to colonic injury

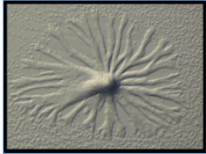
The microbiota can influence normal homeostasis of the gut aside from traditional innate immune responses

Question:

How the normal microbiota mechanistically interacts with the epithelia is not well understood

How the microbiota can influence epithelial growth and proliferation is not well understood

Influence of Reactive Oxygen Species (ROS) on cell proliferation/differentiation



Bloomfield G, et al. **Superoxide signaling required for multicellular development of *Dictyostelium***. 2003, *J. Cell Sci.* 116:3387



Tsukagoshi T et al. **Transcriptional regulation of ROS controls transition from proliferation to differentiation in the root**. 2010, *Cell.* 143:606



Owusu-Ansah E, Banerjee U. **ROS prime *Drosophila* haematopoietic progenitors for differentiation**. 2009, *Nature.* 461:537

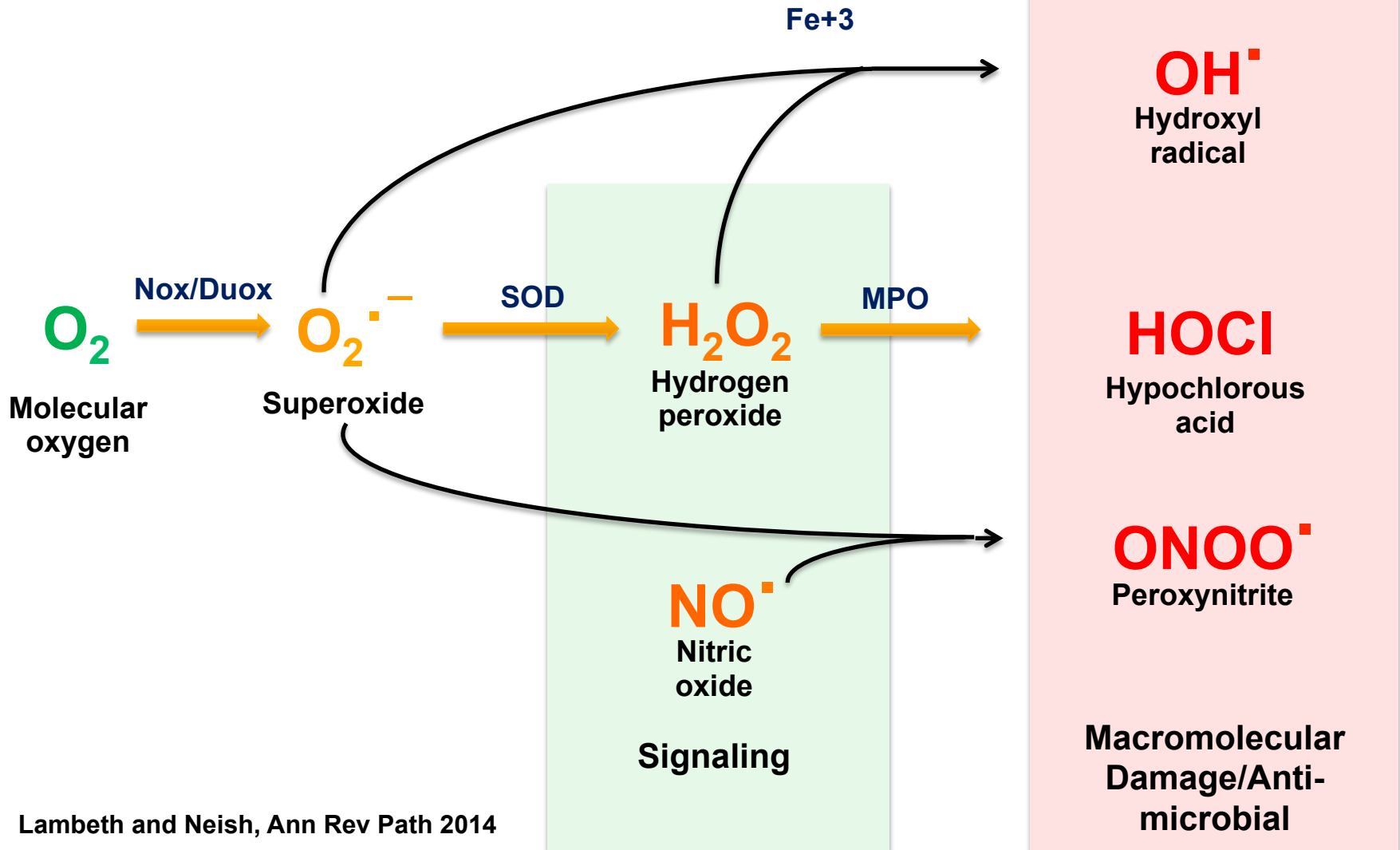


Love S, et al. **Amputation-induced ROS are required for successful *Xenopus* tadpole tail regeneration**. 2013, *Nat Cell Biol.* 15: 222



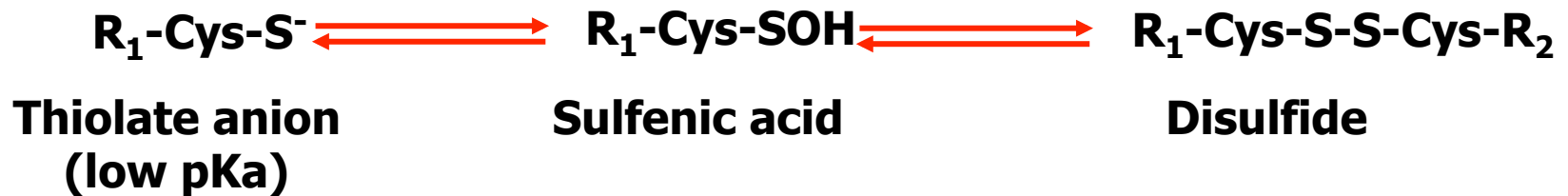
Morimoto H, et al. **ROS Are Required for Mouse Spermatogonial Stem Cell Self-Renewal**. 2013, *Cell Stem Cell.* 12:774

Reactive Oxygen Species



Redox regulation of enzymes

- Mediated by transient oxidation of low pKa catalytic cysteines
- Rapid, reversible and highly localized
- Known target enzymes:
 - Ub like protein ligases; SUMO, Ubc12
 - Dual specificity protein kinases
 - LMW-PTPases
 - Keap1

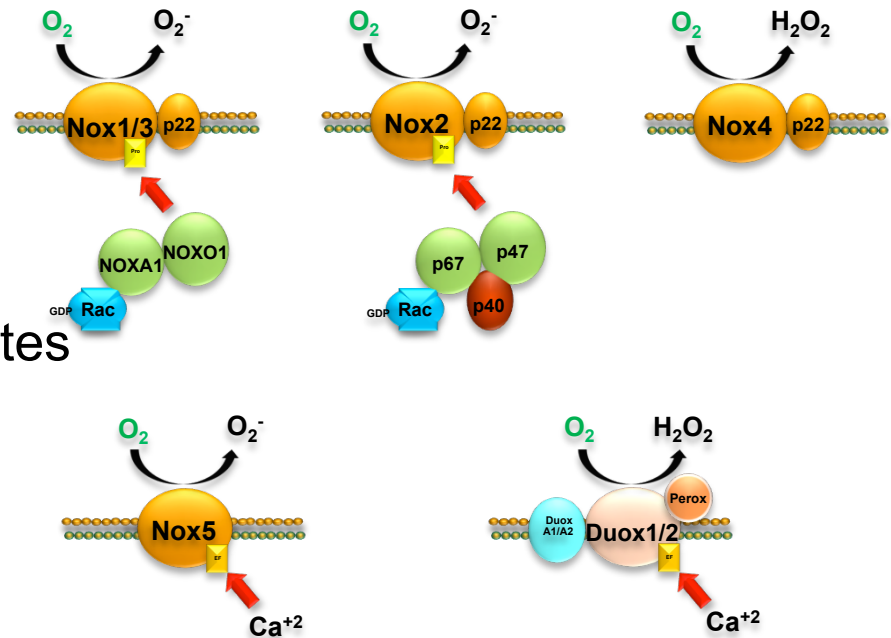


Oxidant stress



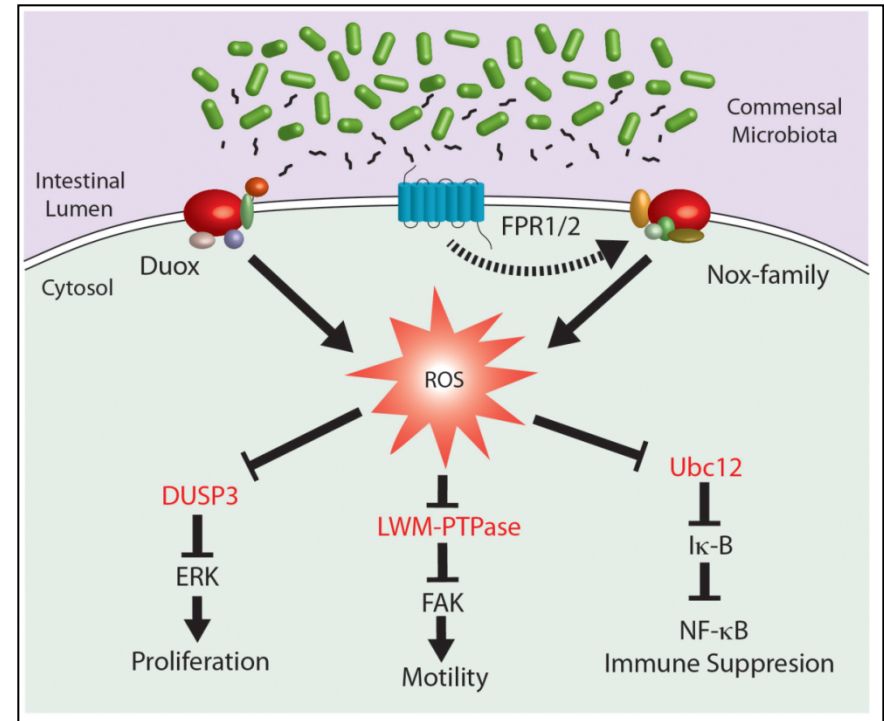
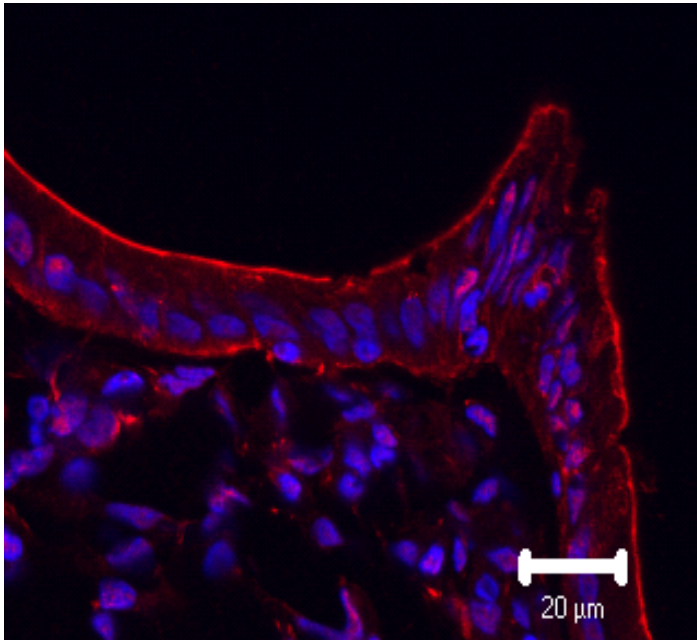
Induced ROS as a response to microbes

- Plants
 - Defensive: Immediate response, lignification response
 - Root development
- *C. elegans*
 - Defensive: anti-microbial, ceDuox/BLI-3
 - Cuticle development
- *Drosophila*
 - Anti-microbial (dDuox)
 - Gut epithelial homeostasis (Nox)
- Mammalian “professional” phagocytes
 - Microbiocidal oxidant burst (Nox2)
- Mammalian barrier epithelia
 - Signaling and homeostatic function (Nox1)



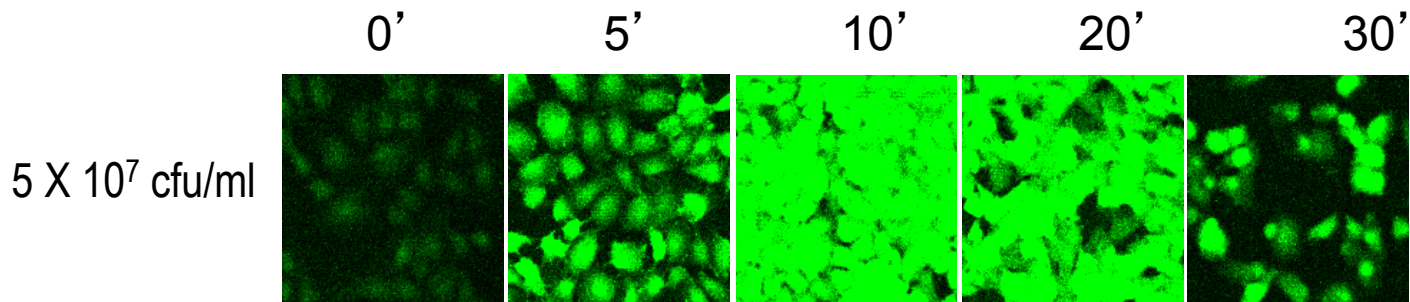
- **Strong conservation of NADPH generating enzymes (Nox' s)**

The Formyl Peptide Receptor pathway



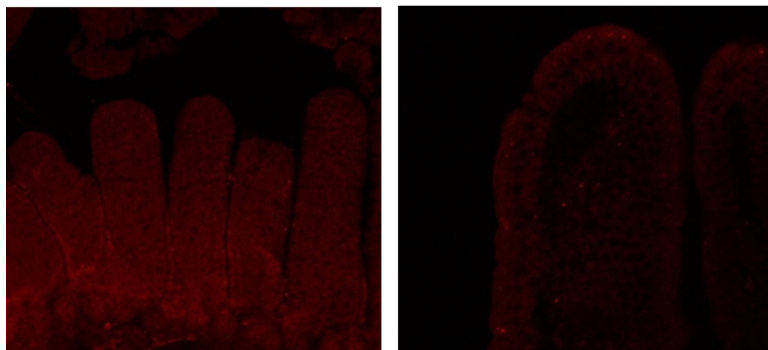
Prokaryotic regulation of epithelial responses by inhibition of IκB-α ubiquitination, *Science*. 2000; 289, 1560-1563.
 Bacterial modulation of epithelial signaling via deneddylation of Cul-1. *J. Immunol.* 2005; 175:4194-4198.
 Commensal bacteria repress cullin dependant-signaling via generation of reactive oxygen species. *EMBO J.* 2007;26:4457-66.
 Commensal-Epithelial Signaling Mediated via Formyl Peptide Receptors. *A J. Path*, 2010; 177:2782-90.
 Enteric commensal bacteria potentiates epithelial restitution via ROS-mediated inactivation of FAK phosphatases. *Proc. Natl. Acad. Sci.* 2011;108:8803-8
 Enteric commensal bacteria induced ERK via FPR dependent redox modulation of DUSP3. *J. Biol. Chem*, 2011; 286:38448-38455.
 Annexin A1- FPR1-Nox1 dependent redox signaling promotes epithelial wound repair. *J Clin Invest*, 2013; 123(1):443-54

Commensal bacteria induce rapid epithelial ROS generation *in vitro* and *in vivo*



Lactobacillus colonized onto IEC-6 monolayers prior to DCF detection of ROS

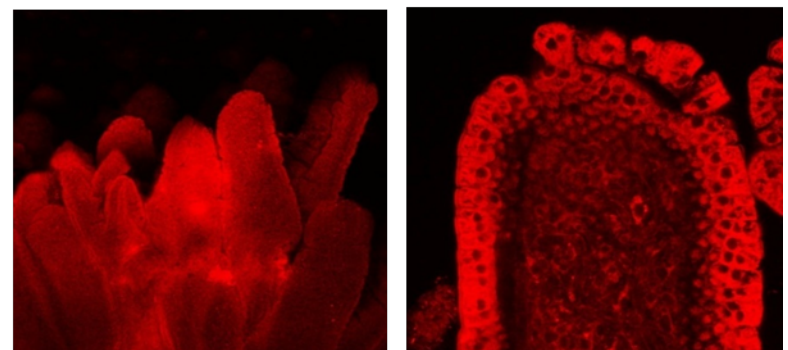
Control Ileum



10x

40x

Lactobacillus, 60'

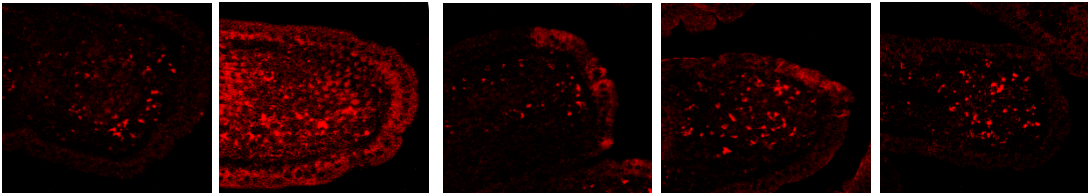
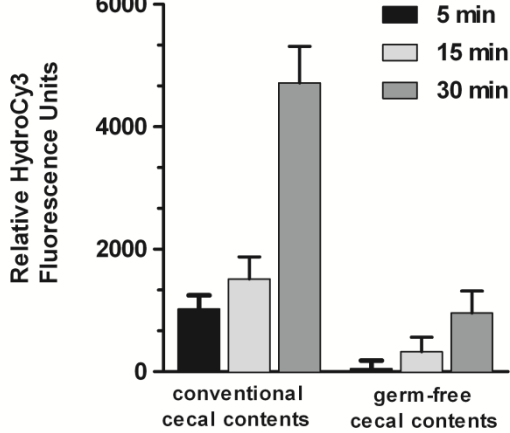
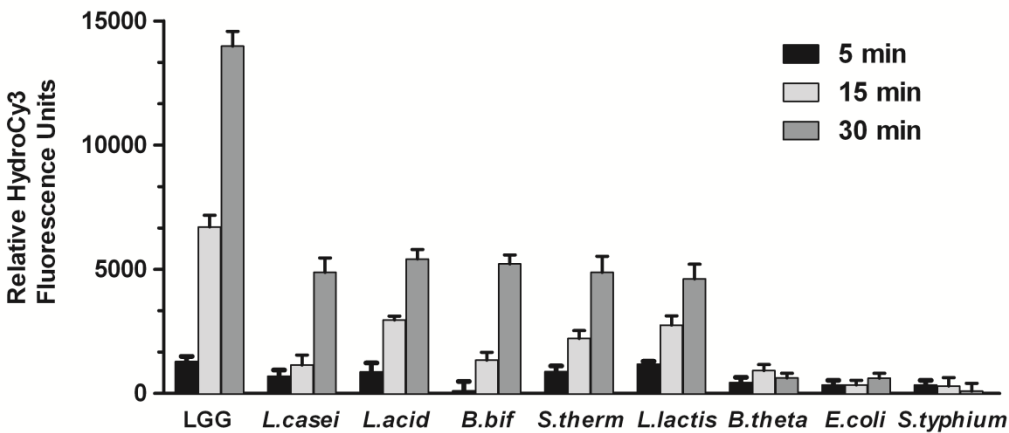


10x

40x

Conventional 10 week female mice were fed cultures of *Lactobacillus* or sterile PBS control for 60 minutes. Hydrocyanine-Cy3 reagent dosed i.p. 20 min prior.

Differential ROS induction by distinct taxa of commensal bacteria *in vitro* and *in vivo*

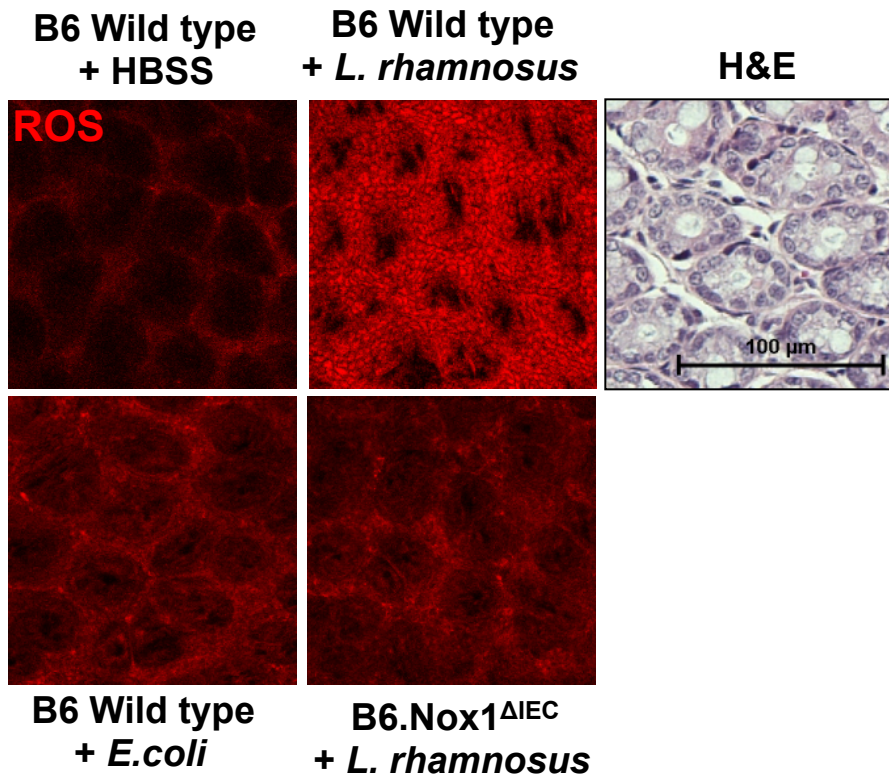


HBSS *L. rhamnosus* *L. casei* *S. themophilus* *E. coli*

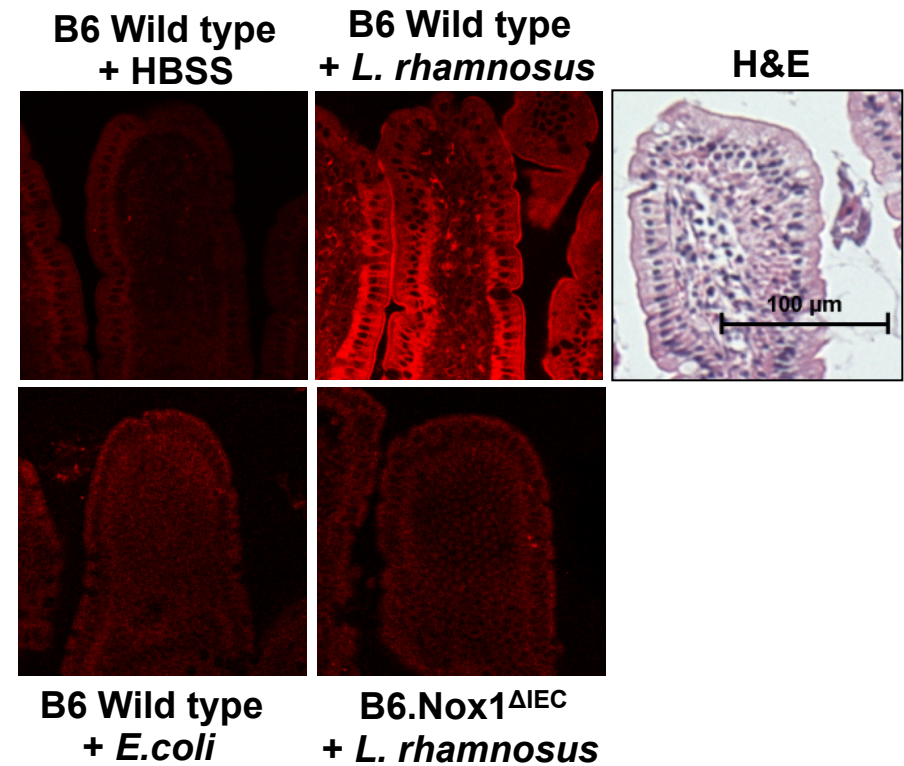
Mouse jejunum

Colonization of intestine with *Lactobacillus* induces Nox1-dependent generation of cellular ROS

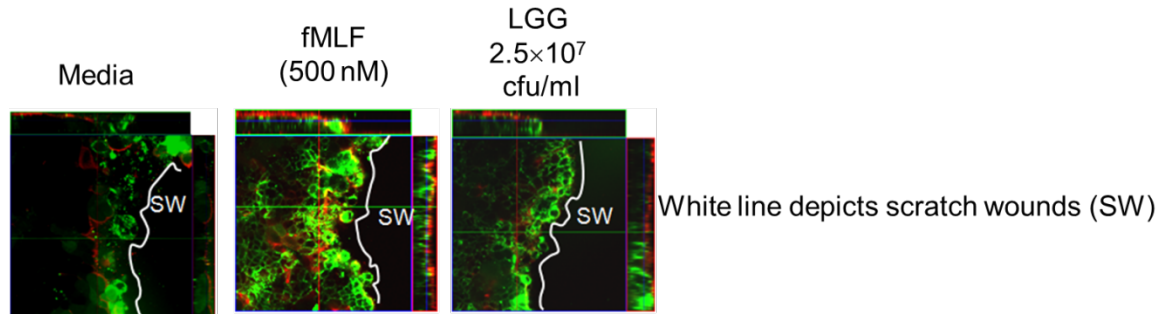
6-week-old colonic tissue



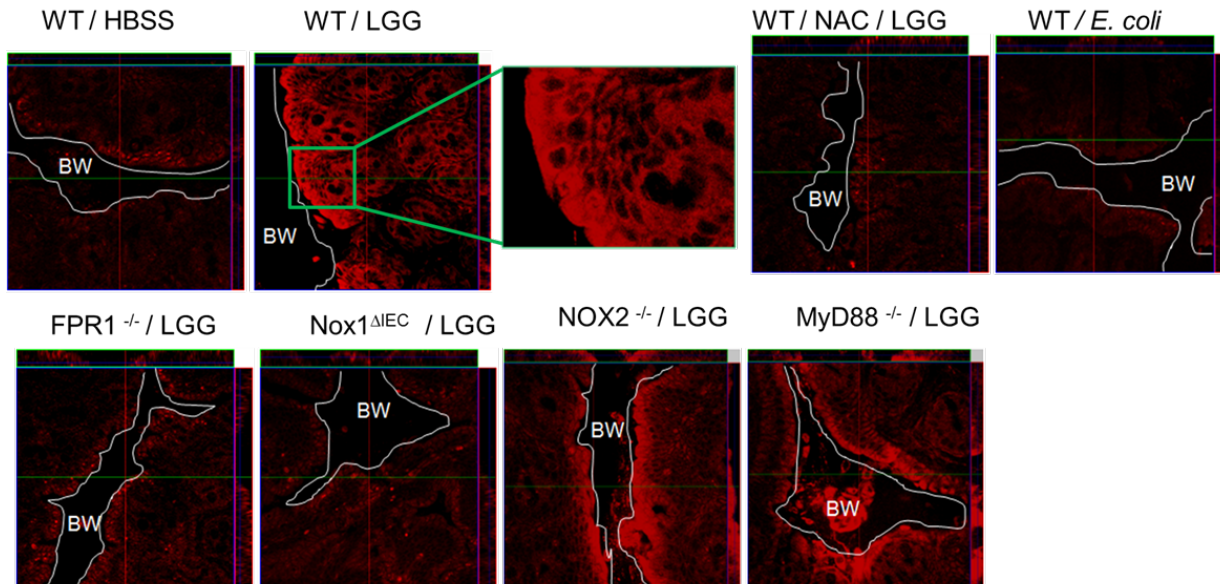
6-week-old distal small intestine tissue



Lactobacilli induces rapid epithelial ROS generation *in vitro* and *in vivo* at wound edges



■ ROS (CM-H2DCF-DA) ■ Wheat Germ Agglutinin



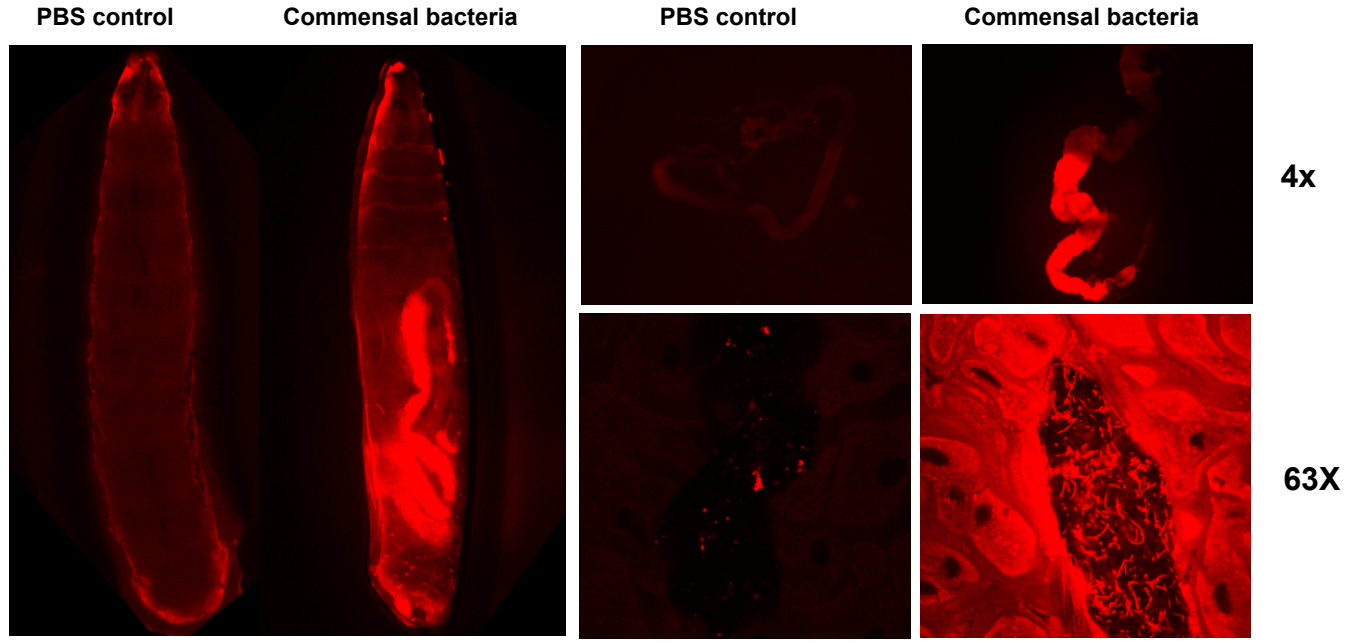
■ ROS (Hydrocyanine3)

White line depicts biopsy wounds (BW)

Commensal bacteria induce rapid epithelial ROS generation *in vivo* in the Drosophila gut

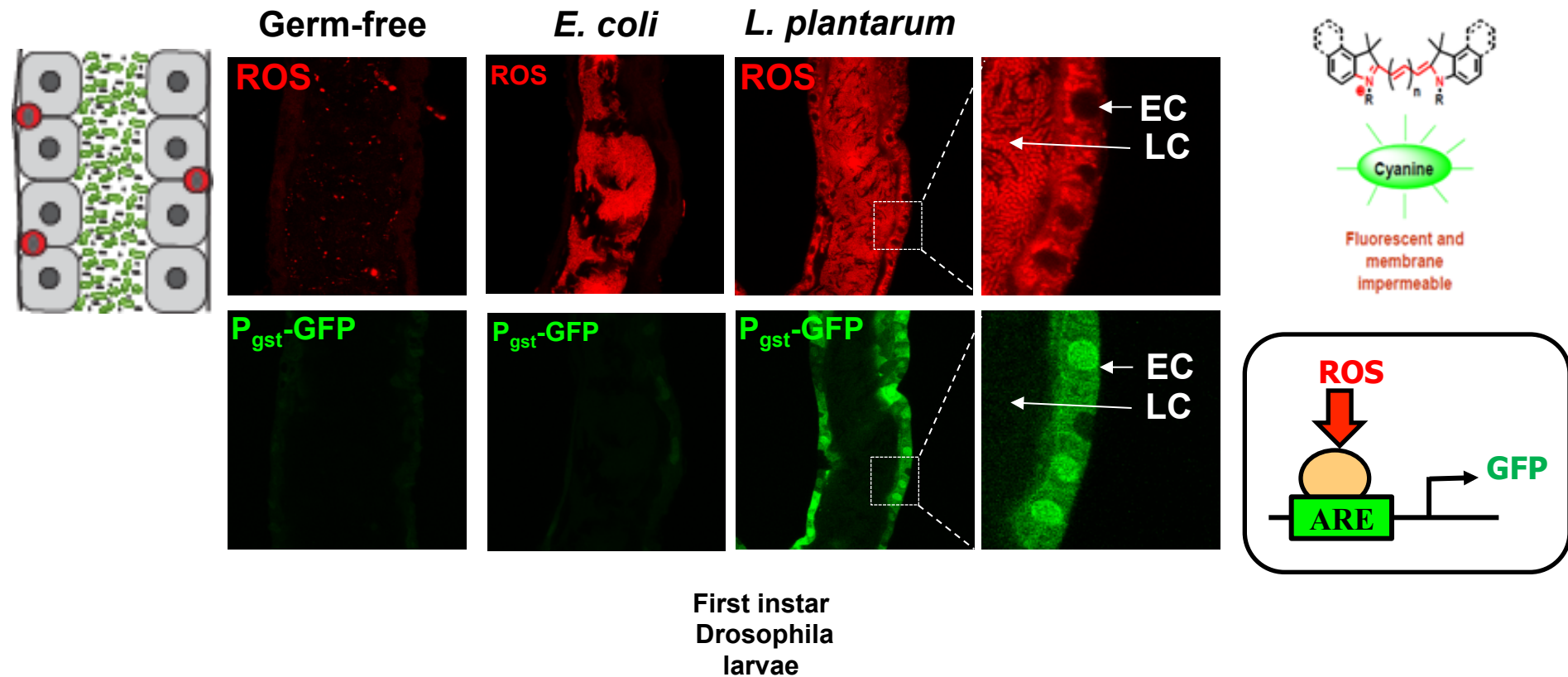


Third instar Drosophila larvae



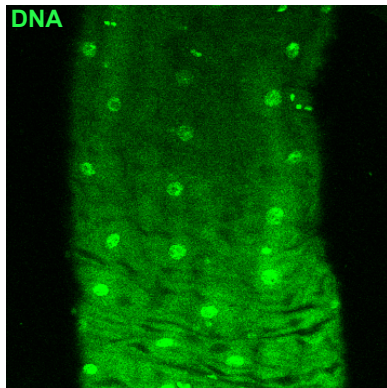
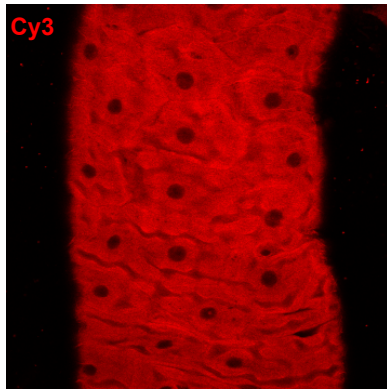
Axenic third instar larvae were fed cultures of *Lactobacillus plantarum* or sterile PBS control for 30 minutes Both *L. plantarum* and PBS cultures included hydrocyanine-Cy3 reagent.

Colonization of the first instar *Drosophila* midgut by *Lactobacillus plantarum* induces cellular ROS generation

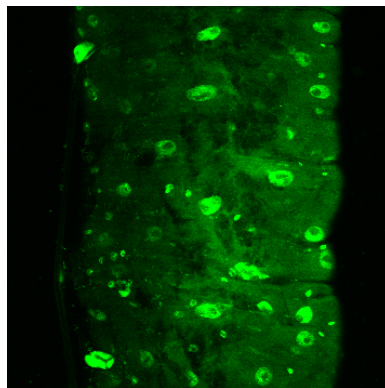
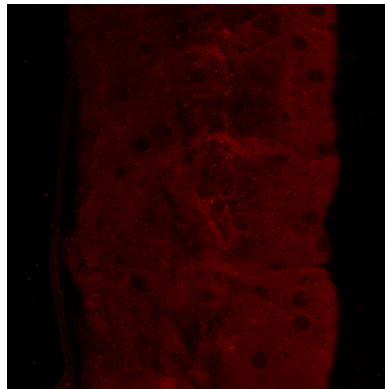


Fly dNox is required for ROS generation

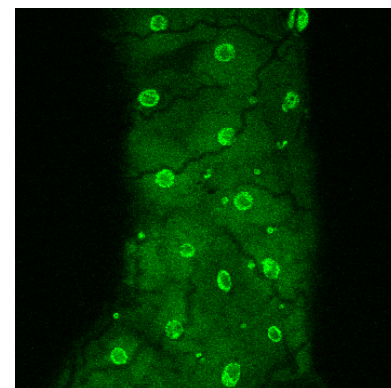
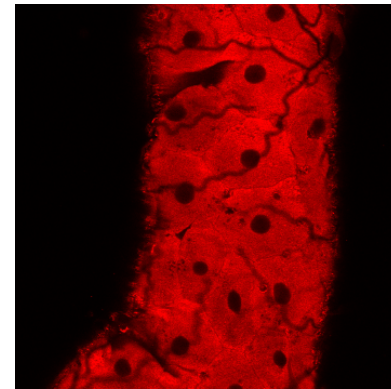
**Myo-GAL4
w1118**



**Myo-GAL4
UAS-Nox-RNAi**

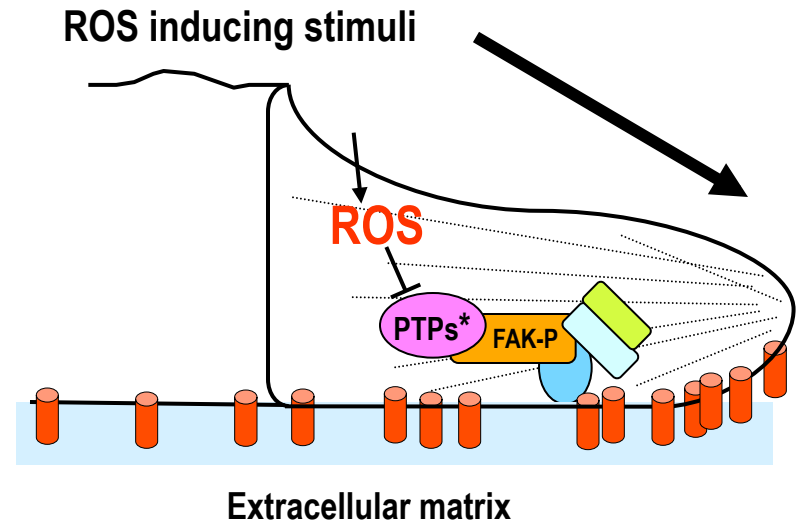
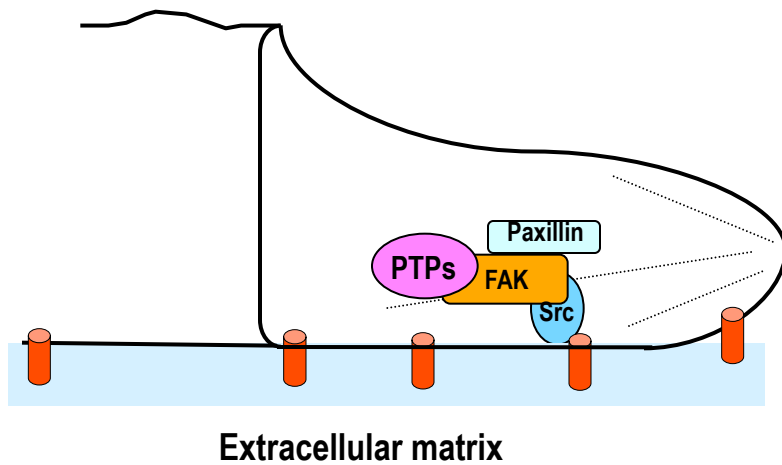


**Myo-GAL4
UAS-Duox-RNAi**

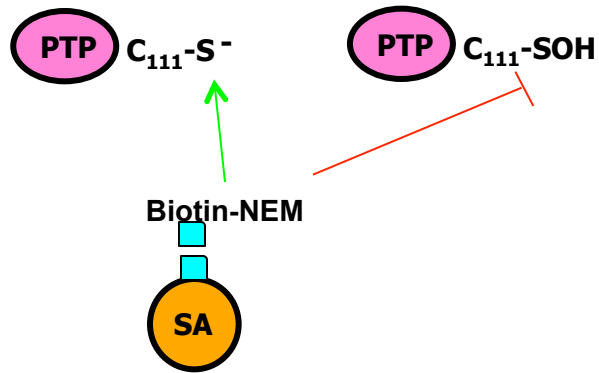


Larval guts fed LGG for 30min with hydro cy3

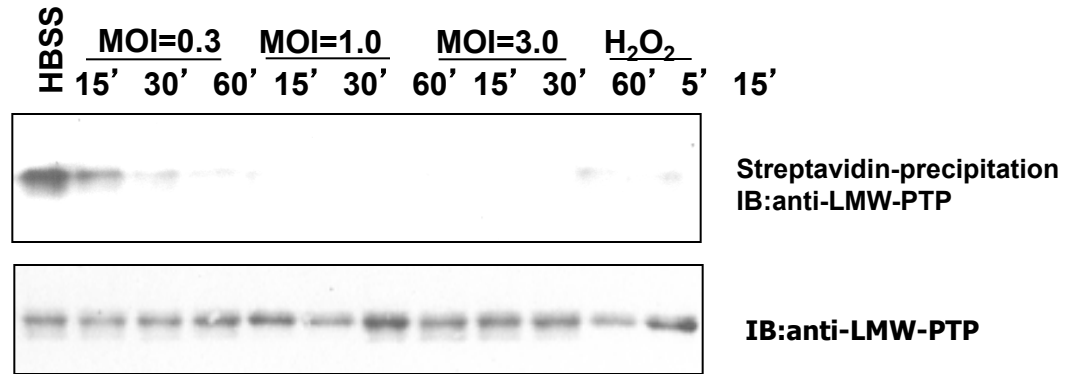
ROS signaling stimulates epithelial movements



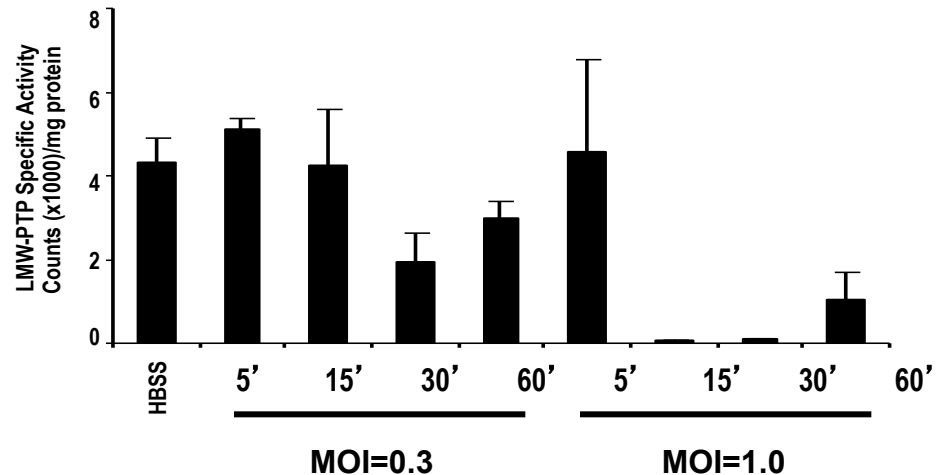
Commensal colonization elicits oxidation and inactivation of FAK phosphatase, LMW-PTPase



Precipitate, immunoblot/assay for PTPase

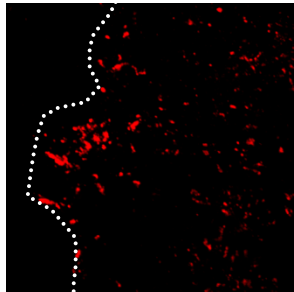


Biotin-n-ethyl-maleimide



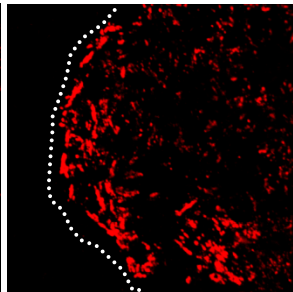
Commensal colonization elicits rapid phosphorylation of FAK

pFAK861

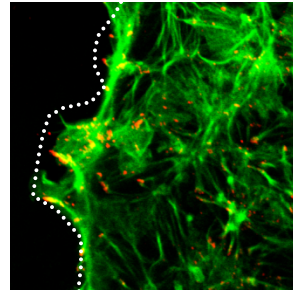


Control

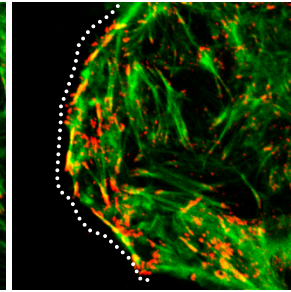
pFAK861, F-actin



Lacto



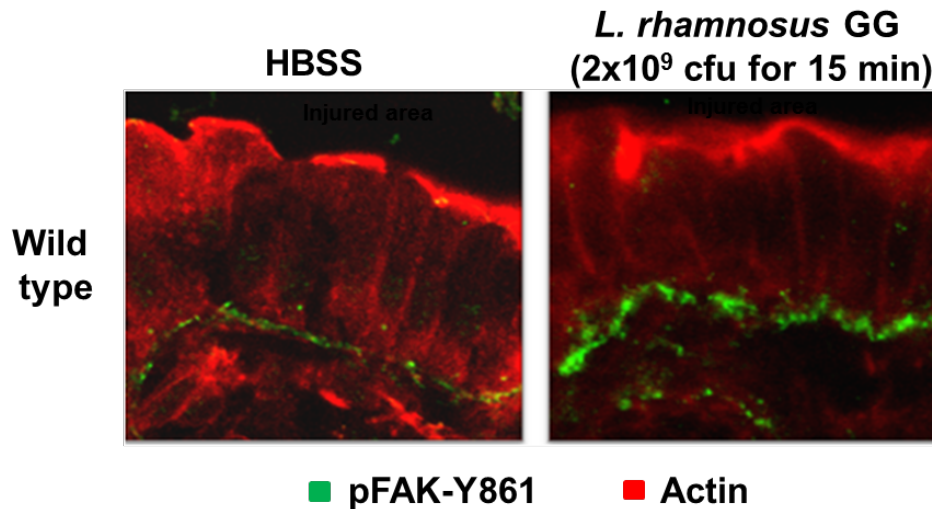
Control



Lacto

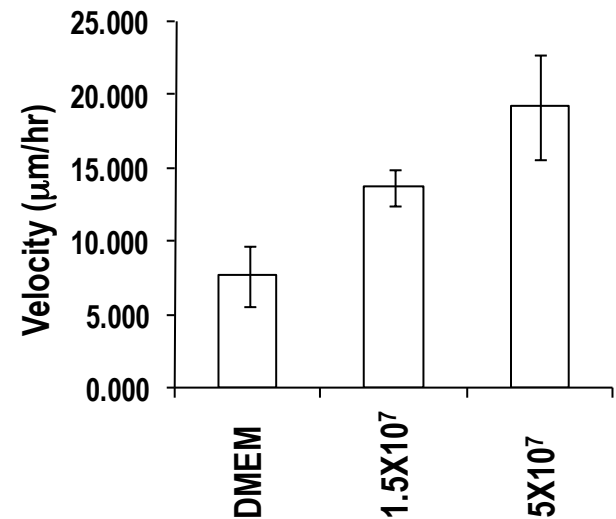
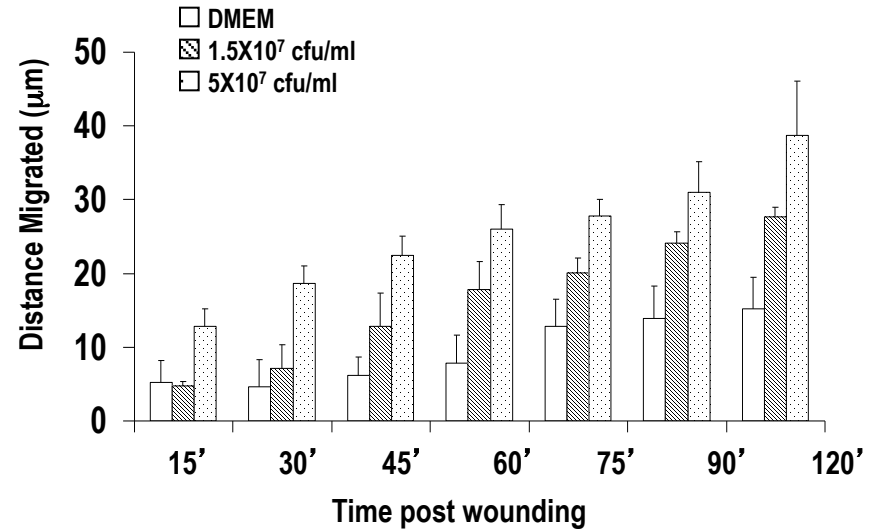
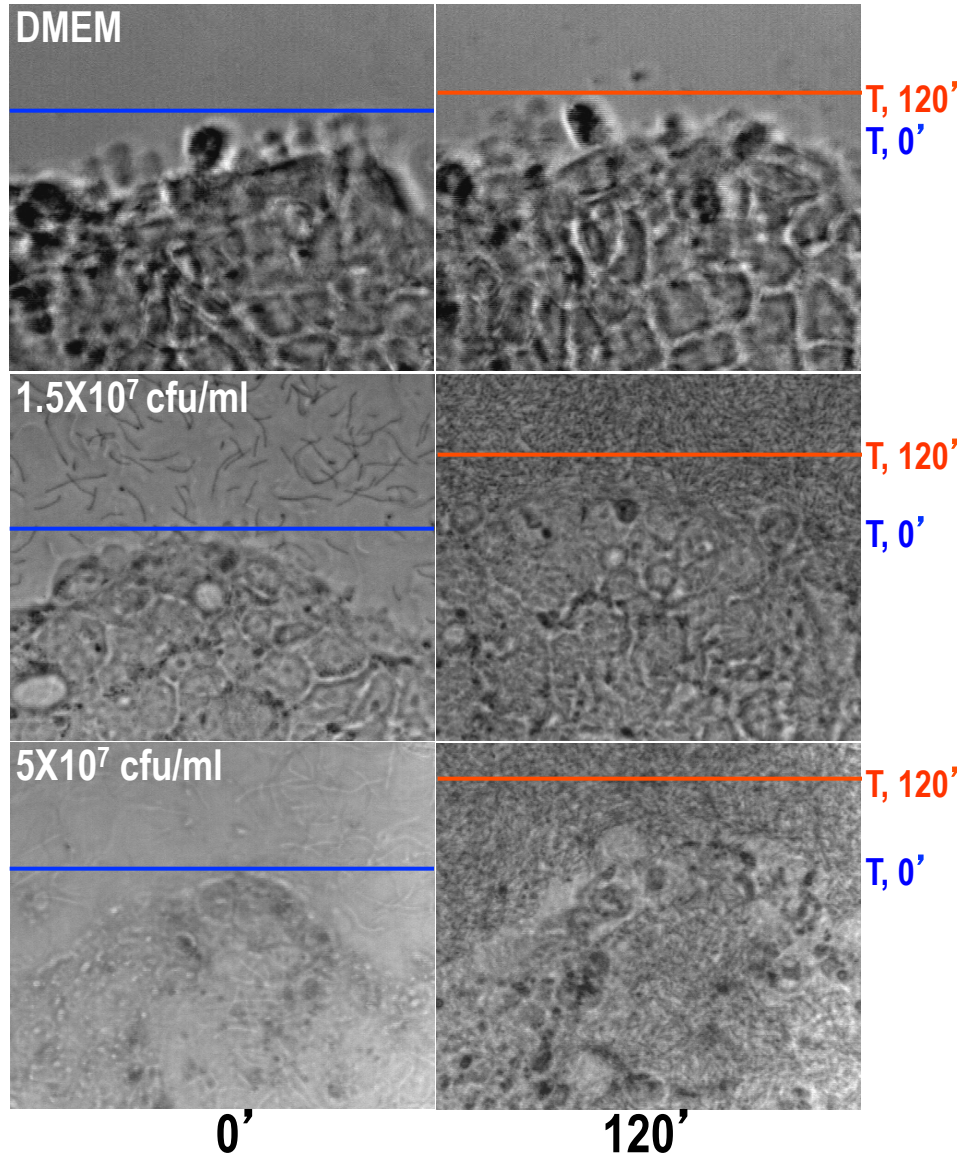
5'

Immunofluorescence of F-actin and phosphorylated FAK-Tyr-861 reorganization induced by LGG.

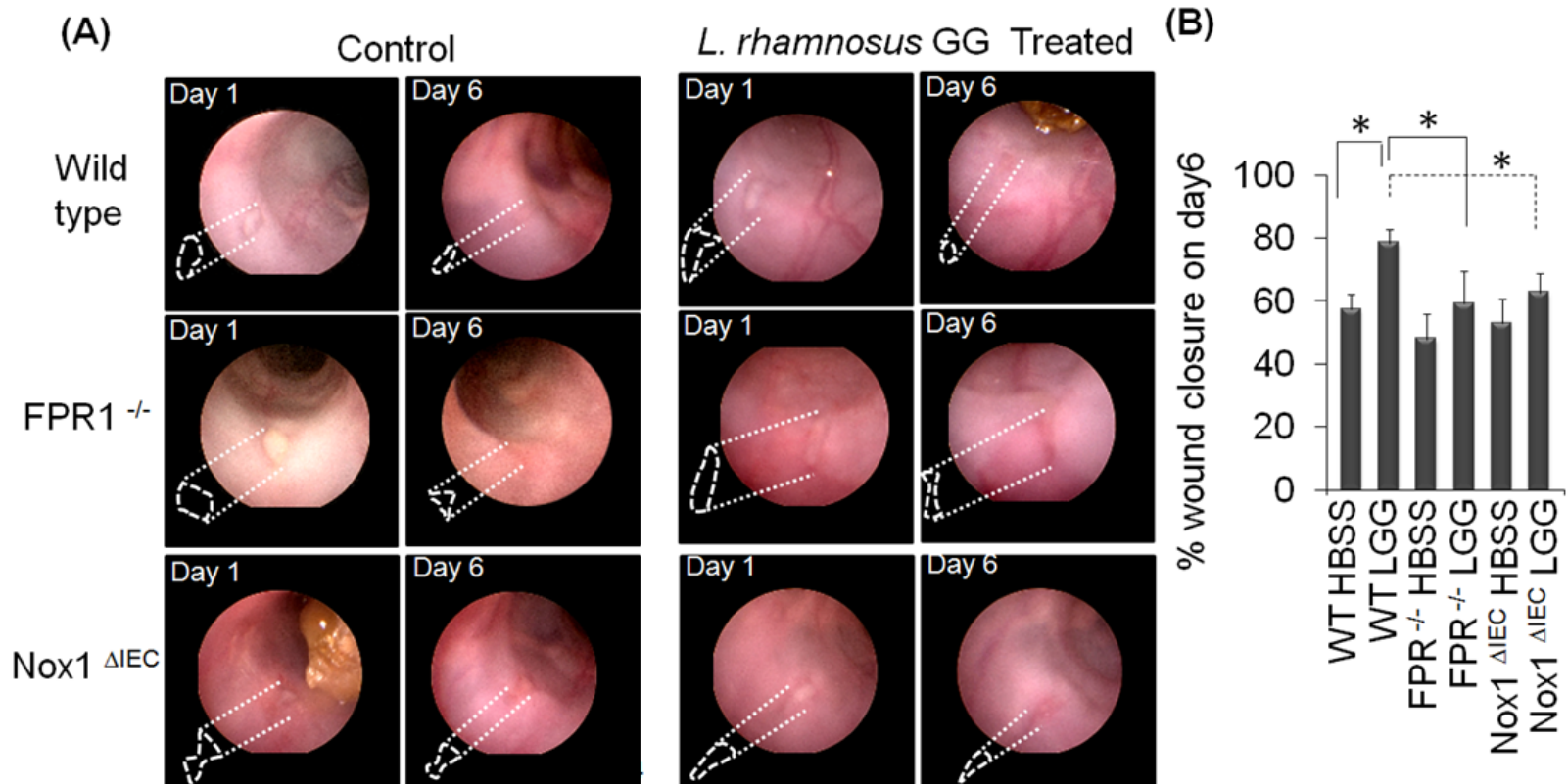


En face image of the wound bed

Commensal colonization elicits increased migration of model epithelia

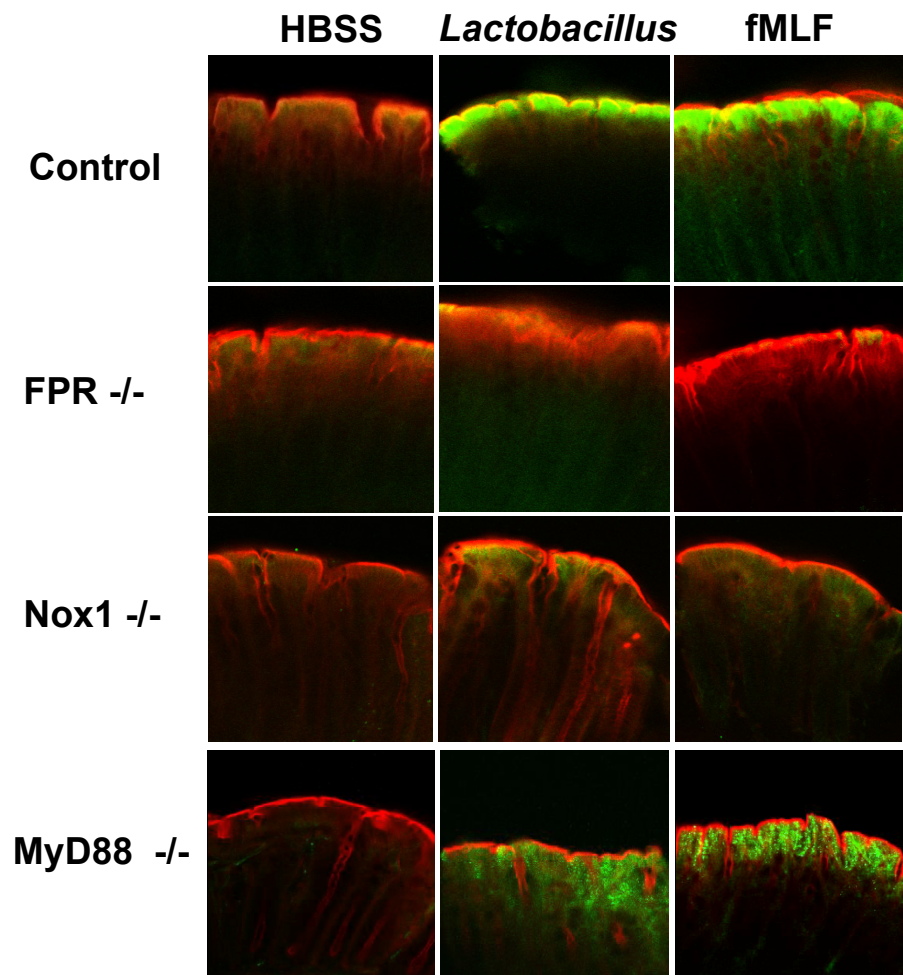


Lactobacilli enhances epithelial wound healing *in vivo*



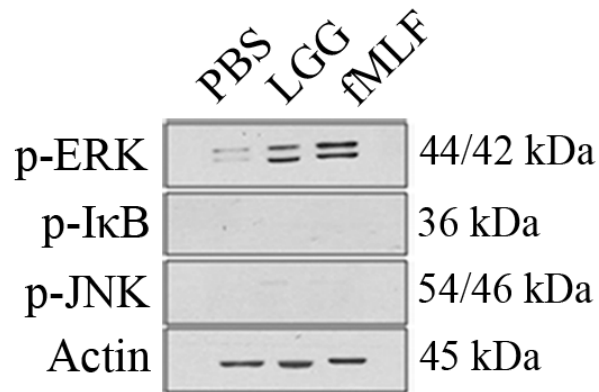
Swanson et al. **Enteric commensal bacteria potentiate epithelial restitution via ROS-mediated inactivation of focal adhesion kinase phosphatases**. PNAS. 2011 24;108(21):8803-8.

Microbiota stimulates phosphorylation of ERK in vivo in a FPR and Nox1 dependent manner



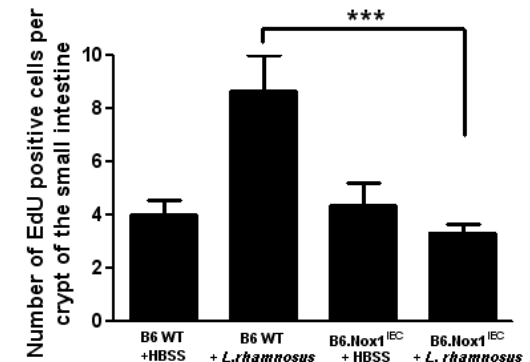
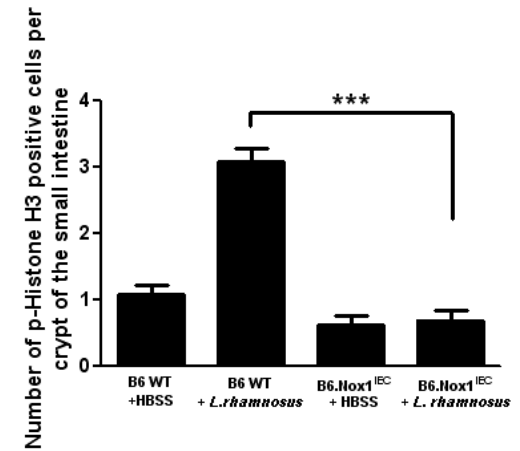
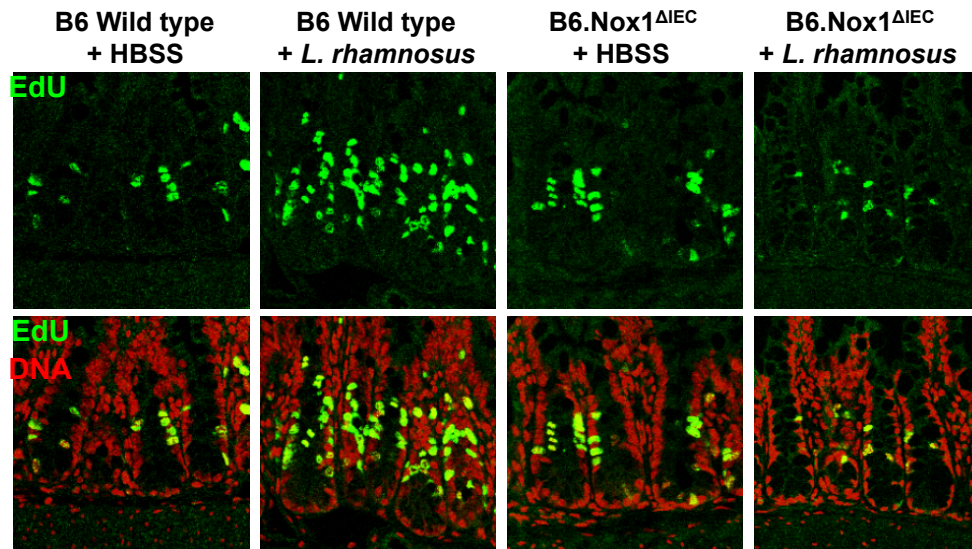
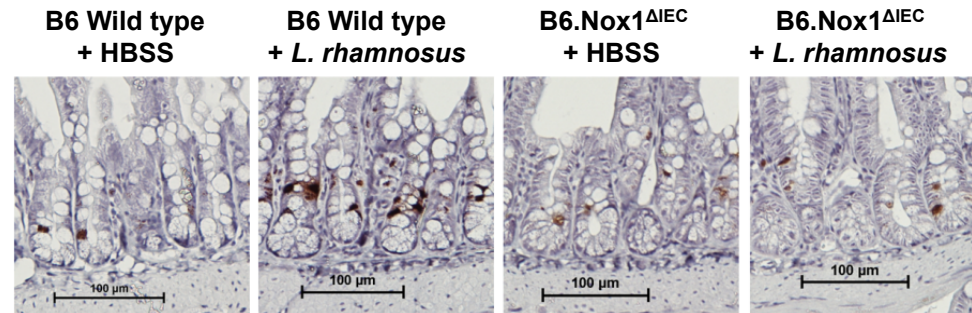
■ Actin
 ■ Phospho-ERK

Intraluminal *Lactobacillus* activates ERK in murine colon in vivo within 15 min

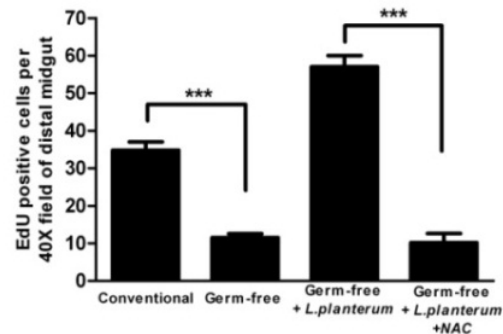
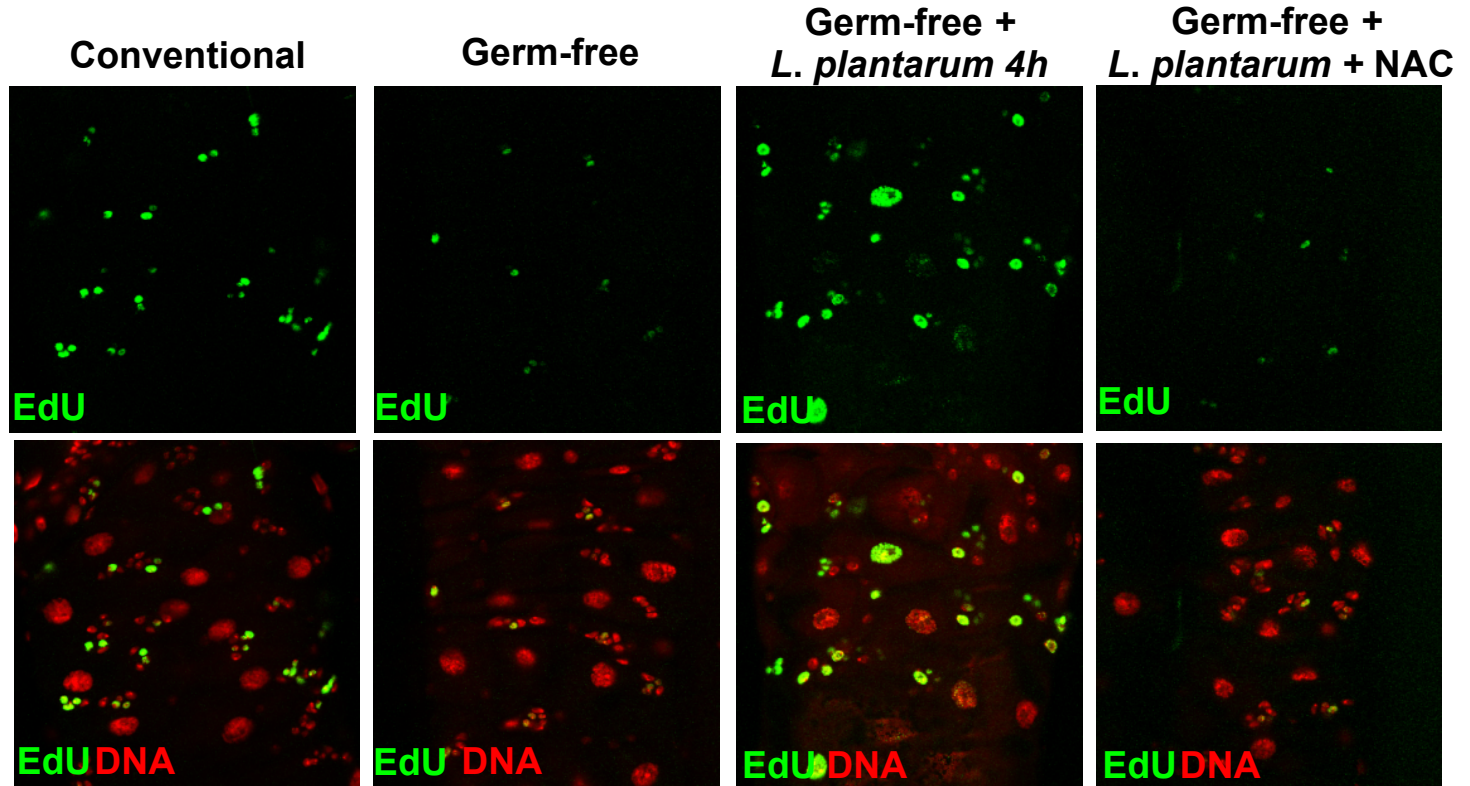
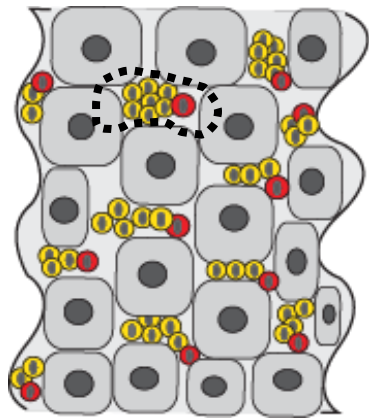


Lactobacilli stimulates crypt proliferation in mice in a Nox1 dependent manner

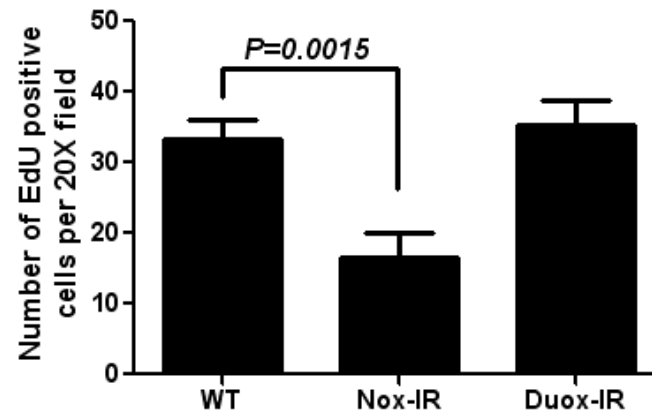
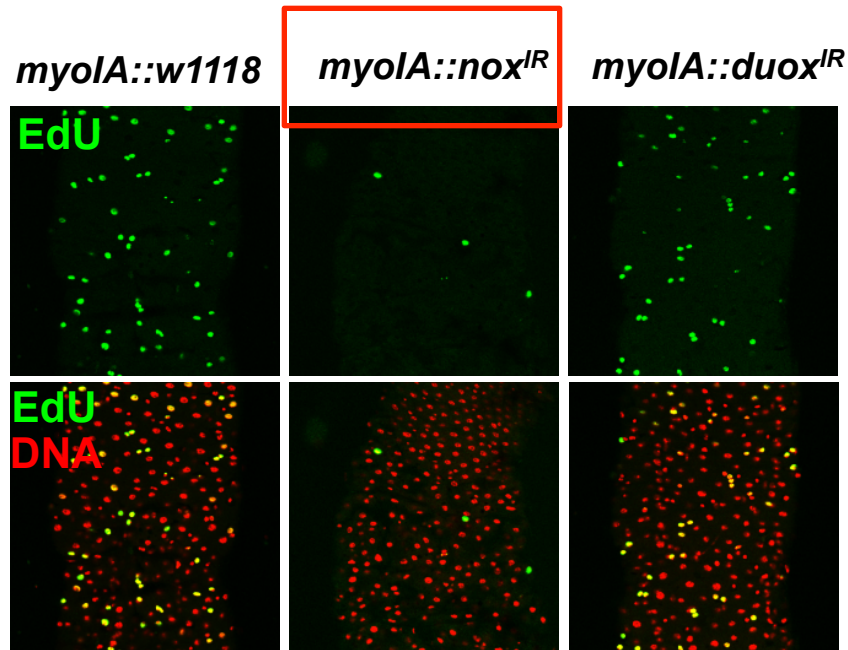
6-week-old small intestine



Lactobacilli induces ROS-dependent cellular proliferation in the *Drosophila* intestine

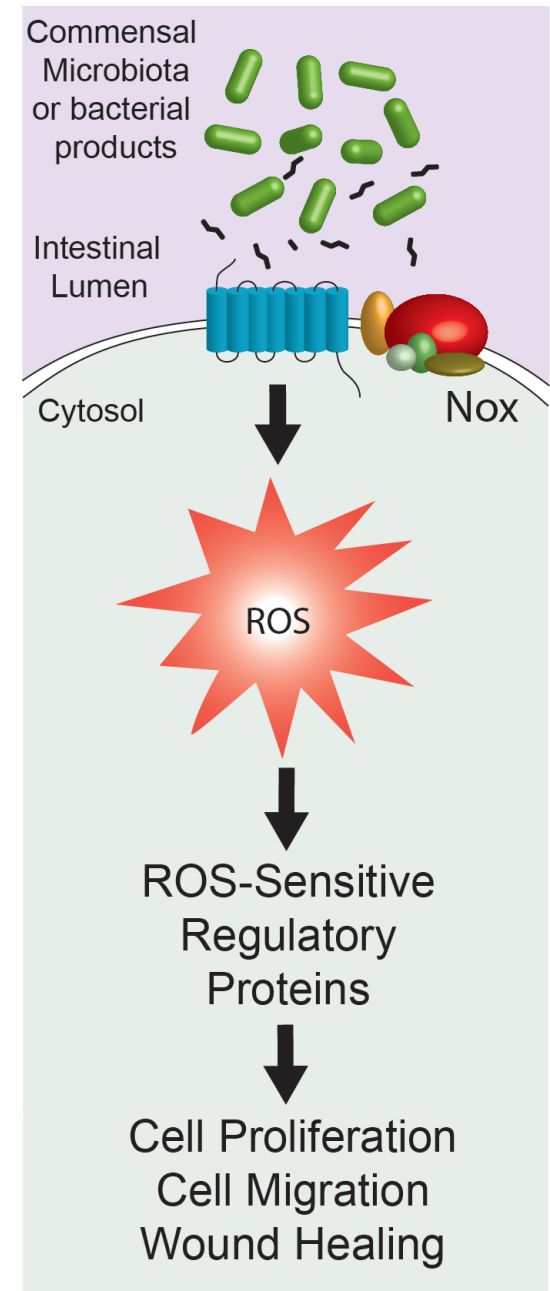


Proliferation in the *Drosophila* epithelia is Nox dependent



Conclusion :

Nox dependent generation of physiological levels of ROS by *Lactobacillus* (and likely other bacteria) is a novel signaling mechanism for transducing bacterial signals into host regulatory events that mediate intestinal homeostasis, proliferation and restitution



Future Challenges

- Novel ROS sensitive proteins/pathways
 - ROS dependent signaling eg Nrf2/ARE pathway
- Correlation with innate immunity and PRR signaling
- Correlation with microbial determinants and bacteria taxonomy
- Role in normal gut development
- Role in wound healing
- Role in oncogenesis

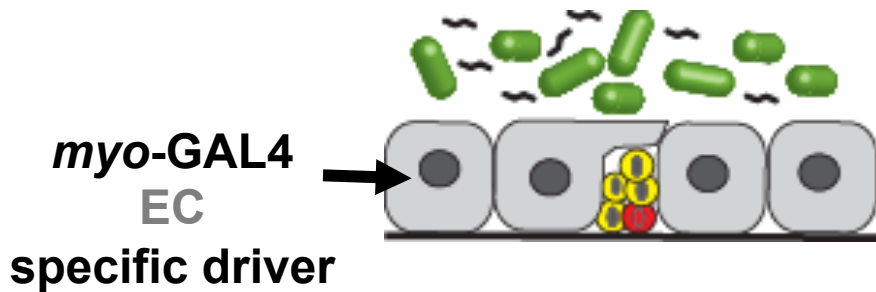
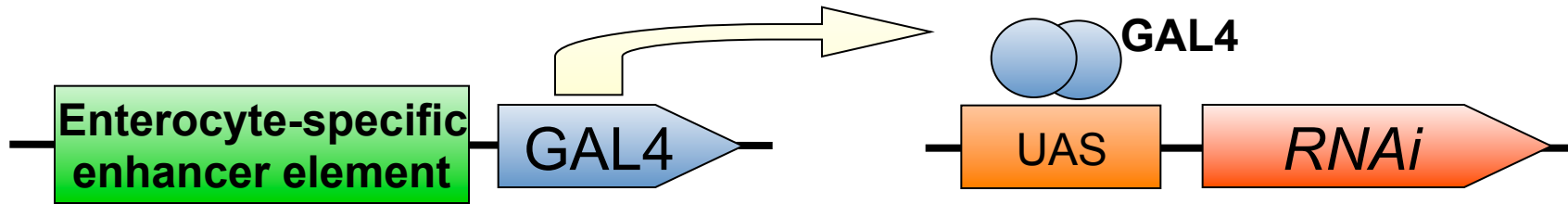
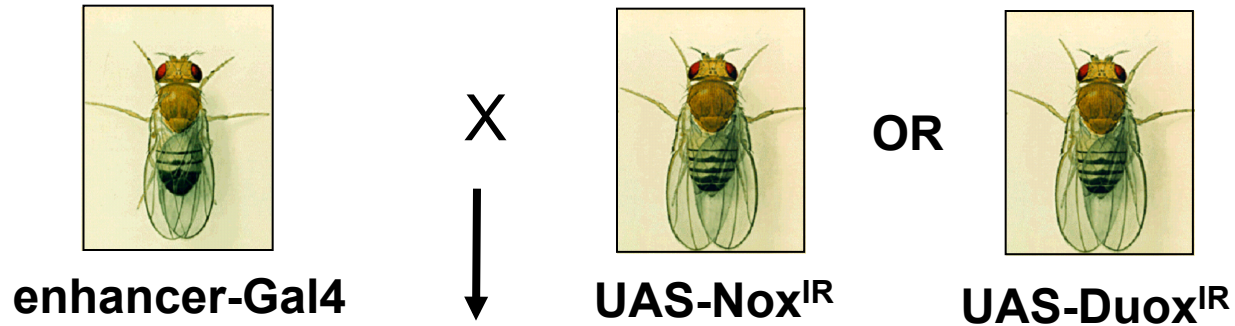
Gaps

- Epithelial cell biology as a host-microbial system
- Cooperation between model systems workers
- “Comparative metagenomics”

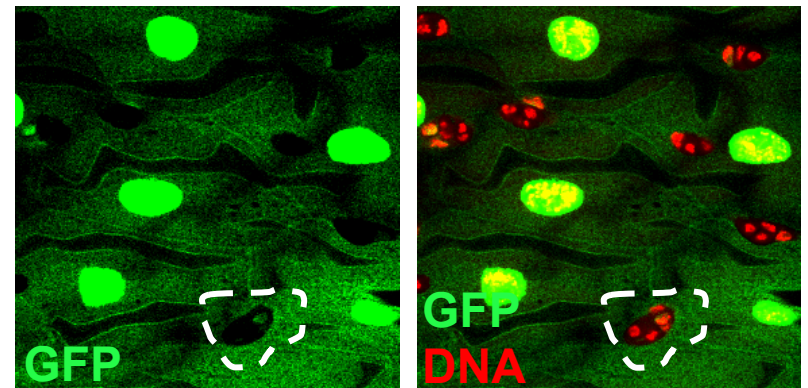
Acknowledgments

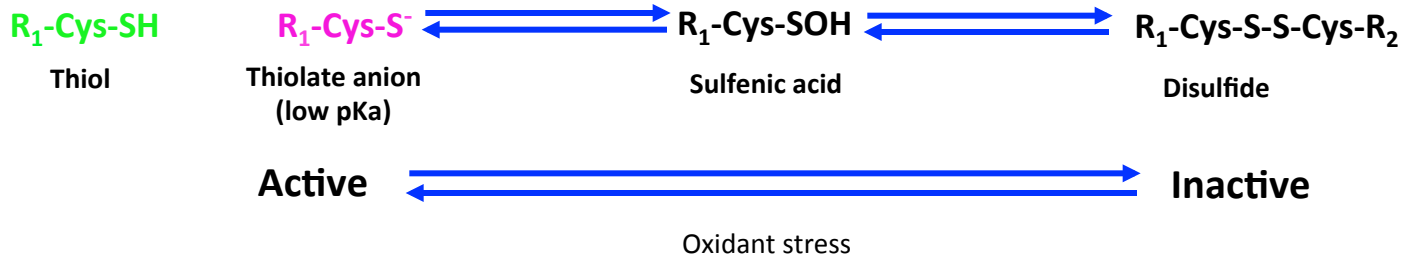
- Neish Lab
 - Ashfaq Alam
 - Huixia Wu
 - Chirayu Desai
 - Jeff Mercante
 - Phil Swanson
- Rheinallt Jones
- Liping Luo
- Asma Nusrat
- Giovanna Leoni
- Supported by NIAID RO1 AI 64462 and NIDDK R01 DK 89763

Expression of RNAi against Nox and Duox in fly enterocytes



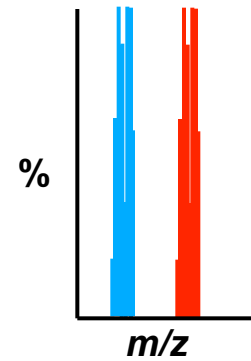
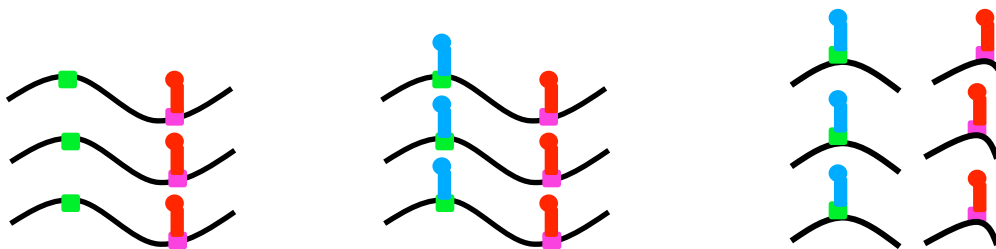
Myo-GAL4, UAS-nlsGFP





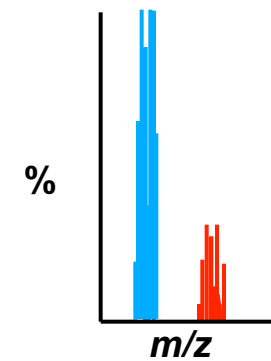
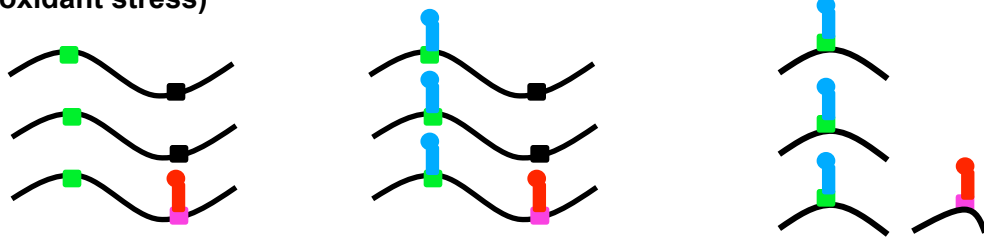
Non-reactive cysteine ■
 Reactive cysteine ■
 Oxidized cysteine ■

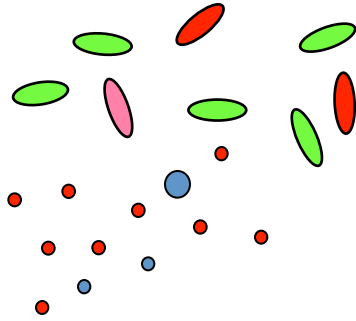
Normal



Heavy ^{14}C ICAT ■ → Reduce with TCEP → Light ^{12}C ICAT ■ → Trypsin digestion → LC/MS

Colonized (oxidant stress)





Extracellular

Identify, characterize
redox sensitive regulatory
targets

Characterize and validate
redox sensitive
physiologic pathways and
processes

1. Epithelial cell culture
2. Drosophila
3. Mice

Identify, characterize ROS
inducing commensals,
bacterial products, small
molecules

Cytoplasm

ROS production

(Hydro Cy3, etc)

Oxidation of regulatory proteins

BIAM/IAA (candidate) ICAT (high throughput)

Experimental validation

Enzymatic, signaling assays

Homeostatic and biochemical alterations

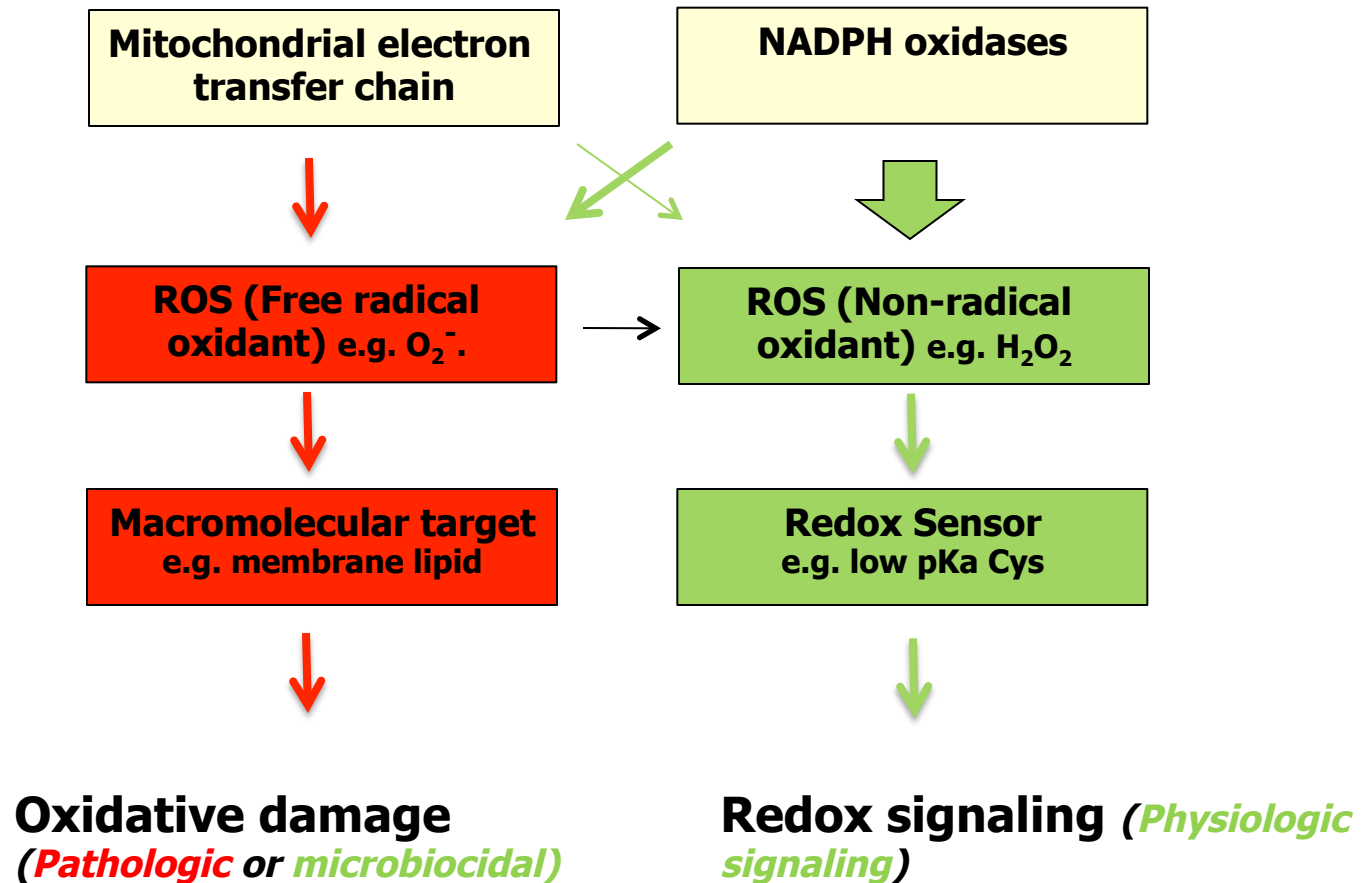
Metabolic and phenotypic analyses

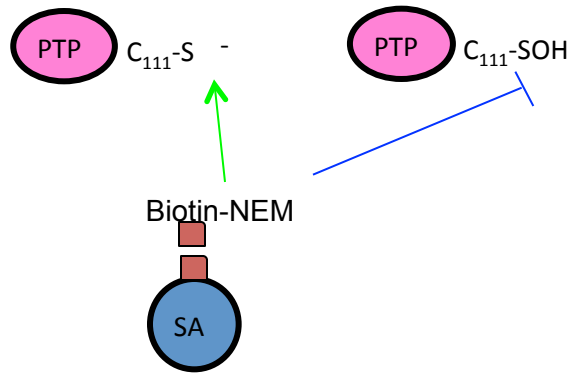
Systems analysis

Redox proteomic database, redox pathway maps

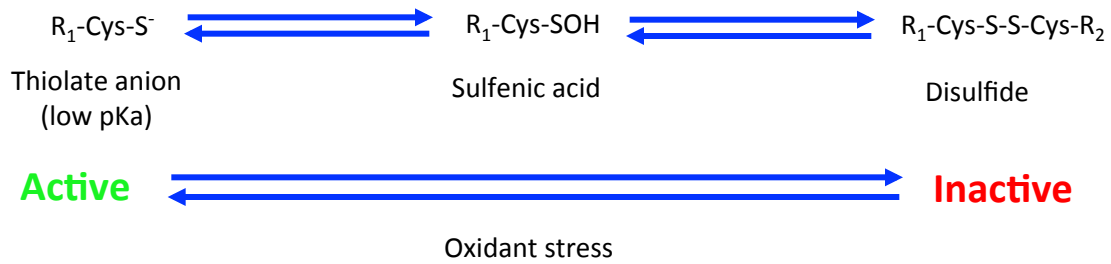
ROS effects on the cell

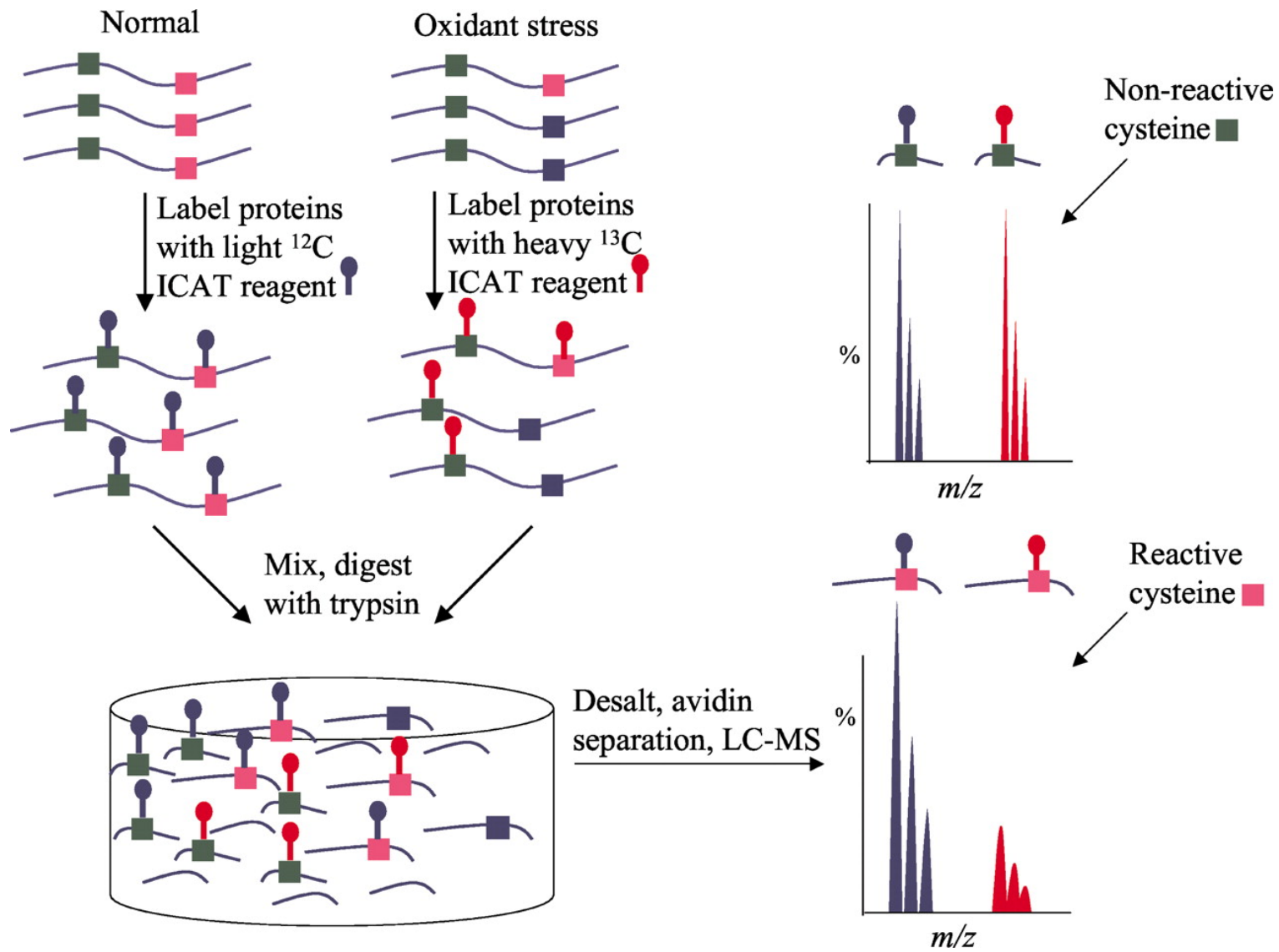
Upstream signals/stress



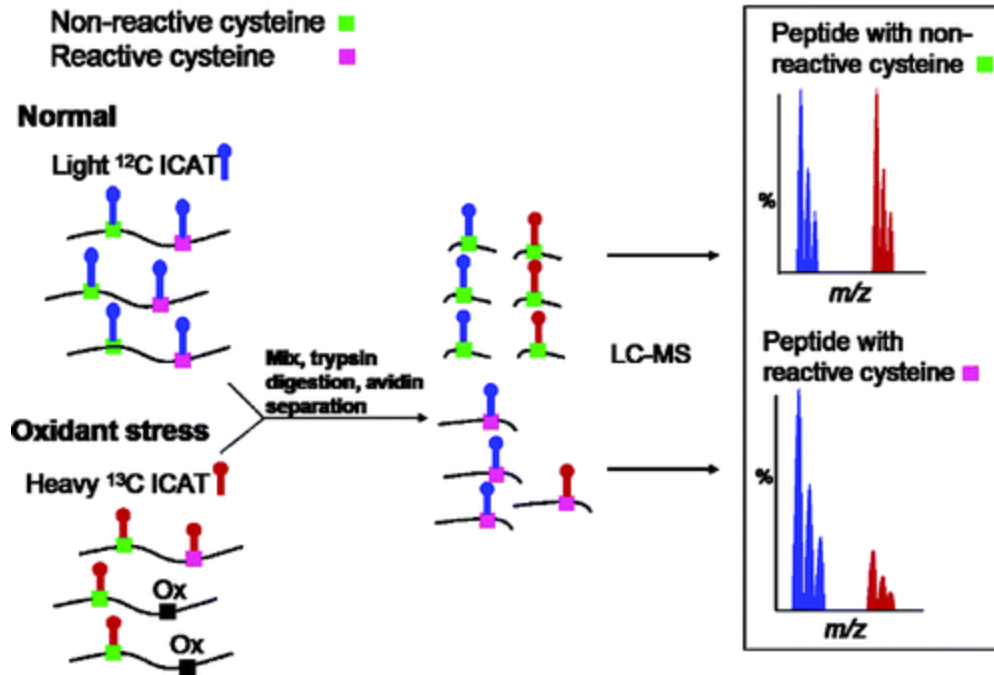


Precipitate, immunoblot/assay for PTPase





ICAT Approach to Redox Proteomics



(a)

Cell extract
 ↓
 Modify thiols with H-ICAT
 ↓
 Reduce with TCEP
 ↓
 Modify thiols with L-ICAT
 ↓
 Treat with trypsin
 ↓
 LC²-MS/MS to determine % reduced for Cys peptides

(b)

Protein Name
Protein metabolism
 Elongation factor1- α
 Elongation factor-2
 Glu-tRNA synthetase
 Ubiquitin-activating E
 Calreticulin
Signal transduction
 Protein phosphatase
 G-protein, β subunit
Detoxification
 Peroxiredoxin 6
 GSH-S-transferase
Stress Response
 Hsp 70
 Hsp 90
Cell Structure/Motility
 Cofilin
 β -Tubulin
 α -tubulin

(c)

Database
chemical

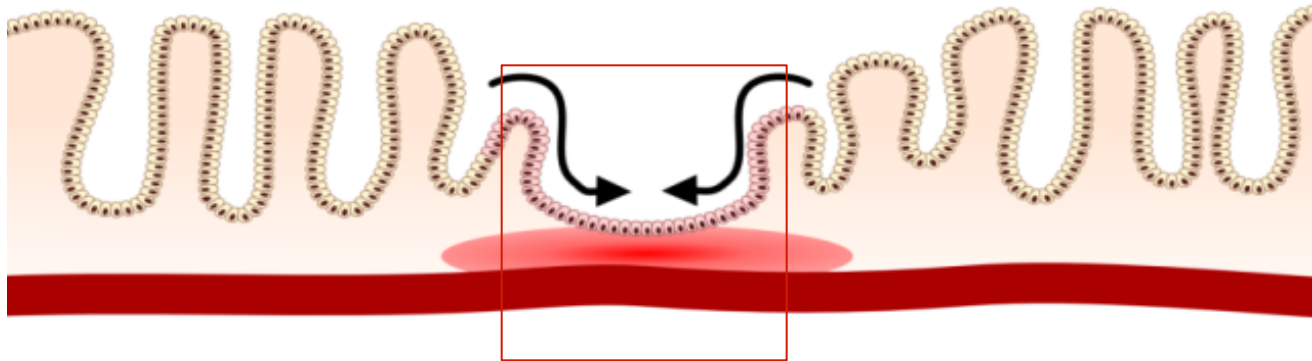
Redox prote

Thiol redox
circuitry maps

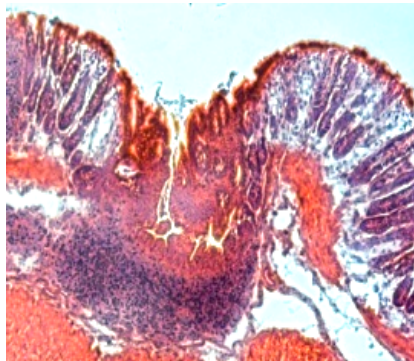
Conclusions

- These data suggest an intriguing role for intestinal bacteria in the modulation of the cellular homeostatic pathways via ROS generation
- Possible mechanism for commensal “crosstalk” with the host
- Possible mechanism for probiotic effects

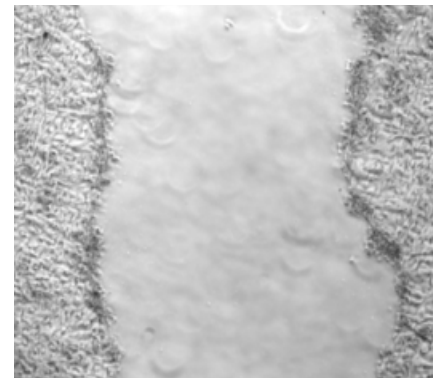
Restitution: Involves coordinated epithelial proliferation and migration



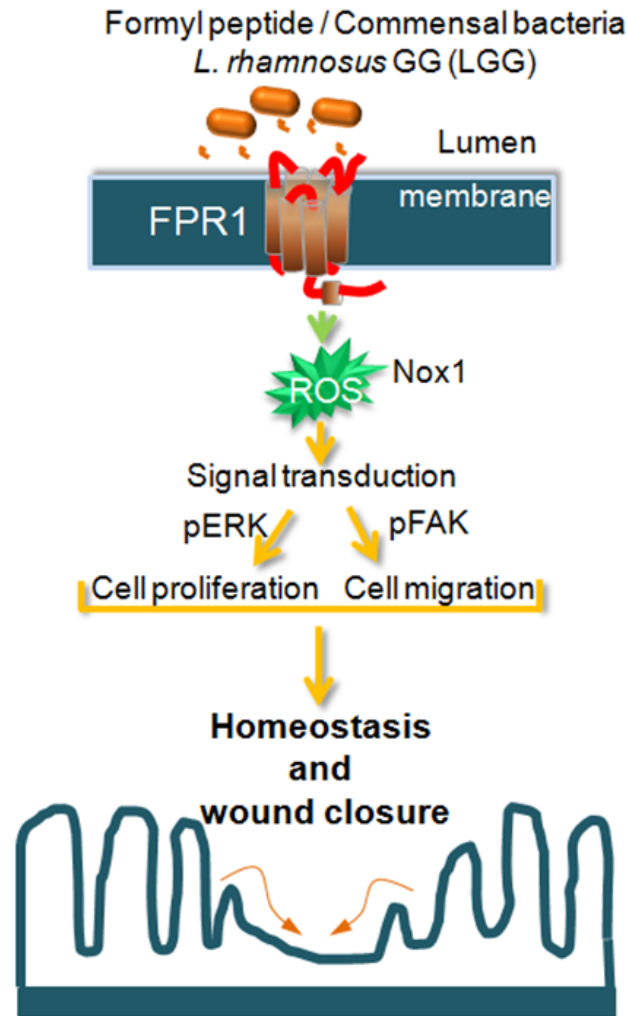
In vivo



In vitro



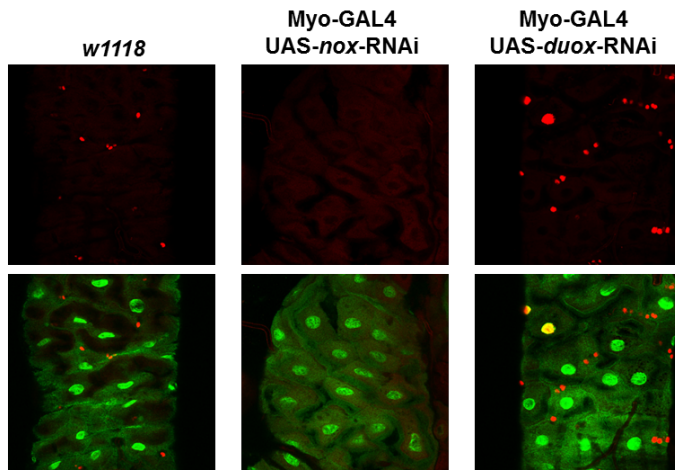
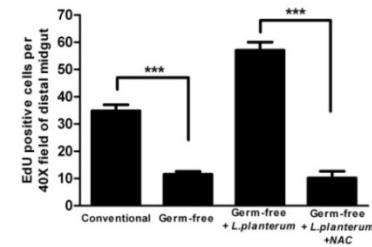
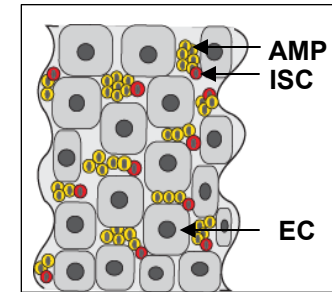
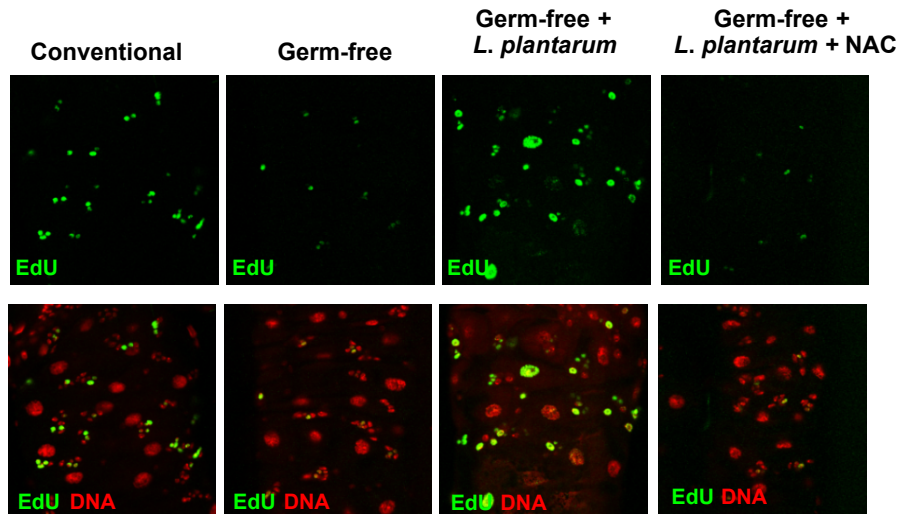
ROS mediated signaling regulates multiple homeostatic pathways



Summary

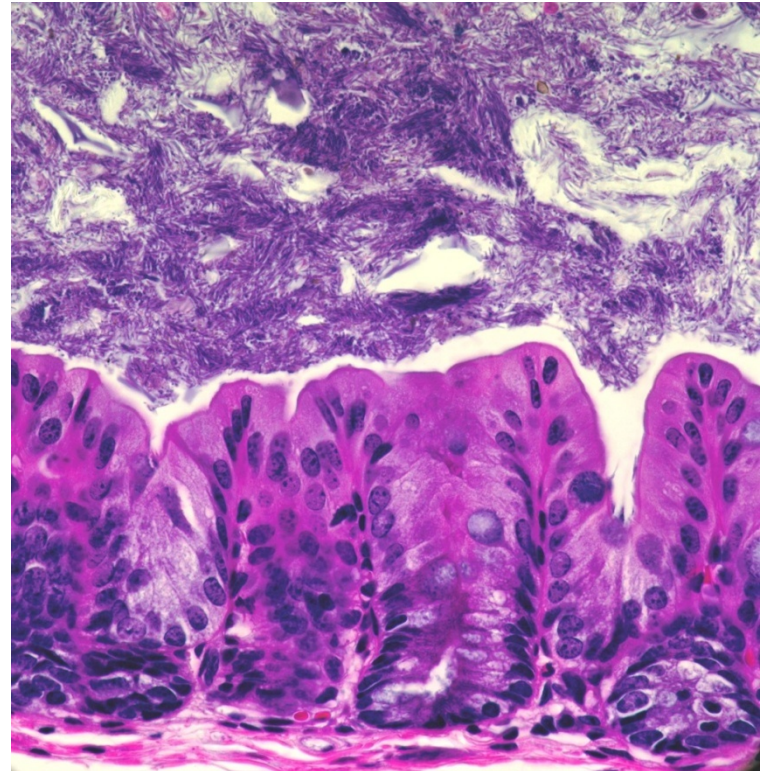
- Specific commensals (and commonly used probiotics) can stimulate non proinflammatory signaling in the gut
- Signaling results in increased epithelial proliferation and migration
- Physiological “deliberate” generation of ROS is involved in these processes
- Possible mechanism for beneficial homeostatic effects of probiotics

Microbiota stimulates crypt proliferation in flies in a Nox1 dependent manner



The Microbiota

- 10^{14} organisms, mostly in colon
 - 10x more than humans cells!
- Acquired at birth
- Approximately 500 genera represented
 - Most in two Divisions, *Bacteroidetes* and *Firmicutes*
- Diverse metabolic abilities
 - Alternate energy sources and fermentative products
- **Symbiotic with host**



ROS generation as a response to bacteria

- Oxidant or “respiratory burst” in phagocytes in response to FPR signaling
- High degree of evolutionary conservation in wide variety of animals and plants
- Strong conservation of NADPH generating enzymes (Nox’ s)

