

Supporting information for

**A facile method for monitoring sphingomyelin synthase activity in HeLa cells using liquid chromatography/mass spectrometry**

Punith M.Sundaraswamy<sup>1#</sup>, Yusuke Minami<sup>2#</sup>, Jayashankar Jayaprakash<sup>1#</sup>, Siddabasave Gowda B. Gowda<sup>1,3\*</sup>, Hiroyuki Takatsu<sup>4</sup>, Divyavani Gowda<sup>3</sup>, Hye-Won Shin<sup>4</sup>, and Shu-Ping Hui<sup>3\*</sup>

1. Graduate School of Global Food Resources, Hokkaido University, Kita-9, Nishi-9, Kita-Ku, Sapporo 060-0809, Japan
2. Graduate School of Health Sciences, Hokkaido University, Kita-12, Nishi-5, Kita-ku, Sapporo 060-0812, Japan
3. Faculty of Health Sciences, Hokkaido University, Kita-12, Nishi-5, Kita-ku, Sapporo 060-0812, Japan
4. Graduate School of Pharmaceutical Sciences, Kyoto University, Kyoto 606-8501, Japan

#Equally contributing authors

\*Correspondences:

1. Siddabasave Gowda B. Gowda\*

Faculty of Health Sciences, Hokkaido University,  
Kita-12, Nishi-5, Kita-ku, Sapporo 060-0812, Japan.

E-mail: [gowda@gfr.hokudai.ac.jp](mailto:gowda@gfr.hokudai.ac.jp)

2. Shu-Ping Hui

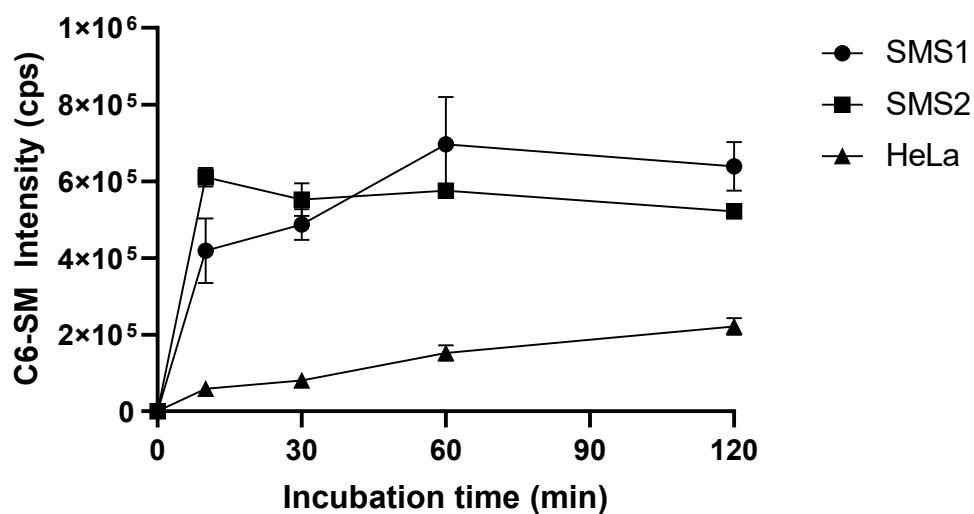
Faculty of Health Sciences, Hokkaido University,  
Kita-12, Nishi-5, Kita-ku, Sapporo 060-0812, Japan.

E-mail: [keino@hs.hokudai.ac.jp](mailto:keino@hs.hokudai.ac.jp) Tel: +81-11-706-3693.

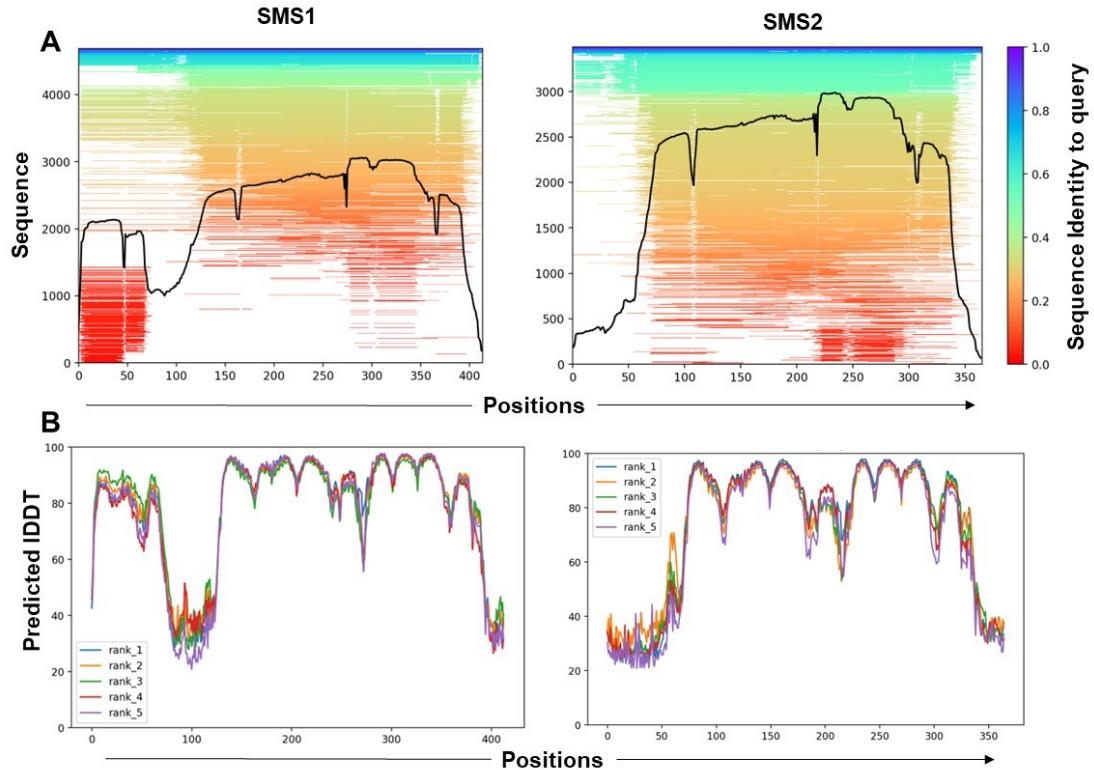
**Table S1.** Method validation for C6-Cer and C6-SM lipid species using cell lysates (n=5). LOD: limit of detection, LOQ: limit of quantification, RSD: relative standard deviation. Low concentration (Low): 5 fmol and High concentration (High): 50 fmol.

Lipids	Detection limits		Extraction recovery (%)		Matrix effect (%)		Intra-day (RSD, %)		Inter-day (RSD, %)	
	LOD (fmol)	LOQ (fmol)	Low	High	Low	High	Low	High	Low	High
C6-Cer	0.05	0.5	105.8	103.7	68.7	73.2	11.4	4.8	5.2	2.4
C6-SM	0.05	0.5	120.5	116.5	72.2	71.1	12.3	5.0	5.4	1.9

**Figure S1:** Production of C6-SM in control (HeLa), HeLa/SMS1, and HeLa/SMS2 over different incubation time at a protein concentration 0.1  $\mu$ g/  $\mu$ L (n=3).



**Figure S2:** (A) Comparison of sequences of SMS1 and SMS2 (B) Predicted Local Distance Difference Test (p lDDT) score of SMS1 and SMS2.



**Figure S3:** Root Mean Square Deviation (RMSD) of protein-ligand Complex of SMS1.

