SNP (control)	Gene	Metabolite	P_{ref}	Spectrum	Cohort	P_C	Rank
rs37369	AGXT2	3-aminoisobutyrate	3.2×10^{-75}	multi-peak		1.0×10^{-46}	
rs8101881	SLC7A9	lysine	1.0×10^{-8}	multi-peak	SHIP	1.5×10^{-15}	1
rs17279437	SLC6A20	dimethylglycine	5.5×10^{-10}	multi-peak	SHIP	4.6×10^{-10}	3
rs11884776	ALMS1	N-acetylated compounds	4.1×10^{-19}	multi-peak	MolPAGE	1.0×10^{-123}	3, 14, 21
rs4539242	PYROXD2	trimethylamine	2.8×10^{-23}	single peak	MolPAGE	1.7×10^{-128}	44
rs830124	<i>PSMD9</i>	2-hydroxyisobutyrate	1.6×10^{-15}	single peak	SHIP	6.6×10^{-11}	65
rs1495743	NAT2	formate	1.4×10^{-10}	single peak	SHIP	6.0×10^{-7}	222 (9)