



# <u>Sample preparation for Permethylated GSL profiling</u> <u>by MALDI/TOF-MS and NSI-MSn</u>

# Date written: 10/8/18

### Authors: Kazuhiro Aoki\* and Mayumi Ishihara

Complex Carbohydrate Research Center, University of Georgia, Athens, GA 30602 \*To whom correspondence should be addressed; Kazuhiro Aoki (<u>kaoki@ccrc.uga.edu</u>)

### **Reagents:**

- Methanol (MeOH)
- α-dihyroxybenzoic acid (DHBA)
- HPLC Water
- Sodium Hydroxide Solution, 50% w/w
- Iso-propanol (2-PrOH)
- n-propanol (1-PrOH)

## **Equipment/Apparatuses:**

- Glass culture tubes (round bottom)
- 500 uL Microcentrifuge tube or glass inserts for GC autosampler vial
- GC glass autosampler vial (conical bottom)
- 1.5 mL Microcentrifuge tube
- Glass pipette
- Plastic pipette
- Glass syringe
- Centrifuge
- Vortex

### **Procedures:**

## MALDI/TOF-MS analysis

MALDI/TOF-MS analysis of permethylated glycosphingolipids is performed in the linear or reflector positive ion mode using  $\alpha$ -dihyroxybenzoic acid (DHBA, 20mg/mL solution in 50% methanol: water) as a matrix. The procedures are described in detail below.

1. Prepare Matrix solution; 20mg α-dihyroxybenzoic acid (DHBA) in 1mL of 50% methanol: water in a 1.5 mL Microcentrifuge tube

- 2. Dissolve the permethylated sample with 20~μL (Adjust the volume depends on your sample amount) of methanol using a clean glass micro syringe (rinsed with Methanol).
- 3. Take  $1 \sim 2 \mu L$  of sample solution using a clean glass micro syringe (rinsed with Methanol), and then transfer into the bottom of a small vial.
- 4. Wait until the solution volume becomes approximately  $1\mu L$  (~3 min).
- 5. Take  $1\mu$ L of matrix solution using a micro pipet and then add into the sample solution.
- 6. Immediately mix them by pipetting the solution several times.
- 7. Take  $1\mu$ L of mixture and then put on a well of a MALDI plate. Immediately, take another 1ul of mixture then put on another well.
- 8. Wait until the mixtures on the plate dry.

### Note:

- The matrix solution is preservable at -20°C. Defrost the matrix solution before use.
- A good crystallized sample is necessary to get a strong signal. However, making a good crystal is not always possible, hence, we recommend making at least two crystals per sample.

### NSI-MSn analysis

Mass analysis by NSI-MS are performed by direct infusion of permethylated glycosphingolipids dissolved in infusion buffer the instrument at a constant flow rate of  $0.4 \sim 1 \mu L/min$  via a syringe pump. The procedures are described in detail below.

- 1. Prepare infusion buffer solution: Methanol/2-Propanol/1-Propanol/1 mM NaOH (16:3:3:2 by volume)
- Dissolve the permethylated GSLs with 20~µL (Adjust the volume depends on your sample amount) of methanol using a clean glass micro syringe (rinsed with methanol).
- 3. Take 10  $\mu$ L of sample solution using a clean glass micro syringe (rinsed with methanol), and then transfer into a small vial (rinsed with methanol).
- 4. Add known concentration of heavy MeI permethylated Dp4 as a reference standard if quantification is desired (Optional).
- 5. Take  $40 \sim 65 \mu$ L of the infusion buffer and then add to the sample solution.
- 6. Screw the cap, mix them by vortexing, and briefly centrifuge the vial.









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