### Acid Sphingomyelinase (aSMase) Activity Assay Kit

K-3200 (96 tests)

Support: echelon@echelon-inc.com

**Materials Provided** 

Catalog #	Description	Amount
K-3205	Standard	2 vials
K-3202	aSMase Substrate	2 vials
K-3203	Substrate Buffer	1 bottle
K-3204	Stop Buffer	1 bottle
	96 well plate	1 plate
	Microtiter Plate Seal	2 seals

#### Additional Materials Provided by User

- Fluorescence microtiter plate reader capable of reading at 360 nm excitation and 460 emission.
- 37 °C Plate Shaker / Incubator
- 70 °C Heat block

### Quick Protocol

Prepare standards and samples.									
Load standards and samples to the 96-well plate.									
Prepare the aSMase Substrate (K-3202).									
Add prepared aSMase substrate to the 96 well plate. Incubate for 3 hrs at 37 °C.									
Add Stop Buffer (K-3204) to the 96 well plate. Incubate for 30 min at room temperature.									
Read plate at 360 ex/460 em.									

#### <u>Storage</u>

Upon receipt, the kit should be stored at -20 °C. Under proper storage conditions, the kit components should remain stable for at least 6 months from date of receipt. Allow the reagents to warm to room temperature before opening vials. Substrate Buffer (K-3203) and Stop Buffer (K-3204) can be stored at room temperature after thawing.

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#### <u>Background</u>

Sphingomyelinase catalyzes the hydrolysis of sphingomyelin into ceramide and phosphoryl choline; and is involved in programmed cell death (apoptosis), cell differentiation and cell proliferation. Sphingomyelinases are classified into five categories: acid sphingomyelinase (aSMase), secretory sphingomyelinase (sSMase), neutral Mg2<sup>+</sup>-dependent sphingomyelinase (nSMase), neutral Mg2<sup>+</sup>-independent sphingomyelinase and alkaline sphingomyelinase. Acid sphingomyelinase was the first described and best characterized of the sphingomyelinases. A deficiency of lysosomal acid sphingomyelinase leads to rapid neurodegeneration and death due to excessive accumulation of sphingomyelin (Niemann-Pick disease).

#### <u>Assay Design</u>

Echelon's Acid Sphingomyelinase Activity Assay Kit uses a fluorogenic substrate, specific for Acid Sphingomyelinase, to provide a sensitive and homogenous method to measure the activity of aSMase in vitro from cell lysates or tissue homogenates. The kit provides all necessary reagents to measure the acid sphingomyelinase activity of 40 samples run in duplicate.

#### <u>Health Hazard Data</u>

The aSMase Substrate (K-3202) and Substrate Buffer (K-3203) contain highly toxic sodium azide and should be handled with caution. Sodium azide can be absorbed into the body by inhalation, ingestion and through the skin causing irritation to the eyes, skin and respiratory tract.

#### Assay Kit Notes

The assay is not compatible with some common lysis buffers decomponents (Table 1). Sonication or freeze-thaw protocols are precommended for preparing cell lysate samples. Avoid in non-compatible components if a lysis buffer is used. See support protocol for Cell lysis samples at the end of document. 1.

<u> Table 1, Incompatible</u>	<b>Buffer Components</b>
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Buffer Component	Concentration
EDTA	≥1 mM
EGTA	≥1 mM
Na+ Pyrophosphate	≥2.5mM
Glycerophosphate	N/A
Na+ vanadate	≥1 mM
Triton X-100	≥1.0% v/v
Deoxycholate	≥0.5% w/v
Igepal CA-630	≥1% v/v
SDS	≥0.1% v/v

 Cell lysis and Tissue homogenate samples should be titrated in the assay for optimal performance. As a starting point use 10 to 30 µg total protein per data point. Sample protein concentra-

<u>Table 2,</u>	Cell Types	and Lysis	<u>Methods</u>
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Lysis Method used	Cell Line # of Cells Used		Protein/well	aSMase Activity (pmol/hr/µg)	
Sonication	NIH 3T3	1 x 10 <sup>6</sup>	1.910 µg	21.753	
Sonication	MDA-MB-231	1.45 x 10 <sup>6</sup>	0.811 µg	28.569	
Sonication	MDA-MB-468	1.45 x 10 <sup>6</sup>	0.558 µg	17.842	
Freeze-thaw	MDA-MB-231	1.45 x 10°	2.834 µg	19.441	
Freeze-thaw	MDA-MB-468	1.45 x 10 <sup>6</sup>	4.833 μg	7.999	

tion should be adjusted depending on the acid sphingomyelinase activity within the sample.

- 3. Tested cell lines and the lysis method used can be found in Table 2.
- 4. Cell lysate and Tissue Homogenate samples should be prepared on the same day as the assay. Samples with freeze defrost cycles or prolonged storage have not been tested.
- 5. Stop Buffer is necessary for fluorescence detection.
- 6. Minimum of 25  $\rho \text{mol}$  hydrolyzed substrate is needed for fluorescence detection.
- 7. The plate can be read multiple times with no significant loss in signal.
- If a lower sensitivity is required (> 200 ρmole/hour) a 2 hour incubation is sufficient.
- Overnight incubation (17 hours) will result in greater sensitivity (< 3.125 ρmol/hour). However, higher coefficient of variation might occur.
- 10. To avoid matrix effects keep the buffer compositions between the standards and samples the same.
- 11. This assay can be adapted for use with 384 well plates. When working with 384 well plates, add 20 uL substrate, standards, and stop solution to each well at the respective steps.

#### Assay Procedure

Please read this entire section, Assay Notes, and relevant support protocols before beginning the assay. This protocol has been developed for duplicate reaction points. If singlet or triplicate points are required, the protocol will need to be adjusted accordingly.

- Turn on a plate shaker / incubator to 37 °C. Warm a heat block, with 1.5 mL tube block, to 70°C. Bring the Substrate Buffer (K-3203), Stop Buffer (K-3204), and Standard (K-3205) to room temperature before use. Place the aSMase Substrate (K-3202) on ice until use. Once defrosted, verify the Substrate Buffer (K-3203) and the Stop Buffer (K-3204) is clear. If precipitation is visible, heat at 37°C until clear. Bring back to room temperature before use.
- 2. Prepare the fluorescent standard curve by adding 400 μL of Substrate buffer (K-3203) to vial of room temperature Standard (K-3205). Vortex 10 sec. Place at room temperature. This is the 32.8 μM standard. This is stable at room temperature for at least 2 hours. Placing on ice may affect solubility. Serial dilute the 32.8 μM standard 2-fold, 6 times for a total of 7-concentrations using the table below.

**Table 3, Preparation of Fluorescent Standard** 

μΜ	pmol/well	μL of Substrate buffer (K-3202)					
32.8	1,640	150 uL of 32.8 μM standard					
16.4	820	150 uL of 32.8 μM standard	150 μL				
8.2	410	150 uL of 16.4 μM standard	150 μL				
4.1	205	150 uL of 8.2 µM standard	150 μL				
2.05	102.5	150 uL of 4.1 µM standard	150 μL				
1.025	51.25	150 uL of 2.05 μM standard	150 μL				
0.5125	25.625	150 uL of 1.025 μM standard	150 μL				
0	0		150 μL				

3. Prepare your samples using Substrate Buffer (K-3203). Please see assay notes for non-compatible buffers, matrix effects, and sample prep suggestions.

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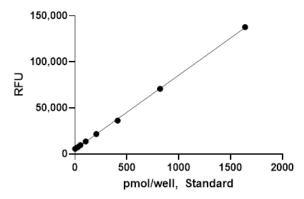
- Load 50 μL/well standards (step 2) or samples (step 3) to the provided 96-well plate using the suggested template as a guide (Table 4).
- Thaw the aSMase Substrate (K-3202) in heat block at 70 °C for 2 min. Mix well. If substrate is not clear, repeat until clear. Precipitation will severely reduce enzyme activity.
- Dilute the aSMase Substrate at 1:40 (80 μL per vial). For the entire plate, add 150 μL aSMase Substrate (K-3202) to 6 mL Substrate Buffer (K-3203). Mix well and keep at room temperature until use. This preparation is stable at room temperature for at least 5 hours.
- Add 50 μL/well of the diluted substrate (step 6) to the 96-well plate. Cover plate with acetate plate seal and incubate at 37 °C for 3 hours with shaking. See assay notes (7-9) for other incubation options.
- Add 50 μL/well Stop Buffer (K-3204) to the 96-well plate. Incubate for 30 minutes at room temperature with shaking. Protect from light. Read plate at 360 nm excitation and 460 nm emission.

#### **Quantification of Samples**

Generate a best fit curve for the fluorescent standards and interpolate relative sample values. We use Graphpad Prism software for sample analysis. The standard curve can be analyzed using a linear curve. Figure 1 shows a 7-point standard linear curve. Determine the activity of your sample by comparing the RFU of your sample to the fluorescent standard curve. For tissue homogenate and cell lysate samples the samples should be normalized with grams of total protein or tissue.

#### Figure 1

#### Acid Sphingomeylinase Activity Assay Flourescent Standard curve



	1	2	3	4	5	6	7	8	9	10	11	12
А	1640	1640	Sample									
~	pmol/well	pmol/well	1	1	9	9	17	17	25	25	33	33
в	820	820	Sample									
В	pmol/well	pmol/well	2	2	10	10	18	18	26	26	34	34
с	410	410	Sample									
L C	pmol/well	pmol/well	3	3	11	11	19	19	27	27	35	35
D	205	205	Sample									
	pmol/well	pmol/well	4	4	12	12	20	20	28	28	36	36
Е	102.5	102.5	Sample									
E	pmol/well	pmol/well	5	5	13	13	21	21	29	29	37	37
F	51.25	51.25	Sample									
Г	pmol/well	pmol/well	6	6	14	14	22	22	30	30	38	38
G	25.625	25.625	Sample									
6	pmol/well	pmol/well	7	7	15	15	23	23	31	31	39	39
н	0	0	Sample									
	pmol/ hr	pmol/ hr	8	8	16	16	24	24	32	32	40	40

Table 4, Suggested Detection Plate Layout

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### Support Protocol

#### Preparation of Cell Lysate

We suggest sonication or freeze-thaw protocols for lysing cells. These protocols are described in further detail below. Cell lysate buffers (see Assay Note 1) can interfere with the detection of aSMase activity. Prior to running the aSMase Activity Assay, total protein should be determined on each sample and the samples diluted to reflect the same concentration before addition to the assay.

#### Sonication:

- 1. Add 500  $\mu L$  1 mM PMSF & scrape cells.
- 2. Sonicate in ice water bath for 10 minutes 3 times.
- 3. Vortex between each sonication.
- 4. Centrifuge 10 mintues at 14000 x g.
- 5. Collect supernatant.
- 6. Follow steps 1-9 outlined in the Protocol: for the detection of aSMase Activity.

#### Freeze-thaw:

- 1. Add 500 μL 1 mM PMSF & scrape cells.
- 2. Freez-thaw 3 times in liquid nitrogen.
- 3. Vortex between each freez-thaw cycle.
- 4. Centrifuge 10 minutes at 14000 x g.
- 5. Collect supernatant.
- 6. Follow steps 1-9 outlined in the Protocol: for the detection of aSMase Activity.

#### Preparation of Tissue Homogenate

This protocol was developed using mouse brain tissue (WT and KO aSMase mice) by Isidora Rovic, M.Sc. at the University of Toronto, Faculty of Medicine; PI: Dr. Andrea Jurisicova. The samples were run in the aSMase Activity Assay at 7  $\mu$ g total protein. Prior to running the aSMase Activity Assay, total protein should be determined on each sample and the samples diluted to reflect the same concentration before addition to the assay. Since there are no protease inhibitors, samples should be kept on ice, at all times, until they are added to the activity assay.

- 1. In a 2ml Eppendorf tube place ~50mg of frozen brain tissue.
- Add 8x volume of ddH20 (ex. For 50 mg tissue add 400µl ddH20). Keep samples on ice.
- 3. Immediately homogenize tissue. Sonicate 3x for 15 seconds each, on medium-high power. Keep samples on ice.
- 4. Freeze-thaw the homogenate once on dry ice.

- Immediately begin sonication of tissue. Sonicate at medium-high power for 30 seconds, allow 10 second break, and resume for 30 more seconds (total 1 minute). Keep samples on ice. Longer sonication may overheat lysates.
- 6. After sonication place tissue on ice.
- Follow steps 1-9 outlined in the Protocol: for the detection of aSMase Activity. For mouse brain lysates, samples were diluted in the sample lysates to 14 µg total protein, and then diluted 1:1 in Substrate Buffer (K-3202) before adding to 96-well plate.

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Products	Catalog Number		Products	Catalog Number
Assays and Reagents			Lipids	
Neutral Sphingomyelinase Assay Kit	K-1800		N-Stearoyl Ceramide 1-phosphate	S-5018
Sphingomyelin Beads	P-B0SM		N-Biotin Ceramide 1-phosphate	S-500B
Ceramide Beads	P-BCer		N-NBD Ceramide 1- Phosphate	S-500N6
SphingoBeads Sample Pack	P-B00Ss		Biotin Sphingomyelin	S-400B

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