

Supplementary Materials: Potential Metabolic Biomarkers to Identify Interstitial Lung Abnormalities

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Table S1. Identified pathways associated with common metabolites.

<i>n</i>	Canonical Pathways	$-\log(p\text{-Value})$	Molecules
1	Phospholipases	5.52E+00	1-Acylglycerophosphocholine, phosphatidic acid, phosphatidylcholine
2	Triacylglycerol Biosynthesis	5.26E+00	1-Acylglycerophosphocholine, phosphatidic acid, phosphatidylcholine
3	RhoA Signaling	4.95E+00	Phosphatidic acid, phosphatidylcholine
4	p70S6K Signaling	3.78E+00	Phosphatidic acid, phosphatidylcholine
5	mTOR Signaling	3.78E+00	Phosphatidic acid, phosphatidylcholine
6	Phospholipase C Signaling	3.51E+00	Phosphatidic acid, phosphatidylcholine
7	Gαq Signaling	3.51E+00	Phosphatidic acid, phosphatidylcholine
8	Choline Biosynthesis III	3.40E+00	Phosphatidic acid, phosphatidylcholine
9	Endothelin-1 Signaling	2.94E+00	Phosphatidic acid, phosphatidylcholine
10	Phosphatidylethanolamine Biosynthesis	1.83E+00	Phosphatidylethanolamine
11	Choline Degradation I	1.73E+00	Betaine aldehyde