



Log₂ fold changes of probes
[Fatty acid biosynthesis; n = 21]

		$\frac{LL_3}{DD_3}$	$\frac{LD_3}{LL_3}$	$\frac{DD_3}{LD_3}$	$\frac{LL_6}{DD_6}$	$\frac{LD_6}{LL_6}$	$\frac{DD_6}{LD_6}$	
K01962 accA; acetyl-CoA carboxylase carboxyl transferase subunit alpha [EC:6.4.1.2 2.1.3.15]								SAR116_1916 Acetyl-CoA carboxylase alpha subunit
K02160 accB; acetyl-CoA carboxylase biotin carboxyl carrier protein								SAR116_1039 acetyl-CoA carboxylase
K02160 accB; acetyl-CoA carboxylase biotin carboxyl carrier protein								SAR116_1773 biotin/lipoyl attachment
K01961 accC; acetyl-CoA carboxylase, biotin carboxylase subunit [EC:6.4.1.2 6.3.4.14]								SAR116_1040 acetyl-CoA carboxylase
K01961 accC; acetyl-CoA carboxylase, biotin carboxylase subunit [EC:6.4.1.2 6.3.4.14]								SAR116_1774 acetyl CoA carboxylase
K01963 accD; acetyl-CoA carboxylase carboxyl transferase subunit beta [EC:6.4.1.2 2.1.3.15]								SAR116_1258 Acetyl-CoA carboxylase carboxyl transferase
K02078 acpP; acyl carrier protein								SAR116_0639 acyl carrier protein
K00645 fabD; [acyl-carrier-protein] S-malonyltransferase [EC:2.3.1.39]								SAR116_0641 acyl-carrier-protein S-malonyltransferase
K00648 fabH; 3-oxoacyl-[acyl-carrier-protein] synthase III [EC:2.3.1.180]								SAR116_0172 probable 3-oxoacyl-acyl-carrier-protein synthase iii
K00647 fabB; 3-oxoacyl-[acyl-carrier-protein] synthase I [EC:2.3.1.41]								SAR116_1573 Beta-ketoacyl synthase
K09458 fabF; 3-oxoacyl-[acyl-carrier-protein] synthase II [EC:2.3.1.179]								SAR116_0638 Beta-ketoacyl synthase
K00059 fabG; 3-oxoacyl-[acyl-carrier protein] reductase [EC:1.1.1.100]								SAR116_0623 3-oxoacyl-acyl carrier protein reductase
K00059 fabG; 3-oxoacyl-[acyl-carrier protein] reductase [EC:1.1.1.100]								SAR116_0640 3-oxoacyl-(acyl-carrier-protein) reductase
K00059 fabG; 3-oxoacyl-[acyl-carrier protein] reductase [EC:1.1.1.100]								SAR116_1718 short-chain dehydrogenase/reductase SDR
K01716 fabA; 3-hydroxyacyl-[acyl-carrier protein] dehydratase / trans-2-decenoyl-[acyl-carrier protein] isomerase [EC:4.2.1.59 5.3.3.14]								SAR116_1572 3-hydroxydecanoyl-ACP dehydratase
K02372 fabZ; 3-hydroxyacyl-[acyl-carrier-protein] dehydratase [EC:4.2.1.59]								SAR116_0508 (3R)-hydroxymyristoyl-[acyl carrier protein] dehydratase
K00208 fabI; enoyl-[acyl-carrier protein] reductase I [EC:1.3.1.9 1.3.1.10]								SAR116_1574 Enoyl-acyl-carrier-protein reductase
K00208 fabI; enoyl-[acyl-carrier protein] reductase I [EC:1.3.1.9 1.3.1.10]								SAR116_1062 Enoyl-acyl-carrier-protein reductase
K01897 ACSL; long-chain acyl-CoA synthetase [EC:6.2.1.3]								SAR116_0193 Putative long-chain-fatty-acid--CoA ligase (Long-chain acyl-CoA synthetase)
K00432 gpx; glutathione peroxidase [EC:1.11.1.9]								SAR116_2363 Phospholipid hydroperoxide glutathione peroxidase
K10804 tesA; acyl-CoA thioesterase I [EC:3.1.2.- 3.1.2.2 3.1.1.2 3.1.1.5]								SAR116_1417 lipolytic enzyme