

Biofiles

Volume 5, Number 7

SIGMA-ALDRICH

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Life Science

Lipid Metabolism



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Coming Next Issue: Cell Based Assay

Life scientists have long realized the value in analyzing biological processes based on its most fundamental unit the cell. This issue will touch on a wide variety of topics ranging from ADMETox to stem cell analysis to the most basic methods of analyzing and staining cells. The issue will announce the addition of 1400 more ECACC cell lines available in the US, Canada and Mexico.

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Introduction

Roland Wohlgemuth

Senior Product Specialist

roland.wohlgemuth@sial.com



After sequencing the human genome, life scientists are now in the era “beyond the genome” and confronting the challenges of bringing this information to life. For lipid researchers, unraveling the complexities of lipid metabolism will bring answers and opportunities in fields as diverse as medicine and biofuels. Today, lipid research addresses all aspects of cell biology including cell structure, energy storage and generation, and cell signaling. The synergies between classical lipid research, systems biology, and the advancement of lipid detection technologies have led to a revolution in lipid biology and the emergence of lipidomics as a branch of metabolomics.

Lipid-based diseases are a growing and expensive challenge to health care systems. As a population ages, chronic conditions associated with aging such as cardiovascular disease, neurodegenerative disorders, and metabolic disorders take increasing tolls in terms of morbidity and mortality. Oxidation of lipids and lipid

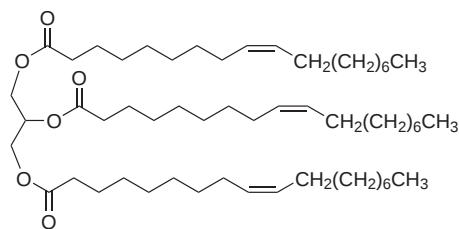
metabolites has been linked to disorders of aging like osteoporosis and vascular calcification. Additionally, research is now trying to explain how dysregulations in lipid metabolism may underlie diseases such as Alzheimer’s, cancer, and asthma. Lipid synthesis pathways, such as the methyl-D-erythritol 4-phosphate (MEP) pathway of isoprenoid synthesis, are being investigated as potential targets for antibacterial therapies and drug targets. As changes in lipid profiles can mark developmental stages or more ominously, pathological states, there is great potential for the use of lipids as biomarkers. Experimental quantification of the levels of lipids and lipid metabolites is therefore essential for enhancing the understanding of different influences such as genetic, nutritional, behavioral, and environmental factors. As part of our commitment to metabolomics research, Sigma offers an expanding range of lipids and lipid metabolites for the mapping and study of healthy and diseased cells.



Natural Abundance Lipid Metabolites

Natural Abundance Lipid Metabolites

Glycerolipid Metabolism

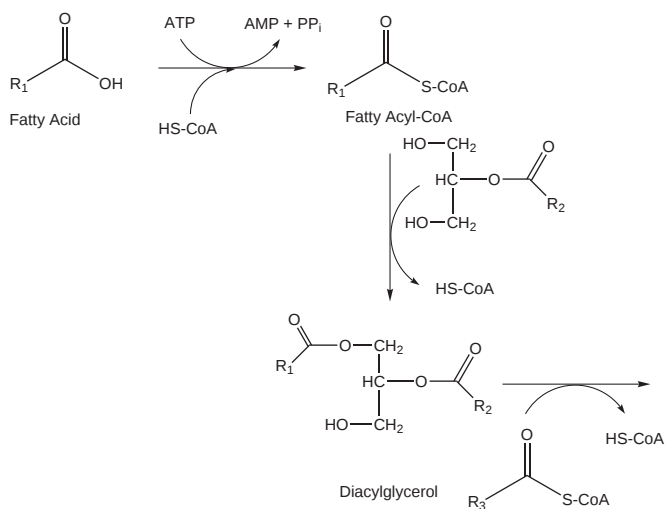


Glycerol esterified with one, two, or three fatty acids make up monoacyl-, diacyl- and triacylglycerols, with a chiral center at carbon-2 of the glycerol moiety. Fats and oils from plants and animals are triacylglycerols, while diacylglycerols are intermediates and cellular messengers, and monoacylglycerols, formed by hydrolysis, are surfactants and intermediates. Because triacylglycerols are insoluble in water, combination or emulsification with other lipids, cellular compounds, or proteins is required before transport and metabolism can occur. Complete or partial lipase-catalyzed hydrolysis yields monoacylglycerols, glycerol, and fatty acids that can be transported and utilized for energy production or biosynthetic pathways of metabolism.

Biosynthesis of triacylglycerols is achieved in a three-step sequence from 2-monoacylglycerols and fatty acids. First, the fatty acid is activated by acyl-CoA synthetase catalyzed conversion to the corresponding fatty acyl thioester with coenzyme A. The fatty acyl-CoA is then coupled with a 2-monoacylglycerol by the catalytic action of a monoacylglycerol transferase to yield a diacylglycerol. The final triacylglycerol is obtained by coupling of fatty acyl-CoA with diacylglycerol through the action of diacylglycerol transferase.

Name	Structure	Cat. No.
Acetylcholine chloride		A6625-10MG A6625-25G A6625-100G A6625-500G
Choline chloride		C7017-10MG C7017-5G C7017-25G C7017-100G
Choline chloride		26978-25G 26978-100G

Name	Structure	Cat. No.
1-Deoxy-D-xylulose-5-phosphate sodium salt		13368-1MG 13368-5MG
1,3-Dihydroxyacetone dimer		D107204-5G D107204-100G D107204-500G
Dihydroxyacetone phosphate dilithium salt		D7137-5MG D7137-10MG D7137-25MG D7137-100MG D7137-250MG
1,2-Dimyristoyl- <i>sn</i> -glycero-3-phosphate monosodium salt		P3650-25MG P3650-50MG P3650-250MG
1,2-Dioctanoyl- <i>sn</i> -glycero-3-phosphate sodium salt	-	P3591-50MG
1,2-Dipalmitoyl- <i>sn</i> -glycerol		D9135-25MG D9135-100MG
1,2-Dipalmitoyl- <i>sn</i> -glycero-3-phosphate sodium salt		P4013-25MG P4013-100MG
D-(+)-Glyceraldehyde, viscous		49800-1G 49800-5G
D-Glyceric acid calcium salt dihydrate		367494-1G 367494-5G
Glycerol, anhydrous		49767-100ML 49767-250ML 49767-1L
<i>rac</i> -Glycerol 1-myristate		M1890-100MG M1890-1G
<i>rac</i> -Glycerol 1-phosphate disodium salt hexahydrate		G2138-10MG G2138-5G G2138-25G G2138-100G G2138-500G
<i>sn</i> -Glycerol 3-phosphate bis(cyclohexylammonium) salt		G7886-250MG G7886-1G G7886-5G



Caption Text?

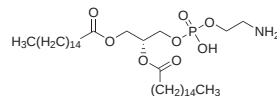
Name	Structure	Cat. No.
Glyceryl 1,3-dipalmitate		D1639-1G
Glyceryl 1,3-distearate	-	D8269-10MG D8269-100MG
Glyceryl triacetate		T5376-500ML T5376-1L
Glyceryl tributyrat		T8626-25ML T8626-100ML
Glyceryl tridecanoate		T7517-1G T7517-5G
Glyceryl tridodecanoate		T4891-100MG T4891-5G
Glyceryl trielaidate	-	T7379-1G
Glyceryl triheptadecanoate	-	T2151-100MG T2151-1G
Glyceryl trilinoleate		T9517-50MG T9517-100MG T9517-500MG T9517-1G T9517-5G
Glyceryl trilinolenate		T6513-100MG

Name	Structure	Cat. No.
Glyceryl trimyristate		T5141-1G T5141-5G
Glyceryl trinonadecanoate	-	T4632-100MG T4632-1G
Glyceryl trioctanoate		T9126-100ML T9126-500ML T9126-1L
Glyceryl trioleate		T7140-500MG T7140-1G T7140-5G T7140-10G T7140-50G
Glyceryl trioleate		92860-5ML 92860-10ML
Glyceryl tripalmitate		T5888-100MG T5888-500MG T5888-1G T5888-5G
Glyceryl tripalmitelaidate	-	T5909-5MG
Glyceryl tripalmitoleate	-	T2630-100MG T2630-1G
Glyceryl tristearate		T5016-5G T5016-25G
Glyceryl tritricosanoate	-	T1412-25MG

Name	Structure	Cat. No.
Glyceryl tritridecanoate	-	T3882-500MG T3882-1G
Glyceryl triundecanoate		T5534-1G
1-Lauroyl- <i>rac</i> -glycerol		M1765-100MG M1765-500MG M1765-1G M1765-5G
1-Linoleoyl- <i>rac</i> -glycerol		M7640-100MG M7640-1G
Lipoteichoic acid from <i>Bacillus subtilis</i>	-	L3265-5MG L3265-25MG
Lipoteichoic acid from <i>Staphylococcus aureus</i>	-	L2515-5MG L2515-10MG L2515-25MG
Lipoteichoic acid from <i>Streptococcus faecalis</i>	-	L4015-5MG L4015-25MG
Lipoteichoic acid from <i>Streptococcus pyogenes</i>	-	L3140-5MG L3140-10MG L3140-25MG
1-Monopalmitoleoyl- <i>rac</i> -glycerol	-	M7890-100MG M7890-1G
1-Oleoyl- <i>rac</i> -glycerol		M7765-25MG M7765-50MG M7765-100MG M7765-1G
Oleoyl-L- α -lysophosphatidic acid sodium salt		L7260-1MG L7260-5MG L7260-25MG
Oleoyl-L- α -lysophosphatidic acid sodium salt		75347
DL- α -Palmitin		M1640-100MG M1640-1G
3-Palmitoyl- <i>sn</i> -glycerol		76184-250MG 76184-1G
3- <i>sn</i> -Phosphatidic acid sodium salt from egg yolk lecithin		P9511-5MG P9511-10MG P9511-50MG P9511-100MG P9511-500MG
Phosphocholine chloride calcium salt tetrahydrate		P0378-10MG P0378-5G P0378-25G P0378-50G
D-(-)-3-Phosphoglyceric acid disodium salt		P8877-10MG P8877-1G P8877-5G

Name	Structure	Cat. No.
O-Phosphorylethanol-amine		P0503-10MG P0503-1G P0503-5G P0503-25G P0503-100G P0503-500G
1,2-Propanediol		12279-1ML-F 12279-5ML-F
1,3-Propanediol		81780-500ML
1-Propanol		96566-5ML-F 96566-10ML-F
Propionaldehyde		538124-250ML 538124-1L
1-Stearoyl- <i>rac</i> -glycerol		M2015-100MG M2015-1G M2015-5G
Tripentadecanoin	-	T4257-100MG T4257-500MG T4257-1G
Tripetroselaidin	-	T5784-25MG T5784-100MG

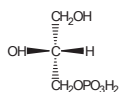
Glycerophospholipid Metabolism



Glycerol is the backbone of the fundamental phospholipids used as the self-assembling units of lipid membranes. It is interesting that both enantiomeric glycerol configurations appear in nature. In eukaryotes and eubacteria, glycerophospholipids are based on the *sn*-3 configuration of the glycerol backbone, while archaeobacteria lipids are based on the *sn*-1-configuration. The metabolic intermediate for all glycerophospholipids is phosphatidic acid (1,2-diacylglycerol-3-phosphate). Lysophosphatidic acid, which lacks the acyl chain in the 2-position, is an important cellular messenger.

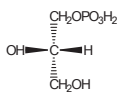
Phosphatidylcholine (lecithin) is the most abundant phospholipid in animal and plant tissues. Phosphatidylethanolamine (cephalin) is an abundant phospholipid in microbial, plant, and animal cells. The corresponding lysophosphatidic acid derivatives, lysophosphatidylcholine and lysophosphatidylethanolamine, are more soluble in water and have surfactant properties, and these are not standard components of cell membranes. Phosphatidylglycerol has important functions in lung surfactants, chloroplasts, and bacterial membranes and is a metabolic intermediate in the biosynthesis of cardiolipin. Phosphatidylserine is involved in biological processes including apoptosis, blood coagulation, and activation of protein kinase C. It can be found in plasma membranes and other membranes of animals, plants, and microorganisms.

A class of phospholipids with a high rate of metabolism is the phosphatidylinositols which have varying degrees of phosphorylation in the polar head group myo-inositol. The metabolic conversion of phosphatidylinositols to diacylglycerols and inositol phosphates is important in the regulation of vital cellular functions such as differentiation, proliferation, and apoptosis, and in anchoring proteins via a glycosyl-bridge to the plasma membrane.



sn-Glycerol 3-phosphate

[L-(glycerol 3-phosphate) = D-(glycerol 1-phosphate)]



sn-Glycerol 1-phosphate

[L-(glycerol 1-phosphate) = D-(glycerol 3-phosphate)]

Name	Structure	Cat. No.
2-Arachidonoyl-1-stearoyl-sn-glycero-3-phosphocholine solution		P1287-5MG P1287-25MG
Cardiolipin sodium salt from bovine heart	-	C0563-10MG C0563-25MG C0563-100MG C0563-500MG
Cytidine 5'-diphosphocholine sodium salt dihydrate		C0256-100MG C0256-500MG C0256-1G
1,2-Didecanoyl-sn-glycero-3-phosphocholine		P7081-25MG P7081-100MG P7081-1G
1,2-Didocosahexaenoyl-sn-glycero-3-phosphocholine		78896-100MG 78896-500MG
1,2-Didodecanoyl-sn-glycero-3-phosphocholine		P1263-25MG P1263-100MG P1263-500MG
1,2-Didodecanoyl-rac-glycero-3-phosphocholine		P1534-100MG
1,2-Dieicosapentaenoyl-sn-glycero-3-phosphocholine		40723-100MG 40723-500MG
1,2-Dierucoyl-sn-glycero-3-phosphocholine		77017

Name	Structure	Cat. No.
1,2-Dierucoyl-sn-glycero-3-phosphoethanolamine		74576
1,2-Dierucoyl-sn-glycero-3-phospho-rac-(1-glycerol) sodium salt		00259
Diethanolamine	HOCH2CH2NHCH2CH2OH	31589-100G 31589-500G
1,2-Dihexadecanoyl-rac-glycero-3-phospho-rac-(1-glycerol) ammonium salt	-	P5650-100MG
1,2-Dihexadecyl-rac-glycero-3-phosphocholine	-	P3777-25MG P3777-100MG
1,2-Di-O-hexadecyl-sn-glycero-3-phosphocholine	-	P1527-100MG P1527-500MG
1,2-Dihexadecyl-sn-glycero-3-phosphoethanolamine		37161-100MG 37161-1G
1,2-Dilinoleoyl-sn-glycero-3-phosphocholine	-	P0537-5MG P0537-25MG
1,2-Dimyrystoyl-sn-glycerol-3-(5'-diphosphocytidine) potassium salt	-	C5430-5MG C5430-25MG
1,2-Dimyrystoyl-sn-glycero-3-phosphate monosodium salt		P3650-25MG P3650-50MG P3650-250MG
1,2-Dimyrystoyl-sn-glycero-3-phosphocholine		P2663-25MG P2663-100MG P2663-500MG P2663-1G
1,2-Dimyrystoyl-rac-glycero-3-phosphocholine		P7930-100MG P7930-250MG
1,2-Dimyrystoyl-sn-glycero-3-phosphoethanolamine		P5693-100MG P5693-250MG P5693-1G
1,2-Dimyrystoyl-sn-glycero-3-phospho-rac-(1-glycerol) ammonium salt		43096-100MG 43096-1G
1,2-Dimyrystoyl-sn-glycero-3-phospho-rac-(1-glycerol) sodium salt		P6412-25MG P6412-100MG P6412-500MG

Name	Structure	Cat. No.
1,2-Dimristoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) sodium salt		90998-100MG 90998-1G
1,2-Dimristoyl- <i>sn</i> -glycero-3-phospho-L-serine sodium salt		80114-100MG 80114-500MG
1,2-Dioctadecanoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) ammonium salt		05717-100MG 05717-1G
1,2-Dioleoyl- <i>sn</i> -glycero-3-phosphocholine		P6354-25MG P6354-100MG P6354-1G
1,2-Dioleoyl- <i>sn</i> -glycero-3-phosphoethanolamine		42490-2.5ML 42490-10ML
1,2-Dioleoyl- <i>sn</i> -glycero-3-phosphoethanolamine		76548-100MG 76548-1G
1,2-Dioleoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) sodium salt		P9664-5MG P9664-25MG P9664-100MG P9664-500MG
1,2-Dioleoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) sodium salt		74384-100MG 74384-1G
1,2-Dioleoyl- <i>sn</i> -glycero-3-phosphoric acid monosodium salt		74304-100MG 74304-500MG
1,2-Dioleoyl- <i>sn</i> -glycero-3-phospho-L-serine sodium salt		P1060-5MG P1060-10MG P1060-25MG
1,2-Dioleoyl-3-linoleoyl- <i>rac</i> -glycerol		D9925-100MG
1,2-Dioleoyl-3-palmitoyl- <i>rac</i> -glycerol		D1782-10MG D1782-100MG
1,3-Dioleoyl-2-palmitoylglycerol		D1657-25MG
1,2-Dipalmitoyl- <i>sn</i> -glycero-3-phosphocholine		P0763-50MG P0763-100MG P0763-250MG P0763-1G P0763-5G
1,2-Dipalmitoyl- <i>sn</i> -glycero-3-phosphocholine		P4329-25MG P4329-100MG P4329-500MG P4329-1G

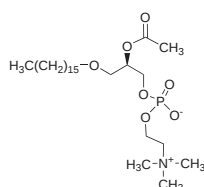
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1,2-Dipalmitoyl- <i>rac</i> -glycero-3-phosphocholine		P5911-100MG P5911-250MG P5911-1G
2,3-Dipalmitoyl- <i>sn</i> -glycero-1-phosphocholine		42566-25MG
1,2-Dipalmitoyl- <i>sn</i> -glycero-3-phosphoethanolamine		P1348-25MG P1348-100MG P1348-250MG P1348-1G
1,2-Dipalmitoyl- <i>rac</i> -glycero-3-phosphoethanolamine		P3275-100MG P3275-500MG
1,2-Dipalmitoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) sodium salt		P9789-25MG P9789-100MG
1,2-Dipalmitoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) sodium salt		50984-100MG 50984-1G
1,2-Dipalmitoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) ammonium salt		42627-100MG 42627-1G
1,2-Dipalmitoyl- <i>sn</i> -glycero-3-phospho-L-serine sodium salt		P1185-10MG P1185-50MG
1,2-Dipalmitoyl- <i>rac</i> -glycero-3-phospho-L-serine		P1902-5MG P1902-25MG
1,3-Dipalmitoyl-2-oleoylglycerol		D2157-10MG D2157-50MG D2157-100MG
1,2-Dipalmitoylphosphatidylinositol 4,5-diphosphate		P7115-100UG
1,2-Dipalmitoylphosphatidylinositol 3,4,5-trisphosphate tetrasodium salt		P4240-100UG

Name	Structure	Cat. No.
1,2-Distearoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) sodium salt		55845-100MG 55845-1G
1,2-Distearoyl- <i>sn</i> -glycero-3-phospho-L-serine sodium salt		43307-100MG 43307-500MG
Ethanolamine		398136-25ML 398136-500ML 398136-2.5L
L-α-Glycerophosphorylcholine	-	G5291-10MG G5291-50MG G5291-100MG
2-Linoleoyl-1-palmitoyl- <i>sn</i> -glycero-3-phosphoethanolamine		62225-25MG
1-O-Octadecyl-2-O-methyl- <i>sn</i> -glycero-3-phosphocholine		O9262-5MG
1-Oleoyl- <i>sn</i> -glycero-3-phosphocholine		L1881-5MG L1881-25MG L1881-100MG
1-Oleoyl-2-palmitoyl- <i>sn</i> -glycero-3-phosphocholine	-	P4142-5MG P4142-25MG P4142-100MG
2-Oleoyl-1-palmitoyl- <i>sn</i> -glycero-3-phosphocholine		P3017-10MG P3017-25MG P3017-100MG
2-Oleoyl-1-palmitoyl- <i>sn</i> -glycero-3-phosphocholine		42773-100MG 42773-500MG
2-Oleoyl-1-palmitoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) ammonium salt		76559-100MG 76559-500MG
2-Oleoyl-1-palmitoyl- <i>sn</i> -glycero-3-phospho- <i>rac</i> -(1-glycerol) sodium salt		63371-100MG 63371-1G
2-Oleoyl-1-palmitoyl- <i>sn</i> -glycero-3-phospho-L-serine sodium salt		51581-100MG 51581-1G
2-Oleoyl-1-stearoyl- <i>sn</i> -glycero-3-phosphocholine	-	P9649-10MG P9649-25MG
1-O-Palmitoyl-2-palmitoyl- <i>rac</i> -glycero-3-phosphocholine	-	P6284-5MG
DL-α-Phosphatidylcholine, distearoyl	-	P8180-100MG P8180-500MG
L-α-Phosphatidylcholine, β-O-methyl-γ-O-octadecyl	-	P1186-1MG P1186-5MG

Name	Structure	Cat. No.
L-α-Phosphatidylethanolamine, dioleoyl	-	P1223-25MG P1223-100MG P1223-500MG
L-α-Phosphatidylinositol dipalmitoyl ammonium salt		P1368-500UG
L-α-Phosphatidyl-D- <i>myo</i> -inositol 3,4-diphosphate, dioctanoyl		P4725-1MG
L-α-Phosphatidyl-D- <i>myo</i> -inositol 4,5-diphosphate, dioctanoyl		P3584-1MG
L-α-Phosphatidyl-D- <i>myo</i> -inositol 3,5-diphosphate, dipalmitoyl		P5713-1MG
L-α-Phosphatidyl-D- <i>myo</i> -inositol 3-monophosphate, dioctanoyl		P5578-1MG
L-α-Phosphatidyl-D- <i>myo</i> -inositol 3-monophosphate, dipalmitoyl		P3953-1MG

Name	Structure	Cat. No.
L- α -Phosphatidyl-D- <i>myo</i> -inositol 4-monophosphate, dipalmitoyl ammonium salt		P7686-1MG
L- α -Phosphatidyl-D- <i>myo</i> -inositol 3,4,5-triphosphate, dioctanoyl		P9953-1MG
1-Stearoyl- <i>sn</i> -glycero-3-phosphocholine	-	L2131-5MG L2131-100MG
Triethanolamine		90278-100ML 90278-500ML

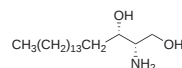
Ether Lipid Metabolism



Phospholipids containing an aliphatic-ether bond or vinyl-ether bond to the carbon-1 of L-glycerol yield 1-alkylglycerols or 1-alkenylglycerols when hydrolyzed. Platelet-activating factor (PAF; 1-alkyl-2-acetyl-*sn*-glycerophosphorylcholine) produces important biological effects such as the aggregation of platelets and induction of hypertensive response at very low concentrations. The metabolic pathway for alkylglycerolipids in the endoplasmic reticulum of tumors and certain healthy cells consist of alkylation of dihydroxyacetone phosphate, ketone reduction, acylation, dephosphorylation, acylation, and/or transfer of phosphorylcholine or phosphorylethanolamine to alkyl glycerol. Pancreatic lipases, phospholipases, and phosphatases do not attack alkyl or 1-alkenyl chains of glycerolipids. Ether-linked phosphatidylcholines and phosphatidylethanolamines containing arachidonic and docosapentaenoic fatty acids are decreased in the blood plasma of hypertensive individuals.

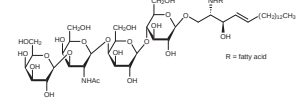
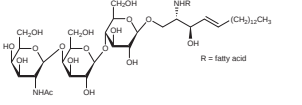
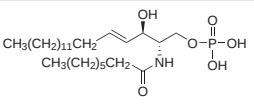
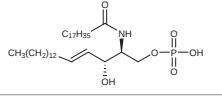
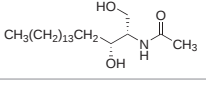
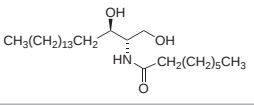
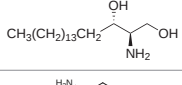
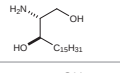
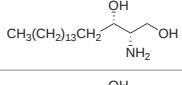
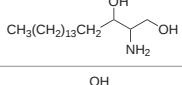
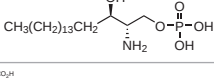
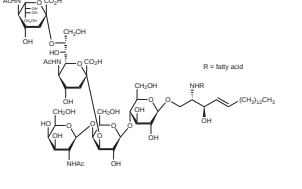
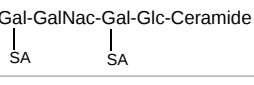
Name	Structure	Cat. No.
β -Acetyl- γ -O-alkyl-L- α -phosphatidylcholine from bovine heart lecithin		P7568-1MG P7568-5MG P7568-10MG
β -Acetyl- γ -O-hexadecyl-L- α -phosphatidylcholine		P4904-1MG P4904-10MG P4904-25MG
1-O-Palmityl-2-acetyl- <i>rac</i> -glycero-3-phosphocholine	-	P1402-10MG
1-O-Palmityl-2-acetyl- <i>sn</i> -glycero-3-phospho-(N,N,N-trimethyl)hexanolamine	-	H8771-250UG
1-O-Palmityl-2-arachidonoyl- <i>sn</i> -glycero-3-phosphocholine	-	P2190-5MG
1-O-Palmityl- <i>sn</i> -glycero-3-phosphocholine		L5016-1MG L5016-10MG
1-O-Palmityl-2-O-methyl- <i>rac</i> -glycero-3-phosphocholine	-	P6034-5MG

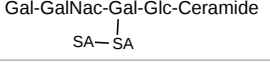
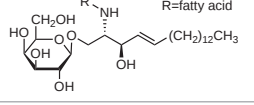
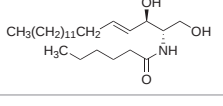
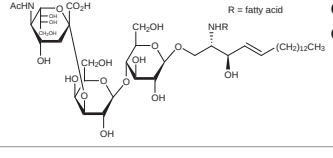
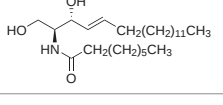
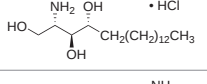
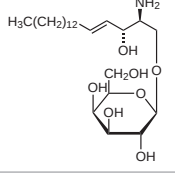
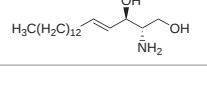
Sphingolipid Metabolism

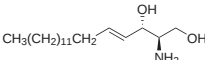
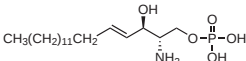
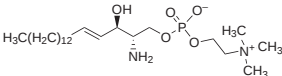


Sphingolipids contain a lipophilic long-chain amino alcohol like sphingosine, dihydrosphingosine, phytosphingosine, or dehydrosphingosine as the characteristic polyalcohol constituent. Sphingosine or a related base is bound to a long-chain fatty acid by an amide linkage and via the terminal hydroxyl group to either a complex carbohydrate or phosphorus-containing moiety. Cerebrosides, sulfatides, ceramide-polyhexosides, and gangliosides are glycosphingolipids, i.e., contain a carbohydrate component instead of phosphorylcholine. The carbohydrate moiety of a glycosphingolipid is oriented towards the exterior cell membrane surface and is characteristic for the cell type, species, growth phase, and differentiated/undifferentiated status.

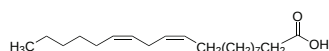
Name	Structure	Cat. No.
N-Acetyl-D-sphingosine		A7191-1MG A7191-5MG

Name	Structure	Cat. No.
Asialoganglioside GM ₁ from bovine brain		G9402-5MG
Asialoganglioside-GM ₂ from bovine brain		G9398-50UG G9398-1MG
Ceramide from bovine brain	-	22244
D-erythro-Ceramide C8 1-phosphate		C8355-1MG
Ceramide 1-phosphate from bovine brain		C4832-1MG
Ceramide phosphorylethanolamine	-	C4987-5MG
Dihydroceramide C2		C7980-5MG C7980-25MG
Dihydroceramide C6	-	C8230-5MG C8230-25MG
Dihydroceramide C8		C8605-5MG C8605-25MG
D-erythro- Dihydrosphingosine		D3314-10MG D3314-50MG
D-threo- Dihydrosphingosine		D4556-1MG
L-threo- Dihydrosphingosine		D4681-1MG
DL-Dihydrosphingosine		D6783-10MG D6783-25MG D6783-100MG
D-erythro- Dihydrosphingosine 1-phosphate		D3439-1MG
Disialoganglioside-GD ₂ from bovine brain, semisynthetic		G0776-1MG
Disialoganglioside GD _{1a} from bovine brain		G2392-1MG G2392-5MG

Name	Structure	Cat. No.
Disialoganglioside GD _{1b} from bovine brain		G8146-1MG
Galactocerebrosides from bovine brain		C4905-10MG C4905-25MG
N-Hexanoyl-D-sphingosine		H6524-1MG H6524-5MG
Lactocerebrosides from bovine brain	-	C3166-1MG C3166-5MG
Monosialoganglioside G _{M3} from canine blood		G5642-5MG G5642-1MG
Monosialoganglioside G _{M1} from bovine brain	-	G7641-1MG G7641-5MG G7641-10MG
Monosialoganglioside G _{M1} from bovine brain	-	G9652-1MG
Monosialoganglioside G _{M2} from bovine brain	-	G8397-1MG
N-Octanoyl-D-sphingosine		O1882-5MG
N-Palmitoyl-D- sphingomyelin semisynthetic from bovine brain sphingomyelin	-	P6778-1MG P6778-10MG
Phytoceramide C2	-	C8105-5MG
Phytosphingosine hydrochloride		P2795-5MG P2795-25MG
Psychosine from bovine brain		P9256-1MG P9256-10MG
Sphingomyelin	-	S7004-5MG S7004-10MG S7004-50MG S7004-100MG S7004-500MG
Sphingomyelin	-	S0756-50MG S0756-100MG S0756-500MG
D-Sphingosine		S7049-5MG S7049-25MG

Name	Structure	Cat. No.
L-erythro-Sphingosine		S7174-5MG
Sphingosine 1-phosphate		S9666-1MG
Sphingosylphosphorylcholine		S4257-10MG S4257-25MG
Sulfatides from bovine brain	-	S1006-5MG S1006-10MG S1006-50MG
Trisialoganglioside-GT _{1b} from bovine brain	-	G3767-1MG G3767-5MG

Fatty Acid Metabolism

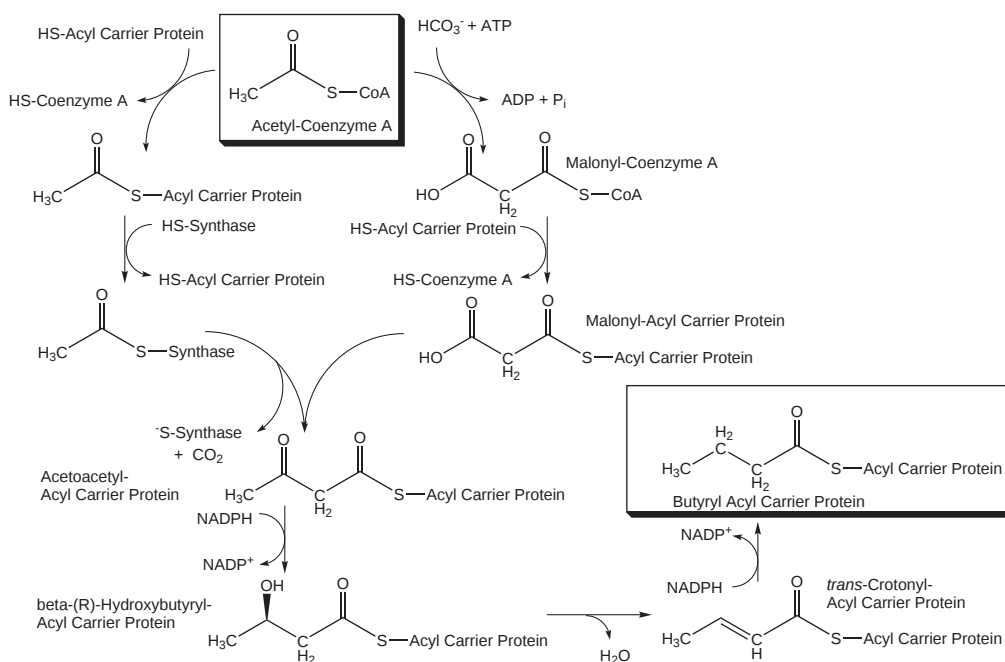


Acetyl coenzyme A is a central metabolite for both the biosynthesis and catabolism of fatty acids. Although the steps in the biosynthetic and catabolic schemes are closely related, there are specific differences. The individual steps are also independent by their spatial separation in biological cells, as biosynthesis occurs in the cytosol while catabolism takes place in the mitochondria. The even-numbered ($[C_2]_n$) common fatty acids result from the sequential two-carbon elongation of acetyl CoA by a series of enzymatic reactions utilizing one or more molecules of acetyl CoA. The two-step

elongation scheme of fatty acid biosynthesis is catalyzed by either discrete enzymes catalyzing specific steps or by a large multi-enzyme complex able to catalyze all steps in the pathway.

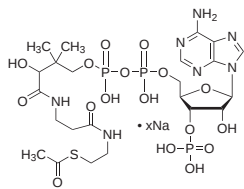
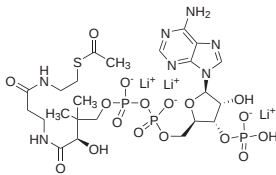
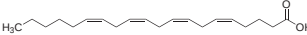
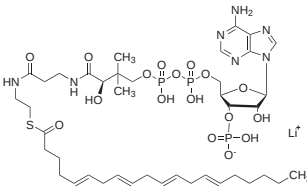
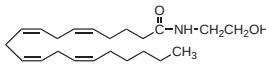
The first step of converting acetyl CoA into the more reactive acetyl-acyl carrier protein (Acetyl-ACP) is catalyzed by ACP transacylase. The acetyl group is then transferred from ACP to a cysteine residue of the synthase complex. In the third step, carboxylation of acetyl CoA is catalyzed by acetyl CoA carboxylase with bicarbonate and ATP to yield malonyl CoA and ADP. In this reaction, biotin acts as a carbon dioxide carrier which transfers carbon dioxide to acetyl CoA. The malonyl group is subsequently converted through a nucleophilic acyl substitution reaction into malonyl-ACP. Thus, both acetyl and malonyl groups are bound to an ACP arm of the synthase complex. The key fifth step consists of the Claisen condensation reaction between the bound acetyl and malonyl groups to give acetoacetyl-ACP. In the sixth step, an enantioselective reduction of the ketone carbonyl group catalyzed by β -ketothioester reductase yields β -hydroxybutyryl-ACP, with a new chiral center in the *R*-configuration. Enzymatic dehydration in step 7 yields *trans*-crotonyl-ACP, which is converted in step 8 by NADPH-dependent reduction of the carbon-carbon double bond to butyryl-ACP.

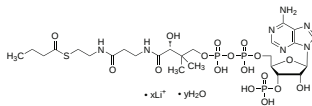
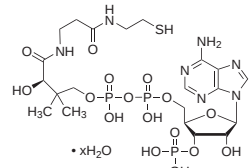
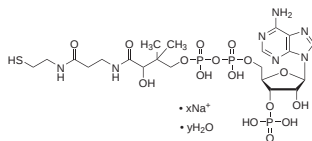
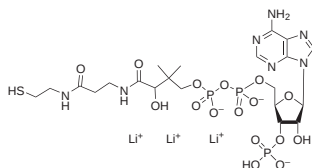
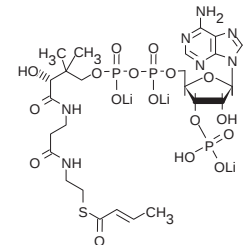
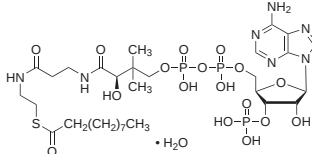
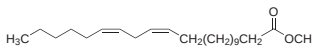
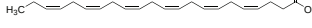
Fatty acid catabolism in the mitochondria of cells provides energy by the β -oxidation pathway and is composed of a repetitive sequence of four enzymatic reaction steps. The fatty acid chain is cleaved from the carboxy-terminus in a stepwise fashion. The first step of the β -oxidation pathway starts with an acyl CoA-dehydrogenase-catalyzed desaturation at C2-C3 using an FAD cofactor to yield an α,β -



Caption Text?

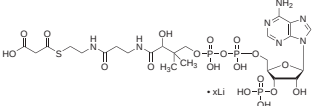
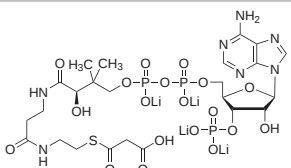
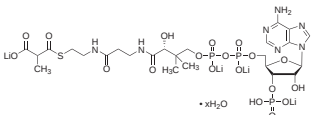
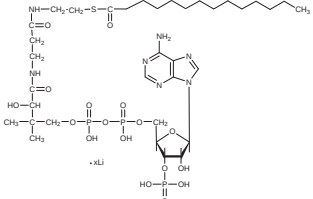
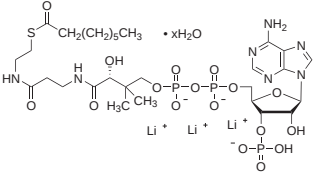
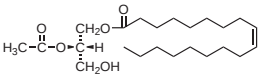
unsaturated acyl-CoA. An enoyl-CoA hydratase-catalyzed *syn*-addition of water to the α,β -unsaturated acyl-CoA in the second step yields 3S-hydroxyacyl-CoA. Next, 3S-hydroxyacyl-CoA is oxidized to the corresponding β -ketoacyl-CoA by NAD-dependent enantioselective 3-hydroxyacyl-CoA dehydrogenases. In the final step, the β -ketoester is cleaved to two CoA esters in a β -ketoacylthiolase-catalyzed retro-Claisen reaction. While the common fatty acids with an even number of carbon atoms produce two molecules of acetyl-CoA in the final passage, fatty acids with an odd number of carbon atoms lead to one molecule of acetyl-CoA and one molecule of propionyl-CoA.

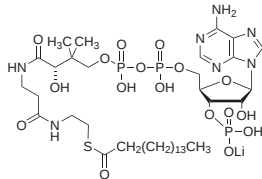
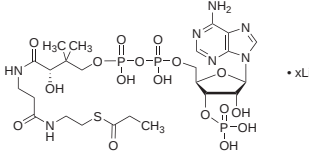
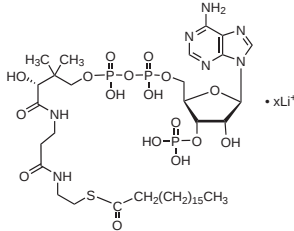
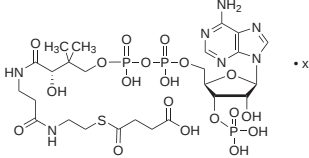
Name	Structure	Cat. No.
Acetoacetyl coenzyme A sodium salt hydrate	-	A1625-5MG A1625-10MG A1625-25MG
Acetyl coenzyme A sodium salt		A2056-1MG A2056-5MG A2056-10MG A2056-25MG A2056-100MG
Acetyl coenzyme A trilithium salt		A2181-1MG A2181-5MG A2181-10MG A2181-25MG A2181-100MG
Arachidic acid	$\text{CH}_3(\text{CH}_2)_{17}\text{CH}_2\text{COOH}$	10930-1G 10930-5G 10930-25G
Arachidonic acid		10931-250MG 10931-1G
Arachidonoyl coenzyme A lithium salt		A5837-5MG A5837-10MG
Arachidonylethanolamide		A0580-5MG A0580-25MG
Behenic acid	$\text{CH}_3(\text{CH}_2)_{19}\text{CH}_2\text{COOH}$	216941-5G
2-Butenoyl coenzyme A lithium salt	-	C6146-5MG C6146-10MG C6146-25MG

Name	Structure	Cat. No.
Butyryl coenzyme A lithium salt hydrate		B1508-5MG B1508-10MG B1508-25MG
Coenzyme A hydrate		C4282-10MG C4282-25MG C4282-100MG
Coenzyme A sodium salt hydrate		C3144-10MG C3144-25MG C3144-100MG C3144-500MG C3144-1G
Coenzyme A trilithium salt		C3019-10MG C3019-25MG C3019-100MG C3019-500MG C3019-1G
Coenzyme A, oxidized lithium salt	-	C2643-5MG C2643-10MG
Crotonoyl coenzyme A trilithium salt		28007-5MG 28007-25MG
Decanoyl coenzyme A monohydrate		D5269-5MG D5269-25MG
3'-Dephosphocoenzyme A	-	D3385-5MG D3385-25MG
cis-13,16-Docosadienoic acid methyl ester		D4034-25MG
cis-4,7,10,13,16,19-Docosahexaenoic acid		D2534-25MG D2534-100MG D2534-1G
cis-4,7,10,13,16,19-Docosahexaenoic acid sodium salt	-	D8768-5MG D8768-25MG

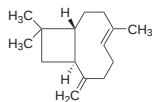
Name	Structure	Cat. No.
all- <i>cis</i> -7,10,13,16,19-Docosapentaenoic acid		D1797-10MG
<i>cis</i> -11,14-Eicosadienoic acid		E3127-25MG E3127-100MG
<i>cis</i> -11,14-Eicosadienoic acid ethyl ester		E7752-25MG
<i>cis</i> -11,14-Eicosadienoic acid methyl ester	-	E7877-25MG E7877-100MG
<i>cis</i> -5,8,11,14,17-Eicosapentaenoic acid		E2011-10MG E2011-25MG E2011-50MG E2011-100MG
<i>cis</i> -5,8,11,14,17-Eicosapentaenoic acid sodium salt		E6627-5MG E6627-10MG E6627-25MG
<i>cis</i> -5,8,11-Eicosatrienoic acid	-	E5888-1MG E5888-10MG
<i>cis</i> -8,11,14-Eicosatrienoic acid		E4504-10MG E4504-100MG
<i>cis</i> -5,8,11-Eicosatrienoic acid methyl ester	-	E6013-1MG
<i>cis</i> -8,11,14-Eicosatrienoic acid methyl ester		E3511-50MG
<i>cis</i> -11,14,17-Eicosatrienoic acid methyl ester	-	E6001-100MG
<i>cis</i> -11-Eicosenoic acid		E3635-100MG E3635-1G
Elaidic acid		E4637-1G E4637-5G E4637-25G
Elaidic acid		45090-10G
Erucic acid		E3385-1G E3385-5G
Glutaryl coenzyme A lithium salt, min. 90%		G9510-5MG G9510-10MG
<i>n</i> -Heptadecanoyl coenzyme A lithium salt		H1385-5MG

Name	Structure	Cat. No.
Hexanoyl coenzyme A trilithium salt hydrate		H2012-5MG H2012-10MG
(<i>R</i>)-3-Hydroxybutyric acid		54920
DL-β-Hydroxybutyryl coenzyme A lithium salt	-	H0261-10MG H0261-25MG
DL-3-Hydroxy-3-methylglutaryl coenzyme A sodium salt		H6132-5MG H6132-10MG H6132-25MG
Isobutyryl coenzyme A lithium salt	-	I0383-5MG I0383-10MG
Isovaleryl coenzyme A lithium salt hydrate		I9381-10MG
Lauroyl coenzyme A lithium salt	-	L2659-5MG L2659-25MG
Lignoceric acid		L6641-100MG L6641-1G L6641-5G L6641-10G
Linoleic acid		L1376-10MG L1376-500MG L1376-1G L1376-5G L1376-10G L1376-25G
Linolenic acid		L2376-500MG L2376-5G L2376-10G
Linolenic acid		62170-10ML-F 62170-50ML-F
γ-Linolenic acid		L2378-10MG L2378-100MG L2378-500MG L2378-1G
Linoleoyl coenzyme A lithium salt, min. 85%	-	L9754-10MG

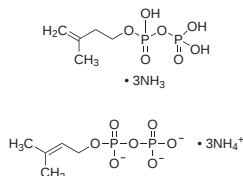
Name	Structure	Cat. No.
Malonyl coenzyme A lithium salt		M4263-5MG M4263-10MG M4263-25MG M4263-100MG
Malonyl coenzyme A tetralithium salt		63410-10MG-F 63410-50MG-F
β-Methylcrotonyl coenzyme A lithium salt	-	M3013-10MG M3013-25MG
Methylmalonyl coenzyme A tetralithium salt hydrate		M1762-1MG M1762-5MG M1762-25MG
Myristoyl coenzyme A lithium salt, min. 80%		M4414-5MG
Nervonic acid	$\text{H}_3\text{C}(\text{CH}_2)_7\text{CH}=\text{CH}(\text{CH}_2)_{13}\text{COOH}$	N1514-100MG
Octanoyl coenzyme A lithium salt hydrate		O6877-5MG O6877-10MG O6877-25MG
1-Oleoyl-2-acetyl-sn-glycerol		O6754-5MG O6754-10MG O6754-25MG

Name	Structure	Cat. No.
Oleoyl coenzyme A lithium salt	-	O1012-5MG O1012-10MG
2-Oleoylglycerol	-	M2787-1MG M2787-5MG M2787-10MG
Palmitoleoyl coenzyme A lithium salt	-	P6775-5MG P6775-10MG
Palmitoyl coenzyme A lithium salt		P9716-5MG P9716-10MG P9716-25MG P9716-100MG
n-Propionyl coenzyme A lithium salt		P5397-5MG P5397-10MG P5397-25MG
Sodium stearate	$\text{CH}_3(\text{CH}_2)_{15}\text{CH}_2\text{COONa}$	S3381-1G S3381-5G S3381-25G
Stearic acid	$\text{CH}_3(\text{CH}_2)_{15}\text{CH}_2\text{COOH}$	S4751-1G S4751-5G S4751-10G S4751-25G S4751-100G
Stearoyl coenzyme A lithium salt		S0802-5MG S0802-10MG S0802-25MG
Succinyl coenzyme A sodium salt, Minimum 85%		S1129-5MG S1129-25MG

Terpenoid Metabolism



The 20,000+ terpenoids found in animals, plants, bacteria, fungi, and archaea are all based on multiples of the 5-carbon isoprene subunit. This is due to biosynthetic pathways that utilize isopentenyl pyrophosphate (IPP) or its isomer dimethylallyl pyrophosphate (DMAPP) as isoprene building blocks for condensation reactions.

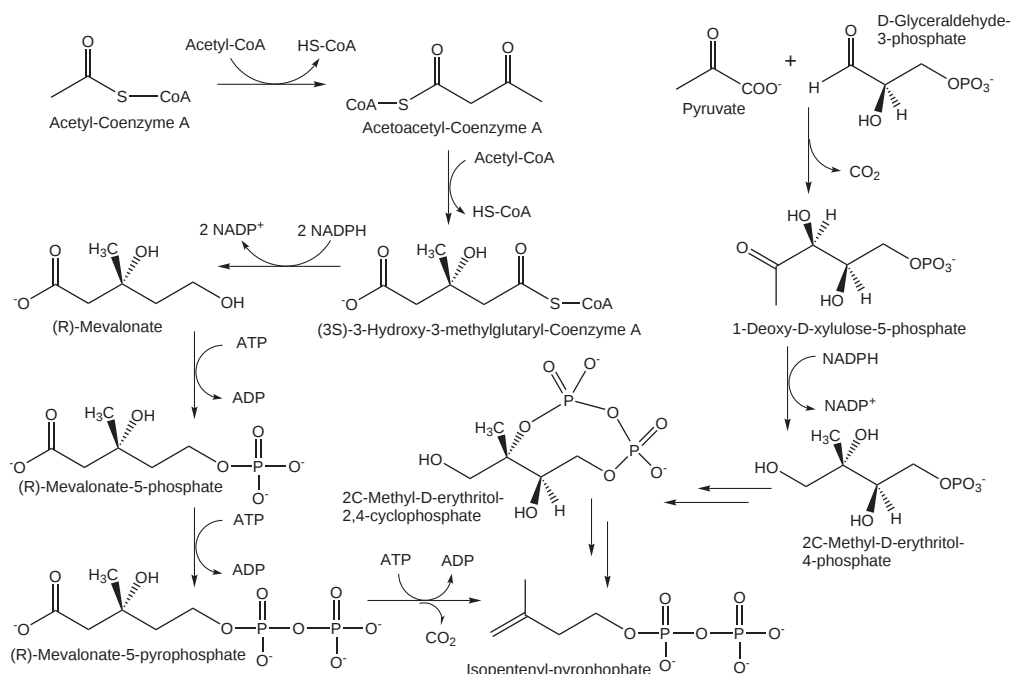


Depending on the biological organism and the terpenoid, two different metabolic pathways exist for the biosynthesis of isopentenyl pyrophosphate:

- the mevalonate pathway and
- the non-mevalonate or methyl D-erythritol 4-phosphate (MEP) pathway.

The mevalonate pathway starts with the formation of acetoacetyl CoA from two molecules of acetyl CoA via Claisen condensation by acetyl-CoA C-acetyltransferase. Acetoacetyl CoA is then condensed with a third molecule of acetyl CoA in an aldol-like reaction catalyzed by hydroxymethylglutaryl-CoA synthase, followed by hydrolysis, to give the metabolite 3-hydroxy-3-methylglutaryl-CoA. In the third step, NADPH-dependent reduction of the thioester group catalyzed by 3-hydroxy-3-methylglutaryl-CoA reductase yields mevalonate. Mevalonate is phosphorylated at the primary hydroxy group by mevalonate kinase to form mevalonate-5-phosphate, which undergoes a second kinase reaction to form mevalonate-5-pyrophosphate. A final phosphorylation at the tertiary hydroxyl group, catalyzed by mevalonate-5-diphosphate decarboxylase, with the simultaneous loss of carbon dioxide and phosphate results in 3-isopentenyl pyrophosphate.

In contrast to the extensively studied and historic mevalonate pathway, the MEP pathway was only discovered in the 1990's and is used by bacteria, algae, and chloroplasts.¹ The first step consists of the 1-deoxy-D-xylulose-5-phosphate synthase-catalyzed coupling of pyruvate with D-glyceraldehyde-3-phosphate to give 1-deoxy-D-xylulose-5-phosphate (DOXP) with concurrent loss of carbon dioxide. Next, an NADPH-dependent rearrangement and reduction reaction is catalyzed by deoxyxylulose-5-phosphate reductoisomerase to produce methyl-D-erythritol 4-phosphate (MEP). MEP is conjugated with cytidine triphosphate (CTP) to produce 4-diphosphocytidyl-2C-methyl-D-erythritol. This intermediate



Caption Text?

undergoes additional phosphorylation at the 2-position of the erythritol moiety (4-diphosphocytidyl-2C-methyl-D-erythritol 2-phosphate) with subsequent enzymatic cyclization and loss of cytidine monophosphate. This metabolite is finally converted to isopentenyl pyrophosphate by the action of two enzymes that reduce the cyclodiphosphate intermediate to a diphosphate and subsequent pyrophosphate.

Another important step in the biosynthesis of terpenoids is the isomerization of isopentenyl pyrophosphate to dimethylallyl pyrophosphate catalyzed by isopentenyl pyrophosphate isomerase. These two fundamental metabolites are joined to form geranyl pyrophosphate (GPP). GPP can be elongated with another molecule of isopentenyl pyrophosphate to farnesyl pyrophosphate (FPP) through sequential condensation by farnesyl pyrophosphate synthetase. Terpenoids with up to 5 isoprenoid units (25 carbons) can be synthesized in this manner, while structures with higher numbers of carbon atoms are formed by dimerization of the corresponding building blocks.

Reference:

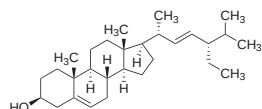
1. Hunger, W.N., et al., Structure and reactivity in the non-mevalonate pathway of isoprenoid biosynthesis. *Biochem. Soc. Trans.*, **31**, 537 (2003).

Name	Structure	Cat. No.
β -Carotene		22040-1G-F 22040-5G-F 22040-25G-F
(-)-trans-Caryophyllene		22075-1ML-F 22075-5ML-F 22075-25ML-F
1-Deoxy-D-xylulose		14764-10MG
1-Deoxy-D-xylulose-5-phosphate sodium salt		13368-1MG 13368-5MG
γ,γ -Dimethylallyl pyrophosphate triammonium salt		D4287-1VL D4287-5VL
trans,trans-Farnesol		277541-1G 277541-10G
Farnesyl monophosphate ammonium salt solution		F1803-5VL
Farnesyl pyrophosphate ammonium salt		F6892-1VL F6892-5VL
Geraniol		163333-25G 163333-100G
Geranylgeraniol	-	G3278-100MG

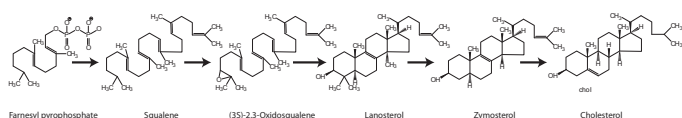
Name	Structure	Cat. No.
Geranylgeranyl monophosphate ammonium salt solution		G2543-5VL
Geranylgeranyl pyrophosphate ammonium salt		G6025-1VL G6025-5VL
Geranyl monophosphate ammonium salt solution		G2293-1VL G2293-5VL
Geranyl pyrophosphate ammonium salt	-	G6772-1VL G6772-5VL
α -Humulene		53675-1ML 53675-5ML
Isopentenyl phosphate dilithium salt		18629
Isopentenyl pyrophosphate triammonium salt solution		I0503-1VL I0503-5VL
Isopentenyl pyrophosphate trilithium salt		00297
(R)-(+)-Limonene		183164-5ML 183164-100ML 183164-500ML
(S)-(-)-Limonene		218367-50G 218367-250G
Lycopene		L9879-1MG L9879-5MG L9879-10MG
(-)-Menthol		15785-100G 15785-1KG
(-)-Menthone		218235-25G 218235-100G
2-C-Methyl-D-erythritol		41707
(R)-Mevalonic acid sodium salt		41288-50MG 41288-10MG
($\pm$)-Mevalonic acid 5-phosphate trilithium salt hydrate		79849-10MG 79849-50MG
(R)-Mevalonic acid 5-pyrophosphate tetralithium salt		77631-50MG 77631-10MG
($\pm$)-Mevalonic acid 5-pyrophosphate tetralithium salt		94259-10MG 94259-50MG
(R)-(-)-Mevalonolactone		68519-100MG 68519-500MG

Name	Structure	Cat. No.
(±)-Mevalonolactone		M4667-1G M4667-5G M4667-10G
<i>cis</i> -Nerolidol		72180-25ML
2,3-Oxidosqualene		41043-1MG 41043-10MG 41043-50MG
Phytol, mixture of isomers (~2/1: <i>trans</i> / <i>cis</i>)		80191-100ML
(-)- α -Pinene		P45702-5ML P45702-100ML P45702-500ML
(<i>R</i>)-(+)-Pulegone		376388-5G 376388-100G
Squalene		S3626-10ML S3626-100ML S3626-500ML S3626-1L
(+)- α -Terpineol		83073-5ML-F

Steroid Metabolism



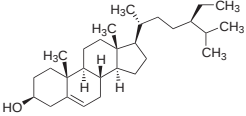
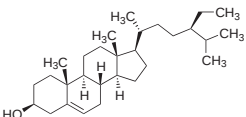
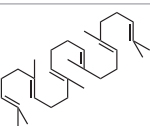
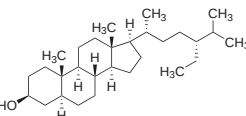
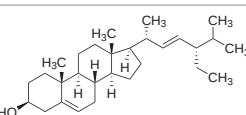
The biosynthesis of steroids originates with farnesyl pyrophosphate through reductive dimerization to squalene by squalene synthase and a subsequent enzymatic conversion of squalene to lanosterol. Enzymatic epoxidation of squalene by squalene epoxidase then yields (3*S*)-2,3-oxidosqualene. Subsequent catalysis by oxidosqualene-lanosterol cyclase generates four rings, six carbon-carbon bonds, and seven chiral centers in one enzymatic reaction and results in a tetracyclic triterpene steroid frame.



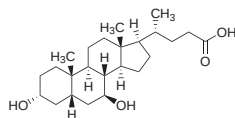
Name	Structure	Cat. No.
Brassicasterol	-	B4936-5MG
Campesterol		C5157-1MG C5157-5MG C5157-10MG
Cholecalciferol		C9756-1G C9756-5G
5 α -Cholest-7-en-3 β -ol	-	C3652-25MG C3652-100MG
Cholesterol		C8667-500MG C8667-1G C8667-5G C8667-25G C8667-100G
Cholesterol		C8503-25G C8503-100G C8503-500G C8503-1KG C8503-5KG
Cholesteryl arachidonate	-	C8753-10MG C8753-25MG C8753-500MG
Cholesteryl hexanoate		C6524-25G C6524-100G
Cholesteryl linoleate		C0289-100MG C0289-250MG C0289-1G
Cholesteryl oleate		C9253-100MG C9253-250MG C9253-500MG C9253-1G
Cholesteryl palmitate		C6072-1G C6072-10G
Corticosterone		C2505-500MG
Cortisone		C2755-250MG C2755-1G C2755-5G

Name	Structure	Cat. No.
7-Dehydrocholesterol		D4429-1G D4429-5G
21-Deoxycortisol		P9521-25MG
1-Deoxy-D-xylulose-5-phosphate sodium salt		13368-1MG 13368-5MG
Desmosterol	-	D6513-5MG D6513-10MG
1α,25-Dihydroxyvitamin D ₃		17936-100UG-F 17936-1MG-F
1α,25-Dihydroxyvitamin D ₂		17944-1MG-F
(24R)-24,25-Dihydroxyvitamin D ₃		17943-100UG 17943-1MG
γ,γ-Dimethylallyl pyrophosphate triammonium salt		D4287-1VL D4287-5VL
Doxercalciferol		D0196-1MG D0196-5MG
Ergocalciferol, activity: ~40000 units/mg		95220-1G 95220-5G
Ergosta-5,7,9(11),22-tetraen-3β-ol	-	E2634-5MG E2634-50MG
Ergosterol		E6510-5G E6510-10G E6510-25G

Name	Structure	Cat. No.
Ergosterol		45480-10G-F 45480-50G-F
trans,trans-Farnesol		277541-1G 277541-10G
Farnesyl pyrophosphate ammonium salt		F6892-1VL F6892-5VL
Geraniol		163333-25G 163333-100G
Geranylgeranyl pyrophosphate ammonium salt		G6025-1VL G6025-5VL
Geranyl pyrophosphate ammonium salt	-	G6772-1VL G6772-5VL
Hydrocortisone		H0135-1MG
22(R)-Hydroxycholesterol	-	H9384-1MG H9384-5MG
22(S)-Hydroxycholesterol		H5884-5MG H5884-10MG
17α-Hydroxypregnenolone	-	H5002-5G
1α-Hydroxyvitamin D ₃		17946-1MG
Isopentenyl pyrophosphate triammonium salt solution		I0503-1VL I0503-5VL
Lanosterol		L5768-1MG L5768-5MG
(±)-Mevalonic acid 5-phosphate trilithium salt hydrate		79849-10MG 79849-50MG
(±)-Mevalonic acid 5-pyrophosphate tetralithium salt		94259-10MG 94259-50MG
2,3-Oxidosqualene		41043-1MG 41043-10MG 41043-50MG
5-Pregnen-3β-ol-20-one		P9129-1G P9129-5G P9129-25G P9129-100G

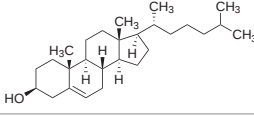
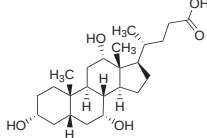
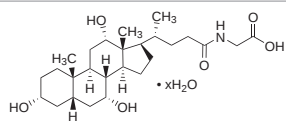
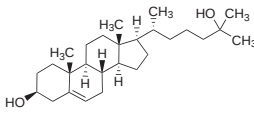
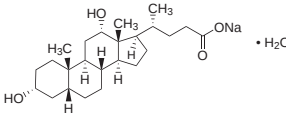
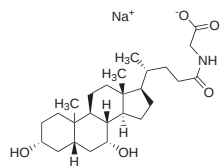
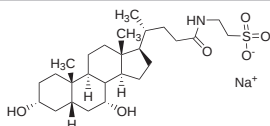
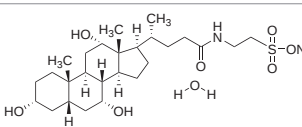
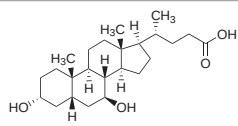
Name	Structure	Cat. No.
β -Sitosterol		S1270-10MG S1270-25MG S1270-100MG
β -Sitosterol		S9889-1MG S9889-5MG S9889-10MG
Squalene		S3626-10ML S3626-100ML S3626-500ML S3626-1L
Stigmastanol		S4297-1G S4297-5G
Stigmasterol		S2424-1G S2424-5G S2424-10G S2424-25G

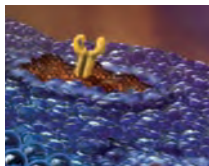
Bile Acid Metabolism



Mammalian cells use multiple biochemical pathways to break down cholesterol into more polar, water-soluble bile acids that can function as emulsifying agents. A series of enzymatic reaction steps involving 7 α -hydroxylations, epimerization of the 3 β -hydroxy group, saturation of the double bond, and side-chain cleavage or oxidation convert cholesterol into the bile acid chenodeoxycholic acid.

7-Deoxycholic acid is formed along a similar pathway, with an additional 12 α -hydroxylation and removal of the 7-hydroxy group. The hydrophobic/hydrophilic balance of the various bile acids facilitate the solubilization of monoacylglycerols, fat-soluble vitamins, and other lipids. This helps digestion and absorption but is also required for solubilization and excretion of lipophilic metabolites such as bilirubin.

Name	Structure	Cat. No.
Cholesterol		20808-100G-R 20808-1KG-R
Cholic acid		C1129-25G C1129-100G C1129-500G C1129-1KG
Glycocholic acid hydrate		G2878-100MG G2878-500MG G2878-1G G2878-5G G2878-25G
25-Hydroxycholesterol		H1015-10MG H1015-25MG H1015-100MG
Sodium deoxycholate monohydrate		30968-25G
Sodium glycochenodeoxycholate		G0759-25MG G0759-100MG G0759-500MG G0759-1G G0759-10G G0759-25G
Sodium taurochenodeoxycholate		T6260-100MG T6260-250MG T6260-1G T6260-5G
Taurocholic acid sodium salt hydrate		T9034-1G T9034-5G T9034-25G
Ursodeoxycholic acid		U5127-1G U5127-5G U5127-25G



Isotopically Labeled Metabolites

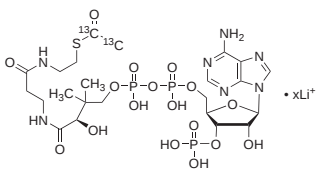
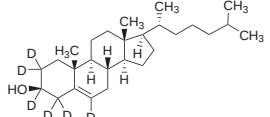
Isotopically Labeled Metabolites

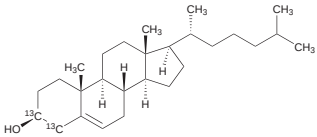
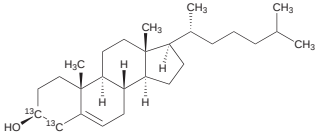
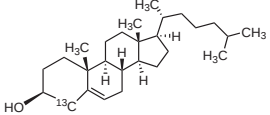
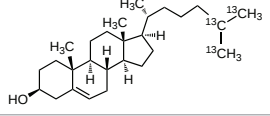
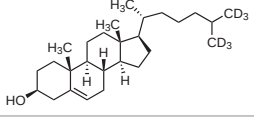
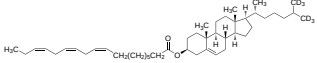
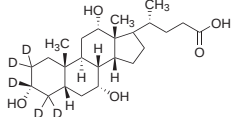
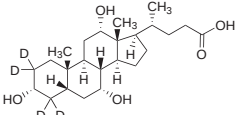
As the body's major source of fuel, the study of fat metabolism is an area of interest and importance. The increased availability of stable isotopically labeled fatty acids and related compounds has aided in the steady accumulation of information surrounding lipid metabolism. Labeled fatty acids have been used in numerous research areas including fatty acid oxidation¹, lipid kinetics², quantitative lipid analyses³, lipid synthesis⁴, and fat absorption.¹ Information gathered from stable isotope tracer studies has added to our understanding of the metabolic pathways and the metabolic fate of certain fatty acids.

The safe labeling of lipids by non-radioactive isotopes has been crucial for advancements in the study of the structural biology of membranes, magnetic resonance imaging and spectroscopy, nutrition science, and metabolomics. The availability of ¹³C- and ¹⁵N-labeled lipids and lipid precursors has enabled selective isotope labeling of membranes by biosynthetic incorporation of a labeled lipid. Lipid metabolism can therefore be studied in several dimensions: at the whole organism level, at the level of individual membranes, or in the small molecule pathways of lipid conversions.

References:

1. Wolfe, R.R. and Chinkes, D.L.; *Isotope Tracers in Metabolic Research, Principles and Practice of Kinetic Analysis*, 2nd ed.; John Wiley & Sons, Inc.: Hoboken, New Jersey, 2005; pp 259-298.
2. Adiels, M., Larsson, T., Sutton, P., Taskinen, M.-R., Borén, J., and Fielding, B.A.; *Rapid Commun. Mass Spectrom.* **2010**, *24*, 586-592.
3. Sun, D., Cree, M.G., and Wolfe, R.R.; *Anal. Biochem.* **2006**, *349*, 87-95.
4. Bergman, B.C., Perreault, L., Hunerdosse, D.M., Koehler, M.C., Samek, A.M., and Eckel, R.H.; *J. Appl. Physiol.* **2010**, *108*, 1134-1141.
5. Jauregui, R.C., Mattes, R.D., and Parks, E.J.; *Gastroenterology* **2010**, in press.

Name	Structure	Cat. No.
Acetyl-1,2- ¹³ C ₂ coenzyme A lithium salt		658650
Cholesterol-2,2,3,4,4,6-d ₆		488577-100MG

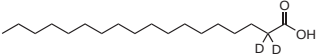
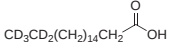
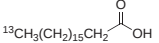
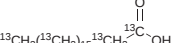
Name	Structure	Cat. No.
Cholesterol-3,4- ¹³ C ₂		488585
Cholesterol-3,4- ¹³ C ₂		662291
Cholesterol-4- ¹³ C		605875
Cholesterol-25,26,27- ¹³ C ₃		707678
Cholesterol-26,26,26,27,27-d ₆		679046
Cholesteryl-26,26,26,27,27-d ₆ linolenate		730238
Cholic-2,2,3,4,4-d ₅ acid		614106
Cholic acid-2,2,4,4-d ₄		614149-500MG

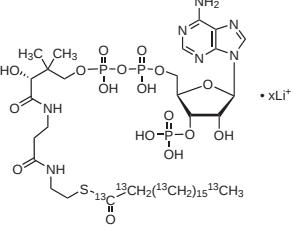
Name	Structure	Cat. No.
Cholic-24- ¹³ C acid		605883
Choline chloride- ¹⁵ N		609269-1G
Choline chloride-1- ¹³ C		605301
Choline chloride-1,1,2,2-d ₄		615544
Choline chloride-trimethyl-d ₉		492051-1G
Cortisone-2,2,4,6,6,9,12,12-d ₈		705586-5MG
1α,25-Dihydroxyvitamin D ₃ (6,19,19-d ₃)		705942-1MG
Ethanolamine- ¹⁵ N		609552
Ethanolamine- ¹³ C ₂		606294
Ethanolamine-2- ¹³ C		606316
Ethanolamine- ¹³ C ₂ hydrochloride		606308-100MG
Glycerol-1,1,2,3,3-d ₅		454524-1G 454524-5G
Glycerol-1,1,2,3,3-d ₅		661473
Glycerol-1,2- ¹³ C ₂		714895
Glycerol-1,3- ¹³ C ₂		492639-250MG
Glycerol-2- ¹³ C		489484
Glycerol-2- ¹³ C		661465
Glycerol- ¹³ C ₃		489476-500MG
Glycerol- ¹³ C ₃		660701

Name	Structure	Cat. No.
<i>rac</i> -Glyceryl-d ₅ -2,3-dioleate-1-palmitate		730076
Glyceryl tri(octanoate-1- ¹³ C)		425893
Glyceryl tri(octanoate-d ₁₃)		617121
Glyceryl tri(oleate-1- ¹³ C)		489514-500MG
Glyceryl tri(oleate-1- ¹³ C)		714771
Glyceryl tri(oleate-1,2,3,7,8,9,10- ¹³ C ₇)		646253
Glyceryl tri(oleate-9,10- ¹³ C ₂)		646245
Glyceryl tri(palmitate-1- ¹³ C)		425907-1G
Glyceryl tri(palmitate-1- ¹³ C)		680842
Glyceryl tri(palmitate-1,2- ¹³ C ₂)		605603
Glyceryl tri(palmitate-16,16,16-d ₃)		615471
Glyceryl tri(palmitate-d ₃₁)		616966
Glyceryl tri(palmitate-d ₃₁), S&P tested		660698
Glyceryl tri(stearate-1- ¹³ C)		492663-250MG
Glyceryl tri(stearate-18,18,18-d ₃)		616117

Name	Structure	Cat. No.
Glycocholic acid-glycyl-1- ¹³ C		605891
Hydrocortisone-1α,2α-d ₂		614157
Hydrocortisone-9,11,12,12-d ₄		705594-5MG 705594-10MG
Linoleic acid- ¹³ C ₁₈		605735-100MG
Linolenic Acid-1- ¹³ C		694940
Linolenic acid- ¹³ C ₁₈		605743-100MG
Malonyl- ¹³ C ₃ coenzyme A lithium salt		655759
Octanoyl-2,4,6,8- ¹³ C ₄ Coenzyme A, lithium salt		703885
Oleic acid-1- ¹³ C		661589
Oleic acid-9,10- ¹³ C ₂		646466
Oleic acid- ¹³ C ₁₈		490431-100MG
Oleic acid-d ₃₄		683582

Name	Structure	Cat. No.
Oleoyl- ¹³ C ₁₈ coenzyme A lithium salt		675768
Palmitoyl-1- ¹³ C coenzyme A lithium salt		658200
Potassium linoleate- ¹³ C ₁₈		605816
1,2-Propanediol-1,2- ¹³ C ₂		603678
1,2-Propane-d ₆ -diol		614416
1,2-Propanediol-d ₈		486272-5G
1,3-Propanediol-1,3- ¹³ C ₂		603554
1,3-Propanediol-2- ¹³ C		491101
1,3-Propanediol- ¹³ C ₃		603562
1,3-Propane-d ₆ -diol		613525
1,3-Propanediol-d ₈		589535
1-Propanol- ¹³ C ₃		640689
Propionaldehyde-1- ¹³ C		603813
Propionaldehyde-2,2-d ₂		588113
Propionaldehyde-2,2,3,3-D ₅		707201
Stearic acid-1- ¹³ C		299162-1G
Stearic acid-1,2- ¹³ C ₂		591602
Stearic acid-2- ¹³ C		591491

Name	Structure	Cat. No.
Stearic acid-2,2-d ₂		493155-1G
Stearic-17,17,18,18-d ₅ acid		615846
Stearic acid-18- ¹³ C		605654
Stearic acid- ¹³ C ₁₈		605581-100MG
Stearic acid-18,18,18-d ₃	-	490393-250MG

Name	Structure	Cat. No.
Stearic-d ₃₅ acid		448249-250MG 448249-1G
Stearoyl- ¹³ C ₁₈ coenzyme A lithium salt		675776



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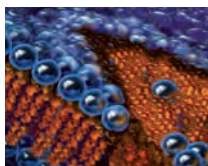
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Characteristic Metabolites for Inborn Errors of Lipid Metabolism

Characteristic Metabolites for Inborn Errors of Lipid Metabolism

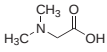
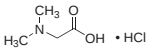
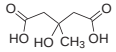
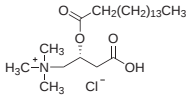
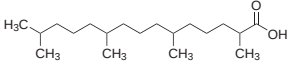
Hereditary disorders in lipid metabolism include Tay-Sachs disease, Gaucher disease, Niemann-Pick disease, metachromatic leucodystrophy, Fabry disease, Refsum disease, and Tangier disease. These lipidoses are characterized by dysfunctional lipid metabolism and result in abnormal metabolite accumulations. One of the first disorders recognized as an inborn error of lipid metabolism was Refsum disease, which produces toxic levels of phytanic acid if untreated.

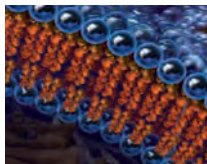
Gaucher disease is a progressive sphingolipid-degradation disease characterized by genetic mutations in the lysosomal enzyme glucocerebrosidase, which leads to decreased enzymatic activity. Measurements of the metabolites methylcholine, phosphatidylcholine, and sphingomyelin are important for studying the pathophysiology of Gaucher disease. The main therapy used to treat Gaucher disease is enzyme-replacement therapy in order to normalize sphingolipid degradation and to prevent tissue damage caused by sphingolipid accumulation. Another promising therapeutic approach to Gaucher disease is to decrease the tissue glucocerebrosidase level to a concentration which can be cleared by the existing glucocerebrosidase.

A deficiency of the lysosomal enzyme α -galactosidase A results in the progressive accumulation of the glycosphingolipids globotriaosylceramide Gb3 and digalactosyl-ceramide in Fabry disease, which can cause early death from cardiac, renal, and cerebrovascular events.

Dimethylglycine dehydrogenase (DMGDH) deficiency is an inborn error of choline metabolism caused by a mutation in the gene *hDMGDH* and results in increased N,N-dimethylglycine concentrations of 100-fold in serum and 20-fold in urine.

Nine inborn errors of bile acid metabolism have been identified that result in enzyme deficiencies and damaged bile acid biosynthesis in infants, children, and adults. Since bile acids have several important physiological functions, such as emulsifying fats and fat-soluble vitamins, and involvement in cholesterol, bilirubin, xenobiotics, and drug metabolites elimination, a failure in the multistep enzymatic conversion of cholesterol to bile acids will accumulate unusual bile acids and metabolic intermediates.

Name	Structure	Cat. No.
O-Acetyl-L-carnitine hydrochloride		A6706-1G A6706-5G
N,N-Dimethylglycine		D1156-10MG D1156-5G D1156-10G D1156-25G
N,N-Dimethylglycine hydrochloride		D6382-5G D6382-25G
Globotriaosylsphingosine from porcine blood	-	G9534-1MG
DL-Hexanoylcarnitine chloride	-	H2132-25MG
3-Hydroxy-3-methylglutaric acid		H4392-100MG H4392-1G
Palmitoyl-L-carnitine chloride		P1645-5MG P1645-10MG P1645-25MG
Palmitoyl-DL-carnitine chloride		P4509-100MG P4509-1G
Phytanic acid, mixture of isomers	-	P4060-5MG P4060-25MG
Pristanic acid solution, mixture of isomers		P6617-5MG



Lipid Metabolites as Biomarkers for the Differentiation of Diseased and Healthy Cells

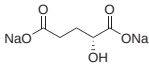
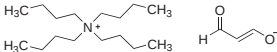
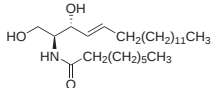
Lipid Metabolites as Biomarkers for the Differentiation of Diseased and Healthy Cells

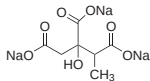
The human body regulates cellular lipid concentrations by location and time, and deregulation is associated with a variety of human diseases. In addition to disorders due to inborn errors of lipid metabolism, diseases like cardiovascular disease, diabetes, cancer, infections, and neurodegenerative diseases are frequently lipid related. A key feature of diabetes, obesity, and atherosclerosis is excessive lipid accumulation in cells. Excess energy is stored as triacylglycerols for later use when nutrients are scarce. These triacylglycerols, along with the precursors to membrane lipids, are stored in cytoplasmic lipid droplets in eukaryotic cells. Host-pathogen interactions in lipid metabolism are also of interest in infectious disease research since pathogenic microorganisms can use the host's lipids as a carbon and energy source, which provides an advantage to the pathogen in persistence, virulence, and overcoming host immune responses. Major drug classes like the statins and cyclooxygenase (COX) inhibitors target enzymes involved in lipid metabolism. Increased concentrations of lysophosphatidylcholines and decreased concentrations of antioxidative ether phospholipids in serum have been associated with acquired obesity. Detailed analysis of the local concentrations of lipids, their precursors, and metabolites as a function of time will aid the understanding of their roles in the vital functions of cells.

Next to adipose tissue, the central nervous system has the highest cellular lipid concentration. Injuries and disorders of the CNS may be influenced by changes in lipid metabolism. The accumulation of LDL-derived lipids in the arterial wall results in atherosclerosis and is a risk factor for stroke. Lipid metabolism is altered by tumor necrosis factor- α (TNF- α) and interleukin-1 (IL-1), which stimulate production of eicosanoids, ceramide, and reactive oxygen species. Reactive oxygen species that exceed the cell's capability for detoxification result in oxidative stress yielding oxidized phospholipids that are metabolized to 4-hydroxynonenal, 4-oxo-2-nonenal, and acrolein. Neurodegenerative diseases, CNS injuries, and brain traumas can be influenced by lipid oxidation because of the high lipid content and oxygen consumption of the brain. Sphingomyelin accumulation

results from acidic sphingomyelinase deficiency in Niemann-Pick disease A and B, while cholesterol accumulation occurs in Niemann-Pick disease C due to mutations in either the *NPC1* or *NPC2* genes. Cholesterol is an important regulator of lipid organization and a precursor for neurosteroid biosynthesis. The gene encoding the apolipoprotein E4 variant of the principal cholesterol carrier protein in the brain (the E4 allele of *APOE*) is a significant risk factor for Alzheimer's disease. Lipid peroxidation due to phospholipase activation may contribute in part to Parkinson's disease.

Name	Structure	Cat. No.
N-Acetyl-D-sphingosine		A7191-1MG A7191-5MG
Acrolein		01679-10ML
Cholesterol		C8667-500MG C8667-1G C8667-5G C8667-25G C8667-100G
Cholesterol		20808-100G-R 20808-1KG-R
Galactocerebrosides from bovine brain		C4905-10MG C4905-25MG
L- α -Glycerophosphorylcholine	-	G5291-10MG G5291-50MG G5291-100MG
N-Hexanoyl-D-sphingosine		H6524-1MG H6524-5MG
HNE-DMA		H9538-2MG H9538-5MG

Name	Structure	Cat. No.
D-α-Hydroxyglutaric acid disodium salt		H8378-25MG H8378-100MG H8378-250MG
Malondialdehyde tetrabutylammonium salt		63287-1G-F 63287-5G-F
N-Palmitoyl-D-sphingomyelin semisynthetic from bovine brain sphingomyelin	-	P6778-1MG P6778-10MG
N-Octanoyl-D-sphingosine		O1882-5MG

Name	Structure	Cat. No.
Sphingomyelin	-	S7004-5MG S7004-10MG S7004-50MG S7004-100MG S7004-500MG
Sphingomyelin	-	S0756-50MG S0756-100MG S0756-500MG
Trisodium (2 <i>RS</i> ,3 <i>RS</i>)-2-methylcitrate		59464-10MG 59464-50MG

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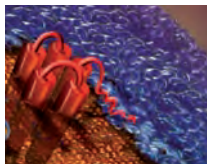
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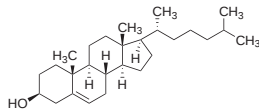
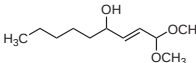
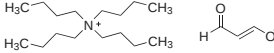
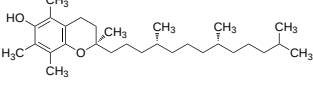
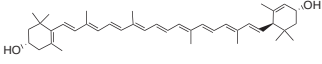


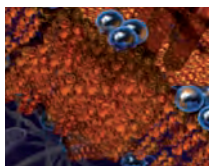
Altered Lipid Metabolites in Aging

Altered Lipid Metabolites in Aging

The development of cognitive and neurodegenerative disorders, cardiovascular diseases, metabolic disorders, bone diseases, certain forms of cancer, and autoimmune diseases with increasing age has been associated with the stability of the cell and biochemical reactions within it. Like DNA and proteins, lipids are sensitive to reactive oxygen species that produce oxidative stress. Endogenous aldehydes, such as malonic dialdehyde and methylglyoxal, are the major initiators of metabolic disorders. Age-related diseases develop when cells cannot control aldehyde formation or abolish the negative methylglyoxal effect on their metabolism. Toxic aldehydes cause cumulative damage over a lifetime if not counteracted by the cell's ability to avoid the formation of endogenous aldehydes or to neutralize their negative effects without damaging vital cellular components. The aging process and the individual's life span is influenced by the balance of the two opposing processes of anti-aging effects and the aging effects of toxic aldehydes.

Cholesterol levels in human serum above 6.5 mmol/L are known to be associated with a higher risk of developing Alzheimer's disease, while a reduction of cholesterol in the serum in midlife lowers the risk of dementia. Bile acids act as regulators of aging that benefit health and longevity and modulate housekeeping longevity assurance pathways. It is likely that the mechanisms for the positive effects of bile acids are evolutionarily conserved. The time-dependent accumulation of age pigments like lipofuscin occurs in all cells, but it is more prominent in cells which are not active in cell division. These age pigments appear together with lipid droplets in the cytoplasm.

Name	Structure	Cat. No.
Cholesterol		C8667-500MG C8667-1G C8667-5G C8667-25G C8667-100G
HNE-DMA		H9538-2MG H9538-5MG
Malondialdehyde tetrabutylammonium salt		63287-1G-F 63287-5G-F
Methylglyoxal solution		M0252-25ML M0252-100ML M0252-250ML M0252-500ML M0252-1L
(+)-α-Tocopherol, activity: ≥1000 IU/g		T1539-25G T1539-100G
Xanthophyll		X6250-1MG X6250-5MG



Lipid Metabolites in Natural Product Biosynthesis

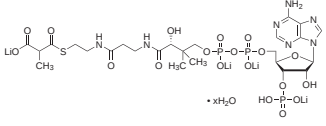
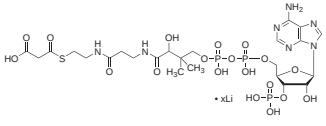
Lipid Metabolites in Natural Product Biosynthesis

Many natural compounds contain tailor-made lipid anchors or lipid-like functional groups intended to localize them on the cellular membrane. Natural products with lipid anchors are of interest for a number of cell migration processes of healthy and diseased cells like wound healing, blood vessel formation, immune system function, and tumor metastasis. Certain lipid components of the cell membrane itself, such as phosphatidylglycerol, cardiolipin, or glycosylphosphatidylinositol, can also serve directly as anchors. These phospholipids are listed in the section on natural abundance lipid metabolites.

Lipids in the cell walls and membranes of pathogenic microorganisms and their related pathways are of particular interest for the development of novel bioactive compounds, especially if they have no eukaryotic counterparts. Antibiotics like the phosphoglycolipids are the only known active site inhibitors of the peptidoglycan glycosyltransferases, which are involved in the biosynthesis of the bacterial cell wall. The lipoteichoic acids of the cell wall of Gram-positive bacteria, anchored by their diacylglycerol moiety, are recognized by the innate immune system, making them an interesting scaffold for new approaches to natural therapeutic products. In Gram-positive bacteria, lipid A is of interest for the development of immunostimulating compounds and vaccine adjuvants.

The biosynthesis of eicosanoids such as prostaglandins, thromboxanes, and leukotrienes begins with arachidonic acid and involves a series of enzymatic reactions including radical formation, reaction with oxygen and cyclization, and reduction. Polyacetylenic natural products contain a unique carbon-carbon triple bond and have a wide variety of biochemical and ecological functions. Initial departure points from primary metabolism include the three fatty acids crepenynic acid, stearolic acid, and tariric acid.

Name	Structure	Cat. No.
Acetyl coenzyme A trilithium salt		A2181-1MG A2181-5MG A2181-10MG A2181-25MG A2181-100MG
Acetyl coenzyme A sodium salt		A2056-1MG A2056-5MG A2056-10MG A2056-25MG A2056-100MG
Coniferyl alcohol		223735-100MG 223735-1G
Farnesyl pyrophosphate ammonium salt		F6892-1VL F6892-5VL
Geranylgeranyl pyrophosphate ammonium salt		G6025-1VL G6025-5VL
Geranyl pyrophosphate ammonium salt	-	G6772-1VL G6772-5VL
Isopentenyl pyrophosphate trilithium salt		00297
Isopentenyl pyrophosphate triammonium salt solution		I0503-1VL I0503-5VL
Malonyl coenzyme A tetralithium salt		63410-10MG-F 63410-50MG-F

Name	Structure	Cat. No.
Methylmalonyl coenzyme A tetralithium salt hydrate		M1762-1MG M1762-5MG M1762-25MG
Malonyl coenzyme A lithium salt		M4263-5MG M4263-10MG M4263-25MG M4263-100MG
Lipoteichoic acid from <i>Streptococcus pyogenes</i>	-	L3140-5MG L3140-10MG L3140-25MG
Lipoteichoic acid from <i>Streptococcus faecalis</i>	-	L4015-5MG L4015-25MG

Name	Structure	Cat. No.
Lipoteichoic acid from <i>Staphylococcus aureus</i>	-	L2515-5MG L2515-10MG L2515-25MG
Lipoteichoic acid from <i>Bacillus subtilis</i>	-	L3265-5MG L3265-25MG
Lipid A, monophosphoryl from <i>Salmonella enterica</i> serotype minnesota Re 595 (Re mutant)	-	L6895-1MG L6895-5MG
Lipid A, monophosphoryl from <i>Escherichia coli</i> F583 (Rd mutant)	-	L6638-1MG L6638-5MG
Lipid A, diphosphoryl from <i>Salmonella enterica</i> serotype minnesota Re 595 (Re mutant)	-	L0774-1MG L0774-5MG
Lipid A, diphosphoryl from <i>Escherichia coli</i> F583 (Rd mutant)	-	L5399-1MG L5399-5MG
L-NASPA	-	P0247-1MG P0247-5MG

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Fax: (+7) 495 621 6037

Singapore

Tel: (+65) 6779 1200
Fax: (+65) 6779 1822

Slovakia

Tel: (+421) 255 571 562
Fax: (+421) 255 571 564

South Africa

Free Tel: 0800 1100 75
Free Fax: 0800 1100 79
Tel: (+27) 11 979 1188
Fax: (+27) 11 979 1119

Spain

Free Tel: 900 101 376
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Tel: (+34) 91 661 99 77
Fax: (+34) 91 661 96 42

Sweden

Tel: (+46) 8 742 4200
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Switzerland

Free Tel: 0800 80 00 80
Free Fax: 0800 80 00 81
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Fax: (+41) 81 755 2815

United Kingdom

Free Tel: 0800 717 181
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Fax: (+44) 1747 833 313

United States

Toll-Free: 800 325 3010
Toll-Free Fax: 800 325 5052
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