

NIH Public Access

Author Manuscript

J Neurosci Res. Author manuscript; available in PMC 2011 October 1

Published in final edited form as:

JNeurosci Res. 2010 October ; 88(13): 2940–2951. doi:10.1002/jnr.22438.

Inhibition of neutral sphingomyelinase-2 perturbs brain sphingolipid balance and spatial memory in mice

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Abstract

The sphingolipid ceramide is a bioactive signaling lipid that is thought to play important roles in modulating synaptic activity, in part by regulating the function of excitatory postsynaptic receptors. However, the molecular mechanisms by which ceramide exerts its effects on synaptic activity remain largely unknown. We recently demonstrated that a rapid generation of ceramide by neutral sphingomyelinase-2 (nSMase2; also known as sphingomyelin phosphodiesterase-3) played a key role in modulating excitatory postsynaptic currents by controlling the insertion and clustering of NMDA receptors (Wheeler et al. 2009). We now demonstrate that nSMase2 plays a role in memory. Inhibition of nSMase2 impaired spatial and episodic-like memory in mice. At the molecular level, inhibition of nSMase2 decreased ceramide, increased PSD-95, increased the number of AMPA receptors and altered the subunit composition of NMDA receptors. Our study identifies nSMase2 as an important component for efficient memory formation and underscores the importance of ceramide in regulating synaptic events related to learning and memory.

Keywords

Memory; NMDA; AMPA; Ceramide; Synapse; Sphingomyelinase

INTRODUCTION

It is becoming increasingly recognized that lipids play critical roles in regulating neuronal excitability. The biophysical properties of lipids regulate shape, endo- and exocytotic events, vesicle and protein trafficking and a considerable number of lipid species also function as second messengers that regulate a multitude of neuronal events (Day and Kenworthy 2009; Owen et al. 2009; Stahelin 2009; Swartz 2008). Considering the potential importance of lipids in regulating neuronal functions, relatively little is known about the mechanisms by which lipids control synaptic events.

Sphingolipids are a class of lipid especially enriched in the central nervous system. The sphingolipid ceramide is a bioactive lipid that modulates a variety of cellular events ranging

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from proliferation to apoptosis (Claus et al. 2009; van Blitterswijk et al. 2003). Ceramide can be rapidly generated in focal compartments by the actions of a family of sphingomyelin catabolizing enzymes called sphingomyelinases. There are three general classes of sphingomyelinase, which based on their pH optima are categorized as neutral, acidic and alkaline sphingomyelinases. Three neutral sphingomyelinases (nSMase2-4) have been identified and cloned (Hofmann et al. 2000; Mizutani et al. 2000; Schneider and Kennedy 1967; Tomiuk et al. 1998). The nSMase2 is highly enriched in the neurons of hippocampus and dentate gyrus (Hofmann et al. 2000; Rao and Spence 1976), suggesting it may play a role in hippocampal dependent memory. Indeed, several lines of evidence suggest that ceramide generated by nSMase2 is important for regulating synaptic activities related to memory. Nerve growth factor (NGF)-induced neurite outgrowth and synaptogenesis are regulated by nSMase2 and its reaction product ceramide (Brann et al. 1999; Hirata et al. 2001; Ito and Horigome 1995), and NGF induced activation of nSMase2 can rapidly enhance the rate of depolarization-induced action potentials (Nicol 2008; Zhang and Nicol 2004; Zhang et al. 2002). Recently we demonstrated that $TNF\alpha$ -induced enhancement of excitatory post-synaptic currents and NMDA-evoked calcium bursts were regulated by nSMase2-mediated generation of ceramide (Wheeler et al. 2009). Similar effects at postsynaptic sites have been produced by direct additions of synthetic cell-permeable ceramide analogs that increase excitatory postsynaptic currents without affecting pairedpulse facilitation (Coogan et al. 1999; Fasano et al. 2003; Furukawa and Mattson 1998). These ceramide-associated enhancements of excitatory currents are often transient and are followed by sustained depression of excitatory postsynaptic currents (Coogan et al. 1999; Davis et al. 2006; Furukawa and Mattson 1998; Tabarean et al. 2006; Yang 2000). Combined, these findings support complex roles for sphingolipids in regulating neuronal excitability that may in part depend on the spatial and temporal production of ceramide. Despite these physiological and biochemical findings that suggest ceramide generated by nSMase2 may be important for synaptic events related to learning and memory, a role for nSMase2 in memory has not been established. Here we provide evidence that nSMase2 is important for spatial and episodic-like memory in mice and uncover roles for nSMase2 in the constitutive regulation of brain ceramide metabolism and postsynaptic structure.

EXPERIMENTAL PROCEDURES

Animals

Mice with a deletion in the gene encoding nSMase2 were kindly provided by Dr. Christophe Poirier. These mice exhibit a complete loss of nSMase2 activity as described previously (Aubin et al. 2005). Since mutation in nSMase2 produces dwarfism and defects of osteogenesis that could interfere with behavioral training (Aubin et al. 2005; Stoffel et al. 2005; Stoffel et al. 2007), the use of these mice was restricted to biochemical analysis of brain sphingolipid content. Since behavioral testing required 14 consecutive days of training, it was not practical to administer GW4869 directly into brain, as this number of repeated injections is very likely to induce local inflammation and tissue damage. Therefore, we used the nSMase2 specific inhibitor GW4869 (Haughey et al. 2004; Kolmakova et al. 2004; Marchesini and Hannun 2004; Peng et al. 2006; Smith et al. 2006; Takahashi et al. 2006; Walton et al. 2006). Male C57BL/6J mice (4 months old) were injected intraperitoneal once daily with GW4869 (1.25 mg/kg, dissolved in saline with 2.5% dimethylsulfoxide, Calbiochem, Gibbstown, NJ) 7 days prior and during behavioral testing for a total of 21 daily injections. An experimental timeline is included in supplementary Fig 1. In preliminary experiments we determined that this dose of GW4869 does not produce any adverse effects, and did not alter the sphingomylein or ceramide content of heart, skeletal muscle or liver (Supplementary Fig 2) and does enter into brain in sufficient quantity to inhibit neutral sphingomyelinase activity (Fig 1). Control mice were-injected with the

vehicle used to dissolve GW4869 (2.5% dimethylsulfoxide). All animals were housed in groups of 5 and maintained on 12 hr light-dark cycle with free access to water and rodent chow. Behavioral training consisted of the following memory paradigms. A classic version of the Morris water maze, Repeated Reversals task, radial water maze and Y maze. On the last day of training all animals were sacrificed and brains rapidly removed. Brains were dissected with a sagittal incision down the center line and one half was immersed in 4% paraformaldahide and stored at 4°C for 5 days, followed by post-fixation in 30% sucrose for 48 hours. The remaining tissue was rapidly dissected into cortex, hippocampus, striatum, cerebellum and snap-frozen. Frozen brain tissues were stored at -80°C. All rodent procedures were conducted according to NIH guidelines for animal care and approved by the JHU Institutional Animal Care and Use Committee.

Pretraining

Mice were handled daily for 5 days to acclimate them to investigator handling. On the sixth day, mice were pre-trained in a small pool (diameter 55 cm) with a platform $(10 \times 10 \text{ cm})$ submerged 1 cm below the water surface as previously described (Savonenko et al. 2005). Mice were placed in the water facing the platform and allowed to swim, climb on the platform and stay there for 10 sec. Mice were then removed from the platform, dried and returned to a dry waiting cage for 10–12 min. A total of 6 trials were performed during the pre-training session.

Classic Morris water maze and Repeated Reversals

A fixed platform version of the water maze paradigm was carried out as previously described (Savonenko et al. 2005). Briefly, mice were trained in a 100 cm diameter pool filled with opaque water ($21 \pm 2^{\circ}$ C) and surrounded by various visual cues. A collapsible platform (10 \times 10 cm) was submerged 1 cm below the water surface in one of the four quadrants and remained in the same position until the end of the training. The starting location for each trial was randomly altered among 3 quadrants. The maximum duration of each trial was 60 sec and mice that failed to find the platform during this time were guided to the platform and then removed. Mice were given 12 trials a day for 5 consecutive days including one probe trail at the beginning and a second probe trial at the end of each training session. The platform was completely submerged during probe trials and the mouse was allowed to swim freely for a variable time interval (30-45 sec). During probe tests, the percent of time spent in annuli (area surrounding the platform, 20 and 40 cm in diameter) was recorded. At the end of the probe trial the platform was raised and the mouse was allowed to escape onto the platform to avoid extinction due to repeated probe trials. Performance in all tasks was recorded using an HVS video tracking system (HVS Image Analysis VP-200; Hampton, UK). After completion of a classic Morris water maze, mice were trained in a Repeated Reversals paradigm for 2 days as previously described (Savonenko et al. 2005). This task differs from the classic water maze by changing the platform location at the start of each day. In this paradigm, mice are required to inhibit memory for the old platform location and learn the new location. Latency (time spent to locate the hidden platform from the start location) and swim speed (average speed during a trial) were measured during the platform trials. In the probe trials, percentage of time spent in the area 20 cm in diameter around the platform was recorded (Annulus 20).

Radial water maze

The radial arm water maze task was conducted as previously described (Savonenko et al. 2005). A six-arm radial maze made of clear plastic was placed into the pool and the platform was submerged at the end of one arm. The platform location was randomly changed between different arms at the beginning of each training day. The start position was located at the end of one arm and was randomly chosen in every trial. Mice were given 6 trials a day for 4

consecutive days. The maximum duration for each trial was 120 sec and mice that did not find the platform were guided to the platform by experimenter, allowed to stay on the platform for 10 sec and then moved to the waiting cage. Latency to find the platform and the number of correct arm entries where the platform was located on the previous day were registered and analyzed.

Y Maze

The Y maze test was conducted in a wooden elevated Y shaped maze according to previously established protocols (Melnikova et al. 2006) with minor modifications. Mice performed a two-trial Y maze task where one arm was blocked during the first trial using a stable barrier. Mice were placed at the end of one arm and allowed to explore the maze for 5 min and the sequence of arm entries was recorded. Following a 30 min rest interval, trial 2 began by placing mice at the end of one arm with all three arms opened and available for exploration. A two-trial Y-maze task represents a classic test for spatial recognition memory. Mice with intact memory have a tendency to explore novel arm more often than the rest of the maze while mice with memory impairment enter in all three arms evenly (Conrad et al. 1996; Dellu et al. 1997). The number of visits to the novel arm was calculated for 5 min duration of testing. Motor activity was assessed on the basis of total number of arms visited during each trial. Spontaneous alternation behavior was calculated as the number of triads containing entries into all three arms divided by the maximum number of possible alternations.

Sphingomyelinase activity assay

The Amplex Red sphingomyelinase assay kit (Molecular Probes, Inc., Eugene, OR) was used to quantify acid- and neutral sphingomyelinase activities according to the manufacturer's protocols. Assays were conducted using 50 μ g of cortical tissue. Cortical homogenates were incubated for 30 min at 37°C in 100 μ l of working solution that consisted of 2U/ml HRP, 0.2 U/ml choline oxidase, 8 U/ml of alkaline phosphatase and 0.5 mM sphingomyelin. Neutral sphingomyelinase activity was determined in working buffer with a pH of 7.4. Acidic sphingomyelinase activity was measured by first adjusting the pH of the working solution to 5.0 during incubation, then increasing the pH to 7.4 to allow for detection of the product with the Ampex Red reagent. The reaction product was measured using a fluorescent plate reader (absorption at 571 nm and emission at 585 nm; SpectraMax M2e, Molecular Devices, CA). Positive controls consisted of sphingomyelinase (40 U/ml) or H₂O₂ (20 mM) added to the working solution.

Immunoblotting

For Western blots, cortical tissues were homogenized in ice-cold buffer containing 20 mM Tris-HCl (pH 7.4), 50 mM NaF, 0.32 M sucrose, 2 mM EDTA, 2 mM EGTA, 0.2 mM sodium orthovanadate, 1 mM PMSF and protein inhibitor cocktail (Roche diagnostics GmbH, Germany). Proteins (10 μ g of total protein per lane) were separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis and transferred to nitrocellulose membranes. Membranes were blocked in Tris-buffered saline (TBS) containing 10 mM Tris-HCl [pH 7.4], 150 mM NaCl and 0.1% Tween 20 (TBS-T) for 1 hr and incubated with primary antibodies overnight at 4°C. Monoclonal antibodies included PSD-95 (1:5000, Oncogene, Cambridge, MA), synaptophysin (1:1000, Sigma, St. Louis, MO), NR1 (1:1000, Calbiochem, Gibbstown, NJ), NR2A (1:1000, Calbiochem, Gibbstown, NJ), GluR1 (1:1000, Millipore, Billerica, MA), GluR2 (1:1000, Cell signaling technology, Danvers, MA) and β -actin (1:1000, Sigma, St. Louis, MO). Following incubations with primary antibodies, membranes were washed in TBS and TBS-T and incubated at room temperature for 1 hr with HRP-conjugated secondary antibodies (1:5000, Goat anti-mouse IgG; 1:5000, Goat

anti-rabbit IgG, Promega, WI). Immunoreactivity was visualized using Pico West SuperSignal (Pierce, Rockford, IL) and optical densities were quantified using NIH ImageJ software.

Immunohistochemistry and immunofluorescent staining for free floating sections

Coronal sections (30 µm) were washed in TBS three times and incubated in 1% hvdrogen peroxide for 30 min to block endogenous peroxidase activity. Tissue sections were then washed three times in T-TBS and blocked for 1 hr in 5% normal goat serum. Rat anti-mouse neutrophil primary antibody was added (1:1000, Serotec, Raleigh, NC) and incubated overnight on shaking platform at 4°C. Tissue sections were then washed in T-TBS and incubated in biotinylated secondary antibody (1:300, Bio-Rad, Hercules, CA) for 1 hr at room temperature and incubated for 30 min in ABC kit reagent mixture in order to form biotin-avidin complexes (Vectastain ABC kit, Vector laboratories, Burlingame, CA). Tissue sections were then briefly rinsed and stained with DAB according to the manufacturer's directions (Vector laboratories, Burlingame, CA). Stained sections were mounted on slides and cover slipped with Permount clear mounting media (Sigma, St. Louis, MO). For immunofluorescent staining the following primary antibodies were used: polyclonal anti-Iba (1:1000, Wako chemicals, Richmond, VA), polyclonal anti-GFAP (1:1000, Dako, Carpinteria, CA). Seconday antibidies were Alexa fluor 488 and Alexa fluor 495 (both 1:500, Invitrogen, Carlsbad, California). Images were obtained using a Zeiss microscope equipped with a Retiga 2000R Fast 1394 camera (QImaging, Surrey, BC, Canada). Fluorescent intensity was quantified using Openlab software (Improvision, Waltham, MA).

Lipid extraction and measurement of sphingolipids using LC-ESI/MS/MS

Total lipids were extracted according to a modified Bligh and Dyer procedure (Shaikh 1994) as described previously (Bandaru et al. 2007). Each sample was homogenized in 10 volumes of deionized water, then in 3 volumes of 100% methanol containing 30 mM ammonium acetate . The mixture was vortexed, four volumes of chloroform added, vortexed and centrifuged at 1,000g for 10 minutes. The chloroform layer was removed, dried and stored at -80°C until use. Samples were re-suspended in 100% methanol for LC/MS/MS. Sphingomyelins and ceramides were detected and quantified using multiple reaction monitoring (MRM). Samples were injected using a PAL autosampler into a PerkinElmer HPLC equipped with a C18 column (100×2 mm, 5 µm) coupled to guard column containing identical packing material (Phenomenex, Torrance, CA). For a typical run, the LC column was first pre-equilibrated for 0.5 min with the first mobile phase consisting of 85% methanol, 15% H₂O, and 5 mM ammonium formate. The column was then eluted with the second mobile phase consisting of 99% methanol, 1% formic acid, and 5 mM ammonium formate at the flow rate of 100.0 μ l/min. The eluted sample was injected into the ion source and the detection and quantitation of each analyte was carried out by LC/MS/MS in MRM mode monitoring the parent compound and products by ion scan. Slight differences in the efficiency of extraction and fluctuations in the efficiency of mass detection were controlled for by normalizing data to the internal standards sphingomyelin C12:0 and ceramdie C12:0 (Avanti Polar Lipids. Alabaster, AL). Sphingomyelin and ceramide standards C16:0, C18:0, C18:1 were purchased from Sigma. Ceramides C20:0, C24:0, C24:1 were purchased from Avanti Polar Lipids (Alabaster, AL). Palmitoyl-lactosyl ceramide C16:0-C16:0, stearoyl-lactosyl-ceramide C16:0-C18:0, lignoceryl-glucosylceramide C16:0-C24:0, lignoceryl-galactosyl-ceramide C16:0-C24:0, and stearoylgalactosyl-ceramide-sulfate C18:1-C24:0 were purchased from Matreya Inc. (Pleasant Gap, PA). Data were collected and processed using Analyst 1.4.2 software.

Statistical analyses

Behavioral data were analyzed using repeated measures or main effect ANOVAs with a minimal level of significance of p < 0.05. Post-hoc Tukey tests were applied to significant differences in repeated measures ANOVAs. One-way ANOVA and non-paired t-tests were used to assess differences in sphingolipid and synaptic protein levels.

RESULTS

Inhibition of neutral sphingomyelinase-2 disrupts brain ceramide-sphingomylein balance

Genetic models of mutation or deletion in nSMase2 result in defects of bone formation (Aubin et al. 2005; Stoffel et al. 2005; Stoffel et al. 2007) and were consequently not appropriate for behavioral testing that requires intact locomotor activity. Therefore, we employed a pharmacological approach using GW4869, a selective inhibitor of nSMase2 (Luberto et al. 2002). We first determined if systemically administered GW4869 entered brain in sufficient amounts to inhibit nSMase2. We found that GW4869 decreased neutral sphingomyelinase activity in cortex by 64.3 % (Fig. 1A). Treatment with GW4869 did not affect acidic sphingomyelinase activity (Fig. 1B), consistent with data from previous reports that this drug specifically targets nSMase2 (Haughey et al. 2004; Kolmakova et al. 2004; Marchesini and Hannun 2004; Peng et al. 2006; Smith et al. 2006; Takahashi et al. 2006; Walton et al. 2006). Since nSMase2 catalyzes the hydrolysis of sphingomyelin to ceramide, we next determined if inhibition of nSMase2 by GW4869 affected brain sphingolipid content. Treatment with GW4869 decreased multiple species of ceramide and increased multiple species of sphingomyelin in all brain regions examined (Fig. 2A-H), with the exception of sphingomyelin C16:0, which was decreased in all brain regions (Fig. 2A, C, E, G). These results are strikingly similar to those obtained by analysis of sphingomyelin and ceramide content in brain tissues from mice containing a deletion mutation in the nSMase2 gene (Supplementary Fig. 2), further confirming the specificity of GW4869 for nSMase2. There were no significant changes of ceramide or sphingomyelin levels in liver, kidney, heart or skeletal muscle of mice treated with GW4869 compared with control mice administered vehicle (data not shown). Unexpectedly, we found that treatment with GW4869 altered the sphingolipid content of serum in a direction opposite from what we observed in brain tissue. In serum, treatment with GW4869 increased multiple ceramides and decreased multiple sphingomyelins with few exceptions (Fig. 3A, B). Based on evidence that ceramide can play important roles in the activation of monocytes (Detre et al. 2006; Sipka et al. 2000), we next we next determined if GW4869 altered the trafficking of monocytes into brain parenchyma or activated brain resident glia. GW4689 did not alter the number or morphology of Iba-1 immunopositive marcophage/microglia, GFAP immunopositive astrocytes or the number of immunopositive neutrophils (Supplementary Fig. 3), indicating that inhibition of nSMase2 did not alter the CNS trafficking of bloodderived monocytes or activation state of brain resident glia.

Inhibition of nSMase 2 impairs early phases of spatial reference memory

Spatial reference memory was tested in a classic version of the Morris water maze beginning 7 days after the initiation of GW4869 treatment. Inhibition of nSMase2 with GW4869 did not affect daily averaged latencies to locate the hidden platform. There was a significant effect of day, indicating that both groups progressively decreased latencies to locate the hidden platform with repeated training (Fig. 4A). There were no group differences in swimming speed (Fig. 4A inset). A trial-by-trial analysis of the first training day showed that while vehicle treated mice progressively decreased latencies to locate the hidden platform, mice treated with GW4869 failed to decrease latencies to locate the platform with repeated trials (Fig. 4B). These differences were not due to an effect of treatment on swimming speed (determined for each trial during the first day of training; data not shown).

An analysis of the time spent in quadrants (target, adjacent left, adjacent right and opposite) during the probe test at the end of the first training day indujcated that mice treated with GW4869 spent less time searching in the target quadrant compared with mice administered vehicle. Furthermore, mice treated with GW4869 spent less time in the area surrounding the hidden platform (Annulus 20) compared with mice administered vehicle (Fig. 4C,D). Both groups of mice spent equal time in the target quadrant (Annulus 20) during the final probe trial performed at the end of the fifth training day (Fig. 4C,E). These data indicate that inhibition of nSMase2 delayed the formation of spatial reference memory in mice.

After 5 days of training in a classic Morris water maze task, mice were trained in a Repeated Reversals paradigm for 2 days to test episodic-like memory. In this setting, the hidden platform location was changed each day and mice were required to quickly learn the new platform location while inhibiting memory for the previous location. Performance in the Repeated Reversals paradigm was not different in mice treated with GW4869 compared with vehicle controls (data not shown), suggesting that inhibition of nSMase2 does not affect episodic-like memory in mice.

To assess episodic-like working memory, a six-arm radial maze enclosure was placed into the pool and the platform location was changed daily so that mice were required to learn a new platform location every day. Latencies to locate the hidden platform during trial 1 of all 4 training days were similar in both groups (Fig. 4F). Treatment with GW4869 significantly increased latencies to locate the hidden platform during trials 2–5 across all 4 training days (Fig. 4F). To confirm that inhibition of nSMase2 interfered with episodic-like memory, we next analyzed the number of correct arm entries. The number of correct arm entries during trial 1 of training day 1 (defined as the arm corresponding to the quadrant where the platform was located on the last day of the Repeated Reversal task), were similar in both groups and were not different from chance (chance level is 16%; Fig. 4G). On training day 2, mice treated with vehicle demonstrated long-term memory for the previous episode of training by visiting the correct arm 44% of the time, while mice treated with GW4869 only visited the correct arm 25% of the time; a frequency not significantly different from chance. These data indicate that inhibition of nSMase2 interfered with the formation of episodic-like working memory.

We next used a Y maze task to assess spatial recognition memory. Performance on this task was not different in mice treated with GW4869 compared with mice administered vehicle (data not shown), suggesting that spatial recognition memory was not altered by inhibition of nSMase2.

Long-term inhibition of nSMase2 disrupts synaptic structure

Since synaptic organization is important for learning and memory, we measured several structural and functional synaptic proteins to determine if inhibition of nSMase2 and dysregulation of brain sphingolipid balance modified synapses. Treatment with GW4869 increased the post-synaptic protein PSD-95, without affecting levels of the pre-synaptic protein synaptophysin (Fig. 5A, B). GW4869 did not alter levels of the NMDA receptor subunit NR1, indicating that inhibition of nSMase2 did not alter the total number of NMDA receptors. In mice treated with GW4869 there were increases in NR2A subunits but not NR2B, suggesting that inhibition of nSMase2 modified the subunit composition of NMDA receptors (Fig. 5C, D). The AMPA receptor subunit GluR1 was increased in mice treated with GW4869 indicative of an increase in the total number of AMPA receptors. Levels of AMPA receptor subunits GluR2 and GluR3 were similar in mice treated with GW4869 and mice administered vehicle (Fig. 5E, F). Combined these data suggest that dysregulation of brain sphingolipid balance following inhibition of nSMase2 modified synaptic structure by increasing PSD95, altering the subunit composition of NMDA receptors and increasing the

number of AMPA receptors. These synaptic changes may account for defects of memory following long-term inhibition of nSMase2.

DISCUSSION

In this study we demonstrate that nSMase2 is important for hippocampal-based memory. At the molecular level, inhibition of nSMase2 increased sphingomylein and decreased ceramide content in brain, increased PSD-95 levels, increased the number of AMPA receptors and modified the subunit composition of NMDA receptors at postsynaptic sites in the hippocampus. Combined, these findings suggest that impairment of memory by inhibition of nSMase2 involves a disruption in sphingomyelin/ceramide balance and dysregulation of postsynaptic glutamatergic signaling.

It is interesting that systemic administration of GW4869 did not alter the ceramide or sphingomylein content of liver, heart or skeletal muscle but did decrease the ceramide content and increase the sphingomyelin content in brain. These findings suggest that nSMase2 may have a constitutive activity in CNS, but is largely an inducible enzyme in peripheral tissues. These drug-induced changes in brain sphingolipid content were extremely similar to those observed in mice with a deletion mutation in nSMase2 gene indicating that GW4869 effectively and specifically targeted nSMase2. The effects of GW4869 on sphingomyelin and ceramide content were most striking in the hippocampus, consistent with the high regional expression of nSMase2 in this brain structure (Hofmann et al. 2000). These findings not only indicate that GW4869 is an effective inhibitor of brain nSMase2, but further suggest that in addition to the inducible actions of this sphingolipid-modifying enzyme (Clarke and Hannun 2006), nSMase2 may be involved in the constitutive regulation of ceramide metabolism in brain.

Inhibition of nSMase2 with GW4869 slowed learning. Mice administered GW4869 did not progressively decrease latency to locate the hidden platform with repeated training trials, suggesting that they had difficulty learning to use spatial cues to navigate the pool. During the first probe test, mice administered GW4869 spent less time in the target area (measured by annulus surrounding the platform) further demonstrating that inhibition of nSMase2 interfered with spatial learning. Starting at day 2 of probe trials, latencies to locate the hidden platform in mice administered GW4869 were similar to vehicle, suggesting that inhibition of nSMase2 slows, but does not abolish learning. Similarly, mice administered GW4869 were able to use the spatial navigation memory acquired in the classic Morris Water Maze in subsequent repeated reversals tasks. In this more challenging procedure, earlier locations of the platform need to be disregarded and memory retrieval must be selective for the most recently encoded location; an 'episodic-like' component of the task. The repeated reversals task has close similarities to working-memory protocols in which the animal acquires information that is only useful for a given period (Steele and Morris 1999). One possibility why inhibition of nSMase2 did not interfere with performance on repeated reversal tasks is that nSMase2- dependent plasticity is important for procedural learning that is intrinsic to the acquisition of the Morris Water Maze (such as learning to use a platform as an escape location), but not critical for episodic-like memory required for repeated reversals tasks. A second possibility is that prior learning with the Morris Water Maze influenced later learning on repeated reversal tasks. Indeed, previous studies have shown that pre-training mice in water maze substantially ameliorated AP5-induced memory deficit despite a complete blockade of LTP (Bannerman et al. 1995; Nakazawa et al. 2004).

Following repeated reversals tasks mice were trained in radial arm water maze. A delay in learning on day 2 of training in the radial arm water maze, similar to what was observed in the classic Morris Water Maze was apparent in mice administered GW4869. Compared with

classic water maze, the radial arm water maze has different geometry and is a more challenging task that invokes a broad range of synaptic activation within hippocampal neuronal networks, that may not involve networks not active during classic water maze or repeated reversals tasks (Schwegler et al., 1996; Crusio and Schwegler, 2005). These findings corroborate observations in the classic water maze that inhibition of nSMase2 slows, but does not abolish spatial learning. We cannot however rule out the possibility that nSMase2 is also important for other types of memory.

Sphingolipids are thought to play important roles in neuronal communication by modulating synaptic strength and connectivity. The nSMase2 belongs to a family of enzymes that cleave the phosphodiester linkage of sphingomyelin to create ceramide and phosphocholine. The rapid kinetics and location of nSMase2 make it a likely candidate to be involved in the modulation of plasticity in the CNS. nSMase2 can be rapidly activated with peak activity in 1-2 minutes and is highly expressed in central nervous system where expression is prominent in large neuronal cells including, Purkinje cells, pyramidal cells, neurons of the dentate gyrus granular laver, and pontine nuclei (Hofmann et al. 2000). At the presynaptic terminal, a rapid generation of ceramide is thought to enhance the rate of synaptic transmitter release by facilitating the trafficking and fusion of synaptic vesicles with the plasma membrane (Bajjalieh et al. 1989; Rogasevskaia and Coorssen 2006; Rohrbough et al. 2004). Preventing rapid ceramide generation by inhibition of sphingomyelinases blocks calcium-induced glutamate release in cerebellar neurons (Numakawa et al. 2003). At the postsynaptic membrane, ceramide appears to play complex roles in modulating neuronal excitability. Ceramide analogs and the endogenous generation of ceramide have been shown to rapidly enhance excitatory postsynaptic currents (Fasano et al. 2003; Furukawa and Mattson 1998; Wheeler et al. 2009; Yu et al. 1999; Zhang et al. 2002), which is often followed by sustained depression of synaptic currents (Coogan et al. 1999; Davis et al. 2006; Furukawa and Mattson 1998; Tabarean et al. 2006; Yang 2000). It has also been demonstrated in vivo that intraperitoneal administration of the ceramide analog PDMP, restored synaptic function and preserved spatial memory in ischemic rats (Inokuchi et al. 1997; Yamagishi et al. 2003). Although the exact molecular mechanisms for the effects of ceramide on neuronal excitability are not yet fully understood, we recently demonstrated that a rapid production of ceramide by nSMase2 was required for TNFα-induced insertion of NMDA receptors into post-synaptic membranes and enhancement.

One mechanism by which sphingolipids may regulate synaptic plasticity is by controlling the location and trafficking of receptors at postsynaptic sites. Sphingomyelin, ceramide, cholesterol and the ganglioside GM1 are key components of specialized membrane domains known as lipid rafts. These highly ordered regions of membrane are thought to play important roles in receptor signaling by compartmentalizing membrane proteins into active focal signaling units and promoting cross-talk between different signaling cascades. In neurons, lipid rafts are important sites for docking and insertion of NMDA and AMPA receptors (Besshoh et al. 2005; Fullekrug and Simons 2004; Wheeler et al. 2009). Disrupting lipid rafts by cholesterol depletion has been shown to reduce NMDA receptor currents and associated calcium flux and increased the basal internalization rate of AMPA receptors (Abulrob et al. 2005; Frank et al. 2004; Hering et al. 2003). Furthermore, pharmacological inhibition or a deletion mutation in nSMase2 prevented TNFa-induced insertion of NMDA receptors into lipid rafts, suggesting that some aspects of receptor trafficking may be regulated by enzymatic control of lipid composition. Our current findings further suggest that nSMase2 is an important component of the neuronal synapse. Long-term inhibition of nSMase2 increased sphingomyelin and decreased ceramide content in multiple brain regions (most notably in hippocampus) and modified synaptic structure. In this setting, PSD-95 increased, the number of AMPA receptors increased and the subunit composition of NMDA receptors altered. PSD-95 is a PDZ-domain containing scaffolding protein that is almost

exclusively located in the postsynaptic density of neurons. The amino terminal region of PSD-95 contains three PDZ domains that are necessary and sufficient to localize PSD-95 to lipid rafts. PSD-95 is an essential synaptic adapter protein that is involved in anchoring NMDA and AMPA receptors. Overexpression of PSD-95 enhances dendritic spine maturation, increases the amplitude and frequency of AMPA receptor-mediated EPSCs and converts silent synapses into functional synapses by driving GluR1 into synapses (Beique and Andrade, 2003; Ehrlich and Malinow, 2004; Stein et al., 2003; Husseini et al., 2000). The PDZ1 domain of PSD-95 also binds to the NR2A subunit of NMDA receptors and is involved in increased insertion and decreased internalization of intact NMDA receptors (Lin et al. 2004). Interaction of PSD-95 and NR2A also regulates downstream signal transduction pathways including CREB interaction with CRE promoter sites on target genes to modulate transcriptional events related to neuronal connectivity and excitability (Deisseroth et al., 1999; Aarts et al., 2002; Gardoni et al., 2006; Sornarajah et al., 2008; Chen et al., 2006). Thus, it is likely that increased PSD-95 was directly associated with increases of GluR1 and NR2A and that these molecular events were involved in the perturbation of memory.

Combined, our findings suggest that nSMase2 is important for memory, by actions that involve regulation of brain sphingolipid metabolism and postsynaptic glutaminergic signaling.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

The authors would like to thank Dr. Christophe Poirier for providing mice with a deletion mutation in neutral sphingomyleinase-2 and Fang Yang for her technical assistance.

These studies were funded in part through NIH grants MH077542 and AA0174078 to NJH and AG24984 to HS and NJH.

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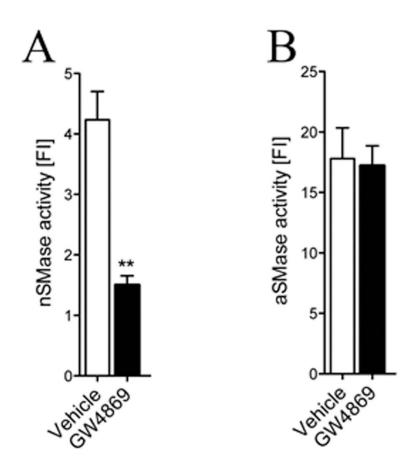
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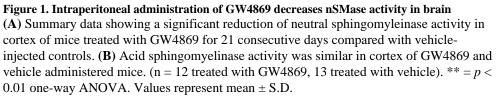
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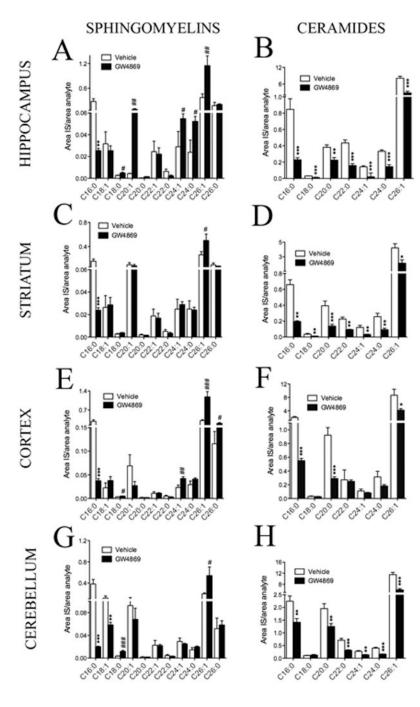


Figure 2. Inhibition of nSMase2 modifies brain sphingolipid content

Brain levels of most sphingomylein species were increased and most ceramides species decreased following long-term inhibition of nSMase2. Summary data showing the effects of inhibiting nSMase2 on sphingomyelin and ceramide content of (**A**, **B**) hippocampus, (**C**, **D**), striatum, (**E**, **F**) cortex and (**G**, **H**) cerebellum (n = 12 treated with GW4869, 13 treated with vehicle). * = p < 0.05; ** = p < 0.01 decreased compared with control, # = p < 0.05; ## = p < 0.01 increased compared with control. ANOVA with Tukey post hoc comparisons. Values represent mean \pm S.D.

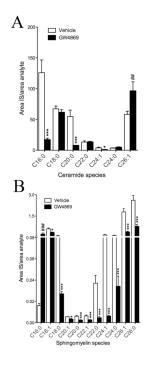


Figure 3. Inhibition of nSMase2 modifies serum levels of ceramide and sphingomyelin Long-term inhibition of nSMase2 decreased ceramide and increased sphingomylein in serum. Figure shows the effects of inhibiting nSMase2 on serum (**A**) ceramides and (**B**) sphingomyelins. (n = 12 treated with GW4869, 13 treated with vehicle)* = p < 0.05; *** = p < 0.001 greater than vehicle, ## = p < 0.01; ### = p < 0.001 less than vehicle. Two-way ANOVA with Tukey post hoc comparisons.

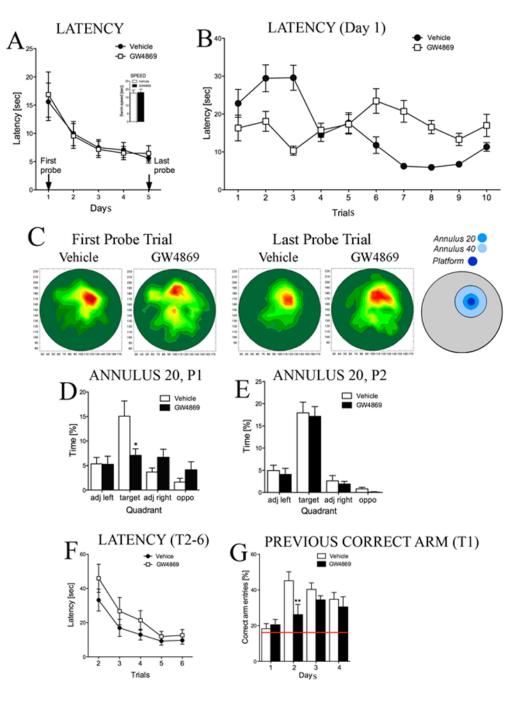


Figure 4. Inhibition of nSMase2 disrupts an early phase of memory

Inhibition of nSMase2 impaired spatial reference and episodic-like memory. (A) Averaged daily latencies to locate the hidden platform in a classic version of Morris water maze were similar in mice treated with GW4869 or vehicle [Two-way repeated measures (Treatment x Day) ANOVA, effect of Treatment F(1, 90) = 0.05, p > 0.8; Day F(4, 90) = 26.6, p < 0.001 and Treatment-by-Day interaction F(4, 90) = 0.27, p > 0.8]. Inset shows that treatment with GW4869 did not affect swimming speed (average of swim speed over all 5 days of training is shown) (t test, p > 0.5). (B) Latencies to locate the hidden platform for each trial of training on day 1 showing that inhibition of nSMase2 impaired the ability of mice to learn the location of the hidden platform. [Two-way (Treatment x Trial) ANOVA for latency on

the first day of training revealed significant effects of Treatment F(1, 230) = 2.49, p < 0.05,Trial F(9, 230) = 2.22, p < 0.05 and Treatment-by-Trial interaction F(9, 230) = 3.63, p < 0.050.001]. (C) Occupancy plots for the probe trials performed at the end of training day 1 and training day 5. Increasing color intensity indicates the region where mice spent the most time during probe trials. Inhibition of nSMase2 with GW4869 significantly decreased the time mice spent in the region surrounding the platform (Annuli 20 and 40) during the first probe trial. Although mice treated with GW4869 were not as accurate in remembering the platform location during the last probe trial, the time they spent in Annulus 20 was not statistically different from mice treated with vehicle. Annuli representing the area of 20- and 40-cm diameters around the platform are indicated in concentric blue circles. (**D**, **E**) Quantitative data showing the time spent in Annulus 20 during the first (P1) and last (P2) probe trials [Two-way (Treatment x Quadrant) ANOVA for the first probe test showed significant effect of Treatment F(1, 92) = 5.35, p < 0.01, Quadrant F(3.92) = 10.73, p < 0.01, p < 0.010.001 and Treatment-by-Quadrant interaction F(3, 92) = 7.79, p < 0.001, Tukey post hoc for Annulus 20, p < 0.01, [Two-way (Treatment x Quadrant) ANOVA for the last probe test did not show significant effect of Treatment F(1, 92) = 0.59, p > 0.4 while there was an effect of Quadrant in both groups F(3,92) = 60.03, p < 0.001 and Treatment-by-Quadrant interaction was not significant F(3, 92) = 0.002, p > 0.9, Tukey post hoc for Annulus 20, p >0.9]. (F) Averaged latencies for trials 2–6 across 4 days of training in a radial water maze. Mice treated with GW4869 showed longer latencies to locate the hidden platform during trials 2–5 across the 4 days of training [Three-way ANOVA revealed a significant effect of Treatment F(1, 23) = 7.01, p < 0.05 and Trial F(4, 92) = 53.80, p < 0.001]. (G) Summary data of correct arm entries in the radial water maze (a correct entry is recorded as entry into the arm where the platform was located on the previous day of training). On training day 2, mice treated with vehicle demonstrated long-term memory for the previous episode of training by visiting the correct arm more often than a chance level (chance level of 16% is indicated with a red line). Mice treated with GW4869 were unable to remember the previous platform location (ANOVA, p < 0.01; Tukey post hoc, p < 0.01).

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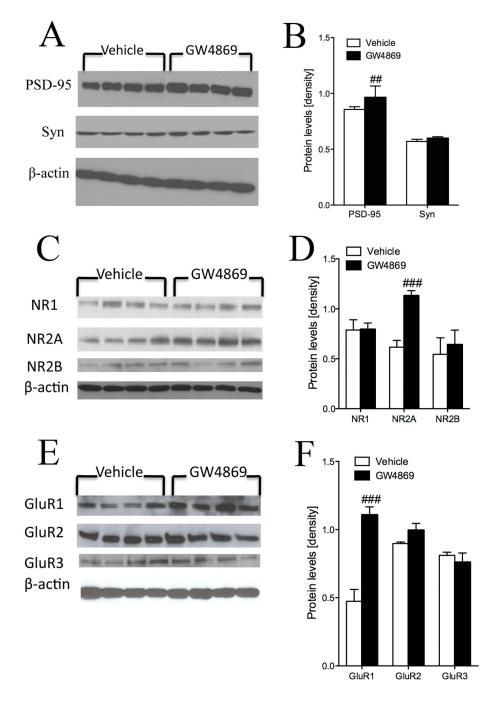


Figure 5. Perturbation of brain sphingolipid content disrupts synaptic integrity

(A, C, E) Representative western blots showing the effects of long-term inhibition of nSMase2 on PSD-95, synaptophysin, NMDA and AMPA receptor subunits. (B, D, F) Quantification of western blot band intensities showing that inhibition of nSMase2 with GW4869 increased PSD-95 levels, increased the amount of NMDA receptor NR2A subunits and AMPA receptor GluR1 subunits. Quantitative data are the mean \pm S.D. from 12 animals/group. ## =p < 0.01; ### =p < 0.001 greater than vehicle. Students t-test.