



Alignment spectra
selected from this
sample treatment class

Manually looked at
individual file from this
class and ID'ed

Name: Methioninesulfoxide; [M+]
RT similarity score: 999.9999
Accurate mass similarity score: 999.6436
Dot product similarity score: 915.0206
Reverse dot product similarity score: 997.7859
Total score: 971.0627

