

# THE EFFECT OF INTRAPARTUM ANTIBIOTIC PROPHYLAXIS ON NEONATAL GUT AND ORAL BACTERIOME

Eliska Pivrcova<sup>1</sup>, Iva Kotaskova<sup>1,2,3</sup>, Lucie Buresova<sup>1</sup>, Lenka Andryskova<sup>1</sup>, Pavel Piler<sup>1</sup>, Petr Janku<sup>4</sup>, Ivo Borek<sup>5</sup>, Jana Klanova<sup>1</sup>, Petra Videnska<sup>1</sup>, Eva Budinska<sup>1</sup>, Petra Borilova Linhartova<sup>1,6</sup>

<sup>1</sup> RECETOX, Faculty of Science, Masaryk University, Kotlarska 2, Brno, Czech Republic; <sup>2</sup> Centre for Cardiovascular Surgery and Transplantation, Brno, Czech Republic; <sup>3</sup> BioVendor MDx, Karasek 1, 621 00 Brno, Czech Republic; <sup>4</sup> Department of Gynecology and Obstetrics, Institution shared with University Hospital Brno and Faculty of Medicine, Masaryk University, Kamenice 5, 625 00 Brno, Czech Republic; <sup>5</sup> Department of Neonatology, Institution shared with University Hospital Brno and Faculty of Medicine, Masaryk University, Kamenice 5, 625 00 Brno, Czech Republic; <sup>6</sup> Clinic of Maxillofacial Surgery, University Hospital Brno, Jihlavská 20, Brno, Czech Republic

## BACKGROUND

Intrapartum antibiotic prophylaxis (IAP) is commonly used in C-section delivery (CS) and in Group B *Streptococcus*-positive (GBS) women before vaginal delivery (VD). IAP may affect neonatal microbial colonization:

- directly by passing into the fetal/neonatal bloodstream
- by passing into breast milk
- by reduced transmission of susceptible bacterial groups from mother to infant

24 %

C-SECTION  
WORLDWIDE  
PREVALENCE



18 %

GBS  
WORLDWIDE  
ESTIMATED  
PREVALENCE

OUR STUDY AIMED TO INVESTIGATE THE EFFECT OF EXPOSURE TO ANTIBIOTICS DURING DELIVERY ON THE NEONATAL MECONIUM AND TRANSITIONAL STOOL AND ORAL MUCOSA BACTERIOME

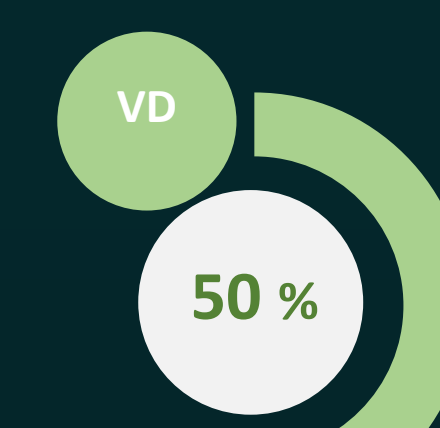
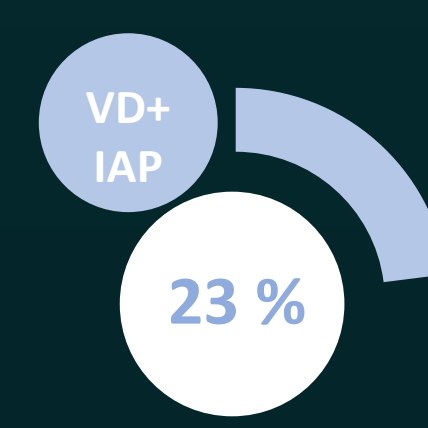
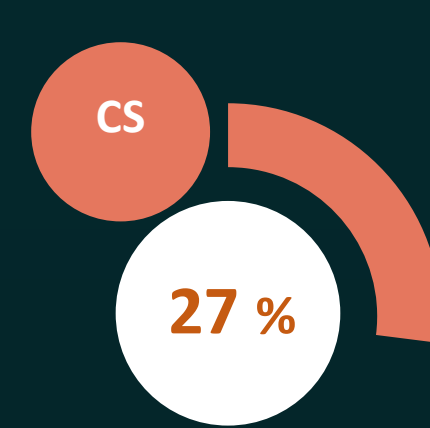
## METHODS

Newborns (n = 66) from the CELSPAC: TNG cohort divided in 3 groups:

CS with IAP

VD with IAP

VD without IAP



Stool swabs from the single-use diaper were collected and evaluated as **meconium** (1-3 days after birth) or **transitional stool** (3-5 days after birth)

Buccal swabs were collected within 48 hours after birth or within 3-5 days

16S rDNA gene sequencing was performed with the MiSeq reagent kit V3 using a MiSeq 2000 instrument.

## RESULTS

### MECONIUM

- No significant differences in bacterial abundance or diversity.

### TRANSITIONAL STOOL

- Significant differences in neonatal transitional stool bacteriome among three studied groups (PERMANOVA;  $q=0.001$ ) (Figure 1B).
- Groups of CS and VD with IAP were significantly associated with ↓ decreased relative abundances of genus *Bifidobacterium* ( $q=0.007$ ) and *Bacteroides* ( $q=0.007$ ).
- Abundance of *Enterococcus* ( $q=0.009$ ) and *Rothia* ( $q=0.012$ ) was significantly ↑ increased in CS neonates compared to VD neonates without IAP (Figure 2).
- ↓ decreased number of observed amplicon sequence variants (ASVs) in CS neonates ( $p=0.02$ ) compared to the VD neonates (Figure 1A).

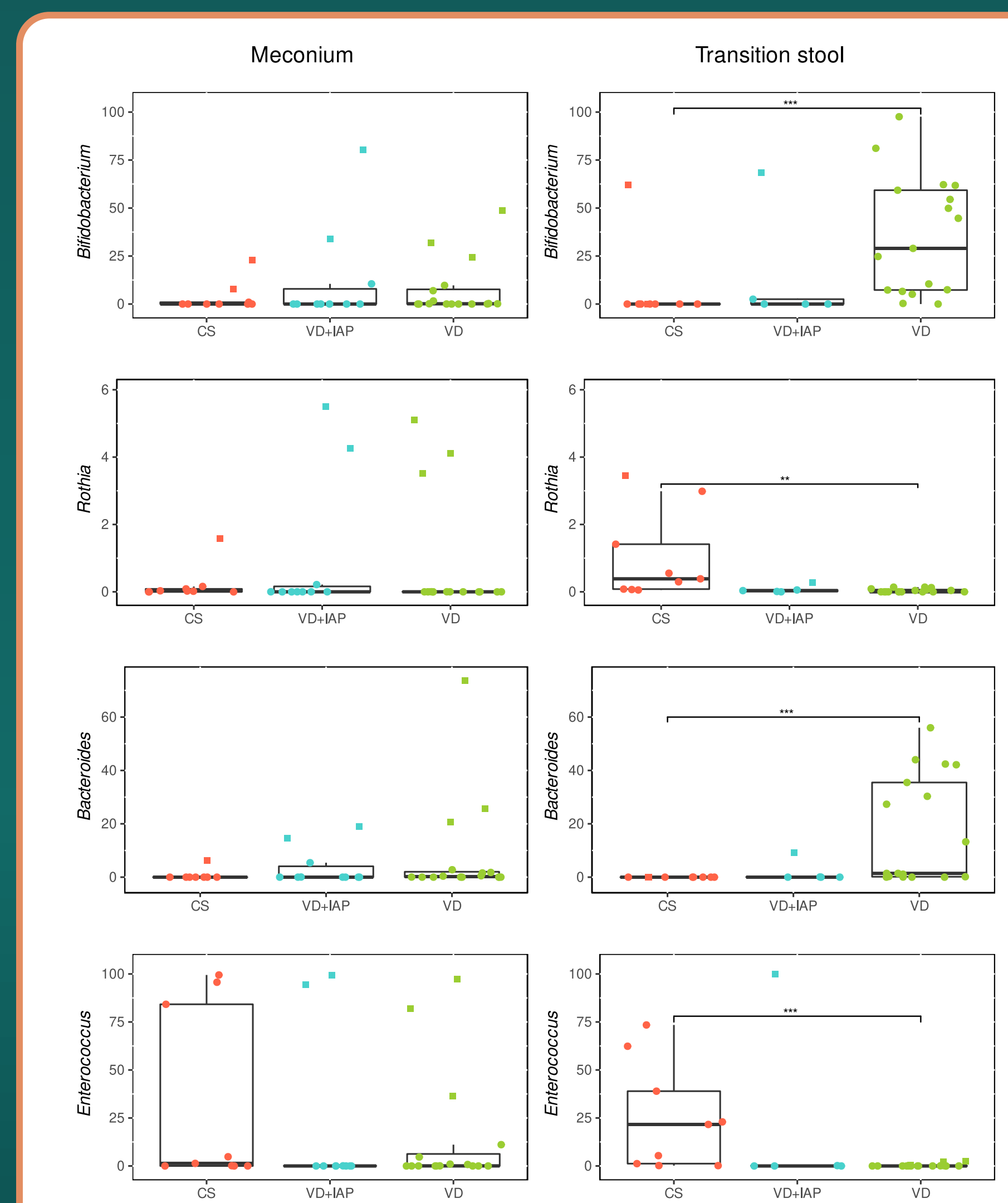


Figure 2. Relative abundance of significantly different bacterial genera in meconium and transitional stool samples (\*\* $p < 0.001$ ; \* $p < 0.01$ ; \* $p < 0.05$ )

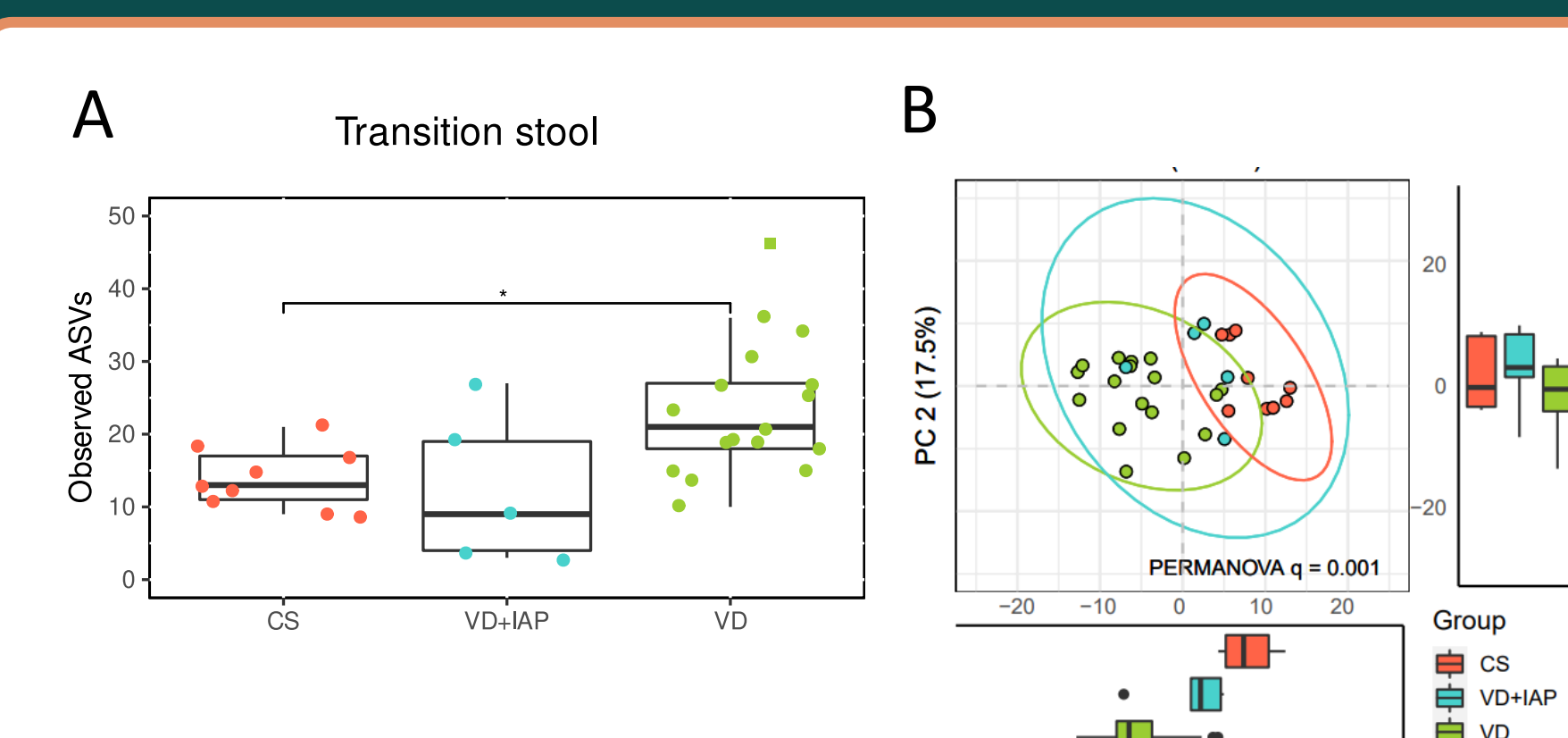


Figure 1. A) Observed ASVs in neonatal transitional stool samples (\* $p < 0.05$ ); B) Principal component analysis of neonatal transitional stool samples



Do antibiotics during labor affect neonatal oral bacteriome as well?

Add your information, graphs and images to this section.

### ORAL MUCOSA

- Differences in alpha diversity (Shannon index,  $p=0.01$ ) and bacterial composition (PERMANOVA,  $p=0.04$ ) were observed in oral samples collected within 48 hours after birth (Figure 3A).
- No significant changes in the bacteriome of oral samples collected after 48 hours.
- Significantly ↓ lower relative abundance of *Gemella* in samples within 48 hours after birth (Figure 3B).

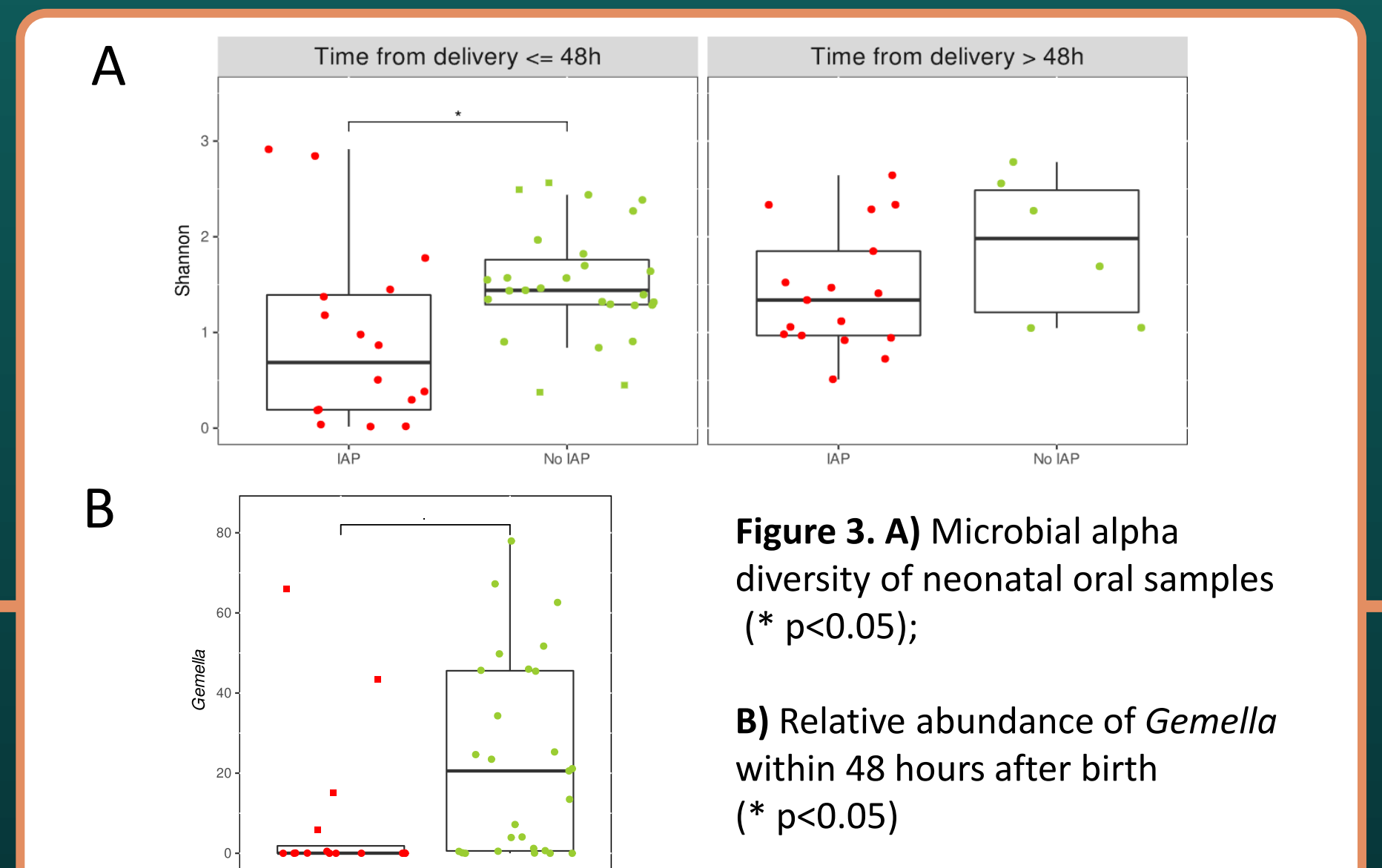


Figure 3. A) Microbial alpha diversity of neonatal oral samples (\* $p < 0.05$ ); B) Relative abundance of *Gemella* within 48 hours after birth (\* $p < 0.05$ )

Does exposure to antibiotics during labor affect neonatal gut bacteriome in the first week of the neonate's life?

## CONCLUSIONS

- The effect of IAP on bacterial abundance and diversity is not reflected in meconium samples yet but is observed in transitional stool samples.
- IAP influences the oral bacteriome of neonates within 48 hours after birth. However, the effect of IAP seems to be suppressed later in the first week of their life.

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