Tool	Result	Ontology	FDR
Metacore	lipid transporter activity	GOFunction	0.000392
Metacore	lipid transport	GOProcess	0.000786
Metacore	lipid localization	GOProcess	0.001043
Metacore	regulation of lipid metabolic process	GOProcess	0.003908
Metacore	Bile acid regulation of lipid metabolism*	Networks	0.006123
IPA	Homeostasis of lipid	DisBio	4.42E-15
IPA	Concentration of triacylglycerol	DisBio	3.61E-14
IPA	Concentration of lipid	DisBio	7.28E-13
IPA	Transport of lipid	DisBio	4.06E-12
IPA	Fatty acid metabolism	DisBio	2.12E-10

*Regulation of metabolism_Bile acid regulation of lipid metabolism and negative FXR-dependent regulation of bile acids concentration