

Supplementary Information

Epigenome-Wide Association Studies of the Fractional Exhaled Nitric Oxide and Bronchodilator Drug Response in Moderate-to-Severe Pediatric Asthma

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Supplementary Table S1. Clinical and demographic characteristics of the individuals with nasal samples.

Characteristics	n	BDR	n	FeNO
Sex (male)	52	34 (65.4)	48	32 (66.7)
Age (years)	52	12.1 (10.4–14)	48	12 (10.4–14)
Ancestry	51		47	
African		2 (3.9)		2 (4.3)
Asian		1 (2)		1 (2.1)
European		43 (84.3)		39 (83)
Latin		3 (5.9)		3 (6.3)
Mixed/other		2 (3.9)		2 (4.3)
Body mass index (z-score)	52	0.6 (-0.3–1.3)	48	0.6 (-0.4–1)
Uncontrolled asthma	52	44 (84.6)	48	41 (85.4)
pre-FEV ₁ (predicted %)	52	91.7 (81–101.7)	48	91.6 (80.9–103.2)
pre-FVC (predicted %)	52	99.8 (91.7–107.8)	48	100.3 (91.6–108.9)
BDR (%)	52	6 (0.6–13)	48	5.2 (0.5–12.3)
FeNO (ppb)	51	13.3 (10.4–45.7)	48	14.8 (10.4–45.7)
SABAs	46	40 (87)	42	36 (90)
LABAs	46	44 (95.7)	42	40 (95.2)
ICS	46	46 (100)	42	42 (100)
LTRA	46	12 (26.1)	42	12 (28.6)
OCS	46	1 (2.2)	42	1 (2.4)
Biological therapy [†]	46	5 (10.9)	42	5 (11.9)

Categorical variables are described as counts (percentage) and continuous variables as median (interquartile range). [†] Biological therapy: Mepolizumab or Omalizumab intake. Abbreviations: FEV₁: Forced Expiratory Volume in the first second; FVC: forced vital capacity; BDR: Bronchodilator drug response; FeNO: Fractional exhaled nitric oxide; SABA: Short-Acting Beta-Agonists; LABAs: Long-Acting Beta-Agonists; ICS: Inhaled corticosteroids; LTRA: Leukotriene Receptor Antagonists; OCS: Oral corticosteroids.

Supplementary Table S2. Summary of the quality control.

Tissue	Blood	Nasal
CpG sites		
Initial number	865,918	866,297
Low quality probes	21,683	46,843
Probes within the sexual chromosomes (X and Y)	18,648	17,893
Multimodal probes	12,168	85,589
Cross-reactive/non-specific probes	40,870	43,254
SNP at the CpG site	24,007	24,208
SNP at single base extension from a CpG site	11,552	11,900
Probes capturing SNPs	55	55
Remaining number of CpGs available for analysis	773,260	636,555
Quality control of individuals		
Initial number	130	62
Outlier or bad quality samples	1	3
Individuals with sex discordance	1	1
Individual with mixed genotype distributions on the control SNP probes	0	2
Related individuals	4	1
Total remaining samples	124	55
Individuals with BDR and methylation data	121	55
Individuals with FeNO and methylation data	109	50

Abbreviations: SNP: Single Nucleotide Polymorphism; BDR: Bronchodilator Drug Response; FeNO: Fractional exhaled Nitric Oxide.

Supplementary Table S3. Sensitivity analysis for the main EWAS findings.

CpG	Chr	Position ^t	Gene	Main Analysis*			Adjusted by ICS dosage*			Adjusted by recruiting center*			Only Europeans*						
				n	Coef	SE	P	n	Coef	SE	P	n	Coef	SE	P	n	Coef	SE	P
FeNO																			
cg12835256	4	110651671	<i>PLA2G12A</i>	109	0.002	2.53x10 ⁻⁹	0.002	92	-0.015	0.002	2.22x10 ⁻⁸	109	-0.015	0.002	1.32x10 ⁻⁸	88	-0.019	0.003	6.73x10 ⁻¹⁰
cg19644580	21	19166676	<i>C21orf91</i>		0.001	1.29x10 ⁻⁷	0.05		0.006	0.001	7.93x10 ⁻⁷		0.006	0.001	3.72x10 ⁻⁷		0.003	0.001	6.96x10 ⁻³
BDR																			
cg06975120	9	139606856	<i>FAM69B</i>		0.006	3.86x10 ⁻⁷	0.01		-0.037	0.007	6.13x10 ⁻⁷		-0.019	0.004	1.89x10 ⁻⁶		-0.037	0.007	2.02x10 ⁻⁷
cg14985321	14	20823915	<i>PARP2</i>	122	0.002	3.71x10 ⁻⁷	0.01	103	-0.010	0.002	1.28x10 ⁻⁵	122	-0.010	0.002	6.02x10 ⁻⁶	97	-0.011	0.002	1.50x10 ⁻⁶
cg26203256	10	111756055	<i>ADD3-AS1</i>		0.004	1.85x10 ⁻⁷	0.01		-0.021	0.004	3.28x10 ⁻⁷		-0.037	0.006	9.06x10 ⁻⁸		-0.020	0.004	3.15x10 ⁻⁶

^tPosition based on GRCh37/hg19 build. *All models were also adjusted by age, sex, ethnicity, and tissue heterogeneity. **Abbreviations.** Coef: Coefficient expressed as log2(fold-change); SE: Standard error.

Supplementary Table S4. Evaluation of the effect of CpGs in nasal epithelial cells.

CpG	Chromosome	Position	Gene	Coef	SE	p-value
Bronchodilator drug response						
cg26203256	10	111756055	<i>ADD3-AS1</i>	NA	NA	NA
cg14985321	14	20823915	<i>PARP2</i>	0.0001	0.004	0.984
cg06975120	9	139606856	<i>FAM69B</i>	0.005	0.009	0.564
Fractional exhaled nitric oxide						
cg12835256	4	110651671	<i>PLA2G12A</i>	-0.015	0.007	0.045
cg19644580	21	19166676	<i>C21orf91</i>	0.006	-0.007	0.406

+Position based on GRCh37/hg19 build. **Abbreviations.** Coef: Coefficient expressed as log2(fold-change); SE: Standard error. NA: not available.

Supplementary Table S5. Summary results of sensitivity analyses stratifying by sex and age.

CpG	Chr	Pos	Gene	All individuals					<13 years old					≥13 years old					Males only			Females only		
				n	logFC	SE	P	FDR	n	logFC	SE	P	n	logFC	SE	P	n	logFC	SE	P	n	logFC	SE	P
FeNO																								
cg12835256	4	110651671	<i>PLA2G12A</i>	109	-0.015	0.002	2.53x10 ⁻⁹	0.002	64	-0.016	0.003	7.30x10 ⁻⁶	45	-0.013	0.004	0.003	67	-0.014	0.004	0.001	42	-0.016	0.003	1.86x10 ⁻⁵
cg19644580	21	19166676	<i>C21orf91</i>		0.006	0.001	1.29x10 ⁻⁷	0.050		0.006	0.002	4.09x10 ⁻⁴		0.004	0.001	0.016		0.001	0.001	0.363		0.008	0.002	8.37x10 ⁻⁵
BDR																								
cg06975120	9	139606856	<i>FAM69B</i>		-0.034	0.006	3.86x10 ⁻⁷	0.099		-0.015	0.009	0.116		-0.053	0.009	9.13x10 ⁻⁷		-0.017	0.007	0.018		-0.082	0.012	2.66x10 ⁻⁸
cg14985321	14	20823915	<i>PARP2</i>	122	-0.010	0.002	3.71x10 ⁻⁷	0.099	71	-0.010	0.003	0.002	51	-0.014	0.003	1.18x10 ⁻⁴	76	-0.011	0.003	1.07x10 ⁻⁴	46	-0.012	0.004	0.009
cg26203256	10	111756055	<i>ADD3-AS1</i>		-0.020	0.004	1.86x10 ⁻⁷	0.099		-0.024	0.005	3.14x10 ⁻⁵		-0.017	0.006	0.008		-0.021	0.005	3.23x10 ⁻⁴		-0.022	0.006	4.92x10 ⁻⁴

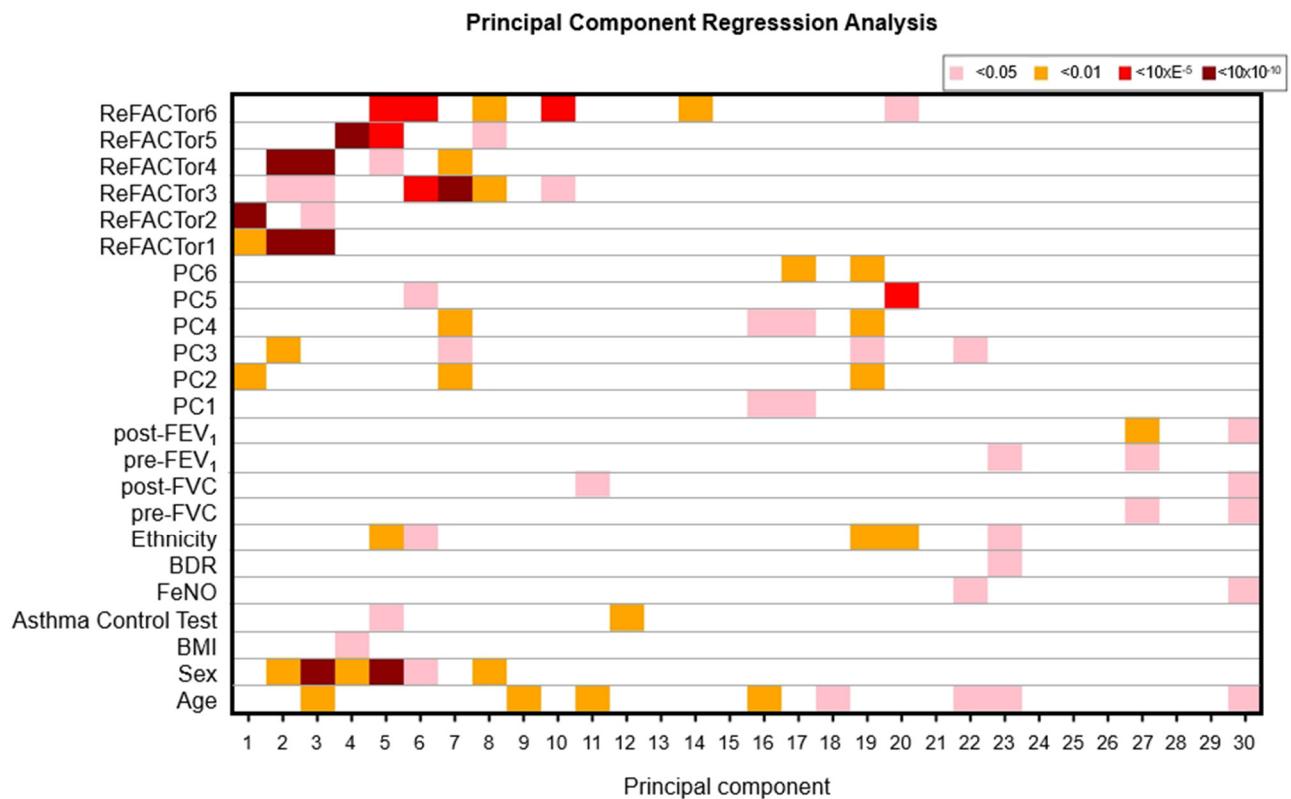
Abbreviations: logFC: log₂(fold-change); SE: standard error; FDR: false discovery rate.

Supplementary Table S6. Summary results of the gene-set enrichment analysis for the EWAS of BDR.

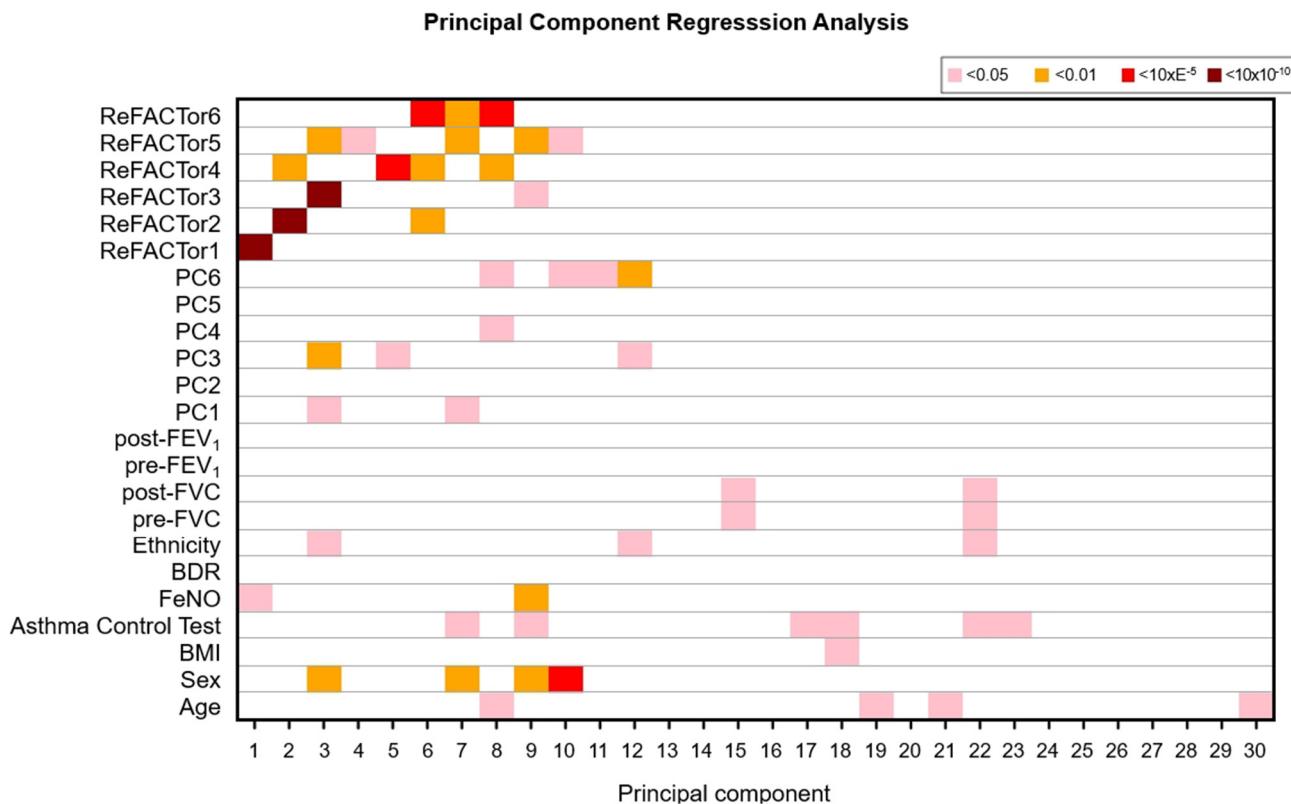
Term	<i>q</i> -value	Odds Ratio	Genes
Pathways			
WikiPathway 2021 Human			
IL-5 signaling pathway WP127	0.011	23	<i>LYN, SYK, KRAS, SOS1</i>
KEGG 2021 Human			
C-type lectin receptor signaling pathway	0.031	10.5	<i>SYK, PRKCD, BCL3, CBLB, KRAS</i>
Fc epsilon RI signaling pathway	0.033	12.9	<i>LYN, SYK, KRAS, SOS1</i>
B cell receptor signaling pathway	0.043	10.7	<i>LYN, SYK, KRAS, SOS1</i>
Biocarta 2016			
Fc Epsilon Receptor I Signaling in Mast Cells <i>Homo sapiens</i>	0.017	23.6	<i>LYN, SYK, SOS1</i>
BCR Signaling Pathway <i>Homo sapiens</i>	0.017	21.9	<i>LYN, SYK, SOS1</i>
IL 2 signaling pathway <i>Homo sapiens</i>	0.042	33.8	<i>SYK, SOS1</i>
Telomeres, Telomerase, Cellular Aging, and Immortality <i>Homo sapiens</i>	0.042	31.2	<i>XRCC6, KRAS</i>
Ontologies			
GO Biological Process 2021			
Positive regulation of protein dephosphorylation (GO:0035307)	0.031	27.6	<i>MAGI2, PRKCD, CAMTA1, DUSP26</i>

Abbreviations: GO: Gene Ontology.

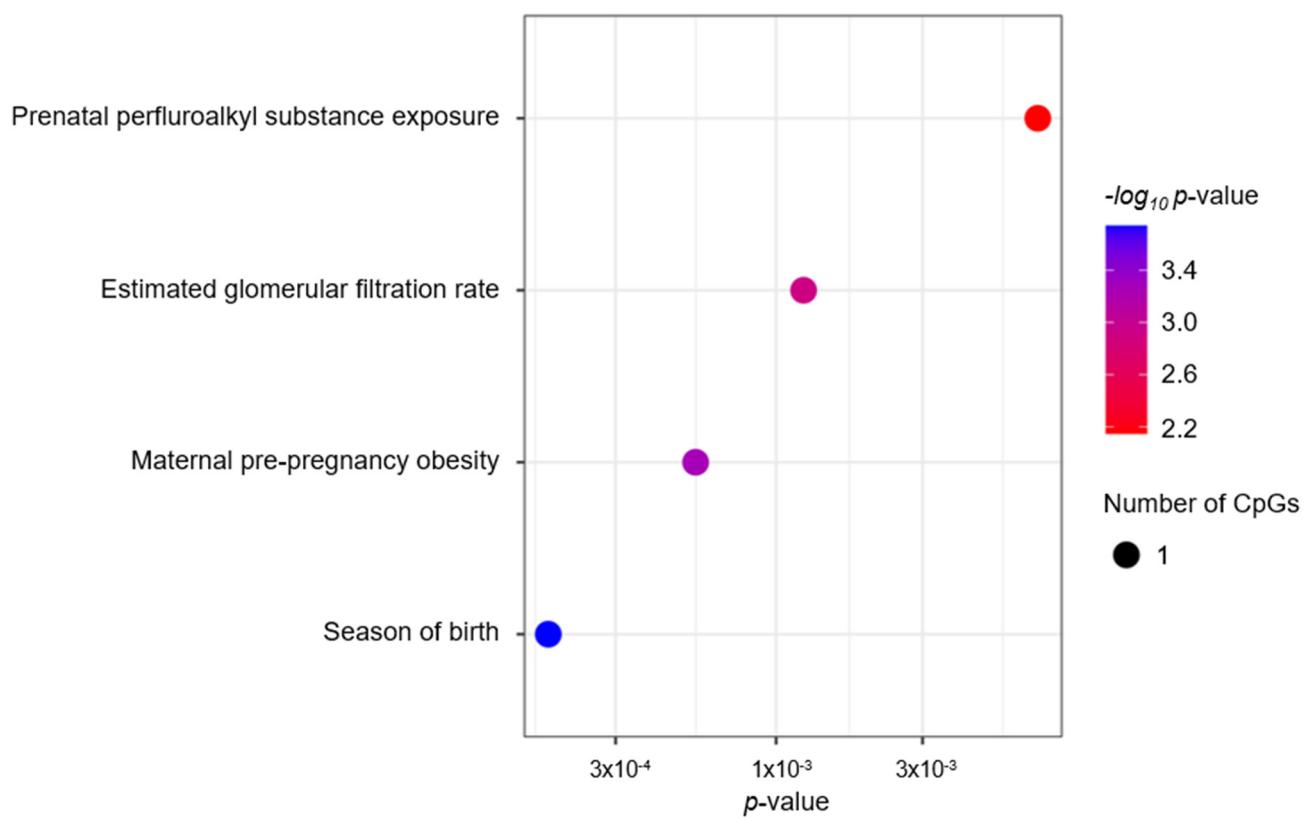
SUPPLEMENTARY FIGURES



Supplementary Figure S1. Principal component regression analysis plot of blood samples. Top principal components are displayed along the X-axis, and covariates are displayed on the Y-axis. It shows with colors different levels of association p-values between principal components (PCs) and covariates. **Abbreviations:** PC: Principal component of genotype data; FEV₁: Forced expiratory volume in the first second; FVC: Forced vital capacity; FeNO: Fractional exhaled nitric oxide; BDR: Bronchodilator drug response; BMI: Body mass index.



Supplementary Figure S2. Principal component regression analysis plot of nasal samples. Top principal components are displayed along the X-axis, and covariates are displayed on the Y-axis. It shows with colors different levels of association p-values between principal components (PCs) and covariates. **Abbreviations:** PC: Principal component of genotype data; FEV₁: Forced expiratory volume in the first second; FVC: Forced vital capacity; FeNO: Fractional exhaled nitric oxide; BDR: Bronchodilator drug response; BMI: Body mass index.



Supplementary Figure S3. Plot summarizing the trait-enrichment analysis for the top 100 CpGs for BDR in EWAS Atlas. *p*-values are displayed on the x-axis and traits on the y-axis. Blue shows the strongest associations, and the size of each point indicates the number of CpGs it correlates with.