

## AMDIS GC/MS Analysis Report

Data: AKS2300.RAW

Library: C:\Program Files (x86)\NISTMS\AMDIS32\libraries\GDM\GMD.MSL

Number of Identifications: 25

| <u>RT(min)</u> | <u>Chemical Name</u>   |
|----------------|--|
| 5.7235         | ? M000413_A105003-101-xxx_NA_1031<br>,31_TRUE_VAR5_ALK_Pyridine, 2-hydroxy- (1TMS) (ID#:NA)<br>RI = 1031.6 RI-RI (lib) = 0.3       |
| 6.0700         | ?? contamination - pyridine/BSTFA (ID#:JCB665-N1012)<br>RI = 1052.4 RI-RI (lib) = -6.6   |
| 7.4758         | ?? M000000_A114002-101-xxx_NA_1131<br>,27_TRUE_VAR5_ALK_NA114002 (classified unknown) (ID#:NA)<br>RI = 1135.2 RI-RI (lib) = 4.0    |
| 9.7261         | M000075_A129001-101-xxx_NA_1262<br>,42_TRUE_VAR5_ALK_Phosphoric acid (3TMS) (ID#:10497-05-9)<br>RI = 1264.8 RI-RI (lib) = 2.4      |
| 10.7773        | ? M000000_A136002-101-xxx_NA_1332<br>,55_TRUE_VAR5_ALK_similar to Lumichrome (2MeOX) (ID#:NA)<br>RI = 1326.4 RI-RI (lib) = -6.1    |
| 13.4964        | ? cis - linalool oxide (furanoid) (ID#:AKS142~1-N1002)<br>RI = 1496.7 RI-RI (lib) = 3.0  |
| 13.8517        | cis - linalool oxide (pyranoid) (ID#:AKS142~1-N1006)<br>RI = 1519.7 RI-RI (lib) = 3.0  |
| 14.1218        | M001212_A155003-101-xxx_NA_1541<br>,53_PRED_VAR5_ALK_similar to Diterbutylphenol (1TMS) (ID#:NA)<br>RI = 1537.2 RI-RI (lib) = -4.3 |
| 18.0201        | ?? contamination_blank (ID#:TAM757-N1008)<br>RI = 1809.6 RI-RI (lib) = -0.4  |
| 18.1055        | ?? contamination_blank (ID#:TAM757-N1010)<br>RI = 1816.3 RI-RI (lib) = -0.5  |
| 18.7104        | ?? M000606_A188004-101-xxx_NA_1863<br>,13_TRUE_VAR5_ALK_Fructose (1MEOX) (5TMS) BP (ID#:NA)<br>RI = 1863.8 RI-RI (lib) = 0.6       |
| 18.8335        | ? RI=1874.2, 18.8427 min AKS2296 (ID#:AKS2296-N1002)<br>RI = 1873.4 RI-RI (lib) = -0.8   |

19.0343            ?? M000633\_A189001-101-xxx\_NA\_1885  
                 ,22\_TRUE\_VAR5\_ALK\_Mannose (1MEOX) (5TMS) BP (ID#:NA)  
RI = 1889.2                      RI-RI(lib) = 4.0

19.5386            IS - 13C6-Sorbitol (ID#:ARP436-N1002)  
RI = 1928.8                      RI-RI(lib) = 6.2

19.8047            M000801\_A196011-101-xxx\_NA\_1945,61\_TRUE\_VAR5\_ALK\_Gallic  
                 acid (4TMS) (ID#:2078-17-3)  
RI = 1949.7                      RI-RI(lib) = 4.1

19.8089            M000801\_A196011-101-xxx\_NA\_1945,61\_TRUE\_VAR5\_ALK\_Gallic  
                 acid (4TMS) (ID#:2078-17-3)  
RI = 1950.0                      RI-RI(lib) = 4.4

20.9469            ? contamination - pyridine/BSTFA - C16:0 acid TMS (ID#:JCB665-  
                 N1044)  
RI = 2042.1                      RI-RI(lib) = -0.8

21.4118            ?? M000060\_A209002-101-xxx\_NA\_2080  
                 ,23\_TRUE\_VAR5\_ALK\_Inositol, myo- (6TMS) (ID#:NA)  
RI = 2081.2                      RI-RI(lib) = 1.0

23.2076            ? contamination - pyridine/BSTFA - C18:0 acid TMS (ID#:JCB665-  
                 N1048)  
RI = 2240.7                      RI-RI(lib) = 1.5

23.2130            ? contamination - pyridine/BSTFA - C18:0 acid TMS (ID#:JCB665-  
                 N1048)  
RI = 2241.2                      RI-RI(lib) = 2.0

27.1469            ?? M000044\_A264001-101-xxx\_NA\_2622  
                 ,87\_TRUE\_VAR5\_ALK\_Sucrose (8TMS) (ID#:NA)  
RI = 2626.6                      RI-RI(lib) = 3.7

27.3416            ? RI=2648.0, 27.3456 min AKS2298\_230228141932 (ID#:AKS229~1-  
                 N1008)  
RI = 2647.6                      RI-RI(lib) = -0.4

29.3721            ?? M000832\_A289005-101-xxx\_NA\_2865  
                 ,56\_TRUE\_VAR5\_ALK\_Catechin (5TMS) (ID#:NA)  
RI = 2871.0                      RI-RI(lib) = 5.4

31.7158            ?? contamination\_blank (ID#:TAM757-N1014)  
RI = 3141.0                      RI-RI(lib) = 7.4

31.7224            ?? contamination\_blank (ID#:TAM757-N1014)  
RI = 3141.7                      RI-RI(lib) = 8.1

## QA/QC:

Instrument type: Ion Trap

Scan Direction None

Highest m/z detected = 799, high m/z setting = 800

High noise level. Median Signal(Noise Level)/Threshold=5.6.

Background (low vs. high retention time):

median low RT S/N=34, high RT S/N=107

Solvent Tailing (m/z 84). Run begins at 5.00 min. Solvent falls below:

S/N=2 before run

S/N=1 at 5.17 min.

Column Bleed (m/z 207):

median low RT S/N=3, high RT S/N=13

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