



A Clinical-Stage Oral β -Lactamase Therapy Prevents
Antibiotic-Mediated Damage of the Gut Microbiome

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Importance of Intestinal Health Has Long Been Recognized



"ALL DISEASE
BEGINS IN
THE GUT!"

-HIPPOCRATES
400 B.C.

Gut Microbiome Involved in:

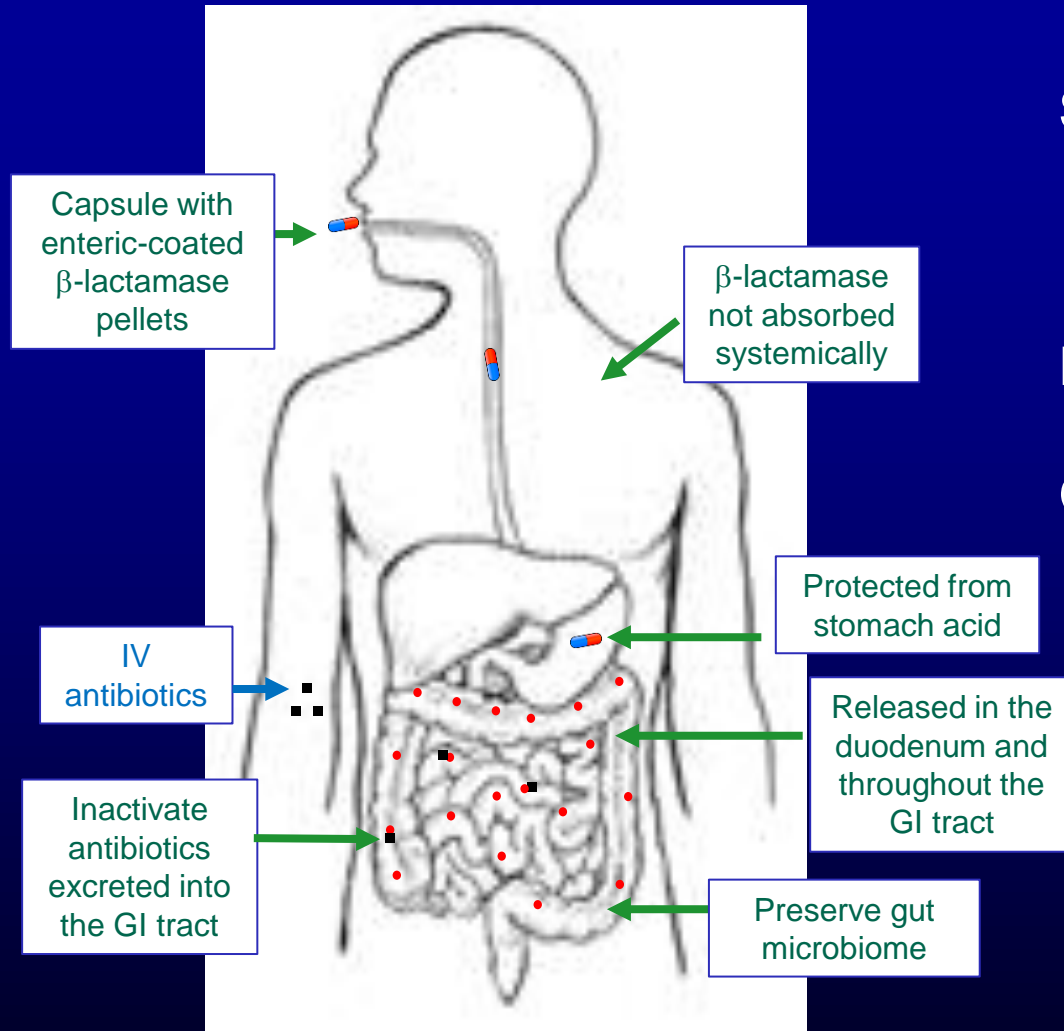
- Digestion
- Nutrient absorption
- Vitamin synthesis
- Bile salt metabolism
- Stimulation of immune system

Disrupted by:

- Disease
- Antibiotics

Synthetic Biologics is developing therapies to protect the gut microbiome from the damage caused by antibiotics

β -Lactamases: From Enemies to Therapies



Strategy: β -lactamase enzyme is intended to degrade residual antibiotics in the GI tract to protect the microbiome

Product: Capsule with enteric-coated enzyme

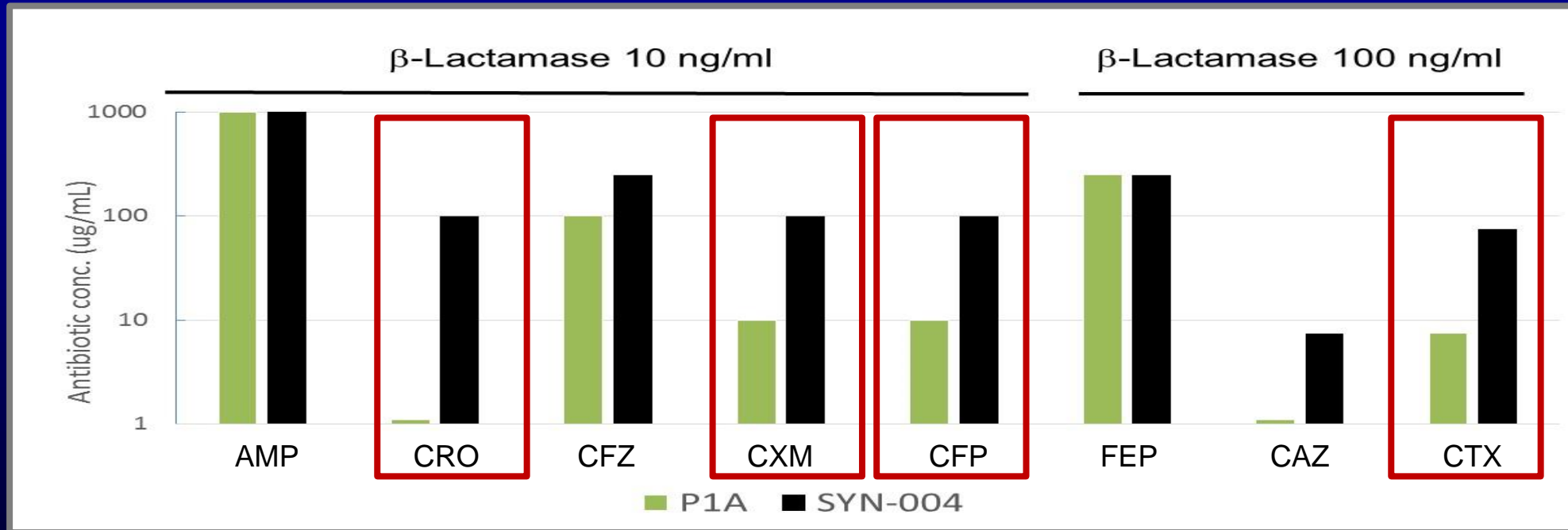
Outcome: Prevention of *Clostridium difficile* infection and antibiotic-associated diarrhea

Orally-delivered β -lactamases intended to degrade residual antibiotics in the GI tract to protect the gut microbiome without affecting antibiotic efficacy

SYN-004 Degrades Cephalosporins

- SYN-004 was engineered from P1A
- P1A is a clinical isolate from *Bacillus licheniformis*

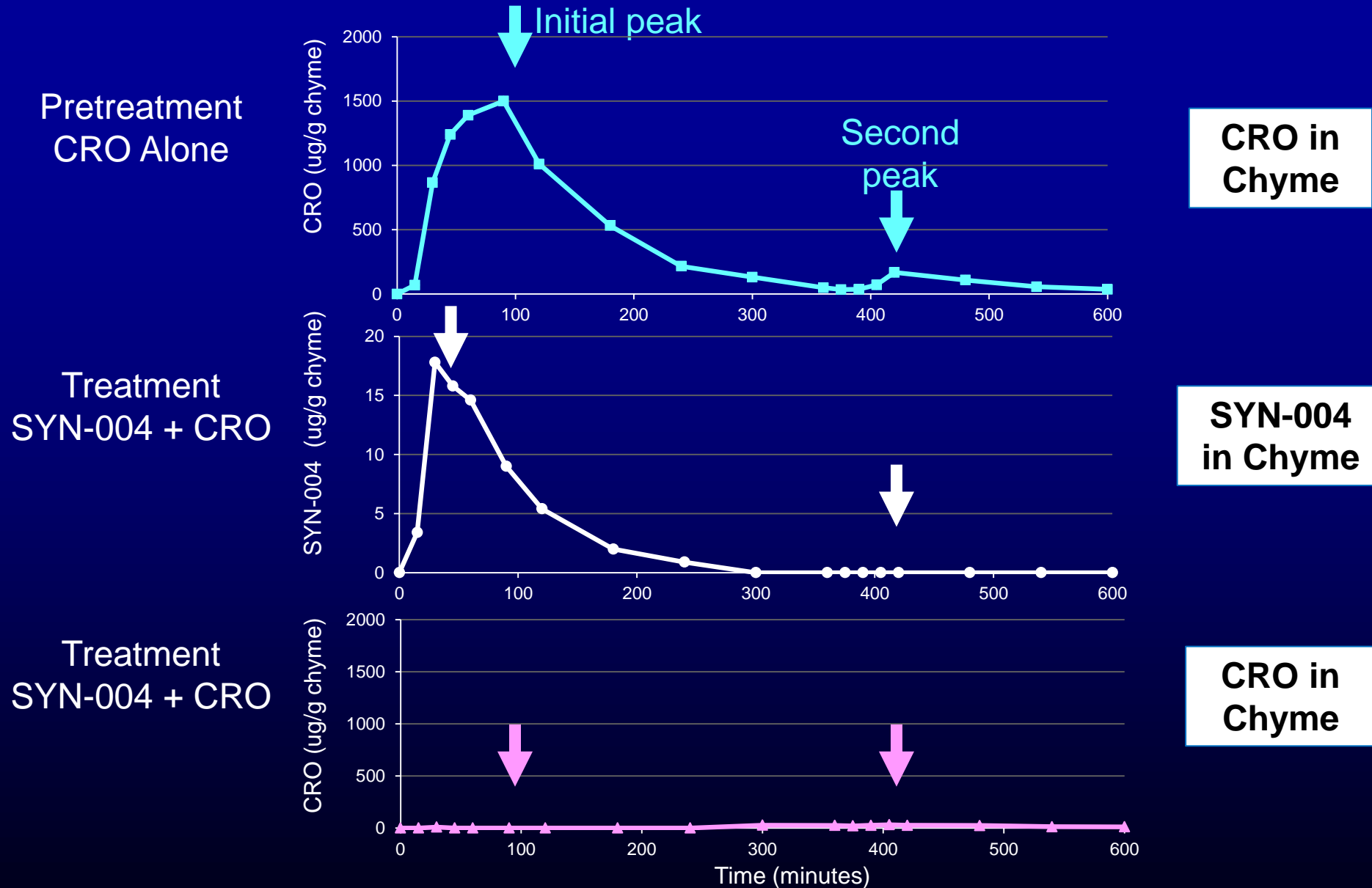
E. coli growth microtiter plate assay



Amp: ampicillin
CRO: ceftriaxone
CFZ: ceftazidime
CXM: cefuroxime
CFP: cefoperazone
FEP: cefepime
CAZ: ceftazidime
CTX: cefotaxime

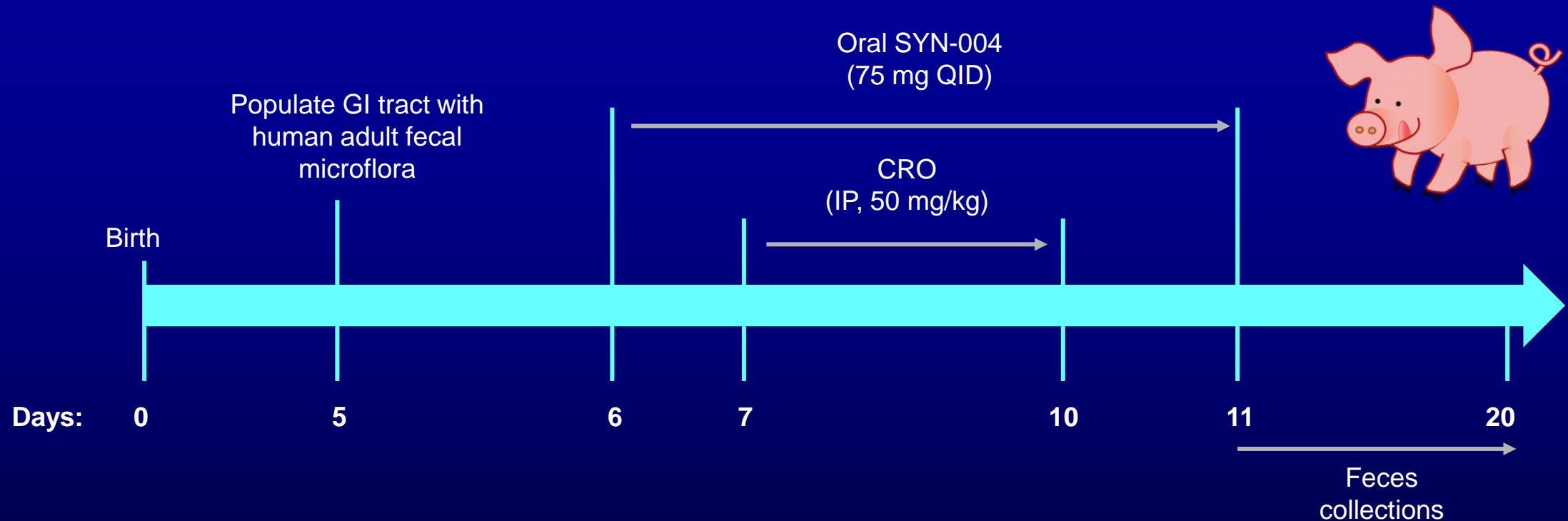
SYN-004 efficiently degrades cephalosporins, including ceftriaxone, cefuroxime, cefoperazone, ceftazidime, and cefotaxime

SYN-004 Degraded Ceftriaxone (CRO) in Dog GI Tract



In the presence of SYN-004 no CRO was detected in chyme

Neonatal Pigs with Human Gut Microflora



Readouts:

- Direct measure of CRO-sensitive bacterial population
- Fecal DNA 16S rRNA V6 region sequence analyses
- Shotgun deep sequencing analyses

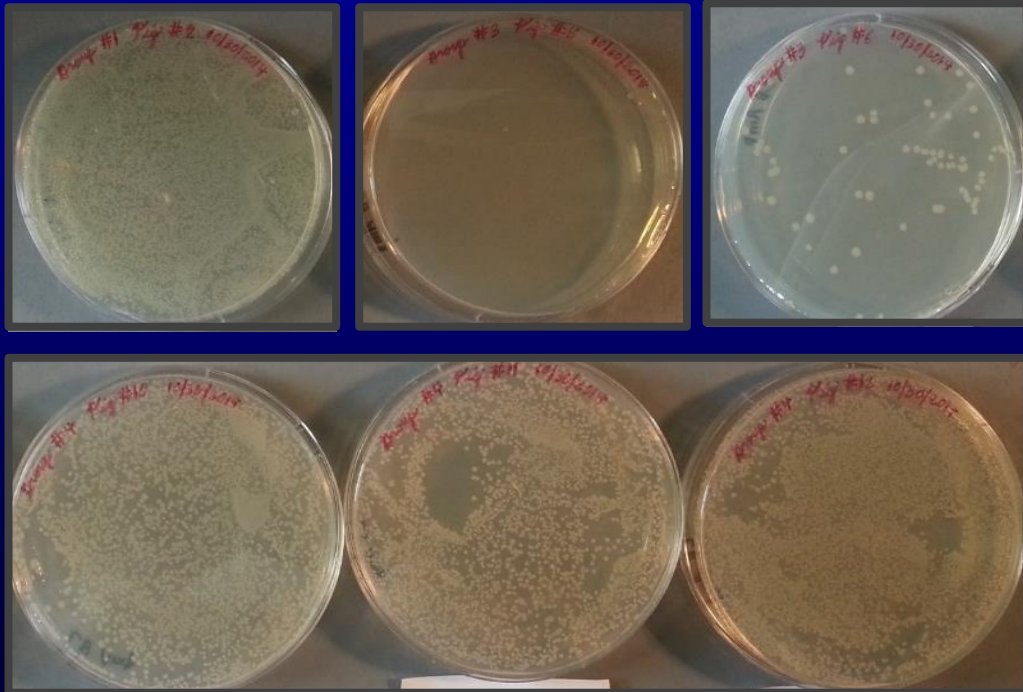
SYN-004 Protected the Gut Microflora in Pigs

- Monitored CRO-sensitive bacterial population, Amp^R aerobes

Bacterial Growth on LB+Amp Plates

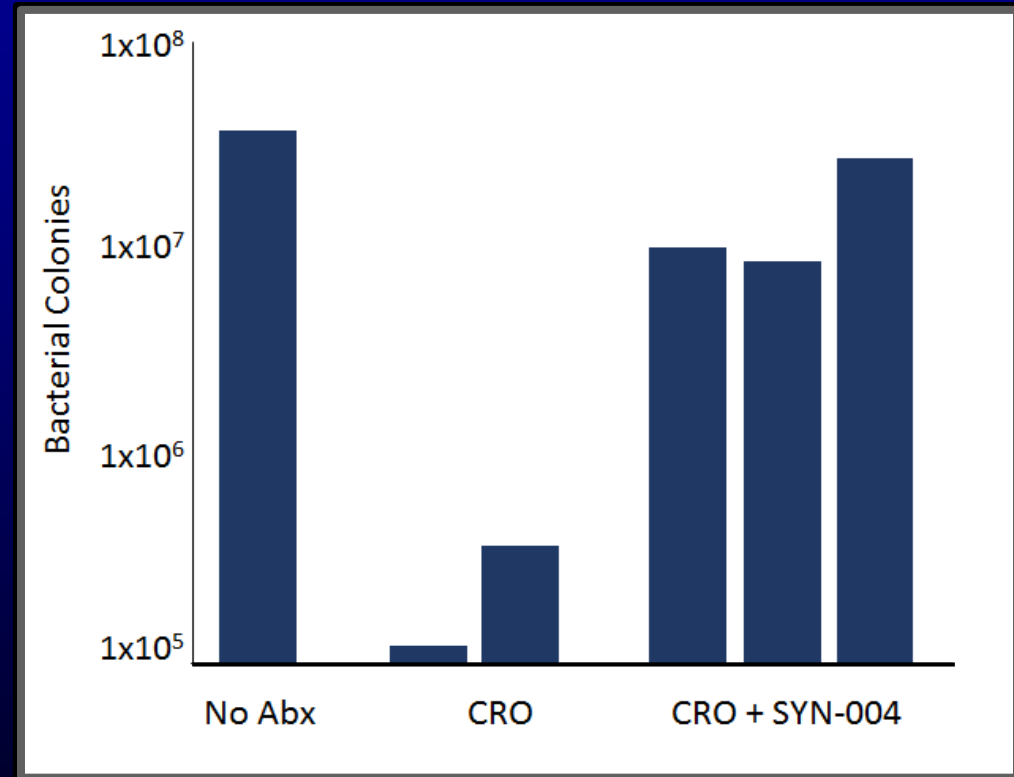
No Abx

CRO



CRO+SYN-004

Bacterial Colonies

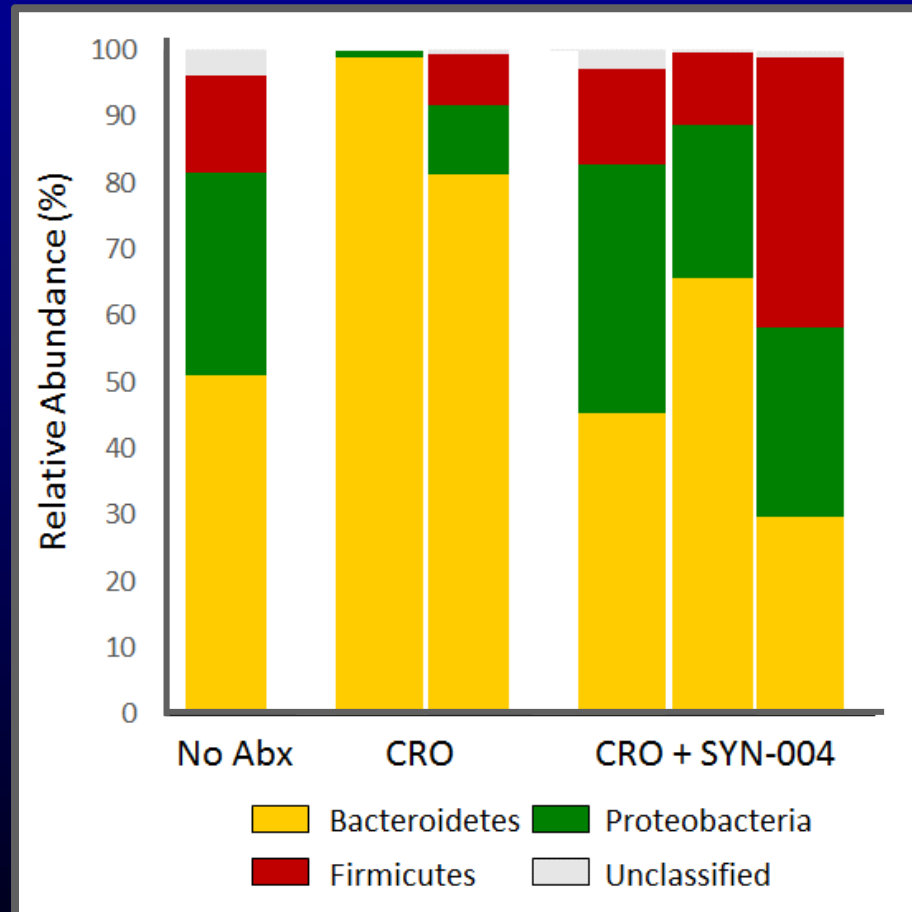


SYN-004 protected the CRO-sensitive fecal bacterial population

SYN-004 Prevented CRO-Induced Dysbiosis

- 16S rRNA V6 region sequence analyses

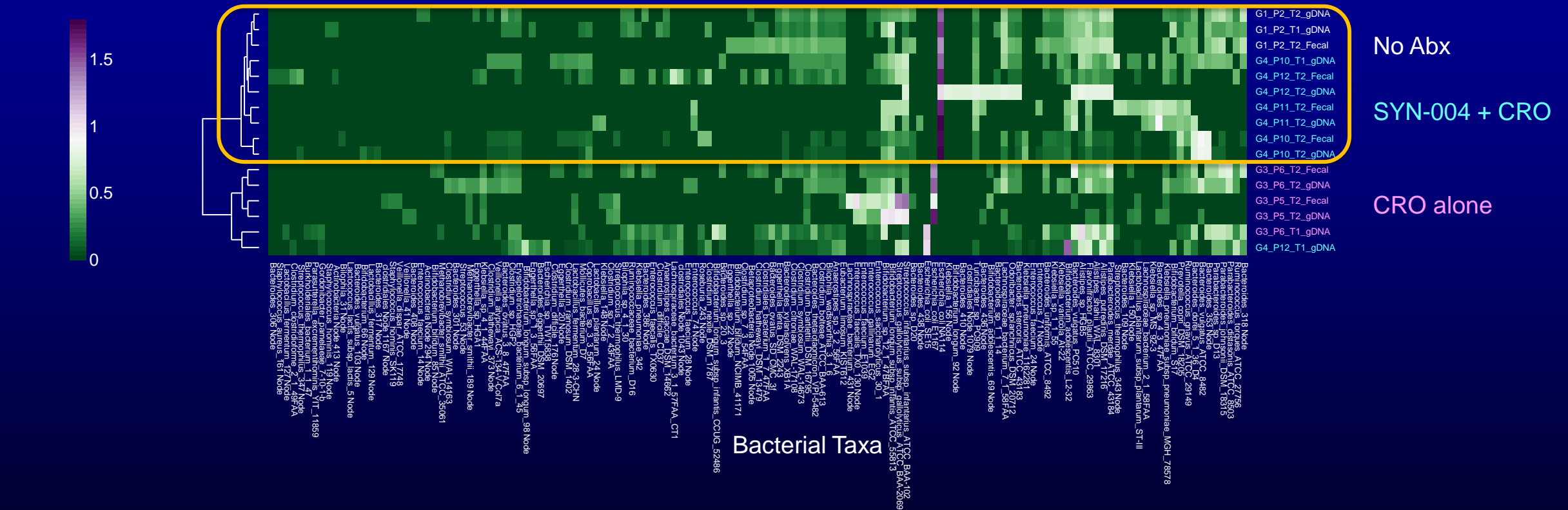
Phylum-Level Taxonomic Classification



SYN-004 protected the
Proteobacteria and Firmicutes
and
prevented overgrowth of
Bacteroidetes in the presence of CRO

SYN-004 Protected the Pig Microbiome

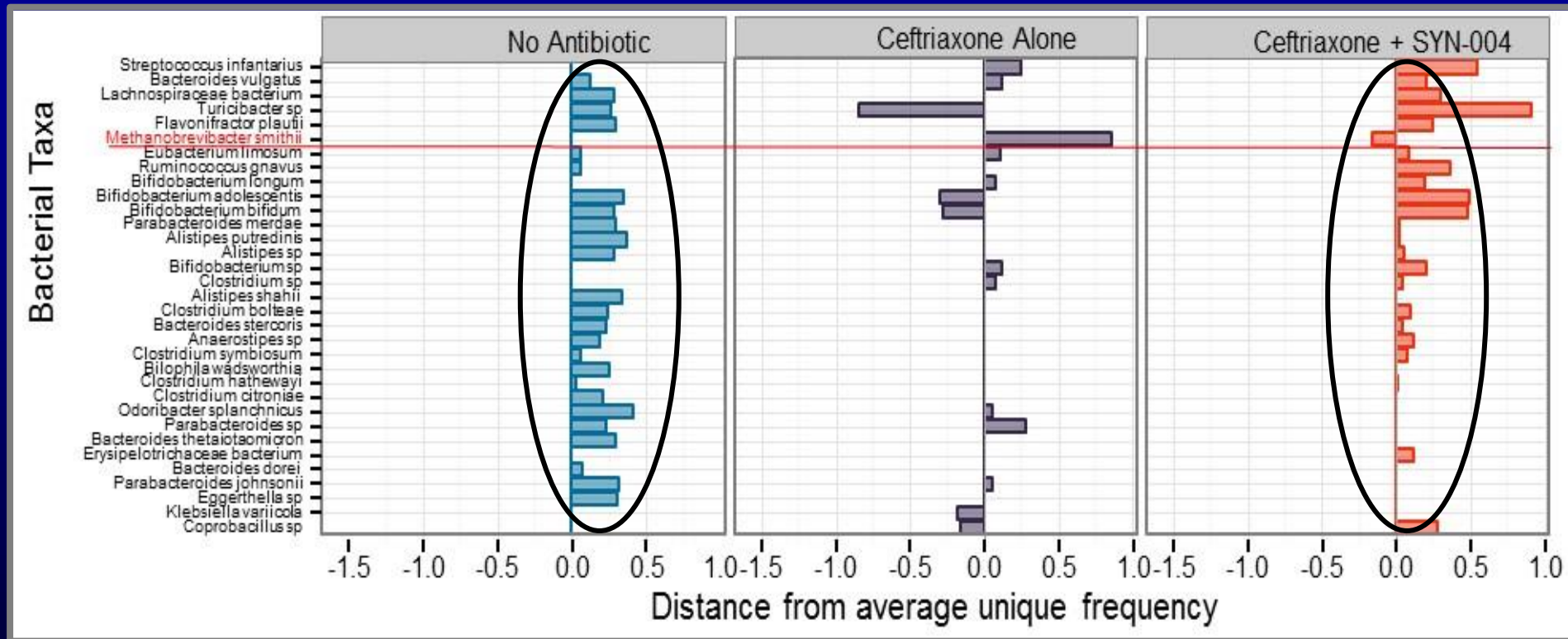
- Whole genome shotgun sequencing and taxonomic profiling
- Heatmap of bacterial strains based on compositional similarity and relative abundance
- Dendrogram clustering samples with similar compositions



The SYN-004-treated animals clustered with the controls demonstrating that SYN-004 protected the GI microflora from the effects of CRO

SYN-004 Prevented Overgrowth of *M. smithii*

- Whole genome shotgun sequencing and analyses
- Nearest shrunken centroid classification of fecal bacterial species



The SYN-004-treated animals were more similar to the no antibiotic controls demonstrating that SYN-004 protected the GI microflora from the effects of CRO

SYN-004 is in Phase 2 Clinical Trials

Preclinical Results

- Safe in two GLP toxicity studies in dogs
- Well tolerated with a NOAEL of 57 mg/kg/day, highest dose tested
- Not detected systemically
- Did not affect ceftriaxone blood levels

Clinical Results

- Phase 1 clinical studies demonstrated SYN-004 safety and tolerability with a single dose of up to 750 mg and multiple doses of 300 mg 4X a day for 7 days
- SYN-004 was not systemically bioavailable
- SYN-004 was not immunogenic
- Phase 2a studies are in progress
- Phase 2b study to initiate 3Q 2015

Conclusions

- SYN-004 degraded ceftriaxone in the dog GI tract
- SYN-004 protected the intestinal microbiome from ceftriaxone in pigs
- SYN-004 prevented the overgrowth of *M. smithii* in antibiotic-treated pigs
- *M. smithii* was reported to be associated with constipation, IBS, and obesity
- SYN-004 is progressing through Phase 2 clinical trials

SYN-004 has the potential to become the first prophylactic therapy designed to prevent antibiotic-mediated microbiome damage including *C. difficile* infection and antibiotic-induced diarrhea

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