

SUPPLEMENTARY MATERIAL

Early-life gut microbiome composition is associated with milk allergy resolution

Supinda Bunyavanich, MD, MPH^{1,2*}, Nan Shen, PhD¹, Alexander Grishin, PhD², Robert Wood, MD³, Wesley Burks, MD⁴, Peter Dawson, PhD⁵, Stacie M. Jones, MD⁶, Donald Leung, MD, PhD⁷, Hugh Sampson, MD⁸, Scott Sicherer, MD⁹, Jose C. Clemente, PhD^{1,8*}

Supplementary Table 1: Baseline characteristics of the participants stratified by age at time of stool collection

Characteristics	All subjects with milk allergy included in the analysis	Age 3-6 months	Age 7-12 months	Age 13-16 months	P-value*
Subjects	226	29 (12.8%)	144 (63.7%)	53 (23.4%)	
Age at last followup (months) – median(IQR)	80.0 (18.0)	72.4 (18.8)	79.4 (22.5)	83.6 (10.8)	
Sex					0.23
Female	77	14 (18.2%)	46 (59.7%)	17 (22.1%)	
Male	149	15 (10.1%)	98 (65.8%)	36 (24.2%)	
Race					0.02
White	174	16 (9.2%)	112 (64.4%)	46 (26.4%)	
Black/African-American	30	8 (26.7%)	16 (53.3%)	6 (20%)	
Asian	19	5 (26.3%)	13 (68.4%)	1 (5.3%)	
Other	3	0 (0.0%)	3 (100.0%)	0 (0.0%)	
Baseline milk IgE level					0.17
<2	91	18 (19.8%)	52 (57.1%)	21 (23.1%)	
≥ 2-10	68	6 (8.8%)	46 (67.6%)	16 (23.5%)	
≥10	67	5 (7.5%)	46 (68.7%)	16 (23.9%)	
Baseline milk SPT response (wheal in mm)					0.0005
<5	63	18 (28.6%)	29 (46.0%)	16 (25.4%)	
5-10	82	7 (8.5%)	56 (68.3%)	19 (23.2%)	
>10	81	4 (4.9%)	59 (72.8%)	18 (22.2%)	
Baseline AD severity					0.01
None	29	4 (13.8%)	15 (51.7%)	10 (34.5%)	
Mild	32	0 (0.0%)	24 (75.0%)	8 (25.0%)	

Moderate	126	14 (11.1%)	83 (65.9%)	29 (23.0%)	
Severe	39	11 (28.2%)	22 (56.4%)	6 (15.4%)	
Breast-feeding at entry					0.15
Never	36	6 (16.7%)	22 (61.1%)	8 (22.2%)	
Yes, currently	115	14 (12.2%)	67 (58.3%)	34 (29.6%)	
Yes, but no longer	75	9 (12.0%)	55 (73.3%)	11 (14.7%)	
Mode of delivery**					0.04
Vaginal	140	23 (16.4%)	90 (64.3%)	27 (19.3%)	
C-section	85	6 (7.1%)	53 (62.4%)	26 (30.6%)	
Solid food intake					2.8x10-7
Yes	219	22 (10.0%)	144 (65.8%)	53 (24.2%)	
No	7	7 (100.0%)	0 (0.0%)	0 (0.0%)	
Antibiotics – lifetime prior to sampling					0.01
Yes	142	11 (7.7%)	95 (66.9%)	36 (25.4%)	
No	84	18 (21.4%)	49 (58.3%)	17 (20.2%)	

**Data N/A for 1 subject

*Fisher's Exact test

Supplementary Table 2: Association between early life exposures and main outcomes of interest

	Milk allergy resolution by age 8 years OR (95% CI)	Sensitization to milk at baseline OR (95% CI)
Mode of delivery	1.08 (0.61-1.94)	1.50 (0.72-3.27)
Breastfeeding status		
Never	reference	Reference
Yes, currently	0.92 (0.42-1.97)	0.72 (0.25-1.83)
Yes, no longer	0.65 (0.29-1.46)	0.74 (0.24-2.00)
Solid food intake	0.51 (0.05-3.20)	10.38 (1.63-112.62)
Antibiotic use	1.05 (0.58-1.87)	1.48 (0.73-2.99)

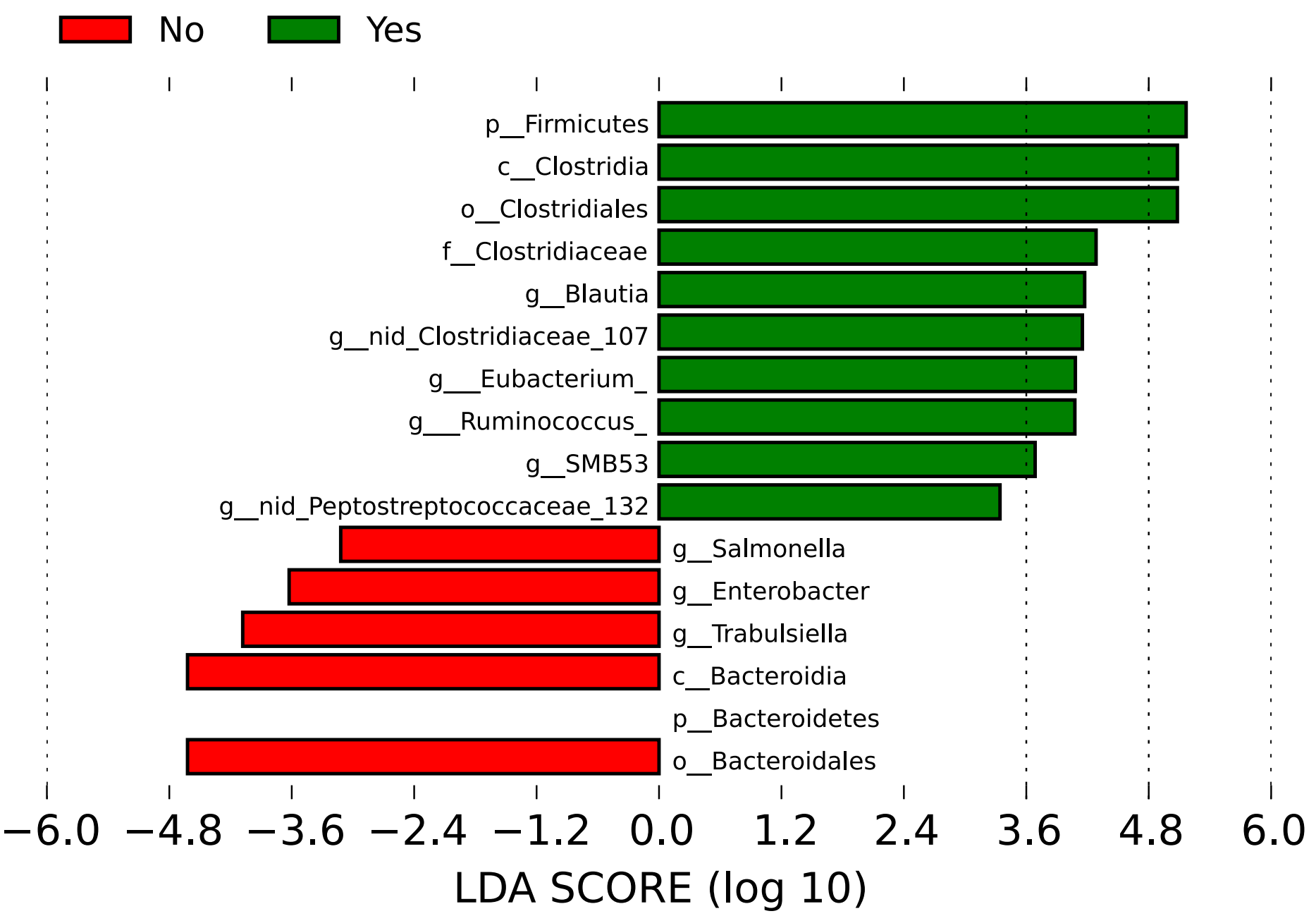
Supplementary Table 3. Results for tests of association between bacterial functional pathways and milk allergy resolution using STAMP. Subjects were stratified by age (3-6 months, 7-12 months, 13-16 months). PICRUSt-predicted KEGG metabolic pathways were utilized as input for STAMP analysis. ANOVA with Tukey-Kramer post-hoc test and Benjamini-Hochberg FDR correction for multiple testing were utilized to test for associations between each pathway and resolution or persistence. Effect size was measured as η^2 . Pathways are represented using standard KEGG conventions, e.g. “Metabolism” at the top level, “Lipid Metabolism” as the second level, and “Fatty acid metabolism” at the lowest level. Degenerate cases where the test statistic could not be calculated are indicated. See file [Bacterial_functional_pathways_milk_resolution_STAMP.xlsx](#).

Supplementary Figure 1. Linear discriminant analysis shows distinct gut microbiome composition associated with milk allergy resolution regardless of covariates. After accounting for specific covariates known to play a role in shaping infant microbiome composition, taxa within *Clostridia* and *Firmicutes* were still enriched in children sampled at age 3-6 months with milk allergy resolution vs. milk allergy persistence by age 8 years, while taxa within *Bacteroidetes* were underrepresented. Linear Discriminant Analysis scores as calculated by LEfSe are presented for antibiotics, breastfeeding, mode of delivery, race, and solid food intake. All taxa were significantly associated with either resolution or persistence of milk allergy ($P < 0.05$). Only taxa with LDA scores >2 are presented.

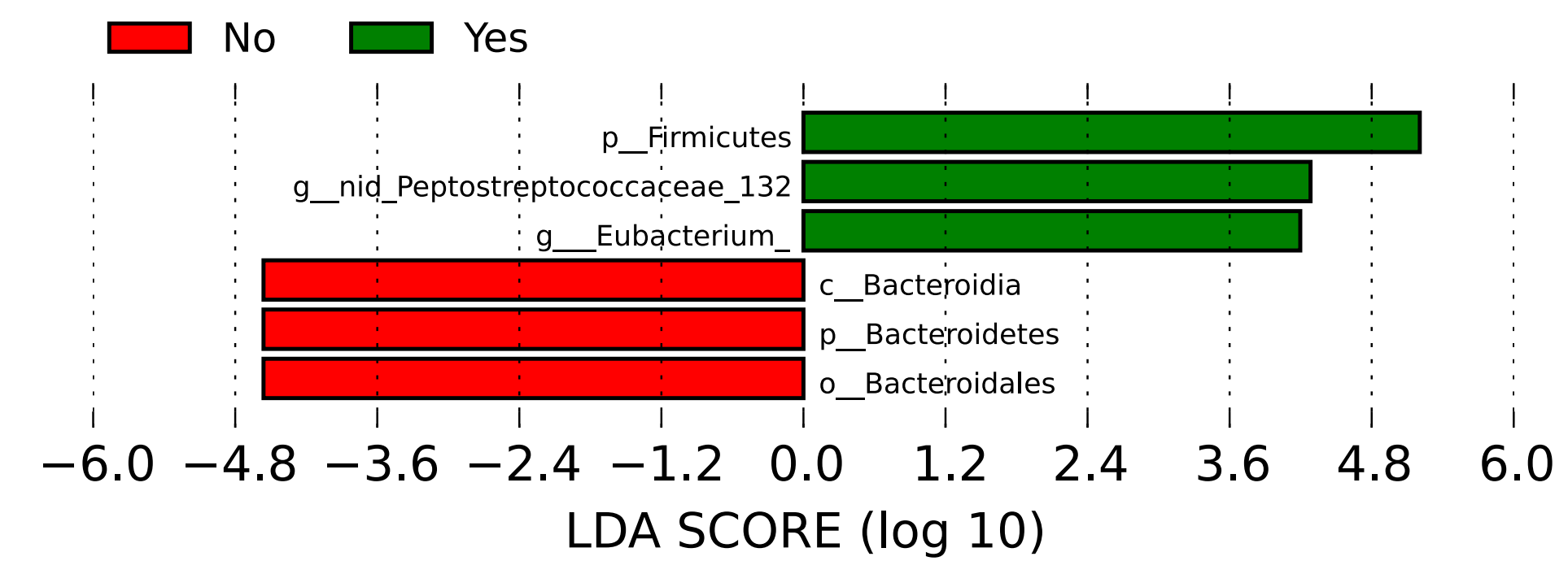
Metabolism Lipid metabolism	Lipid metabolism_Unclassified	0.843860869	276.7863649	0.001462523	0.146009196	0.054308209	0.149148573	0.027385534
Metabolism Biosynthesis of Other Secondary Metabolites	Stilbenoid, diarylheptanoid and gingerol biosynthesis	0.876071468	287.3514416	0.00091718	0.001757368	0.004243124	0.002009992	0.003919988
Metabolism Metabolism of Terpenoids and Polyketides	Prenyltransferases	0.897233102	294.2924575	0.000629185	0.287896653	0.068070199	0.290573069	0.038523696
Metabolism Lipid Metabolism	Lipid biosynthesis proteins	0.923254833	302.8275852	0.000350078	0.527921102	0.049050206	0.526067061	0.047464845
Metabolism Biosynthesis of Other Secondary Metabolites	Streptomycin biosynthesis	0.935087267	306.7086237	0.000250243	0.276899212	0.079469545	0.27882474	0.041941986
Metabolism Biosynthesis of Other Secondary Metabolites	Butirosin and neomycin biosynthesis	0.937216743	307.4070917	0.000234062	0.056446665	0.023119498	0.056992273	0.01247638
Metabolism Amino acid metabolism	Amino acid metabolism_Unclassified	0.93995026	308.3036853	0.000214089	0.219963849	0.067246139	0.221536632	0.040227637
Metabolism Lipid Metabolism	Secondary bile acid biosynthesis	0.954781921	313.16847	0.0001213	0.033027318	0.022249225	0.033374427	0.008613916
Metabolism Biosynthesis of Other Secondary Metabolites	Isoflavonoid biosynthesis	0.979875552	321.3991811	2.40E-05	3.31E-07	1.05E-06	3.44E-07	1.42E-06
Metabolism Xenobiotics Biodegradation and Metabolism	Xylene degradation	0.980292846	321.5360534	2.30E-05	0.067656022	0.022473245	0.067478882	0.014434701
Metabolism Xenobiotics Biodegradation and Metabolism	Ethylbenzene degradation	0.981598092	321.9641742	2.01E-05	0.039260169	0.012778844	0.039123785	0.015865002
Metabolism Lipid Metabolism	Primary bile acid biosynthesis	0.989429314	324.5328149	6.62E-06	0.033665751	0.021984565	0.033745806	0.008467424
Metabolism Biosynthesis of Other Secondary Metabolites	Clavulanic acid biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Enzyme Families	Cytochrome P450	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Glycan Biosynthesis and Metabolism	Glycosaminoglycan biosynthesis - chondroitin sulfate	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Glycan Biosynthesis and Metabolism	Glycosylphosphatidylinositol(GPI)-anchor biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Glycan Biosynthesis and Metabolism	Other types of O-glycan biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Metabolism of Terpenoids and Polyketides	Biosynthesis of 12-, 14- and 16-membered macrolides	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Metabolism of Terpenoids and Polyketides	Biosynthesis of type II polyketide backbone	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Metabolism of Terpenoids and Polyketides	Sesquiterpenoid biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0

Metabolism Metabolism of Cofactors and Vitamins	Ubiquinone and other terpenoid-quinone biosynthesis	0.927171539	304.1122649	5.90E-05	0.155495786	0.088443053	0.156838781	0.085074815
Metabolism Biosynthesis of Other Secondary Metabolites	Clavulanic acid biosynthesis	0.928191245	304.4467284	5.74E-05	3.81E-08	2.98E-07	4.35E-08	3.92E-07
Metabolism Lipid Metabolism	alpha-Linolenic acid metabolism	0.931610292	305.5681757	5.20E-05	0.007491134	0.012577925	0.007316072	0.01157112
Metabolism Carbohydrate Metabolism	Amino sugar and nucleotide sugar metabolism	0.939045387	308.0068868	4.13E-05	1.525691638	0.152603198	1.523874396	0.1296052
Metabolism Biosynthesis and biodegradation of secondary metabolites	Biosynthesis and biodegradation of secondary metabolites_Unclassified	0.943409872	309.438438	3.56E-05	0.068022687	0.036682305	0.067604827	0.033071328
Metabolism Xenobiotics Biodegradation and Metabolism	Polycyclic aromatic hydrocarbon degradation	0.956840584	313.8437115	2.07E-05	0.099518043	0.022695343	0.099320652	0.020519053
Metabolism Biosynthesis of Other Secondary Metabolites	Novobiocin biosynthesis	0.959421363	314.6902072	1.83E-05	0.135456232	0.012427849	0.135351882	0.011810023
Metabolism Amino Acid Metabolism	Amino acid related enzymes	0.969044566	317.8466177	1.06E-05	1.445095238	0.115708433	1.445854257	0.114820108
Metabolism Amino Acid Metabolism	Valine, leucine and isoleucine biosynthesis	0.971387912	318.615235	9.09E-06	0.798994124	0.081837924	0.799505127	0.085451796
Metabolism Biosynthesis of Other Secondary Metabolites	Flavone and flavonol biosynthesis	0.976208377	320.1963476	6.29E-06	0.005054524	0.005178453	0.005080003	0.004918558
Metabolism Metabolism of Terpenoids and Polyketides	Geraniol degradation	0.976411912	320.2631072	6.18E-06	0.040314828	0.04608646	0.040090605	0.043562593
Metabolism Glycan biosynthesis and metabolism	Glycan biosynthesis and metabolism_Unclassified	0.980023833	321.4478174	4.43E-06	0.027993974	0.031267095	0.028119965	0.028343259
Metabolism Energy Metabolism	Photosynthesis proteins	0.99315022	325.7532722	5.21E-07	0.39252015	0.099267099	0.392408706	0.053065188
Metabolism Amino Acid Metabolism	Histidine metabolism	0.999621254	327.8757714	1.59E-09	0.60178324	0.059634292	0.60178832	0.065495458
Metabolism Glycan Biosynthesis and Metabolism	Glycosaminoglycan biosynthesis - chondroitin sulfate	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Glycan Biosynthesis and Metabolism	Glycosylphosphatidylinositol(GPI)-anchor biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Glycan Biosynthesis and Metabolism	Other types of O-glycan biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Metabolism of Terpenoids and Polyketides	Biosynthesis of 12-, 14- and 16-membered macrolides	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Metabolism of Terpenoids and Polyketides	Biosynthesis of type II polyketide backbone	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Metabolism of Terpenoids and Polyketides	Sesquiterpenoid biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0

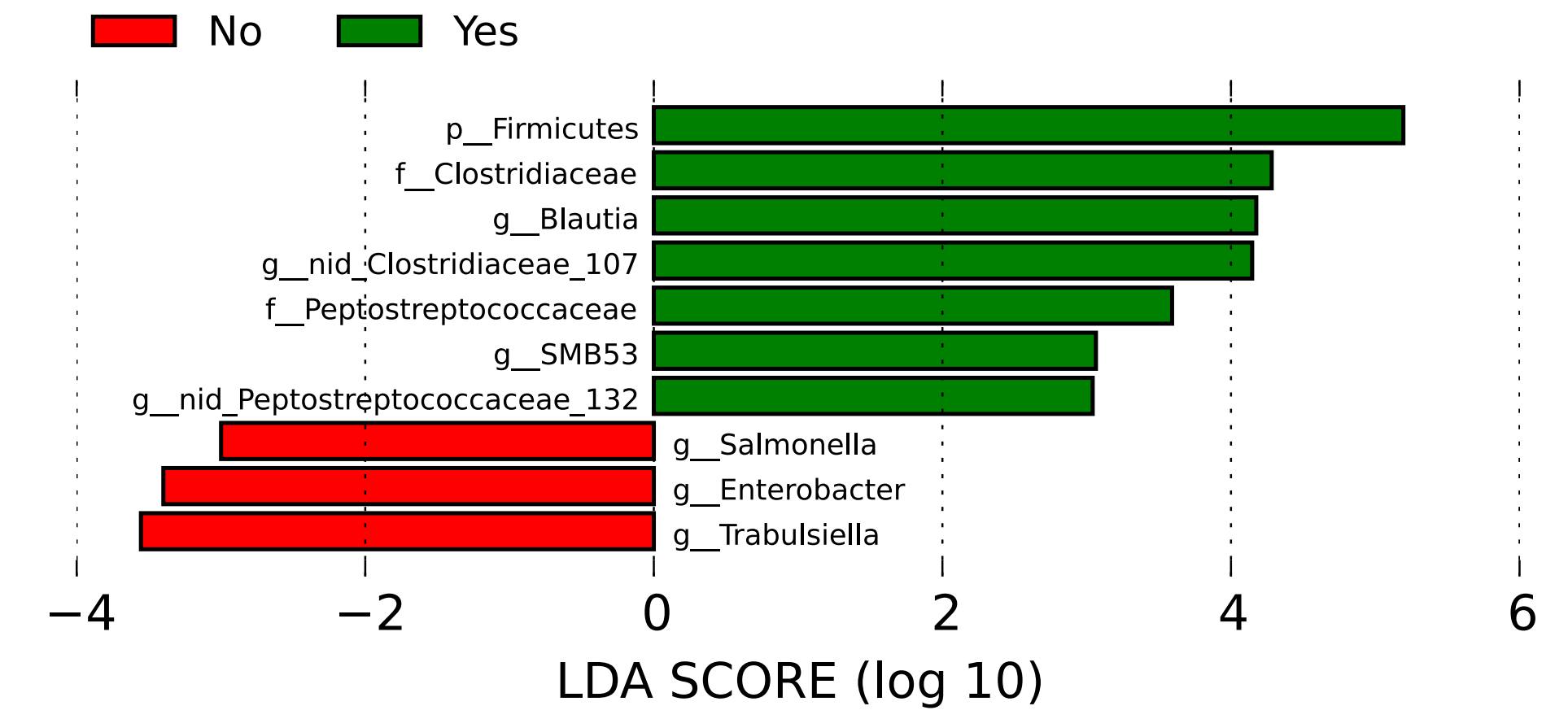
Metabolism Lipid Metabolism	Lipid biosynthesis proteins	0.934867523	306.6365477	0.000132233	0.54893116	0.051213294	0.549974231	0.039488007
Metabolism Biosynthesis of Other Secondary Metabolites	Novobiocin biosynthesis	0.947498573	310.7795321	8.59E-05	0.13426206	0.00986359	0.134438097	0.009057363
Metabolism Energy Metabolism	Sulfur metabolism	0.947562563	310.8005208	8.56E-05	0.278881941	0.035179283	0.278278586	0.029926875
Metabolism Amino acid metabolism	Amino acid metabolism_Unclassified	0.949302819	311.3713247	8.00E-05	0.201792901	0.045628984	0.201025879	0.03991316
Metabolism Biosynthesis of Other Secondary Metabolites	Flavone and flavonol biosynthesis	0.955522814	313.4114829	6.16E-05	0.006963852	0.008258799	0.007084349	0.007071328
Metabolism Metabolism of Other Amino Acids	Phosphonate and phosphinate metabolism	0.959918581	314.8532947	5.00E-05	0.06005206	0.010363232	0.059891407	0.011900241
Metabolism Energy Metabolism	Photosynthesis - antenna proteins	0.976254636	320.2115208	1.75E-05	3.09E-05	6.62E-05	3.03E-05	8.63E-05
Metabolism Energy Metabolism	Carbon fixation pathways in prokaryotes	0.976744082	320.372059	1.68E-05	0.924870865	0.091899017	0.925571139	0.078563156
Metabolism Energy metabolism	Energy metabolism_Unclassified	0.982314928	322.1992965	9.73E-06	0.778388217	0.116936608	0.779094115	0.108358105
Metabolism Lipid Metabolism	Steroid hormone biosynthesis	0.983417245	322.5608563	8.55E-06	0.020566688	0.022671196	0.020433963	0.02235168
Metabolism Biosynthesis of Other Secondary Metabolites	Betalain biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Biosynthesis of Other Secondary Metabolites	Clavulanic acid biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Biosynthesis of Other Secondary Metabolites	Indole alkaloid biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Biosynthesis of Other Secondary Metabolites	Isoflavonoid biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Glycan Biosynthesis and Metabolism	Glycosaminoglycan biosynthesis - chondroitin sulfate	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Glycan Biosynthesis and Metabolism	Glycosylphosphatidylinositol(GPI)-anchor biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
Metabolism Glycan Biosynthesis and Metabolism	Other types of O-glycan biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0
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Metabolism Metabolism of Terpenoids and Polyketides	Sesquiterpenoid biosynthesis	1	328	-1 degenerate case: failed to calculate p-value	0	0	0	0



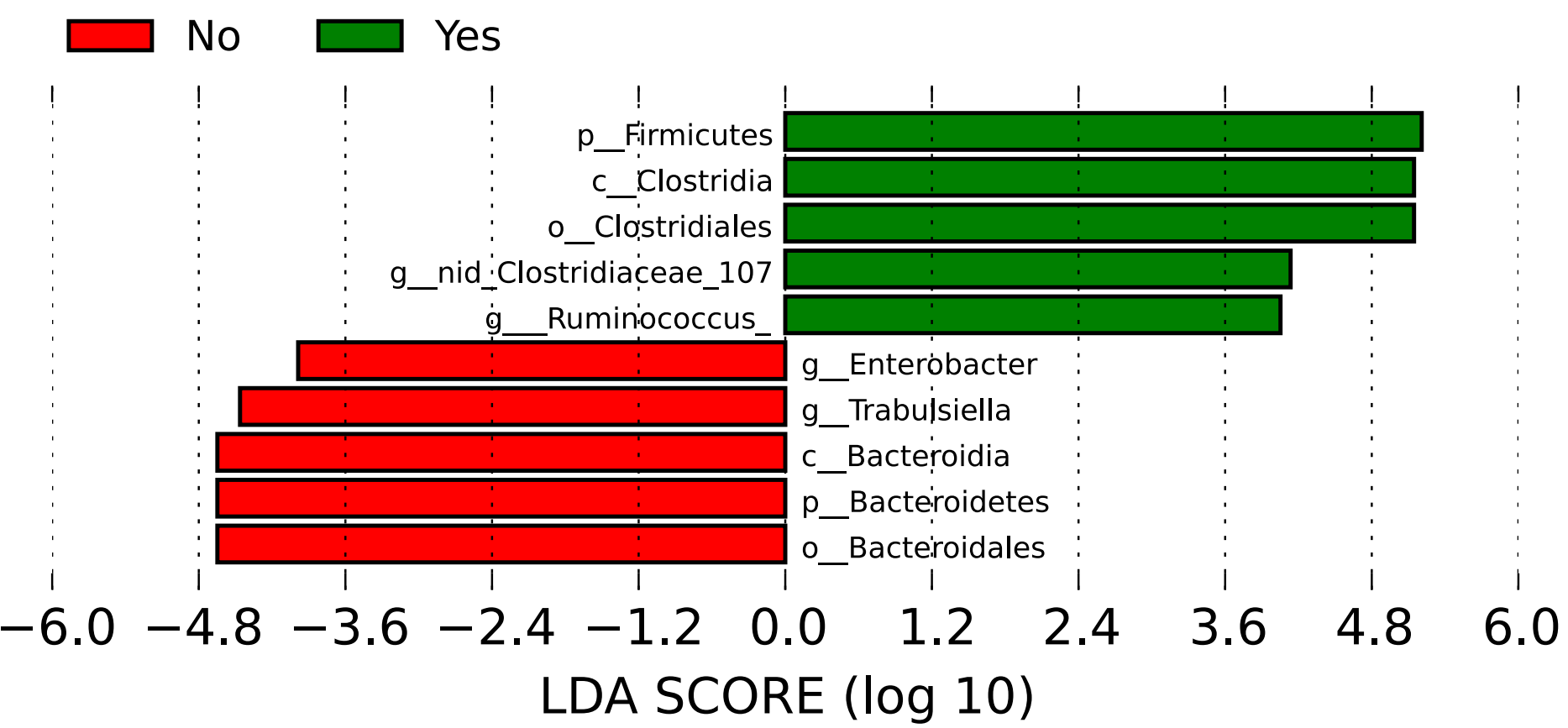
Antibiotics



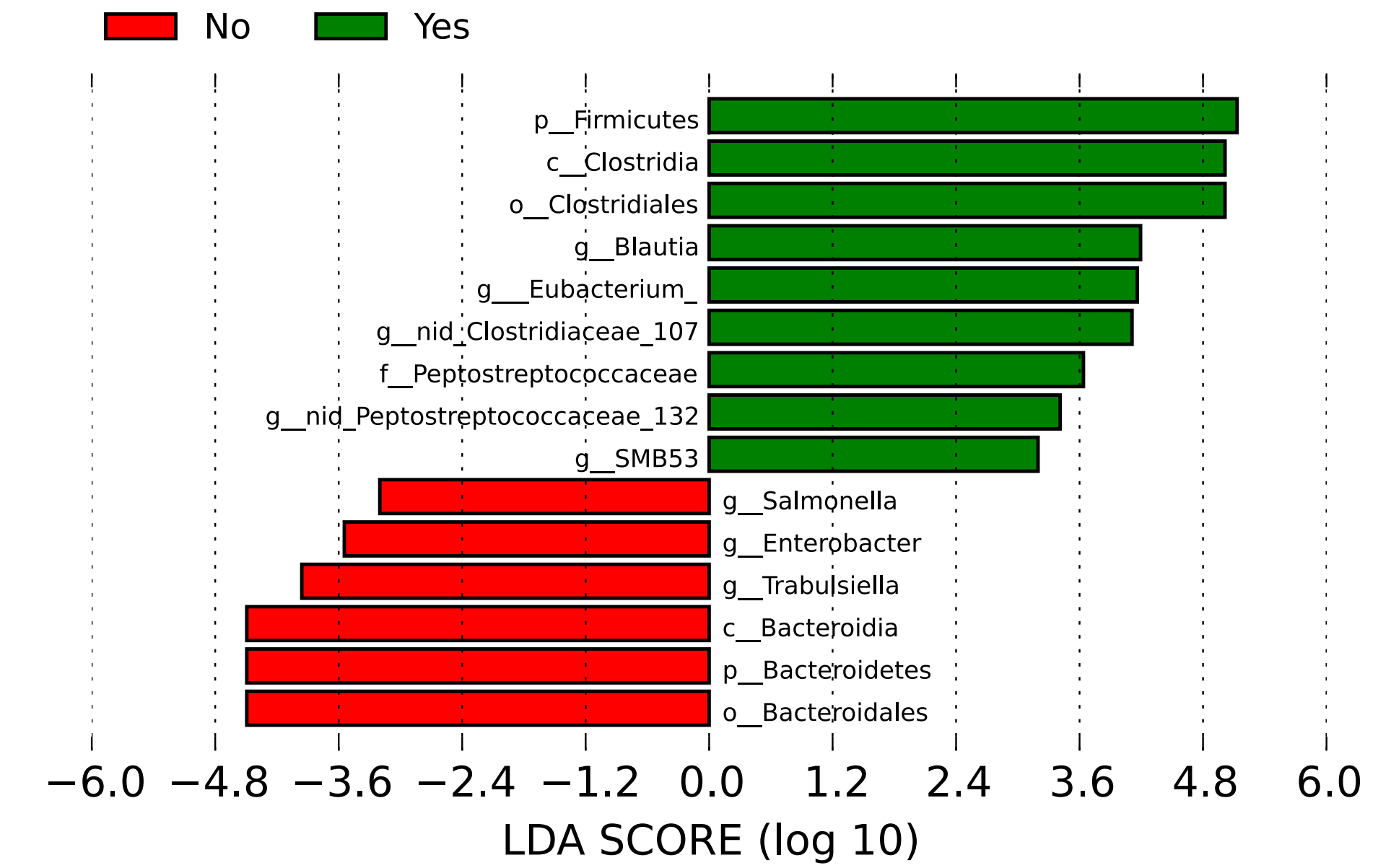
Breastfeeding



C-section



Race



Solid food