

Compound Identification Process for GC-MS Non-Targeted Analysis of JUUL Aerosol using a Custom Mass Spectral Library

Juul Labs Science

Lena N. Jeong¹, Michael R. Noë², Niti H. Shah²,
Saibal Chakraborty², John Miller IV², I. Gene Gillman¹

1. Juul Labs, Inc.
2. Altria Client Services, LLC.

Introduction

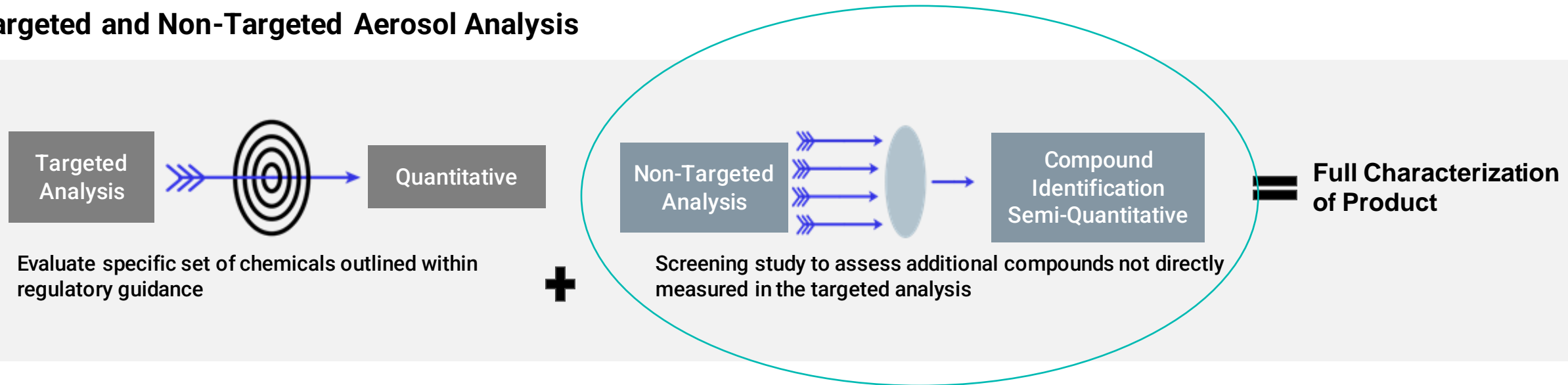
PMTA guidance for ENDS recommends:

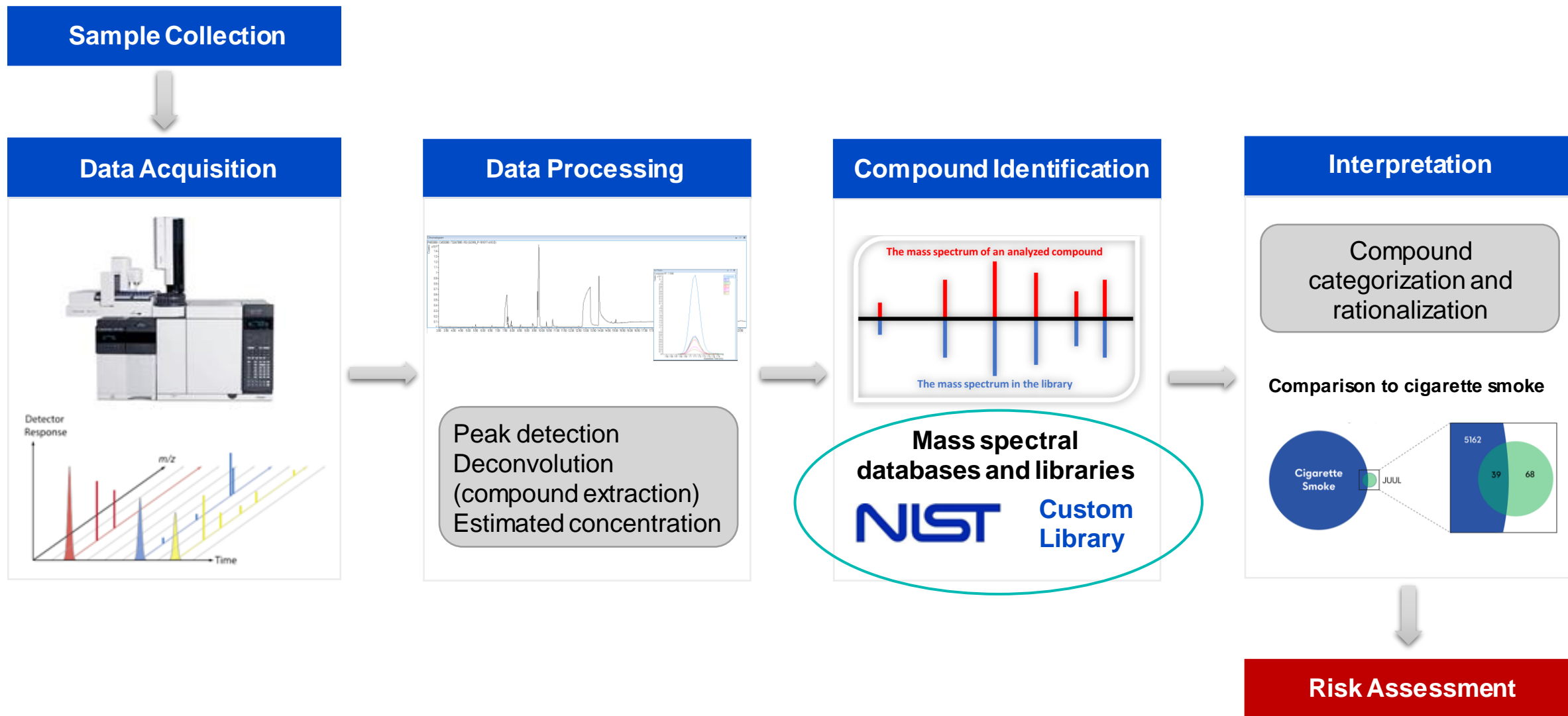
“a list of uniquely identified constituents or chemicals, including ... other toxic chemicals contained within the product or delivered by the product to be provided”

Objective:

- Full Chemical characterization of the ENDS aerosol
- Determination of aerosol constituent emission levels
- Risk assessment

Targeted and Non-Targeted Aerosol Analysis



Non-Targeted Analysis of ENDS by GC-MS

Non-Targeted Analysis: Compound Identification

- Many published NTA studies rely heavily on NIST and/or other pre-established vendor libraries for compound identification
- There are several limitations in solely relying on pre-established vendor databases:
 - Need for a match factor threshold (“good” match/identification ≥ 80)
 - Poor match factor due to matrix effects
 - Not all e-vapor related compounds are included
 - Inconsistent compound identification between samples
 - The NIST database mostly consists of non-deconvoluted mass spectra

Mass Spectral Libraries – Comparison of Match Factor Threshold

NIST

Mass Spectral Database

267,376 compounds

Metabolites, drugs, pesticides, surfactants, etc.

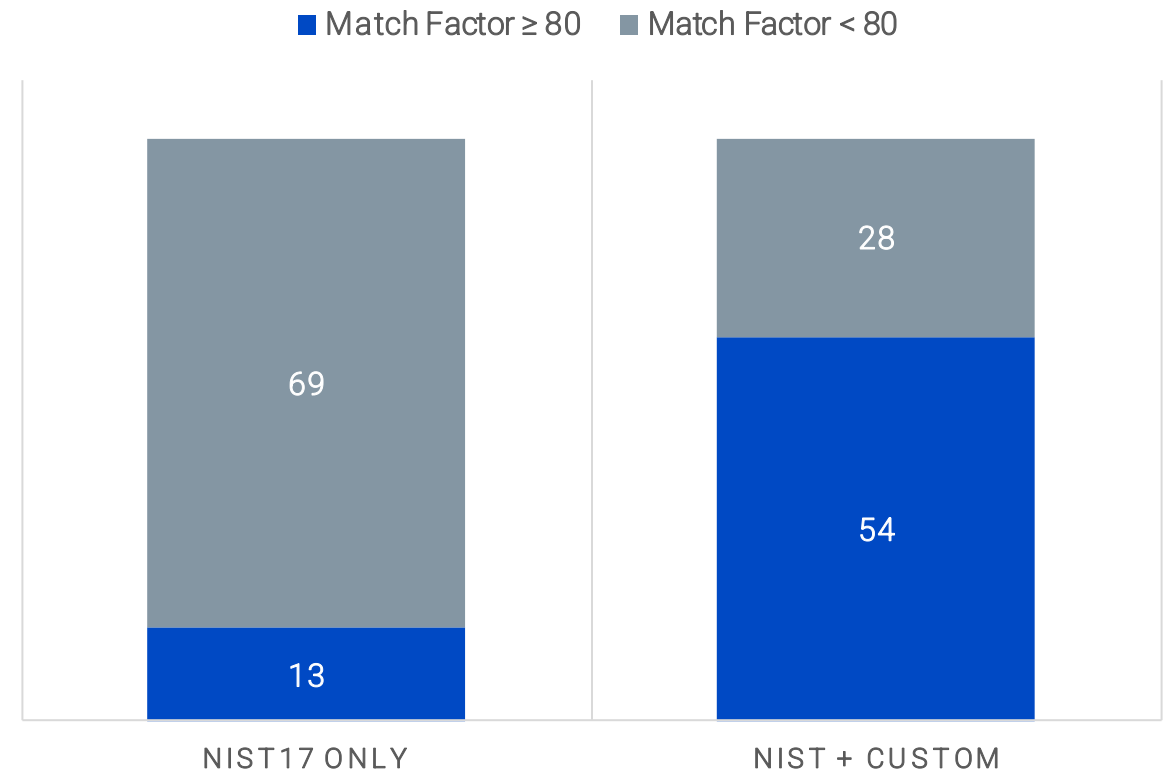
Custom mass spectral library

Reference standard (matrix matched)

Tentative identifications

Previously observed unknown compounds

JUUL Golden Tobacco 5.0% Aerosol NTA Constituents*



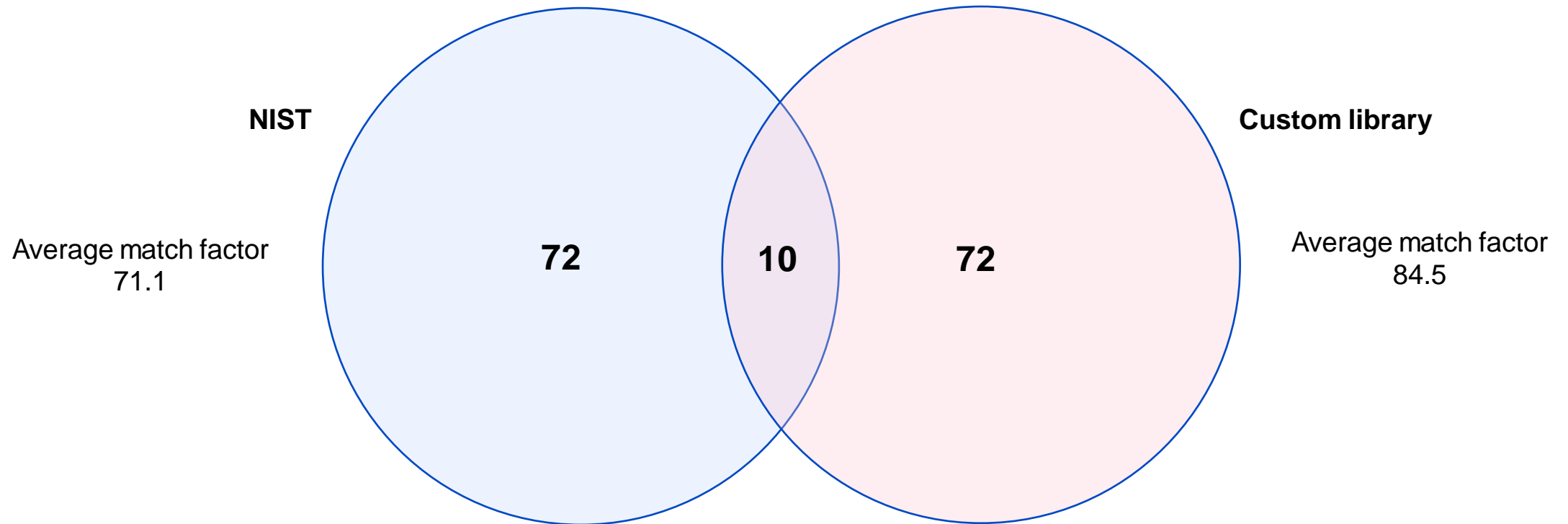
*Primary constituents (propylene glycol, glycerol, nicotine, and benzoic acid) are not included in reporting

Identification Comparison

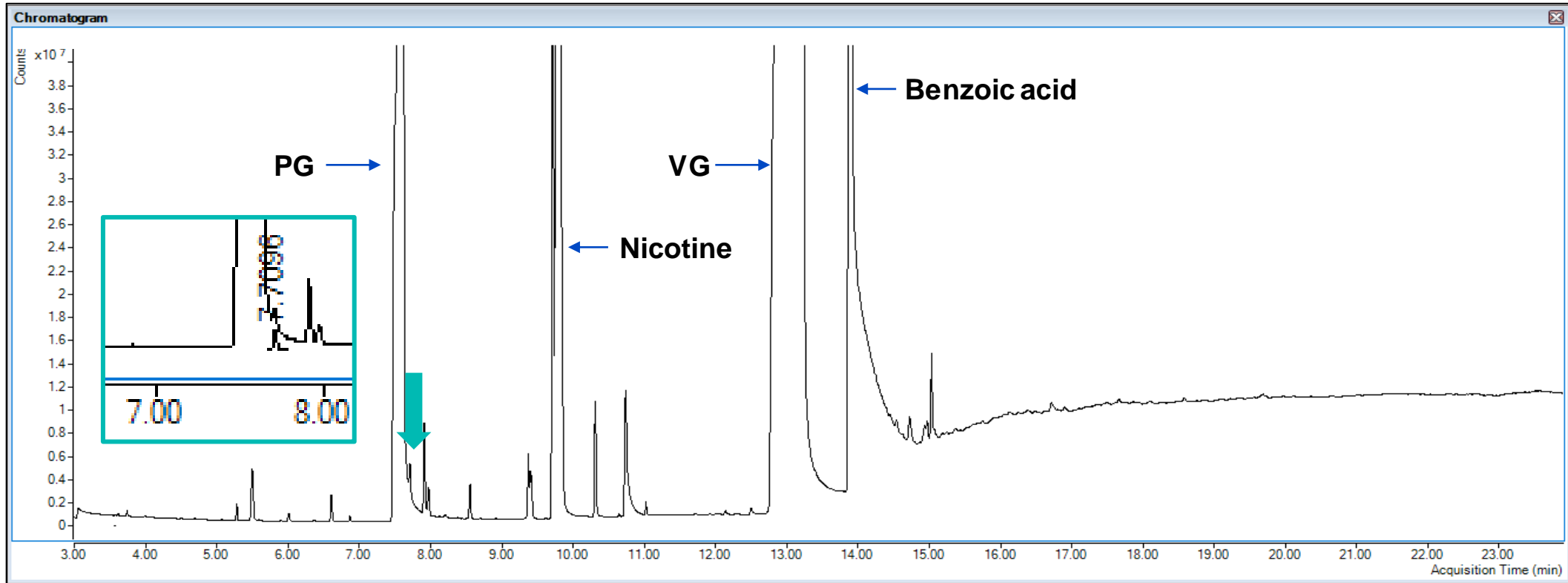
Sample set: Golden Tobacco 5% aerosol

NIST identifications were compared to custom library search results

10 out of 82 (~12%) accurate identifications by NIST (compounds present at relatively high concentrations)

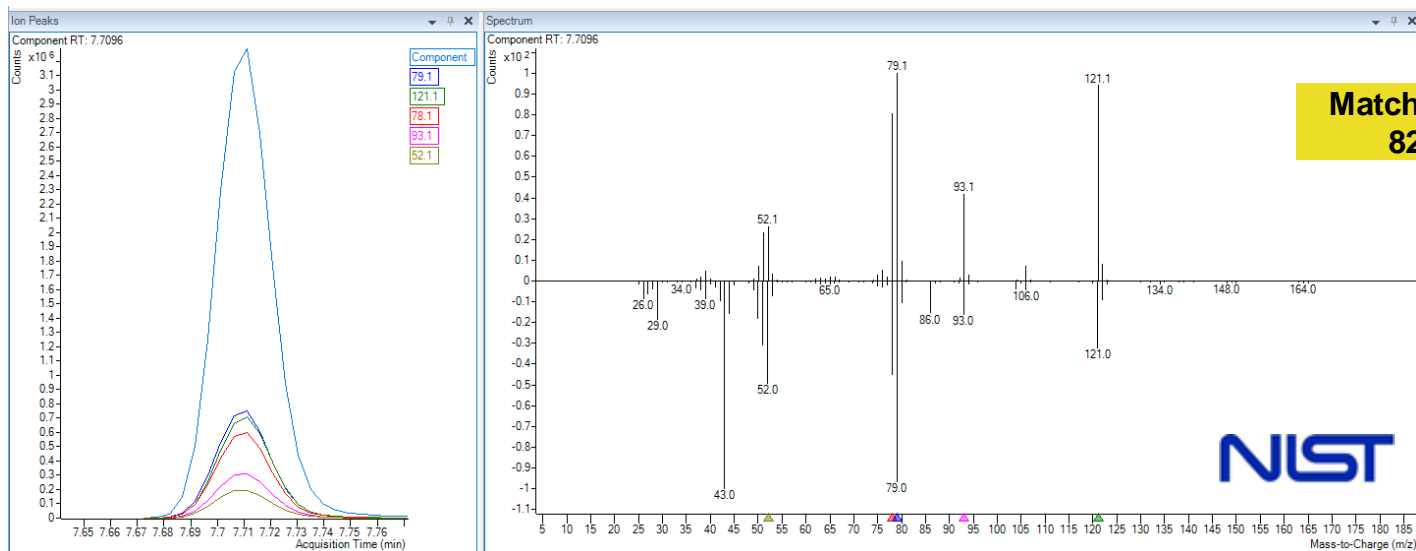


Impact of Matrix Effects on Compound Identification

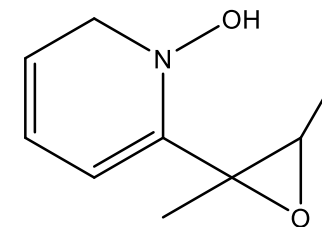


PG – Propylene glycol; VG – Vegetable glycerin

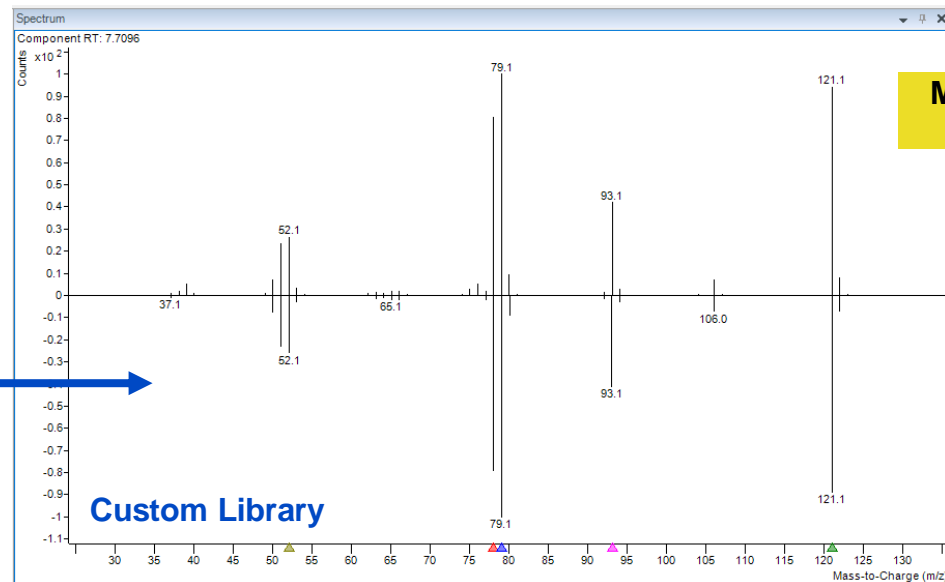
Impact of Matrix Effects on Compound Identification



“Good” match score, but incorrect identification

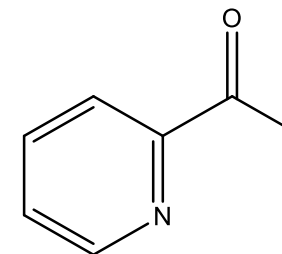


2-(2,3-Dimethyl-oxiran-2-yl)pyridin-1-ol



Matrix matched reference standard

Match Factor 92.65

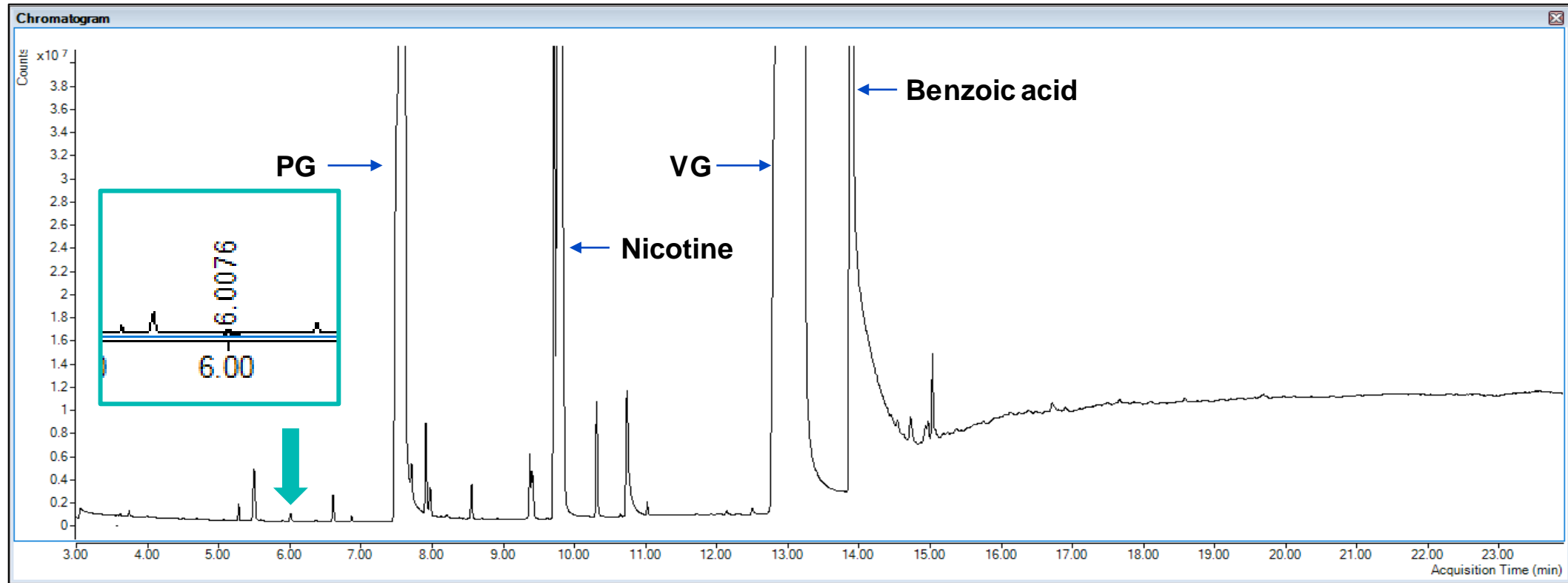


Component RT = 7.70 min
Library RT = 7.68 min

Ethanone, 1-(2-pyridinyl)-

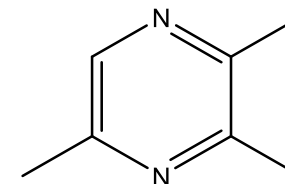
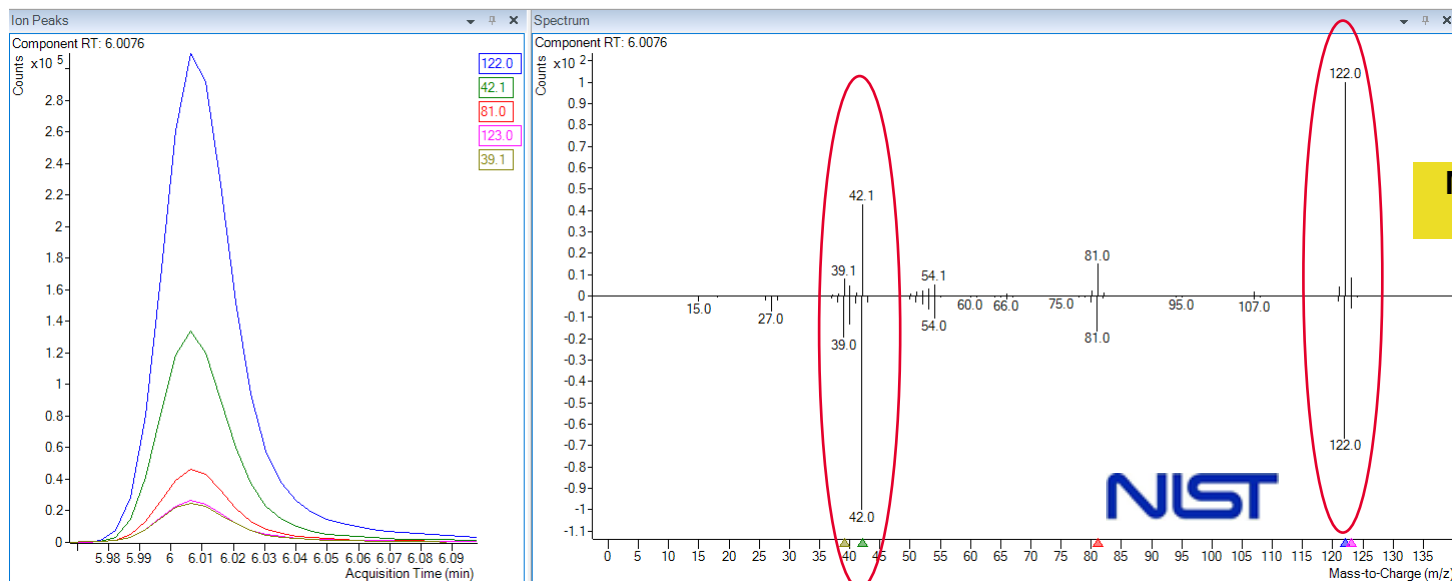
Confirmed by mass spectral match and retention time match

Improved Match Factor



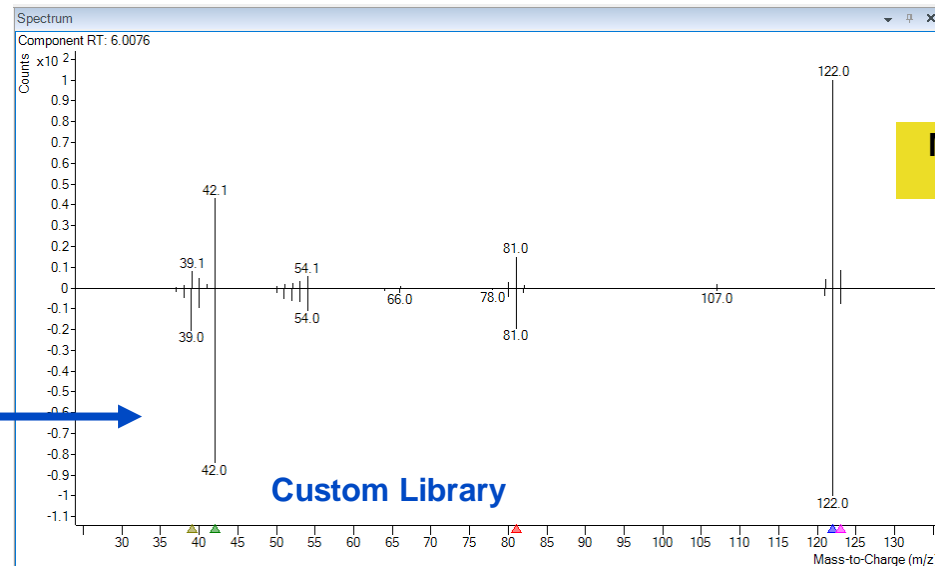
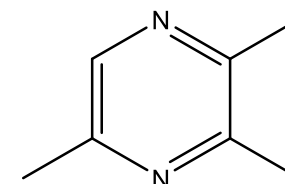
PG – Propylene glycol; VG – Vegetable glycerin

Improved Match Factor



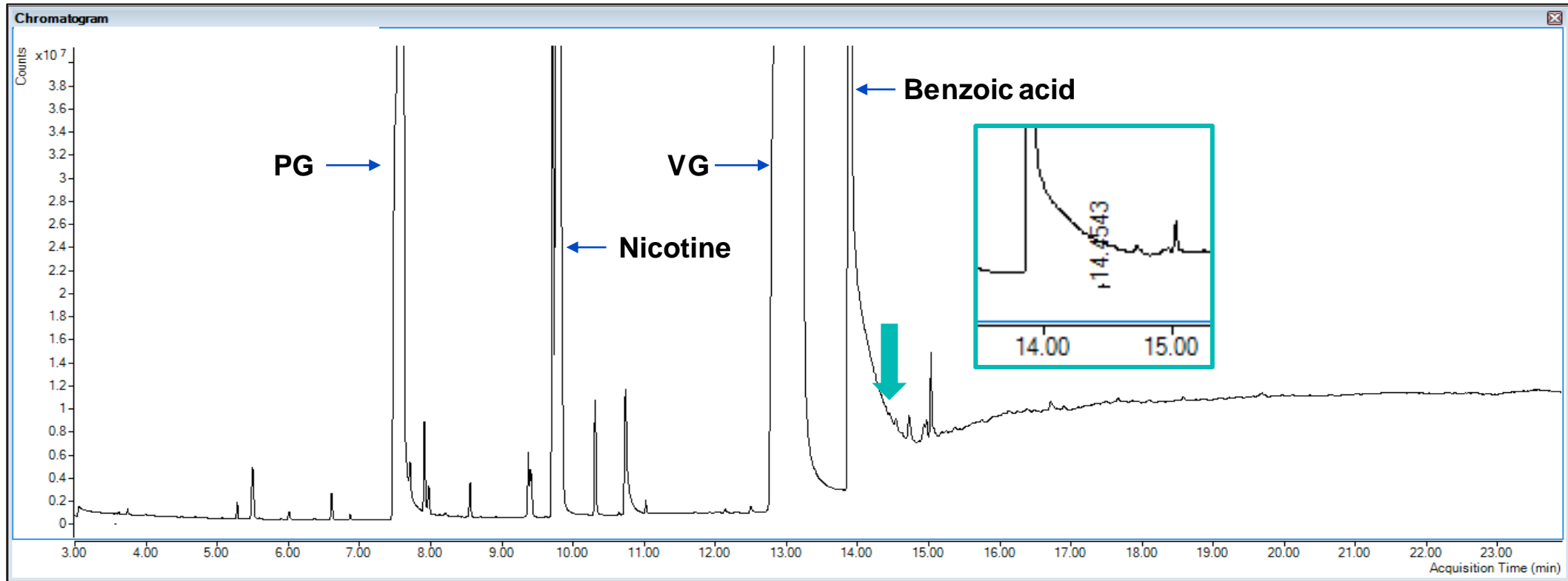
Correct identification

Differences in ion ratios

Matrix matched
reference standard

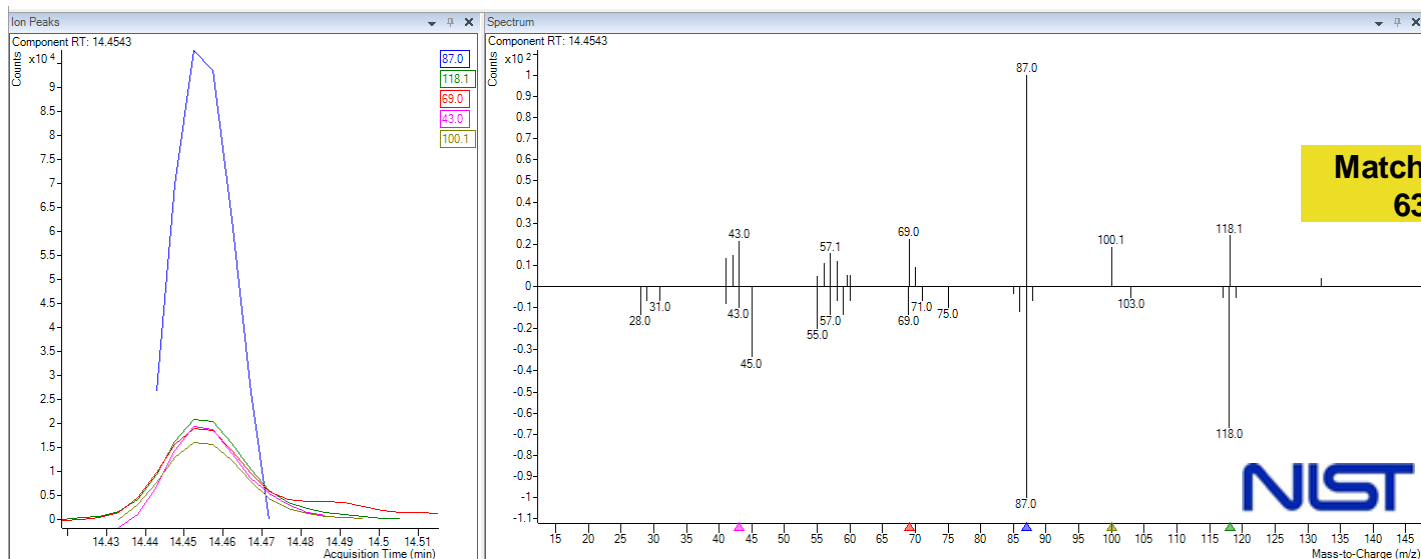
Trimethylpyrazine

Confirmed by mass spectral match and
retention time matchComponent RT = 6.00 min
Library RT = 6.00 min

Absence in NIST Database

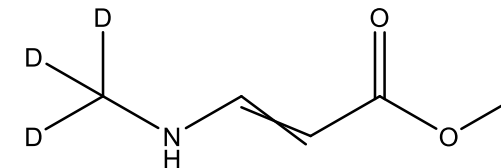
PG – Propylene glycol; VG – Vegetable glycerin

Absence in NIST Database

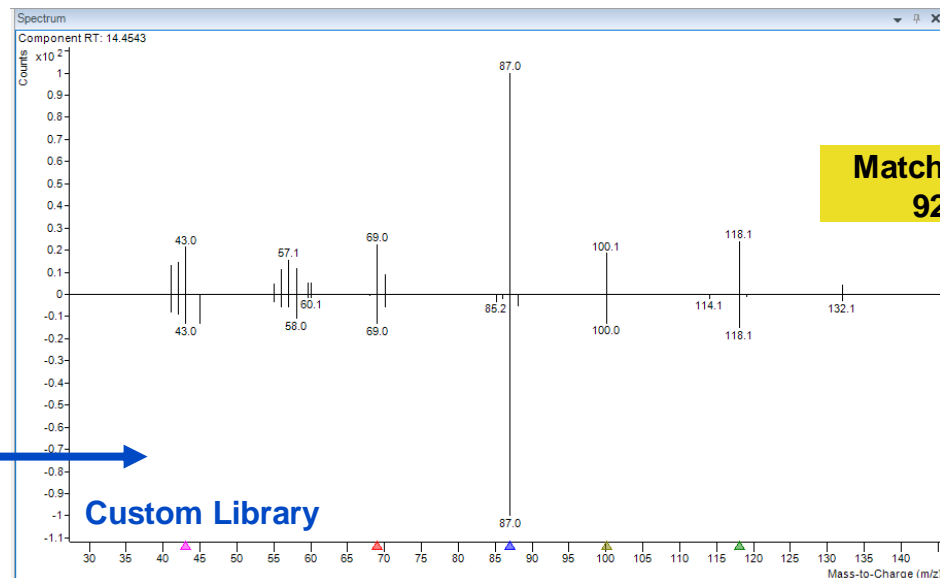


Match Factor
63.46

Incorrect identification due to absence of compound mass spectrum.



Initially labeled as an unknown compound



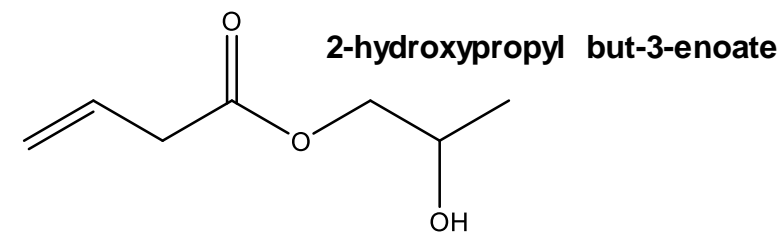
Match Factor
92.10

Component RT = 14.45 min
Library RT = 14.50 min

Custom synthesized
reference standard

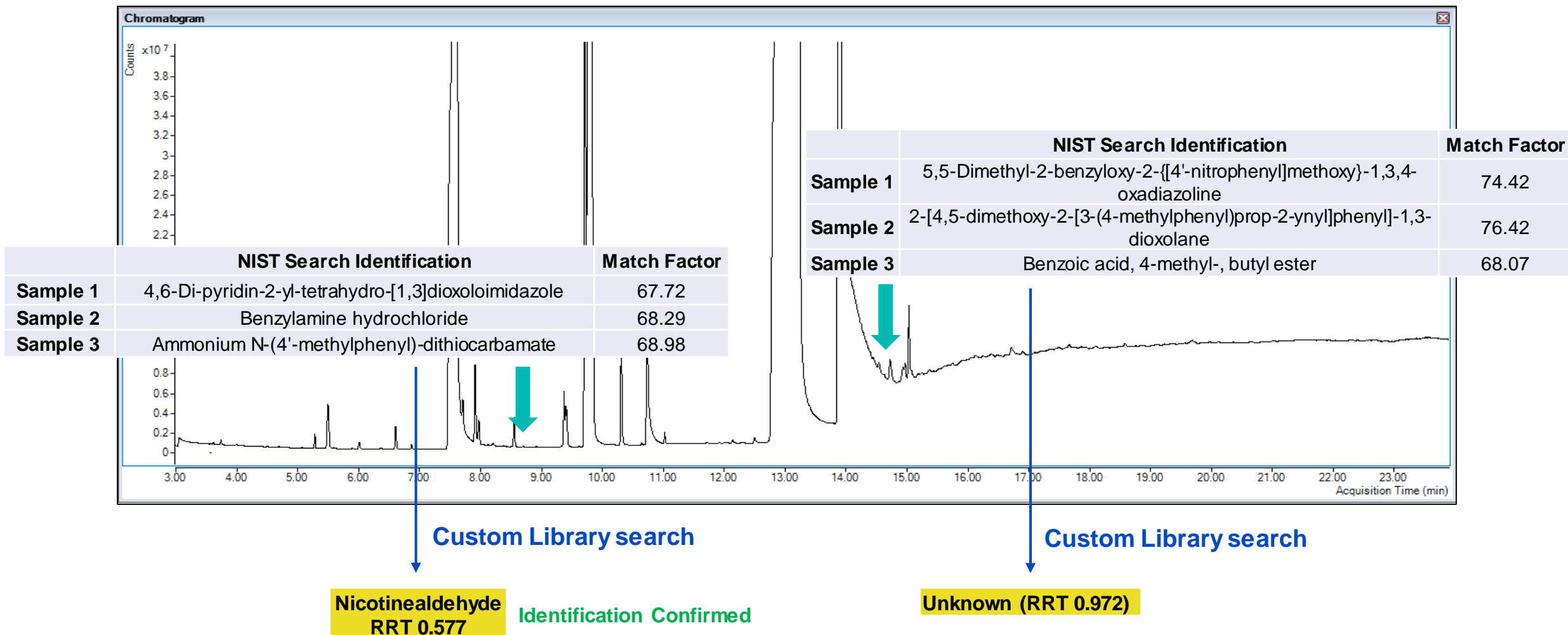


Custom Library



Butyrolactone + PG reaction product

Inconsistent Compound Identification



Relative Retention Time (RRT) = compound RT / internal standard RT

Summary

- Limitations using NIST database only search for chemical identification of ENDS aerosol:
 - < 20 % of the detected compounds had an acceptable tentative identification (match factor ≥ 80)
 - Peaks co-eluting with primary constituents exhibited poor match due to matrix effects
 - Inconsistent identifications between samples
- A comparison of the compound identifications when performing a search including and excluding the custom library combined with the NIST17 database for JUUL aerosol resulted in < 20% common identifications.
- The use of a custom library greatly increased acceptable tentative identifications by consistent detection and peak tracking based on relative retention time (RRT).
- The use of a custom mass spectral library is a valuable addition to the non-targeted assessment approach.