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# Development of a Stability-Indicating Reverse-phase HPLC Method for M1 Peptide Extracts

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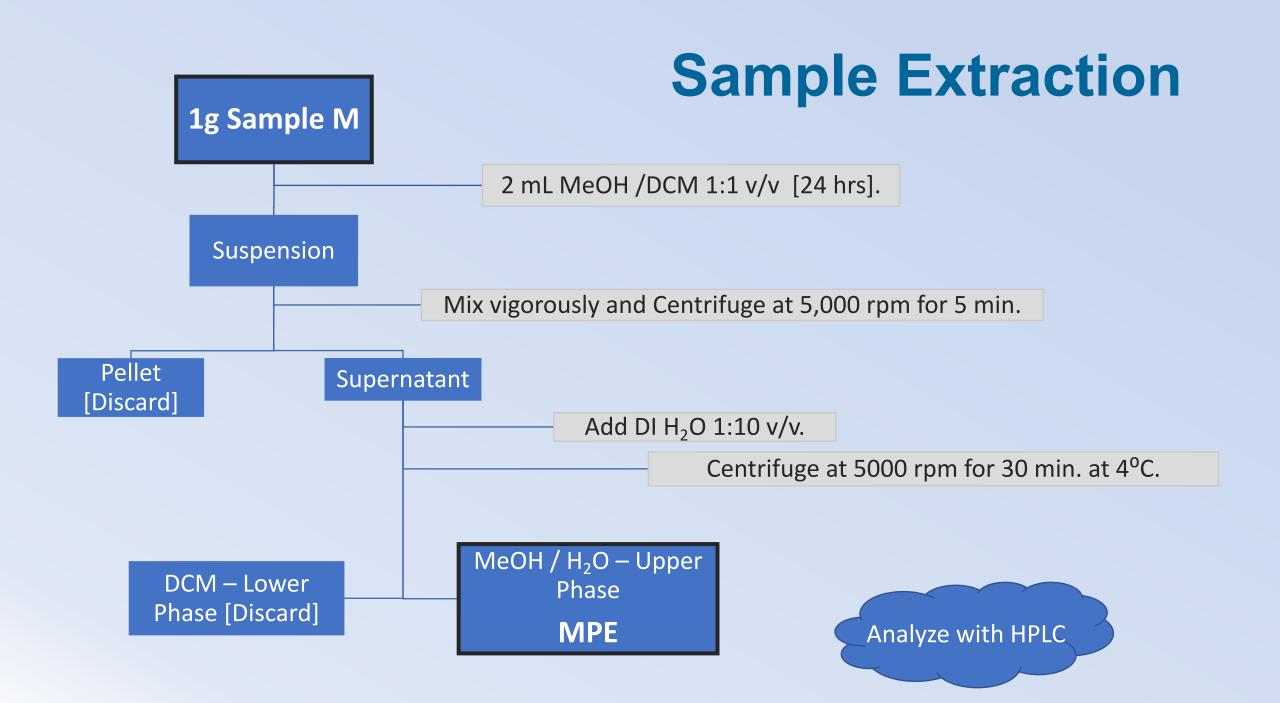
## Introduction

- Cosmonauts are exposed to, and experience radiation induced DNA damage during space travel.
- Tiwari (2019) found this type of DNA damage to be one of the most significant health hazards radiation can create.
  - Necessary to monitor DNA stability during and after travel.
- Recent novel nuclear matrix metalloprotease (nMMP) identification
  - In need of a reliable analysis method.
  - Helena (2018) discusses nMMP presence can be an indicator of both DNA damage and subsequent repair processes.

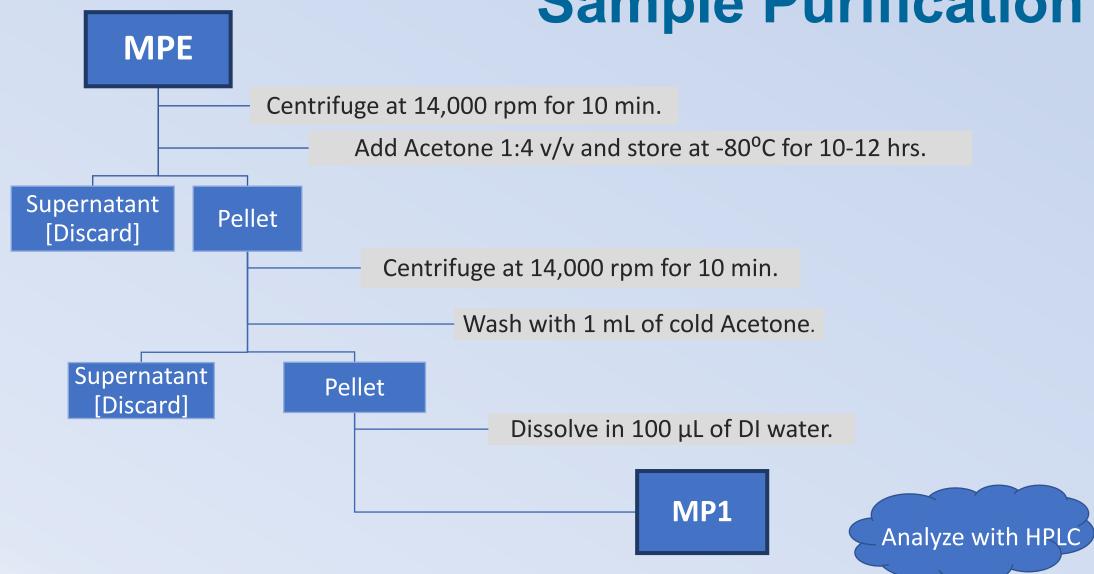
### Introduction



- We have undertaken development of a dependable High-Pressure Lipid Chromatography (HPLC) analyzation method.
  - Our preliminary results are outlined here.
- Only after an acceptable HPLC method is obtained and verified, then sample can be analyzed.
  - Then further research on physicochemical characterization and functional identification of sample M can be done.



# Sample Purification



# **HPLC Method Development**

- Began with literature to get an idea of the HPLC method steps that would work for our sample.
- Gradient elution method as shown in Table 1 below.
- First analyze sample M at the crude state then at the purified state.
- Before any HPLC is ran the lines are first washed, then the column itself is washed.
- A blank sample (DI Water) is then injected at the beginning to equilibrate the system.

#### **HPLC** method details:

Flow rate: 0.5 mL/min Injection volume: 10 μL

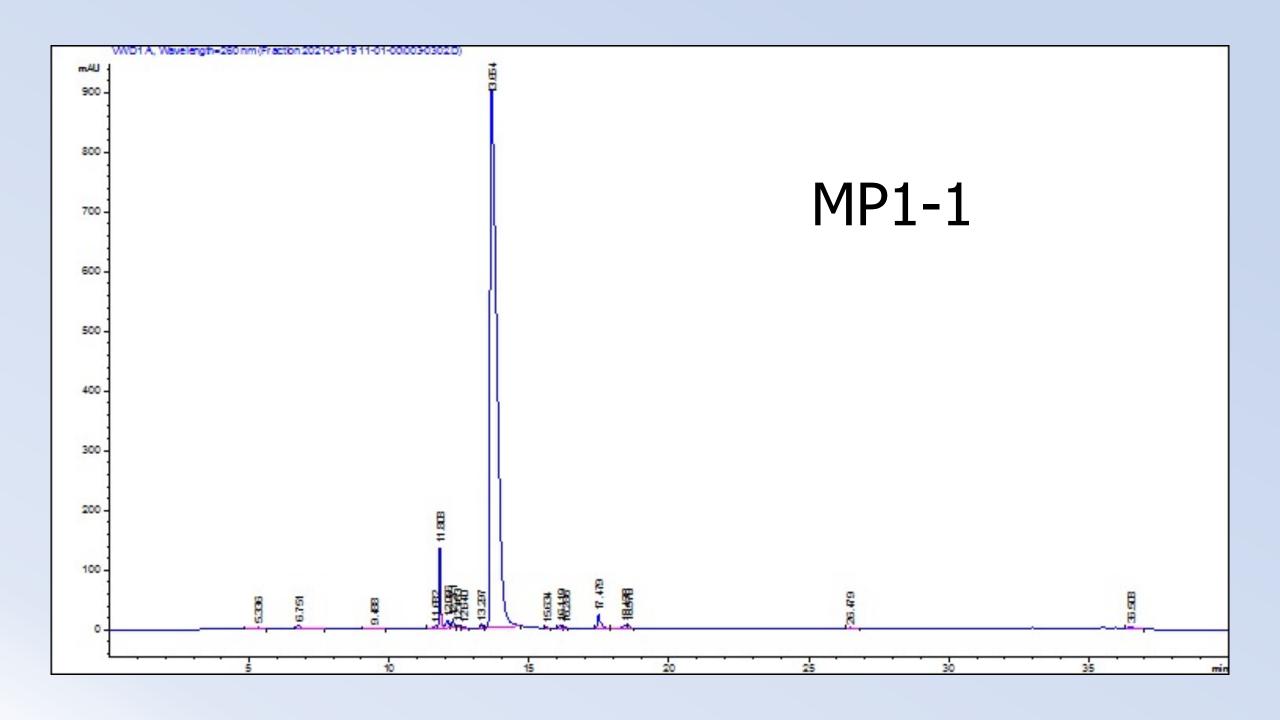
Detection: 260 nm

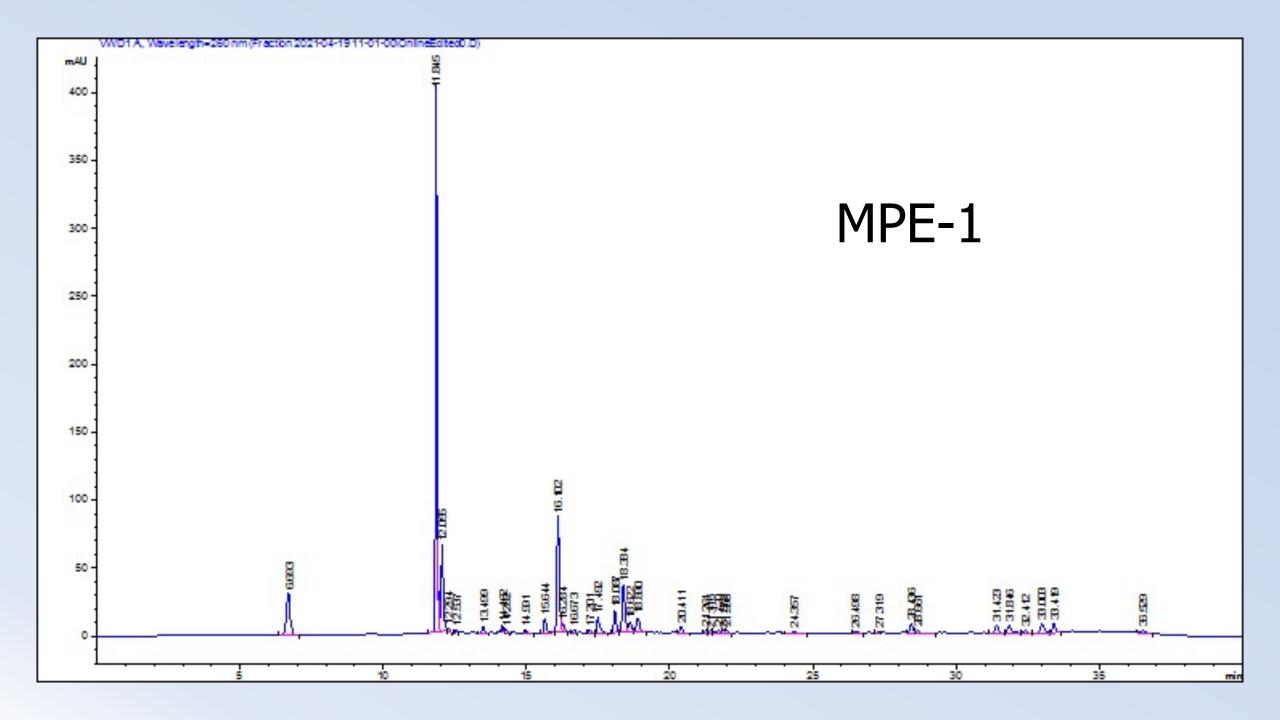
C18 Column

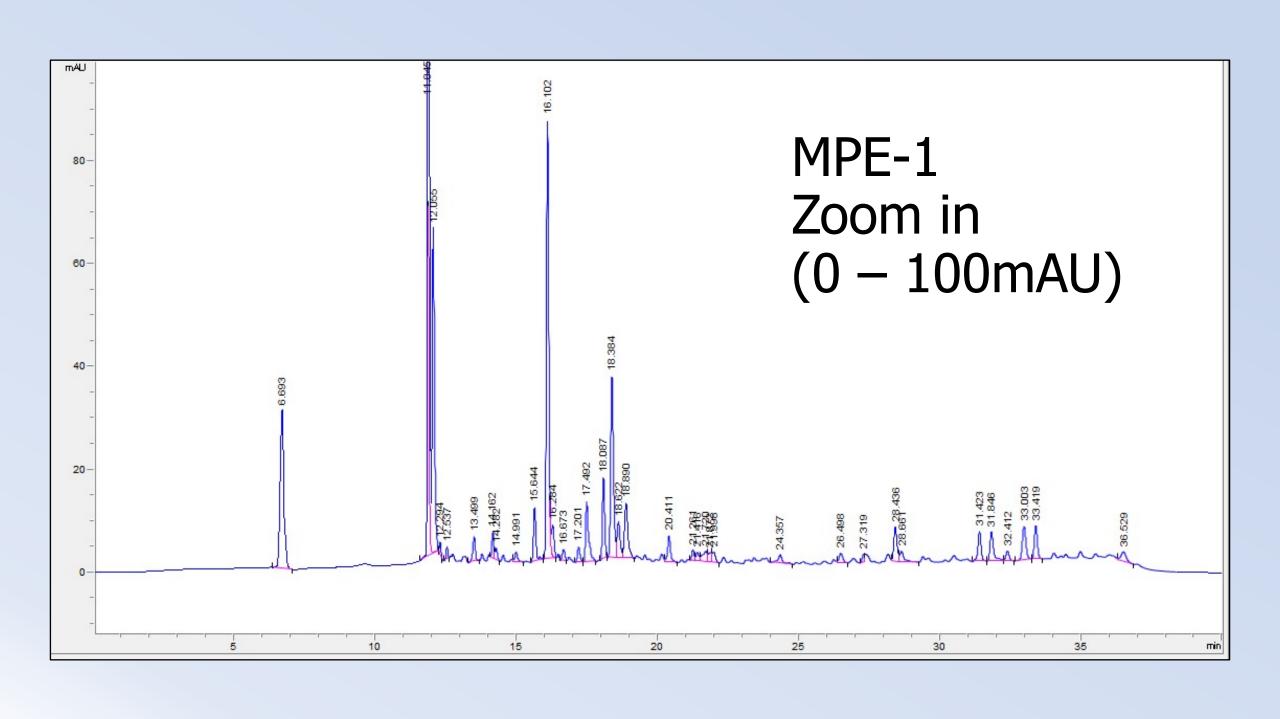
Each sample is run for 30 min. with a 10 min wash of 100% 0.1% formic acid to follow

Total run time: 40 min.

Table 1: Representative HPLC Method		
Time (minutes)	0.1% Formic Acid (%)	Acetonitrile (%)
0	100	0
30	100	0
35	20	80
40	100	0







# **Discussion**

- Repeated HPLC analysis has shown similar elution profiles.
  - Indicate that sample M is relatively pure as there is only 1 major peak.
  - Indicates a lack of stability of the sample throughout the purification process.
- Analysis of sample after 1 week were also performed.
  - Samples saved at room temperature and −20°C and analyzed with HPLC, all showing similar elution profiles.

# **Future Studies**

- After preliminary results of the HPLC method there are new questions to answer.
  - Mass spectrometry could provide further indication of degradation as well as knowing the mw of each peak would help in identification.
  - Introductory use of the HPLC to analyze the sample after a week
    of degradation are promising and the stability of the sample would
    also be a possible pathway of further study.

# Acknowledgements

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## References

- 1.Craik, D.J., Troeira Henriques, S., Mylne, J.S., Wang, C.K. (2012). Cyclotide Isolation and Characterization. *Methods in Enzymology, 156,* 37-62. <a href="https://doi.org/10.1016/B978-0-12-394291-3.00024-1">https://doi.org/10.1016/B978-0-12-394291-3.00024-1</a>
- 2.Helena, J., Margaretha, A., Grobbelaar, S., Magalena Nolte, E., Nel, M., Sean Pepper, M., Coetzee, M., and Mercier A.E. (2018). Deoxyribonucleic Acid Damage and Repair: Capitalizing on Our Understanding of the Mechanisms of Maintaining Genomic Integrity for Therapeutic Purposes. *International Journal of Molecular Sciences*, 19, 1148. 10.3390/ijms19041148.
- 3.Tiwari, V., Wilson III, D.M. (2019). DNA Damage and Associated DNA Repair Defects in Disease and Premature Aging. *The American Journal of Human Genetics*, *105*, 237-257. <a href="https://doi.org/10.1016/j.ajhg.2019.06.005">https://doi.org/10.1016/j.ajhg.2019.06.005</a>.