Supplemental Material

Johswich et al., http://www.jcb.org/cgi/content/full/jcb.200801071/DC1



Figure S1. **Negative-mode MALDI-TOF-MS of S2 cell glycosphingolipids.** Glycosphingolipid extracts of nondepleted S2 cells were analyzed by negative-ion mode MALDI-TOF-MS in the reflectron mode using 2,5-dihydroxybenzoic acid as matrix for sample preparation. A list of newly registered masses and assigned glycosphingolipid structures compared with known GSL structures in *Drosophila* is given in Table S1. b, glycosphingolipid with a Cer composition of C14:1 tetradecasphingenine and C20:0 arachidic acid; a and c, glycosphingolipid with Cer masses that are 28 D (two methylene groups, C₂H₄) lower (a) or higher (c) than for b. Blue circles, glucose; green circles, mannose; yellow circles, galactose; yellow squares, GalNAc; blue squares, GlcNAc; white square, *N*-acetylhexosamine; PE, phosphoethanolamine; m/z, mass/charge.



Figure S2. **MALDI-TOF/TOF-MS fragmentation analysis of two zwitterionic glycolipid species containing lacdiNAc tandem repeats.** (A and B) Zwitterionic glycosphingolipid species with a hexasaccharide glycan moiety (mass/charge = 1,796) with one phosphoethanolamine (PE) modification (A) and the hexasaccharide glycan moiety (mass/charge [m/z] = 1,919) containing two phosphoethanolamine residues (B) were predicted to contain two hexose and four N-acetylhexosamine residues. Fragmentation patterns by MALDI-TOF/TOF-MS (in deprotonated form using 2,5-dihydroxy-benzoic acid as matrix) confirmed the presence of a tetra N-acetylhexosamine sequence with one or two phosphoethanolamine modifications, which can only be interpreted as a repeat of two lacdiNAc units. Blue circles, glucose; green circles, mannose; blue squares, GlcNAc.



Figure S3. **MALDI-TOF/TOF-MS analysis of two zwitterionic glycolipid species.** (A and B) Zwitterionic glycosphingolipid species with five N-acetylhexosamine residues (mass/charge [m/z] = 2,027; A) and six N-acetylhexosamine sugars (mass/charge = 2,487; B) were analyzed by MALDI-TOF/TOF-MS in a deprotonated form using 2,5-dihydroxybenzoic acid as a matrix. Fragmentation patterns confirmed the presence of an additional terminal N-acetylhexosamine residue compared with known structures present in *Drosophila*, probably a GalNAc residue in α -1,4 linkage in analogy to other *Drosophila* zwitterionic glycosphingolipid structures. Blue circles, glucose; green circles, mannose; yellow circles, galactose; yellow squares, GalNAc; blue squares, GlcNAc; white square, N-acetylhexosamine; PE, phosphoethanolamine.

lable S1. Newly registered zwifferionic glycosphingolipia specie	lable S1.	Tak	C	a	2	1		ļ	ł	k	k	0)	l	e	Э	;	S)	I		•			r	J	e	١	N)	1	J	r	e	g	jİ	S	i	e	ł	r	е	•			7	Z	V	N	/Ì		t	e	2	r	(r	Ţ	(C	1	g	J	١	/	C	¢)	S	F)	h	Ţ	ľ	1	g	I	3		Ĩ	P	J	(d	l	1	5	p)(е	C	Ĩ	¢	2	5	
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Zwitterionic GSL	Proposed structure	Registered mass
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Nz6*	GalNAcβ,4(PE-6)GlcNAcβ,3GalNAcβ,4GlcNAcβ,3Manβ,4GlcβCer, GalNAcβ,4GlcNAcβ,3GalNAcβ,4(PE-6)GlcNAcβ,3Manβ,4GlcβCer	1,796.3/1,824.4ª
Nz ₂ 6*	GalNAcß,4(PE-6)GlcNAcß,3GalNAcß,4(PE-6)GlcNAcß,3Manß,4GlcßCer	1,919.4
Nz7*	(HexNAc1-)GalNAcβ,4GlcNAcβ,3GalNAcβ,4(PE-6)GlcNAcβ,3Manβ,4GlcβCer	1,999.5/2,027.5°
Nz ₂ 7*	(HexNAc1-)GalNAcβ,4(PE-6)GlcNAcβ,3GalNAcβ,4(PE-6)GlcNAcβ,3Manβ,4GlcβCer	2,122.5
Nz9*	(HexNAc1-)GalNAcβ,4(PE-6)GlcNAcβ,3Galβ,3GalNAcα,4GalNAcβ,4(PE6) GlcNAcβ,3Manβ,4GlcβCer	2,364.7
Nz ₂ 9*	(HexNAc1-)GalNAcβ,4(PE-6)GlcNAcβ,3Galβ,3GalNAcα,4GalNAcβ,4(PE6) GlcNAcβ,3Manβ,4GlcβCer	2,487.7

GSL, glycosphingolipid. Zwitterionic glycosphingolipids were registered as [M–H]⁻ species and are alternative structures to those described by Seppo et al. (2000). °Glycosphingolipid with a Cer species with a 28-D higher mass, which reflects two additional methylene groups in the Cer moiety.

References Seppo, A., M. Moreland, H. Schweingruber, and M. Tiemeyer. 2000. Zwitterionic and acidic glycosphingolipids of the *Drosophila melanogaster* embryo. *Eur. J. Biochem.* 267:3549–3558.