

University of Dundee

New genetic signals for lung function highlight pathways and chronic obstructive pulmonary disease associations across multiple ancestries

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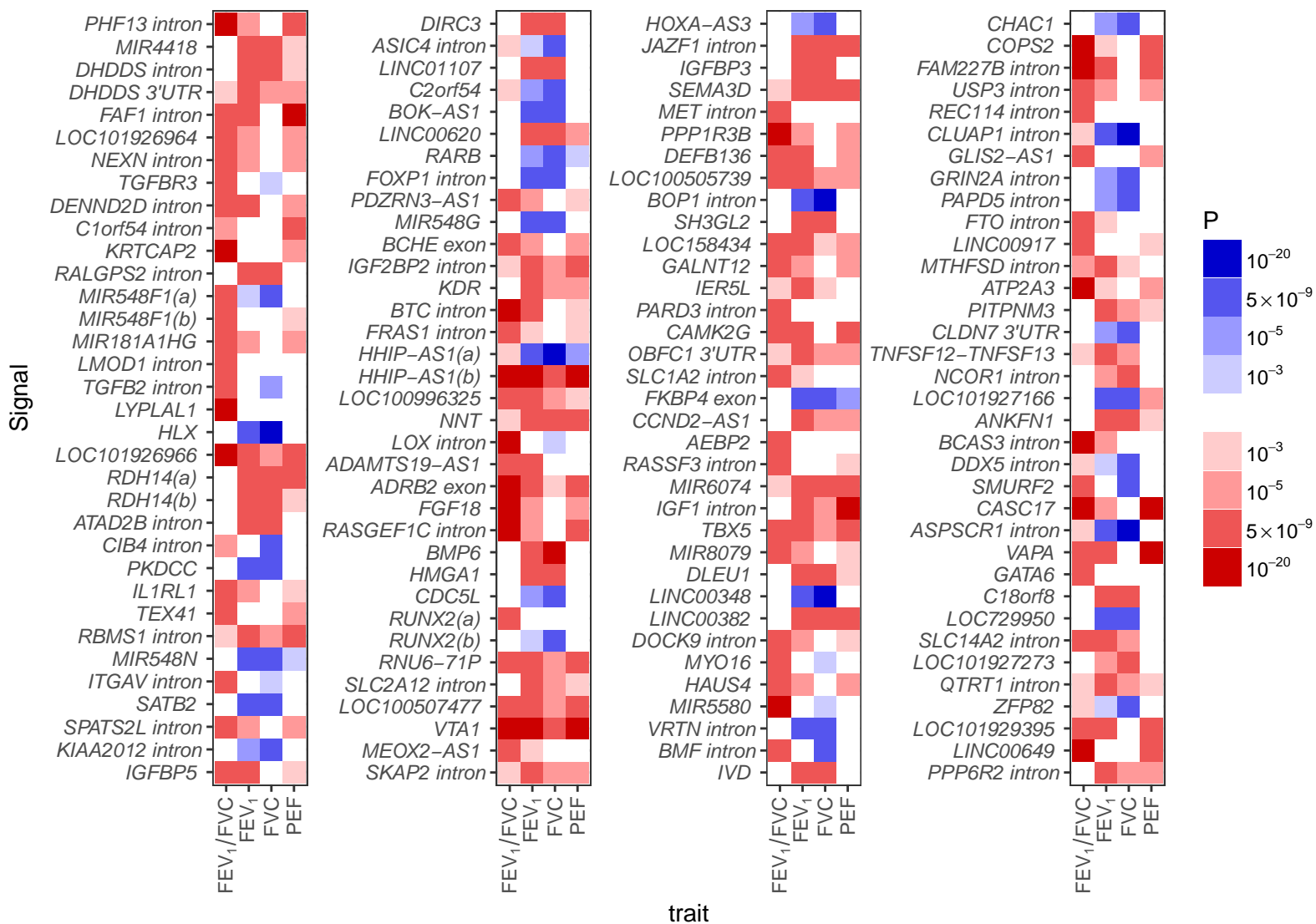
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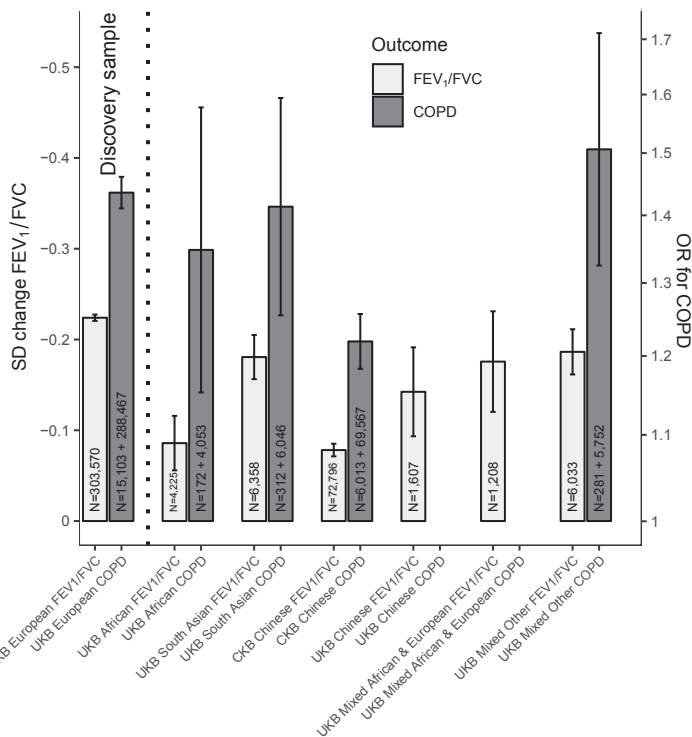
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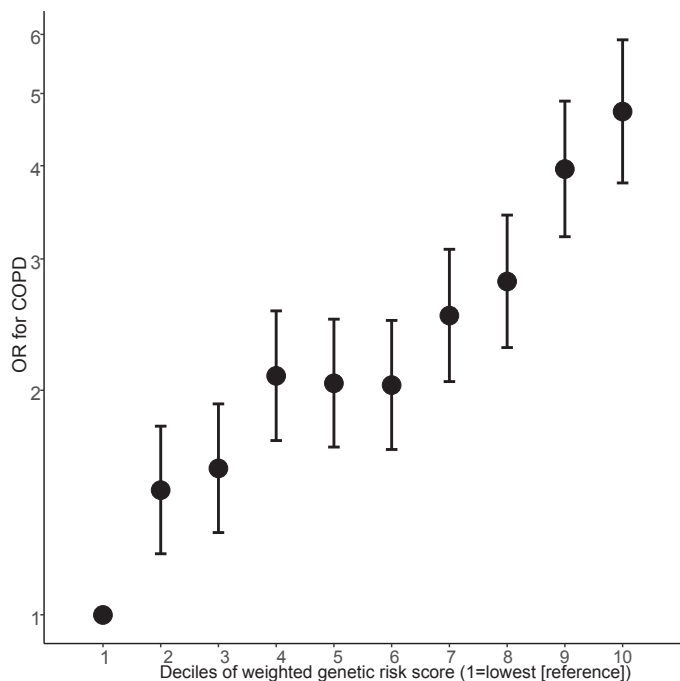


a Weighted risk score associations with FEV₁/FVC and COPD in population-based studies



Ancestral group and phenotype studied in UK Biobank or China Kadoorie Biobank

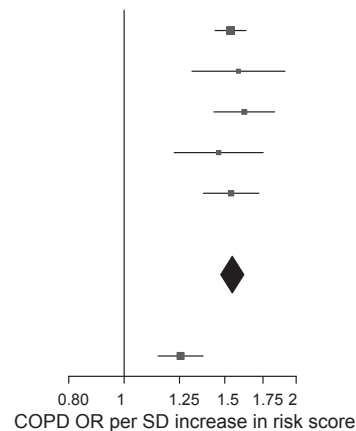
c Odds ratio of COPD per decile increase in the weighted genetic risk score

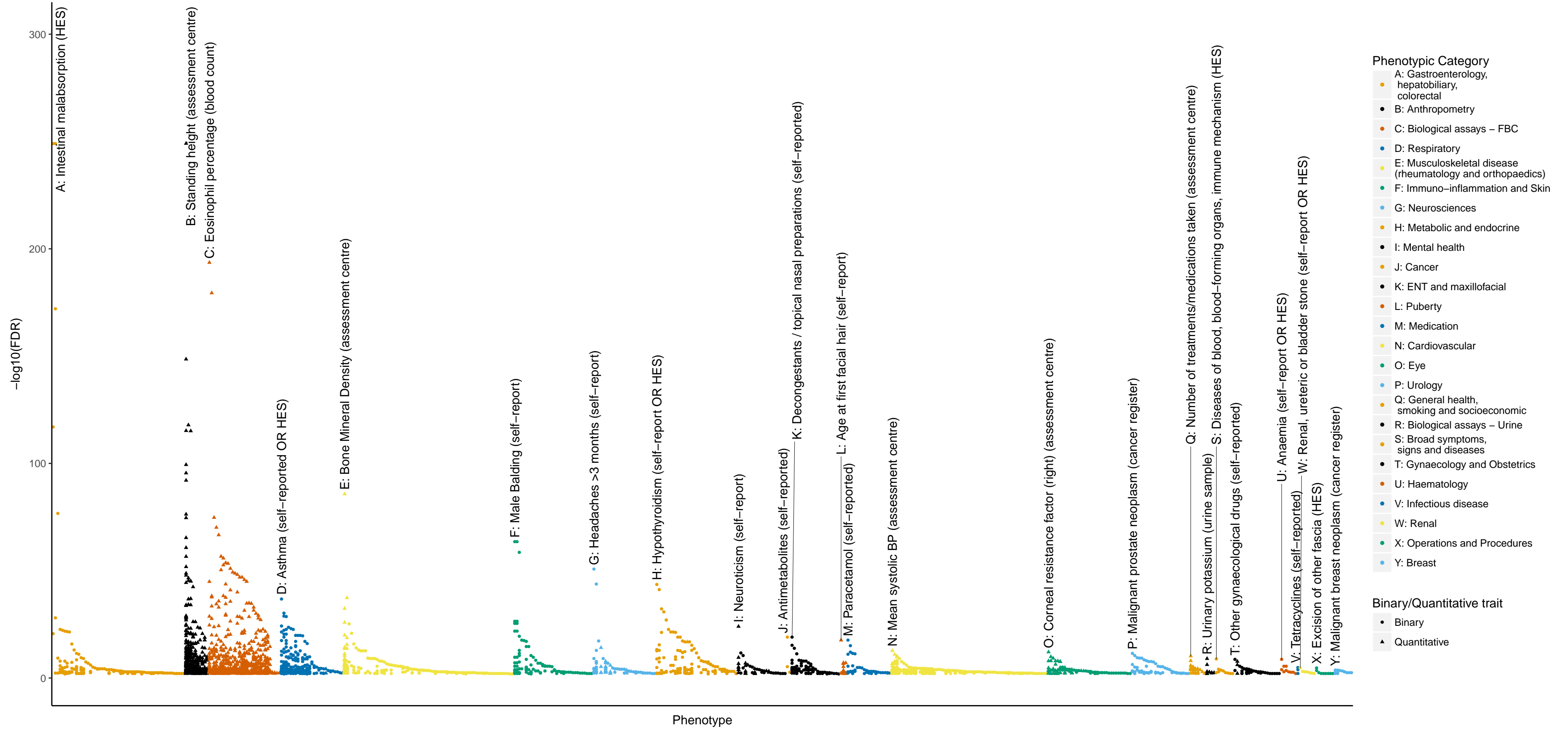


b Weighted risk score associations with COPD susceptibility in COPD case-control studies

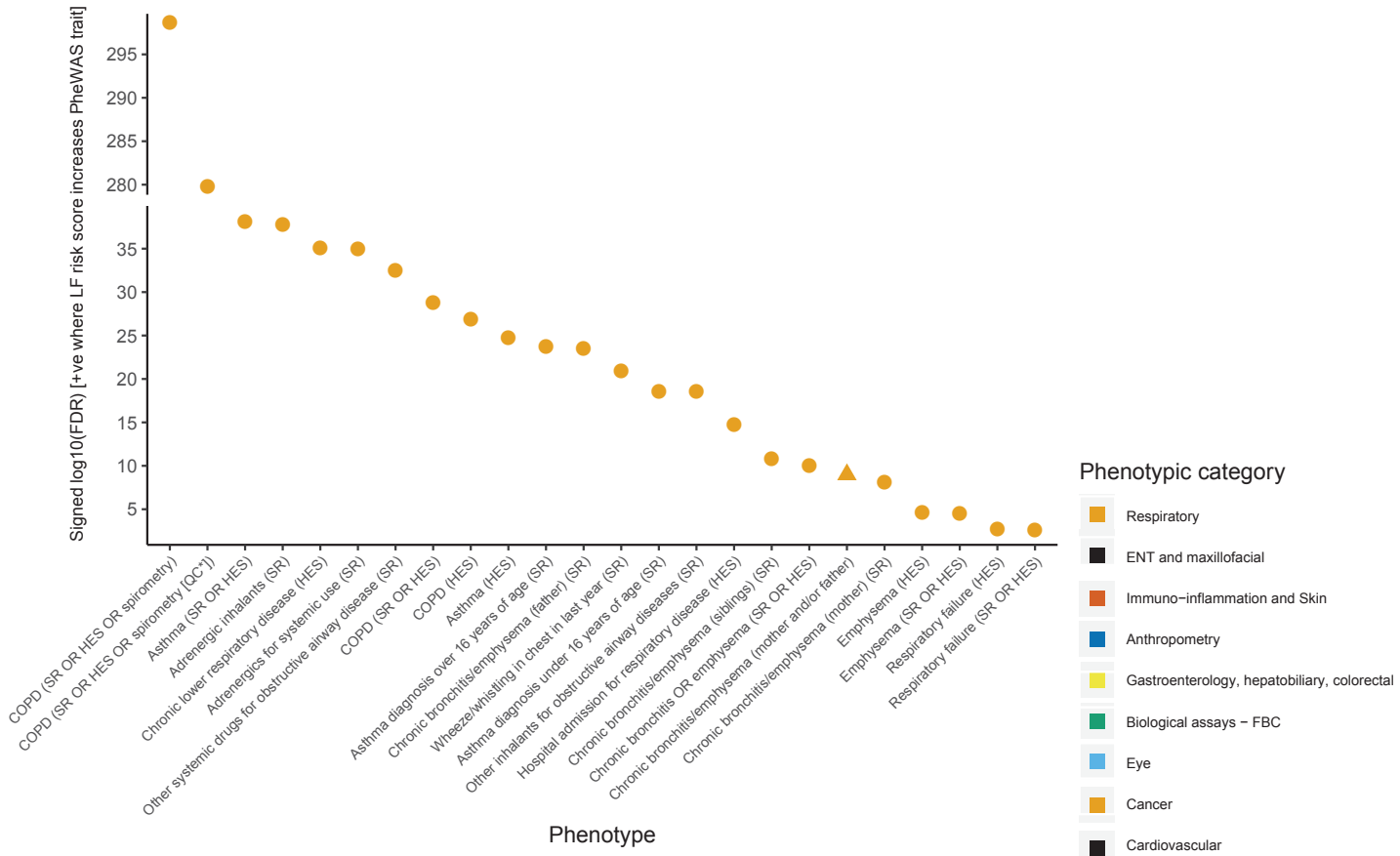
Ancestry	Cohort	OR	95%LCI	95%UCI	P	Cases	Controls
European	COPDGene (EUR)	1.54	1.44	1.63	1.97x10 ⁻⁴¹	3068	2110
	ECLIPSE	1.59	1.31	1.91	1.42x10 ⁻⁰⁶	1713	147
	GenKOLS	1.62	1.44	1.83	8.99x10 ⁻¹⁵	836	692
	NETT-NAS	1.46	1.22	1.75	3.13x10 ⁻⁰⁵	374	429
	SPIROMICS	1.54	1.38	1.72	4.47x10 ⁻¹⁴	988	537
Meta-analysis		1.55	1.48	1.62	1.48x10⁻⁷⁵	6979	3915

African	COPDGene (AFR)	1.26	1.15	1.37	8.36x10 ⁻⁰⁷	910	1556
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a



b

