Table S1. Mass-to-Charge Ratios (m/z) and LC Retention Times (RT) of Lipids that Change during Cytokinesis and at the Midbody, Related to Table 1

m/z	RT	Lipid	Fold increase during	Fold increase at
	(min)		cytokinesis ^(a)	midbody ^(b)
441.3344	~40	sterol derivative	>27	unchanged
661.5191	~46	phosphatidic acid ether/ester	40	unchanged
837.5525	~54	(O-18:0/16:0) phosphatidylinositol (16:0/18:0)	10	unchanged
538.5216	~70	C16 diH-ceramide	>11	unchanged
566.5518	~72	C18 diH-ceramide	accumulated ^(c)	unchanged
594.5830	~73	C20 diH-ceramide	accumulated ^(c)	unchanged
622.6170	~75	C22 diH-ceramide	accumulated ^(c)	accumulated ^(c)
650.6456	~77	C24 diH-ceramide	accumulated ^(c)	accumulated ^(c)
620.5987	~77	C22 ceramide	4	4
648.6366	~79	C24 ceramide	4	14
698.5610	~68	C16 hexosylceramide	>16	13
810.6783	~79	C24 hexosylceramide	unchanged	>36
647.4645	~48	phosphatidic acid (16:0/16:0)	unchanged	14
792.7113	~72	triacylglycerol (16:1,12:0, 18:1)	unchanged	54
810.5278	~49	phosphatidylserine (18:0/20:4)	unchanged	6

For chemical structures of the lipids, see Fig. 1.

^(a) Fold increase in S-phase vs cytokinesis cells is determined by [Abundance _{cytokinesis}] / [Abundance _{S-} phase] for each lipid. Abundance is the total ion count for a given ion. Each ion corresponds to a massto-charge ratio (m/z), which is used to assign the lipid species.

^(b) Fold increase in midbody vs cytokinesis cells is determined by [Abundance midbody] / [Abundance purified lysate] for each lipid.
(c) A numeric value for fold increase could not be calculated due to the low abundance of these ions in

S-phase or cytokinesis cells.

Table S3. Targeted Analysis of the Sphingolipids in siSMPD4-, siGALC-, and siDGAT2-Treated Cells, Related to Figure 3

Fold changes in ceramides, diH-ceramides, hexosylceramides and sphingomyelins were calculated in siRNA treated cells. Fold changes are an average of two independent experiments. For a graphical representation see Fig. S3A.

		siSMPD4	siGALC	siDGAT2	
		Average fold changes relative to control			
	п Оп				
CERAMIDES	n=1 (C14 ceramide)	not detected	not detected	not detected	
	n=2 (C16 ceramide)	4.5x increase	unchanged	unchanged	
	n=3 C18 ceramide)	not detected	not detected	not detected	
	n=4 (C20 ceramide)	not detected	not detected	not detected	
	n=5 (C22 ceramide)	8x increase	unchanged	unchanged	
	n=6 (C24 ceramide)	3.5x increase	unchanged	unchanged	
	н он				
	NH H				
	$\sim \sim [\sim]_{n} \sim \sim ~$				
diH-CERAMIDES	n=1 (C14 diH-ceramide)	not detected	not detected	not detected	
	n=2 (C16 diH-ceramide)	not detected	not detected	not detected	
	n=3 (C18 diH-ceramide)	not detected	not detected	not detected	
	n=4 (C20 diH-ceramide)	not detected	not detected	not detected	
	n=5 (C22 diH-ceramide)	not detected	not detected	not detected	
	n=6 (C24 diH-ceramide)	not detected	not detected	not detected	
AIDES					
HEXOSYLCERAN	n=1 (C14 hexosylceramide)	unchanged	unchanged	unchanged	
	n=2 (C16 hexosylceramide)	3.1x increase	3.2x increase	unchanged	
	n=3 (C18 hexosylceramide)	unchanged	2.1x increase	3.5x increase	
	n=4 (C20 hexosylceramide)	unchanged	unchanged	unchanged	
	n=5 (C22 hexosylceramide)	unchanged	unchanged	unchanged	
	n=6 (C24 hexosylceramide)	2.2x increase	2.1x increase	unchanged	
HINGOMYELINS					
	n=1 (C14 SPM)	unchanged	unchanged	unchanged	
	n=2 (C16 SPM)	unchanged	unchanged	unchanged	
	n=3 (C18 SPM)	unchanged	unchanged	unchanged	
	n=4 (C20 SPM)	not detected	not detected	not detected	
	n=5 (C22 SPM)	unchanged	unchanged	unchanged	
\mathbf{SP}	n=6 (C24 SPM)	unchanged	unchanged	unchanged	