
CHILDHOOD OBESITY,
EARLY CHILDHOOD
CARIES, AND THE ORAL
MICROBIOME:
A SECONDARY
BIOSTATISTICAL
ANALYSIS OF ZOE 2.0

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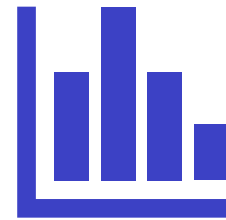
OVERVIEW



Objective



Methods

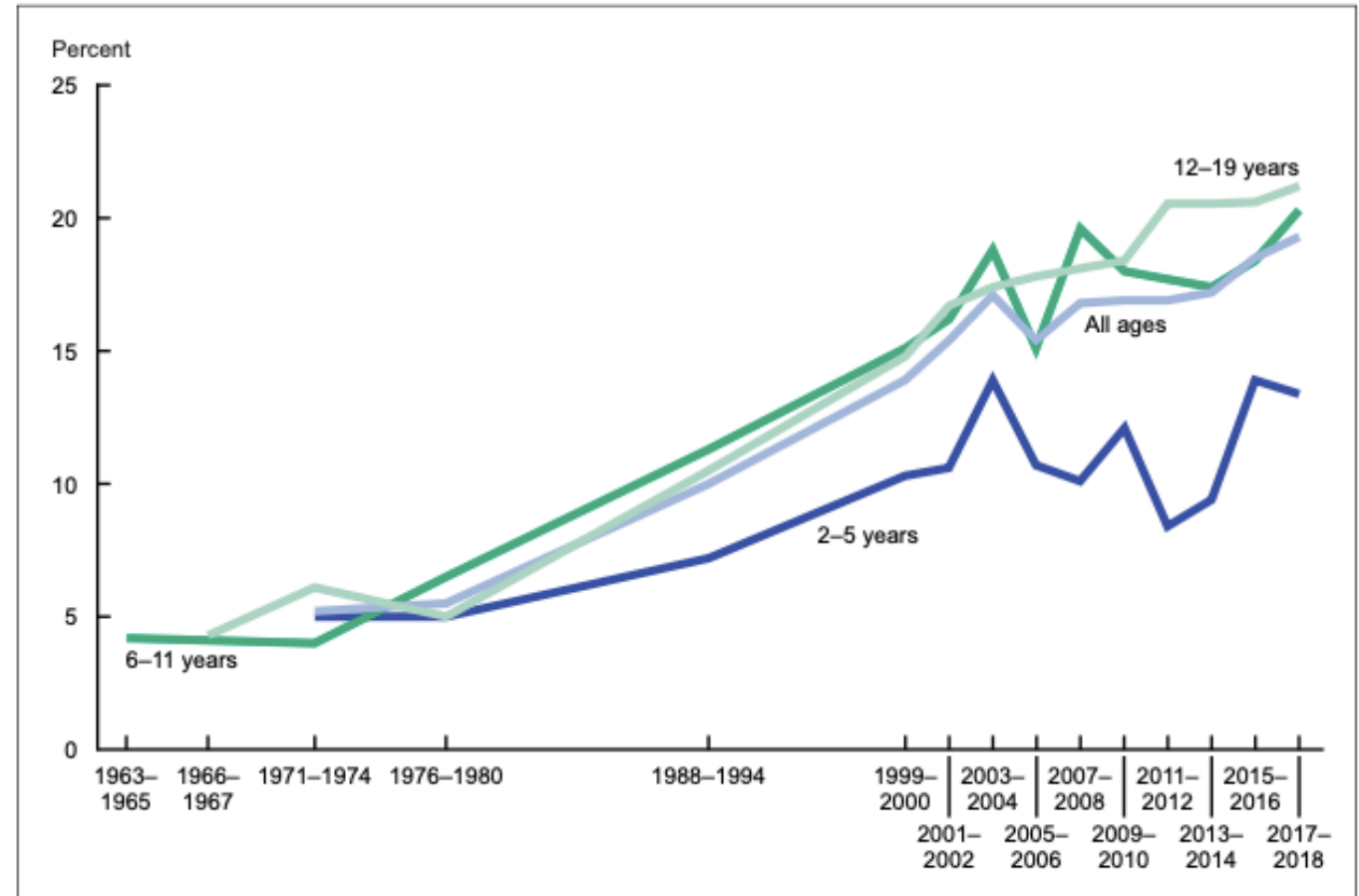


Results

CHILDHOOD OBESITY

- *Cause*: “imbalance between energy intake and expenditure”¹
- *Measure*²: body mass index (BMI), [kg/m²]
 - Obese = 95th percentile among same age and gender group
- *Risks*²
 - Higher risk of chronic diseases in children
 - Higher likelihood of adult obesity

Figure. Trends in obesity among children and adolescents aged 2–19 years, by age: United States, 1963–1965 through 2017–2018



Adopted from: Fryar CD, Carroll MD, Afful J. Prevalence of Overweight, Obesity, and Severe Obesity Among Children and Adolescents Aged 2–19 Years: United States, 1963–1965 Through 2017–2018. *NCHS Health E-Stats*. 2020.

EARLY CHILDHOOD CARIES (ECC)

- *Definition:* “the presence of one or more decayed (noncavitated or cavitated lesions), missing (due to caries), or filled tooth surfaces in any primary tooth in a child under the age of 6”³
 - *Cause:* disrupted balance of demineralization and remineralization at the tooth-biofilm interface, where the microbiome ferments carbohydrates and produces acid⁴
 - *Measure:* International Caries Detection and Assessment System (ICDAS)⁵
 - Sensitive: ICDAS ≥ 1
 - Established/severe: ICDAS ≥ 3
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THE ROLE OF ORAL MICROBIOME

ECC

- Gives rise to dental caries via its metabolic activity⁴

Childhood obesity

- Benahmed et al. (2020)⁶ – modulating effects on the gut microbiome, which plays an important role in the development of obesity
 - Craig et al. (2018)⁷ – stronger indicator of infant weight gain than the gut microbiome
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OBJECTIVE

- i. Explore the potential of oral microbiome as an indicator of childhood obesity
 - ii. Explore the potential of oral microbiome as a common biomarker for childhood obesity and early childhood caries
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ZOE 2.0⁸

- Genetic epidemiological study of oral health in preschool-age (3-5 years) children enrolled in Head Start centers in North Carolina
- Data includes:
 - Family history and demographics
 - Oral health status (dental caries)
 - Environmental factors
 - Some, oral microbiome
- 8,059 enrolled, 6,404 with dental clinical data

STUDY SAMPLE

302 subjects

- 4 subjects excluded from the DNA analysis
- 7 subjects excluded from the RNA analysis

666 core species

- 418 DNA core species
- 632 RNA core species
- 384 overlap

MULTIPLE LINEAR REGRESSION MODELS

Model 1

$zbmi \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

Model 2

$zbmi \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t6 = 0)$

Model 3

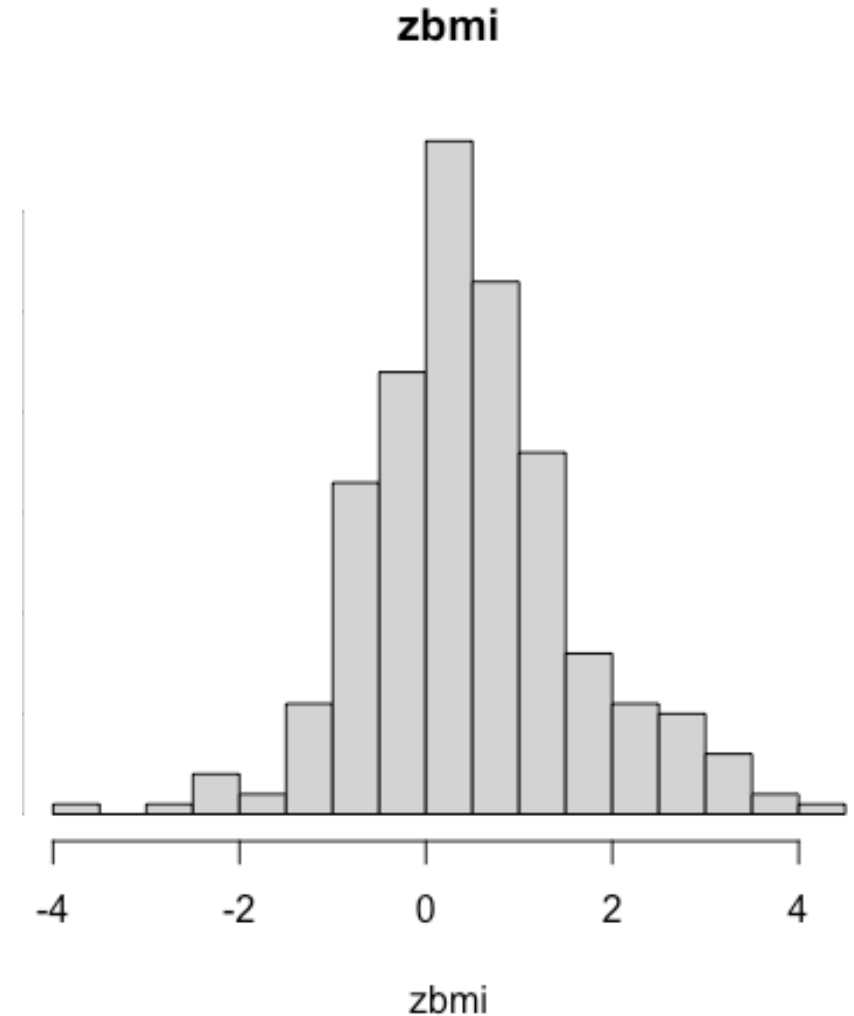
$zbmi \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t6 = 1)$

Model 4

$t6 \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

VARIABLES & DATA DISTRIBUTION

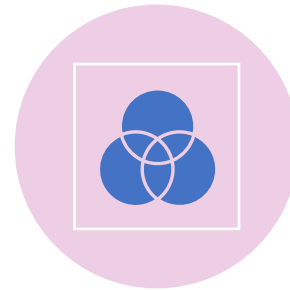
Variable	Type	Description
zbmi	Continuous	Z-score transformed age- and sex- adjusted body mass index (BMI)
species	Continuous	DNA/RNA sequence reads, normalized in transcripts per million (TPM), for the individual bacterial species
age	Continuous	Age in months at the time of exam
gender	Binary	Genetically verified gender (boy/girl)
batch	Categorical	Identification number for each sequencing batch
t6	Binary	Presence of established/severe dental caries, based on international standards (ICDAS ≥ 3)



STATISTICAL ANALYSIS



Each linear regression model run with DNA sequence data and RNA sequence data



P-values and coefficients extracted



P-values adjusted for multiple regression using the FDR method



List of significant species from each model compared

RESULTS

Model	$p < 0.05$	FDR adjusted $p < 0.2$
1	75	5
2	26	0
3	78	29
4	31	11

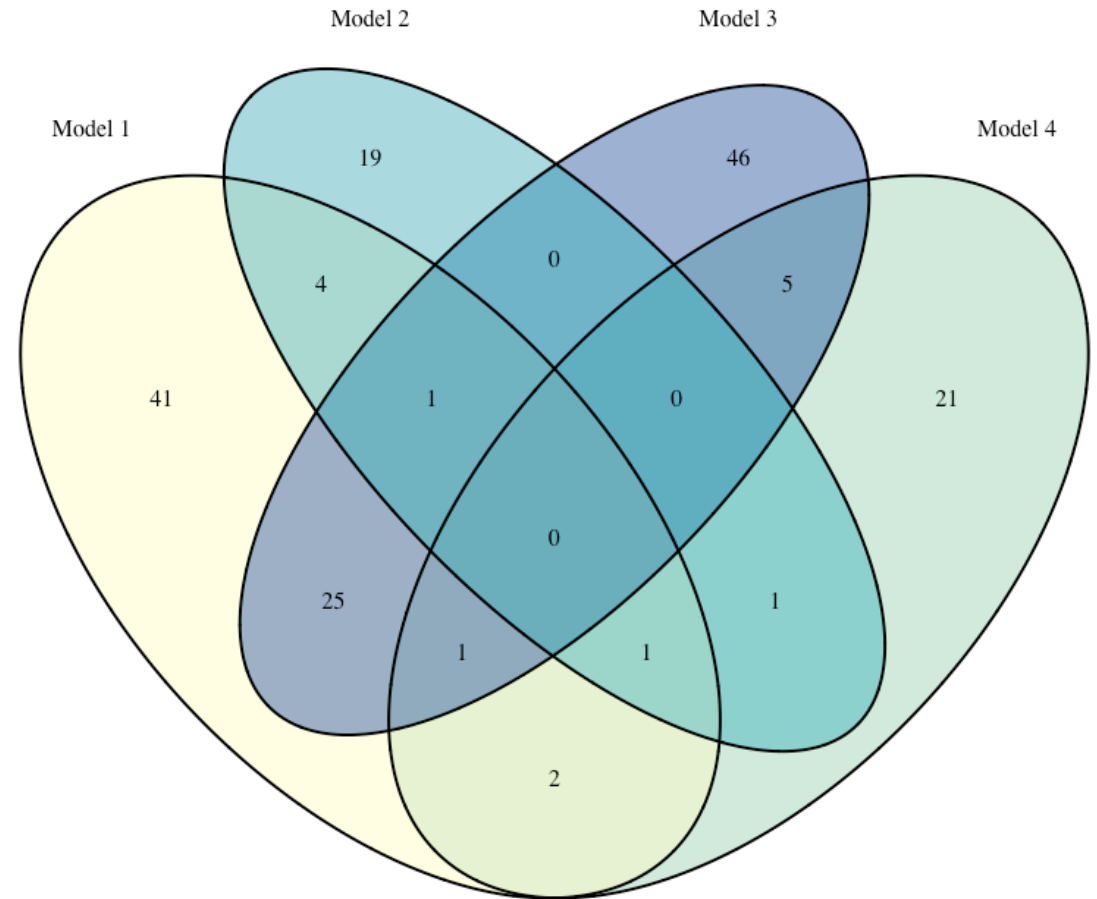
Model 1: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

Model 2: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 0)$

Model 3: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 1)$

Model 4: $t_6 \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

Significant species, nominal $p < 0.05$



Significant species, nominal $p < 0.05$

RESULTS

Total number of significant species
across the 4 models: 167

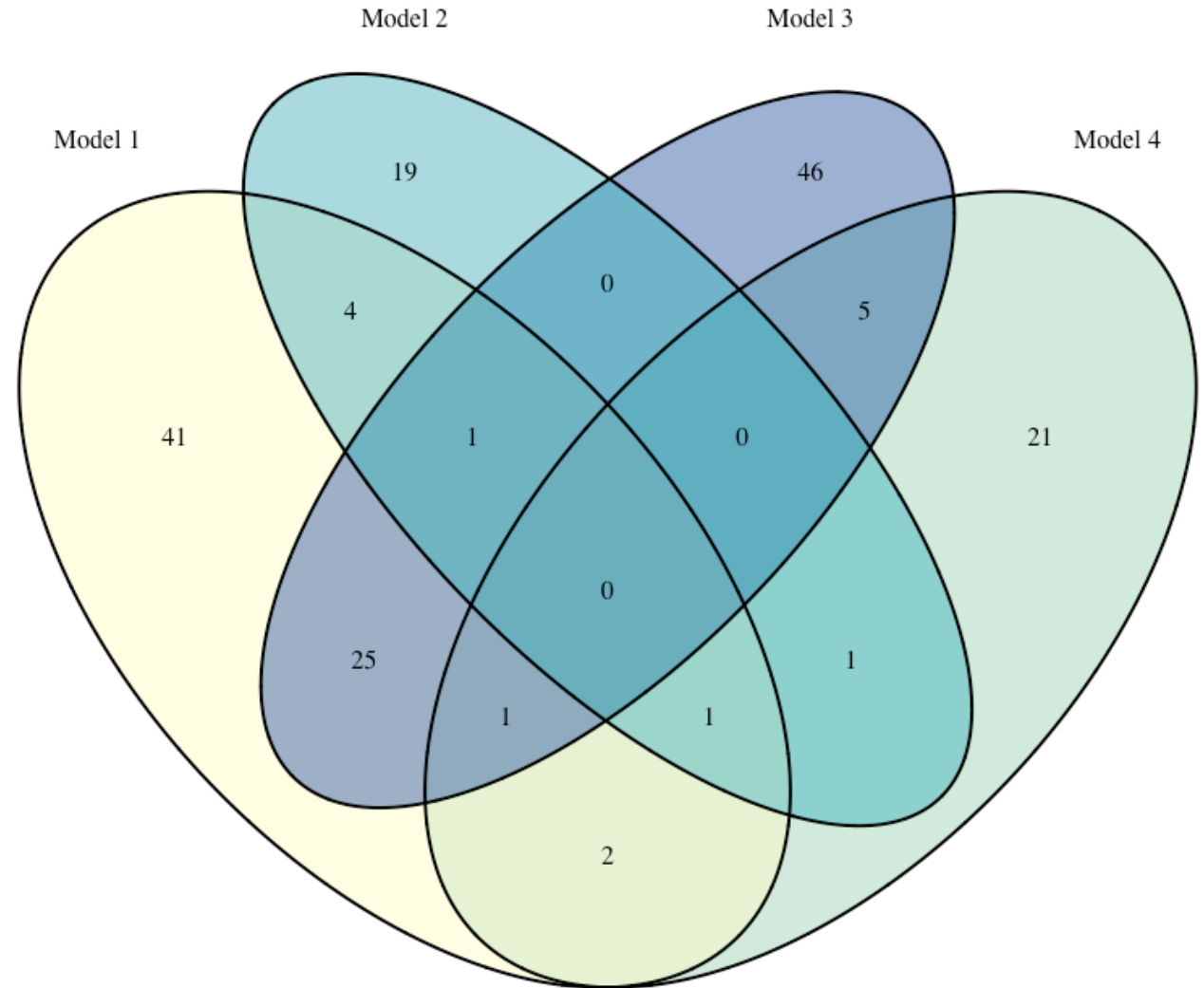
- Obesity associated: 146
- Obesity + ECC associated: 104

Model 1: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

Model 2: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 0)$

Model 3: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 1)$

Model 4: $t_6 \sim \text{species} + \text{age} + \text{gender} + \text{batch}$



Significant species, nominal $p < 0.05$

RESULTS

Associated with obesity only in the absence of dental caries

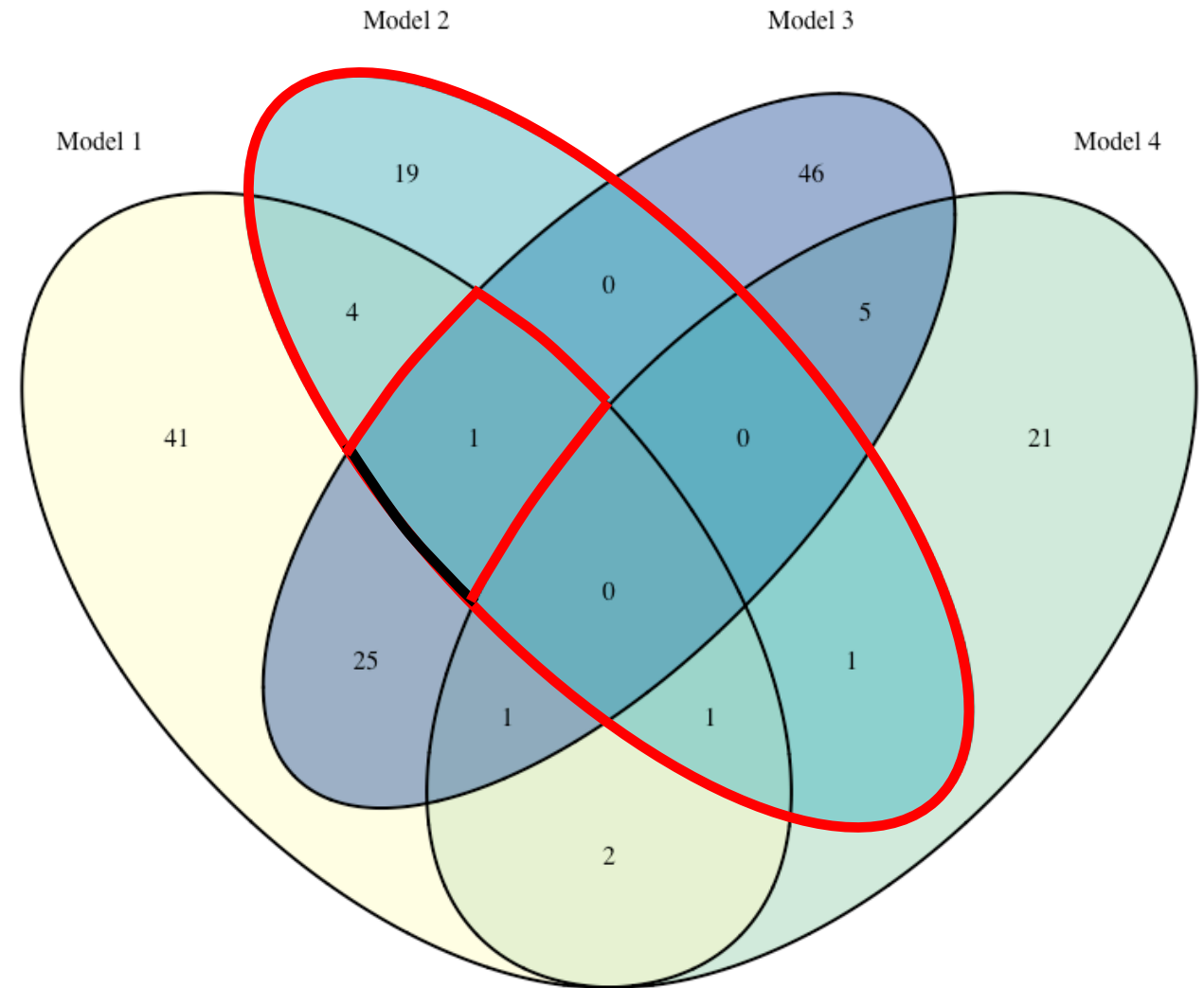
- 25 species

Model 1: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

Model 2: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 0)$

Model 3: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 1)$

Model 4: $t_6 \sim \text{species} + \text{age} + \text{gender} + \text{batch}$



Significant species, nominal $p < 0.05$

RESULTS

Associated with obesity only in the presence of caries

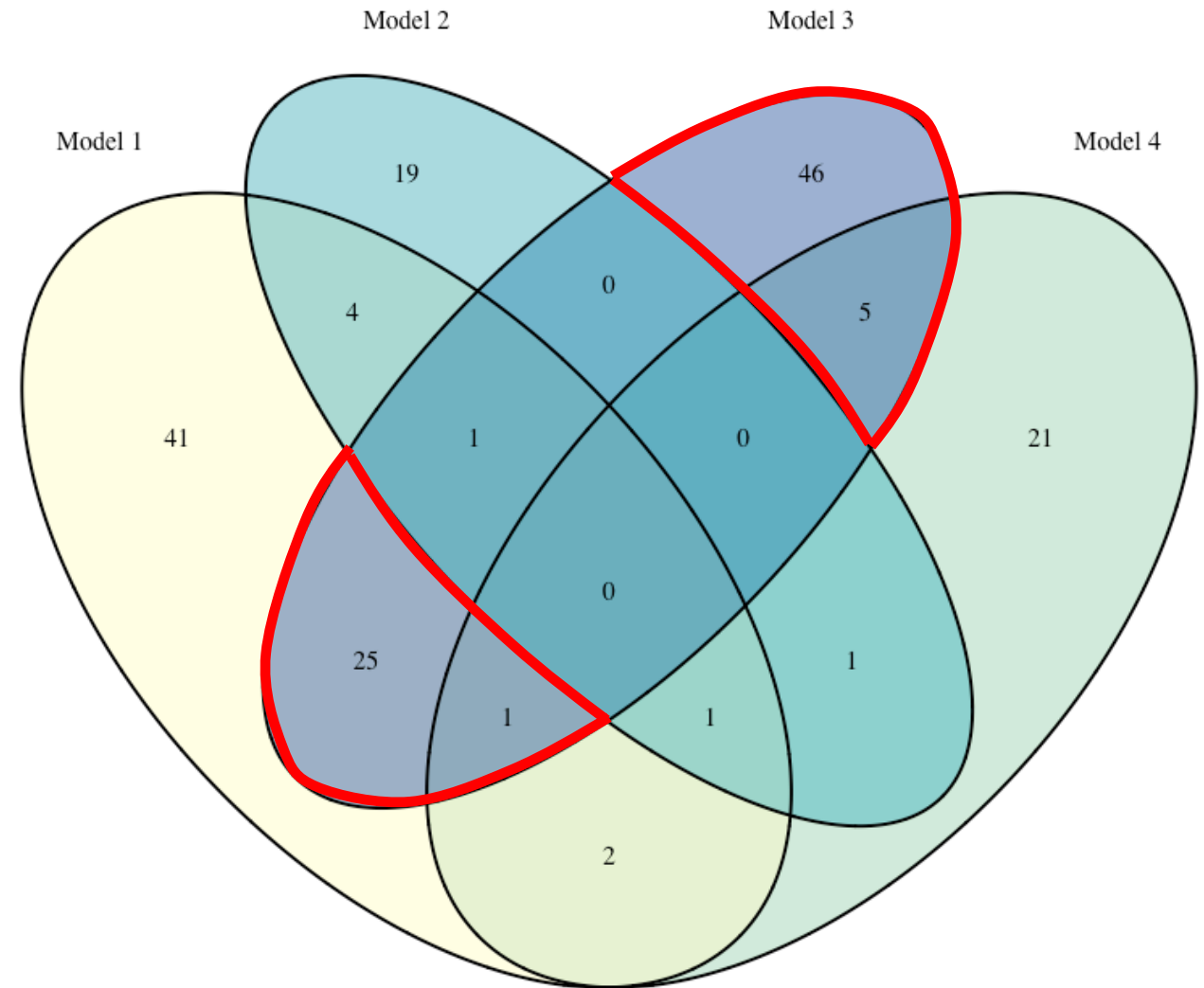
- 77 species
- 24 species with FDR adjusted $p < 0.2$

Model 1: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

Model 2: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 0)$

Model 3: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 1)$

Model 4: $t_6 \sim \text{species} + \text{age} + \text{gender} + \text{batch}$



Significant species, nominal $p < 0.05$

RESULTS

Only associated with obesity, no ECC association

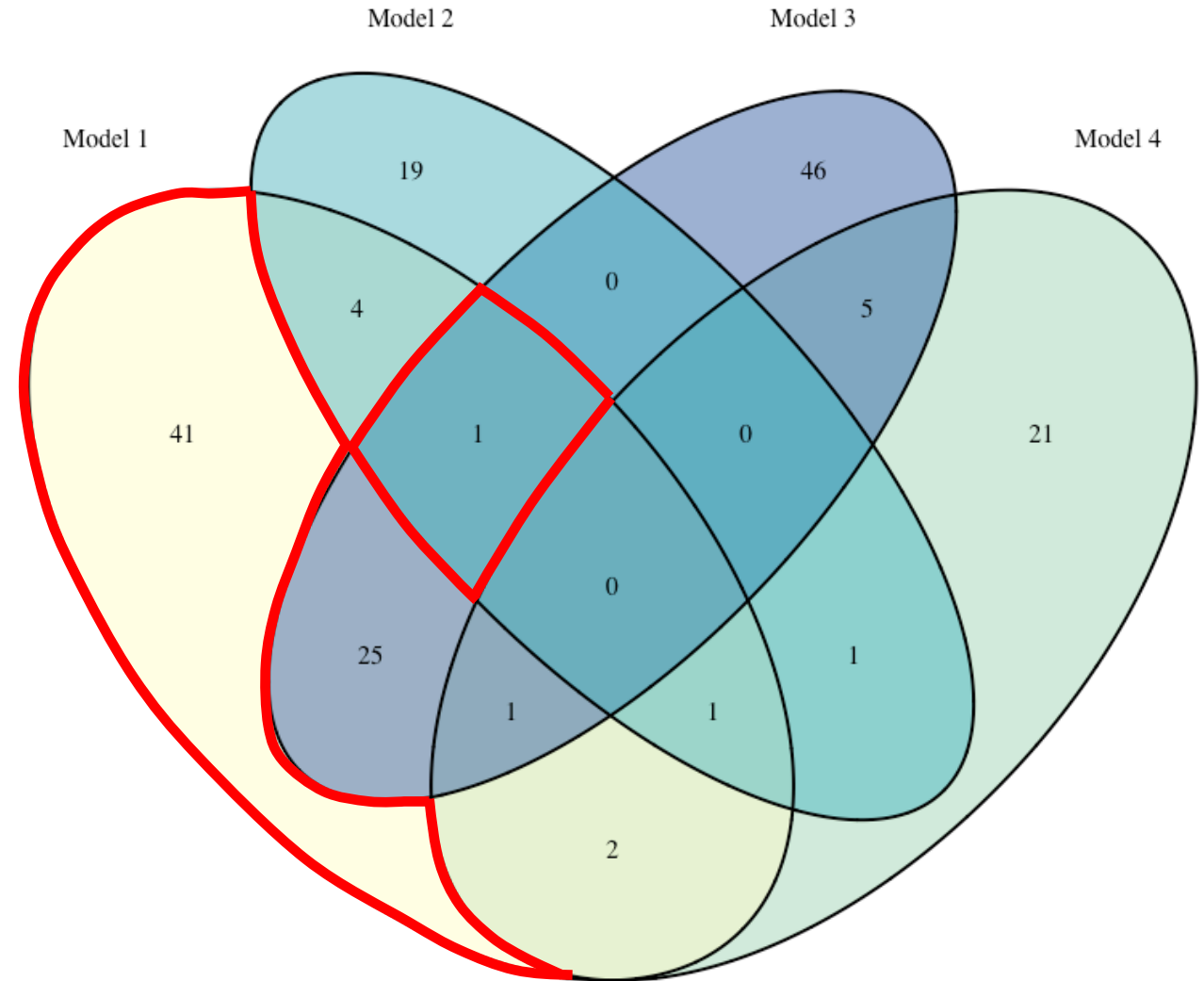
- 42 species
- 2 species with FDR adjusted $p < 0.2$
 - *Glaesserella* sp. 15-814 (-)
 - *Actinobacillus pleuropneumoniae* (+)

Model 1: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

Model 2: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 0)$

Model 3: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 1)$

Model 4: $t_6 \sim \text{species} + \text{age} + \text{gender} + \text{batch}$



Significant species, nominal $p < 0.05$

RESULTS

Associated with obesity and ECC
on separate terms

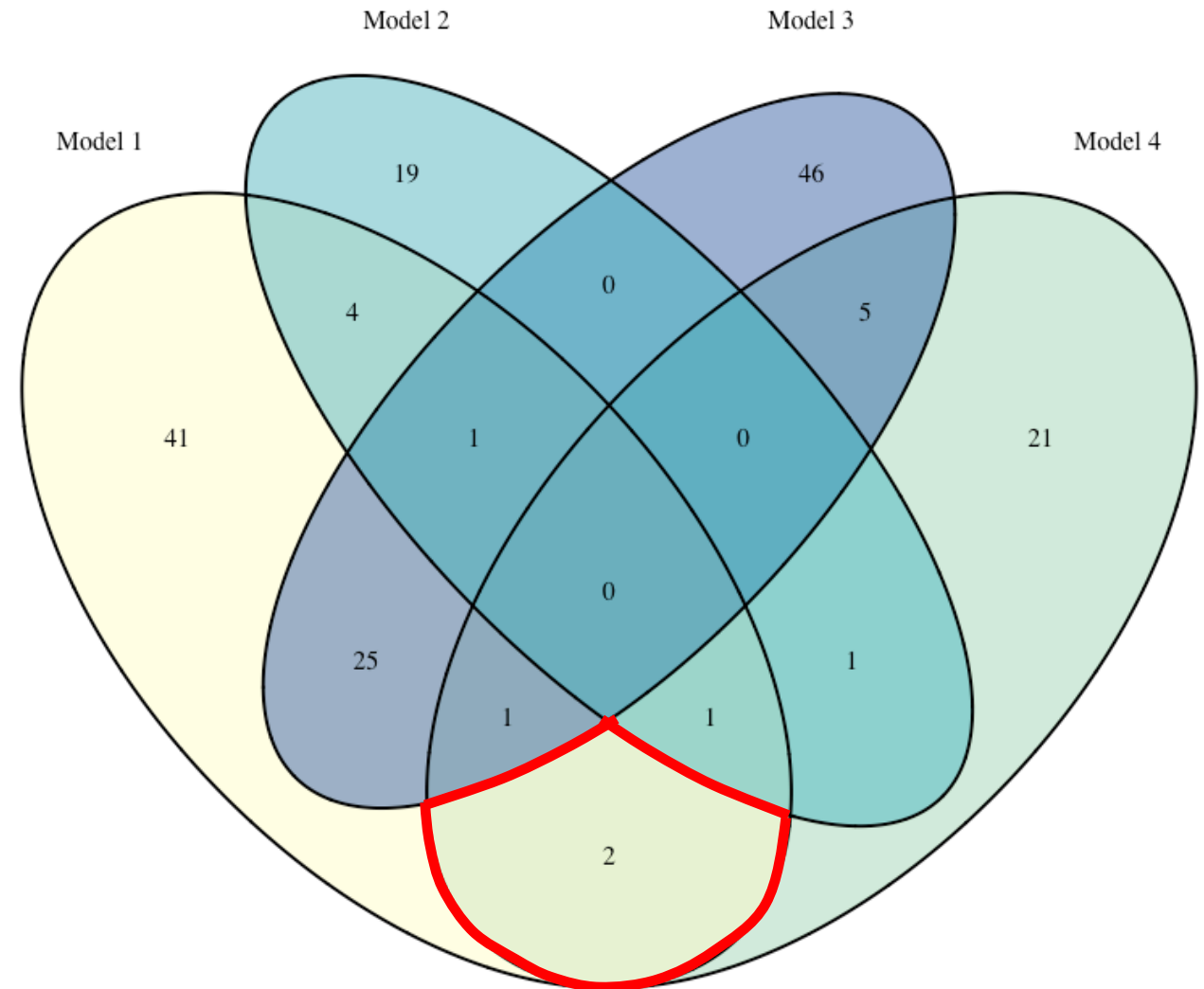
- 2 species
- 1 species with FDR adjusted $p < 0.2$
 - *Veillonella dispar* (-/-)

Model 1: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch}$

Model 2: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 0)$

Model 3: $z_{bmi} \sim \text{species} + \text{age} + \text{gender} + \text{batch} (t_6 = 1)$

Model 4: $t_6 \sim \text{species} + \text{age} + \text{gender} + \text{batch}$



CONCLUSION

- I. *Glaesserella sp. 15-184* and *Actinobacillus pleuropneumoniae* may potentially serve as an indicator of childhood obesity.
 - II. *Veillonella dispar* may potentially serve as a common biomarker for childhood obesity and early childhood caries.
 - III. Other species need further investigation for conclusive results.
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LIMITATIONS + FUTURE DIRECTIONS

- Limitation: limited sample size, resulting in limited testing power
- Future directions
 - 1) Contradicting directionality of correlation from stratified sampling
 - 2) Biological mechanisms of identified species in their roles in childhood obesity and ECC
 - 3) Metabolomics of the oral biofilm with regards to the two conditions

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THANK YOU!

