

## Identification of Microplastics in Water by Pyrolysis Gas Chromatography Mass Spectrometry

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#### From NEMC 2021: How is MP Mass Balance Determined?

Mass Balance = Relative amounts of each polymer in MP sample

- Thermal Extraction GC-MS (TE GC-MS)
- Pyrolysis GC-MS (Py GC-MS)
- Thermal Extraction / Desorption GC-MS (TED GC-MS)



#### Today's Pyrolysis Focus

- Mass Balance of Microplastics by Pyrolysis is Established
- ASTM's D19 and ISO's TC147 groups have methods in draft
- To find mass balance, you have to find the polymer first
- Today the focus is on Polymer ID.



#### The end at the beginning

- Pyrolysis GC-MS is a great tool for the ID and measurment of Microplastics
- Advances in open source data analysis software make analysis more reliable
- The same advances make data work-up faster and largely automated
- The sample sizes are too small, resulting in more manual sample prep work



#### PYROLYSIS and Microplastics: How does pyrolysis GC-MS work?

- Samples are placed in pyrolysis tubes
- Can be particles, punches of filter media, or cryo-milled sediment
- Pyrolysis occurs between 600 and 1000 °C
- Non-volatile sample material is decomposed



PYROLYSIS and Microplastics: How does pyrolysis GC-MS work?

- The sample <u>is fully consumed</u> in the process
- 100% of the decomposition products are introduced to the GC/MS
  - A trap can optionally be used for trapping (focusing) and/or splitting of the sample
- Material(s) in the sample are identified by their pyrolysis fragments

The resulting pyrograms show not what the sample *is*, but what it *became* when heated.

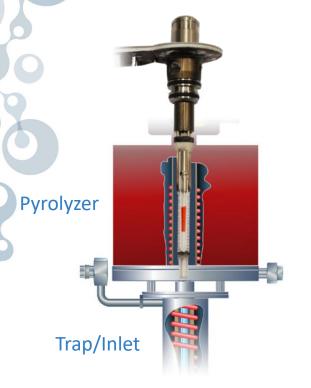


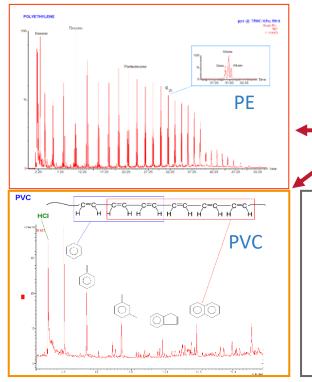


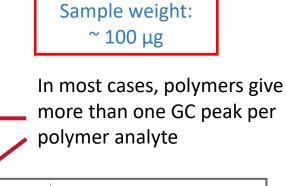


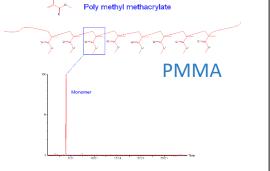


#### **Pyrolysis Examples of Common Plastics**



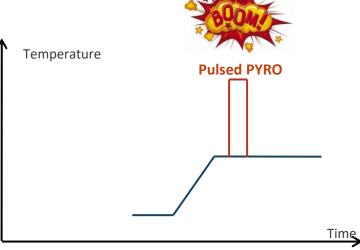








- Fast heating, fast pyrolysis
- <u>Required</u> if going direct to column without a focusing trap
- Same temperature used for ALL polymers in the sample
  - The pulse temp is a compromise
  - Secondary reactions due to overheating add complexity
- Requires method development
  - Works best for known, pure samples
- Not suitable for complex unknown samples or complex mixtures (can't optimize temp)





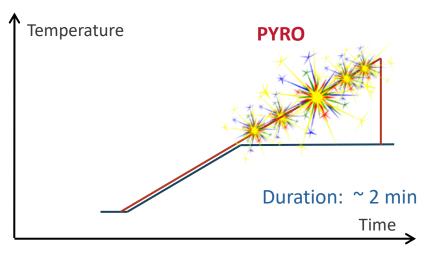


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#### A newer alternative: Smart-Ramped Pyrolysis (SRP)

- Pyrolysis using a temperature ramp
- Trapping the pyrolysates <u>is necessary</u> before introduction to the column
- Polymers are pyrolyzed without overheating and secondary reactions are eliminated
- One combined GC/MS run follows with thermal pyrolysis AND thermal extraction data

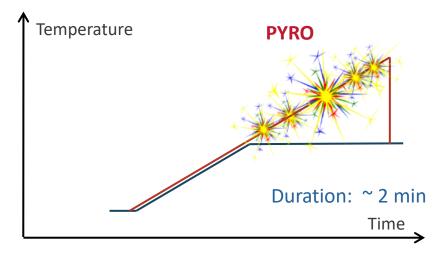


If you can do pulsed pyrolysis, you can also do ramped – the heating rate is a method parameter Ramping does require trapping the pyrolysates on a focusing trap or the column head



#### Smart-Ramped Pyrolysis (SRP): Improvements over Pulsed Pyrolysis

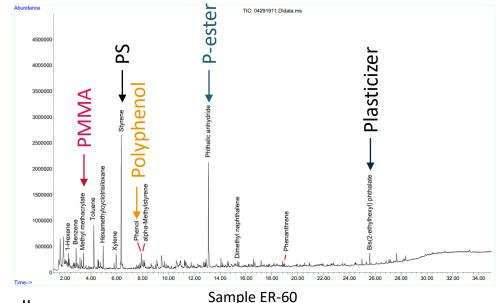
- Better signal and less noise
- Don't have to choose a single pyro temp
- Even unknown samples can be analyzed
- Polymer & additive information in one run



#### Lake Erie River Sediment: Smart-Ramped Pyrolysis



- Several polymers present:
  - Methyl Methacrylate / PMMA
  - Styrene (PS)
  - Phenol (Polyphenol)
  - Phthalic Anhydride (Polyester)
- Monomers can be used as "marker compounds"
- Markers can be used for ID and for quantitation.



 Note the presence of some additives as well (Bis(2-ethylhexyl)phthalate)



- Direct SRP Pyrogram up to 600 °C
  - -Very complex but the use of marker compounds allows the MP polymers to be 'pulled out' (more later)

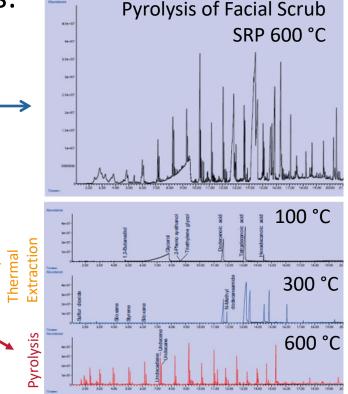
Fractionated Pyrolysis: TE followed by Pyrolysis

A: 100 °C (VOCs) B: 300 °C (SVOCs, additives) C: 600 °C (Well defined pyrogram of MP polymer)

 Fractionated pyrolysis is an easy way to get polymer and additive information from MP's

 Interpretation is simplified, but data in 2-3 (or many more) files – dozens of 'shots' at a sample







#### Using Filters to Extract and Identify MP's: Practical Examples

- One Liter samples of local waters were run through a 10 um PTFE filter w/ vacuum
- All were allowed to dry overnight
- A 1.2 mm diameter punch was taken of each sample and pyrolyzed
- The Street Runoff did not give much signal, so a few milligrams were scraped off & run



Synthetic Grey Water

Street Runoff

Pond Water

**Bottled Water** 

Column:

Pneumatics:

GC:

**Analysis Conditions** 



DB-5MS UI (Agilent) dia = 0.25 mm,  $d_f$  = 0.25 $\mu$ m, L = 30 m

Carrier = Helium Constant flow = 1.0 mL/min

Initial 40 °C (1.0 min) Ramp 15 °C/min to 320 °C (15 min) Detection = 5977 MSD



#### Pyrolysis Conditions PYRO body

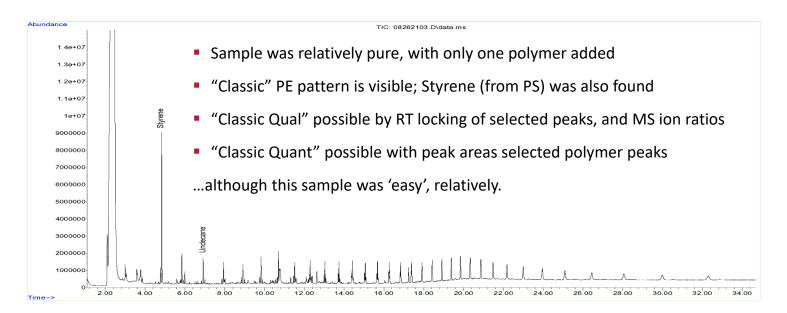
Trap:

Lead Time: Follow up Time: Initial Time: PYRO Program: Splitless: 80 °C (0 min), 300 °C/min, 300 IC (2.17 min) Split 25:1 300 °C isothermal (not trapping) 0.00 min 0.50 min 0.00 min SRP: Initial 300 °C (0 min), ramp 5.0 °C/s to 800 °C (0.0 min)



#### Synthetic Grey Water Filtrate Pyrogram MAKING LABS WORK

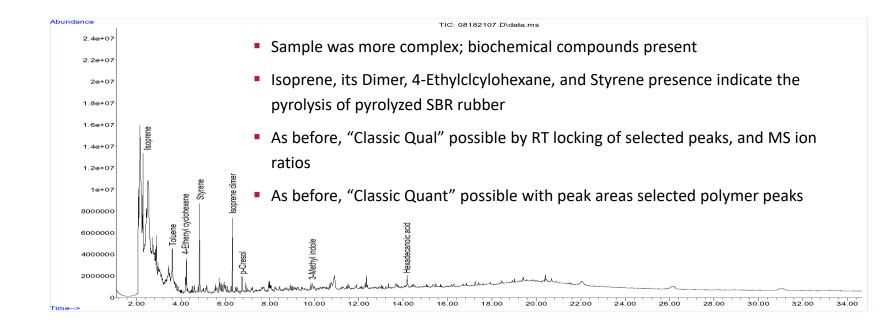
- A facial scrub was spiked with cryomilled PE
- The scrub was then added to pure water, shaken, and then analyzed





#### Street Runoff Filtrate Pyrogram

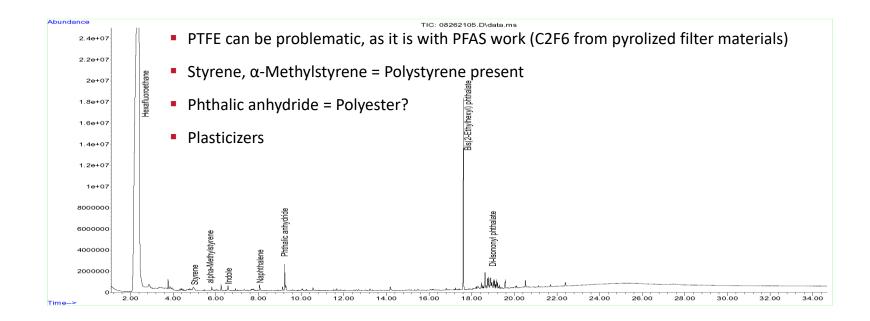
- A punch of the filter did not yield much signal
- Several milligrams of filtrate were scraped off and a 1-2 mg sample of that was run





#### Pond Water Filtrate Pyrogram

- A 1.2 mm diameter punch produced sufficient signal
- Several polymers and plasticizers present





## Problems with the "Classical Approach"

- Polymers usually have more than one GC peak / polymer; we are ignoring information ("Qualifier Peaks" aka Marker Compounds)
- A polymer should also have the right NUMBER of GC peaks, and at RT's that are known
- The library used should then have both GC and MS information in it, and not MS information alone

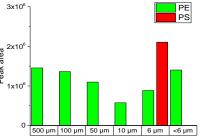
Advances in open-source software has solved these problems

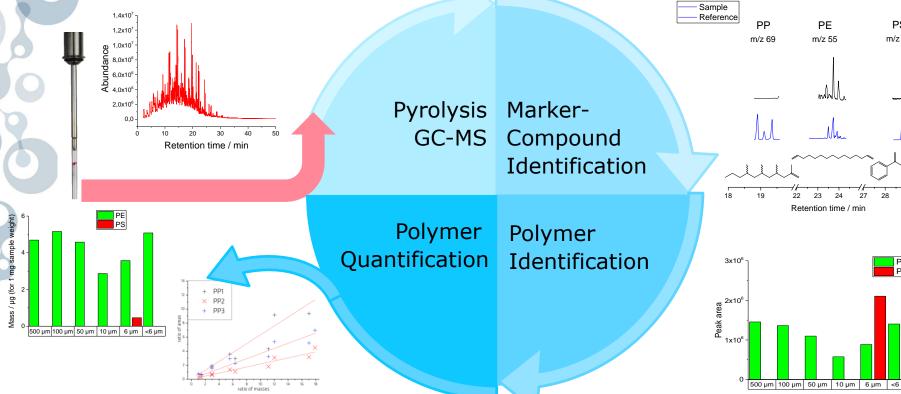
#### Openchrom & ChromIdent: **ID and Quant of Pyrolysis Products**



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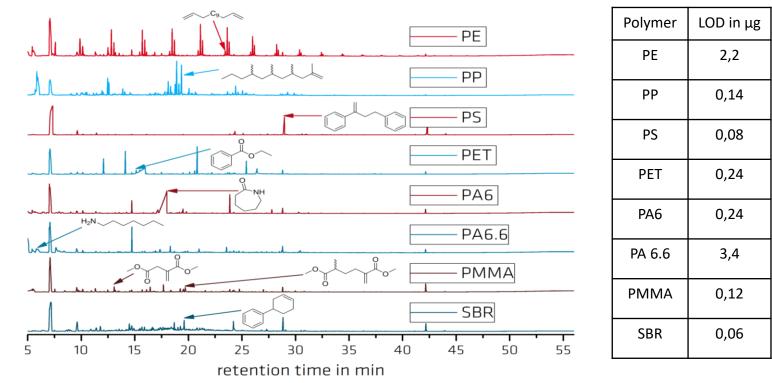
Sample Reference PP ΡE PS m/z 55 m/z 91 m/z 69 MUU 18 19 22 23 24 27 28 29 Retention time / min





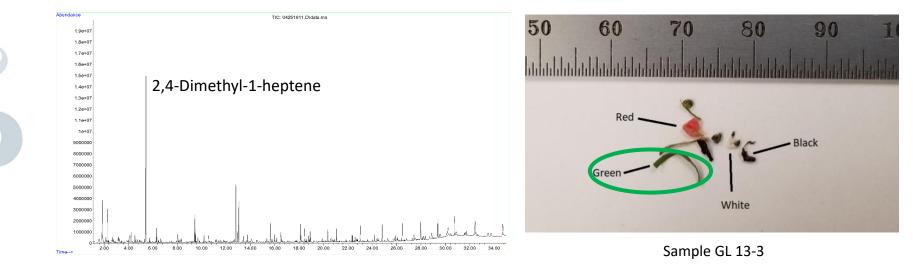


# Eight example Microplastic targets and example marker compounds



Netted Polymer Sample – Lake Erie: Section of Green Portion Only, ~ 0.9 mg



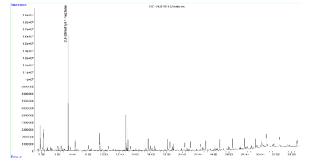


Library	Polymer	Forward	Reverse	Marker	Ambiguous	Unidentified
Entry		Match	Match	Peaks	Peaks	Peaks
P-0001	PP	82.1	81.4	1	55	73
P-0024	PP	89.9	83.8	0	58	71
P-0046	PP	84.1	63.5	14	79	36

#### How ChromIdent "Does It"



- Look for entries with a high number of marker peaks (entry #46 had 14)
- A high Forward match indicates a pure material (84% in this case – relatively pure)
- A low Forward match and high number of unidentified peaks points to a mixed material
- A low Forward match and high number of unidentified peaks <u>but a good Reverse match</u> are an indication that the polymer is present in the mix.

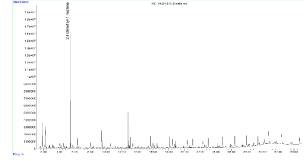


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Non-targeted via ChromIdent



- Look further at the marker peaks and Compare the mass spectra of the marker peaks with the library
- Verify a match by direct comparison of the database entry and sample
- Finally, if more than one entry for a polymer appears in the results list look at results with high numbers of ambiguous peaks
- If no Marker peaks are identified, go through the ambiguous peak identifications to possibly identify the polymer.



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#### Selected PYRO & Microplastics References

- Microplastics (MPs) by Pyrolysis GC-MS (AppNote 212) MPs in Filtered Great Lakes Water and Sediment MPs in Body Care Product (Facial Scrub)
- Identification of Microplastics in Water by Pyrolysis Gas Chromatography Mass Spectrometry (AppNote 232)
   Grey Water, Pond Water, Street Runoff, Bottled Water
- Microplastics from fish stomach content, Texas Gulf Coast ...
  E. Hendrickson et al., Mar. Pollut. Bull., 2018, 137, 91-95
- Microplastics from Western Lake Superior ...
  E. Hendrickson et al., Environ. Sci. Technol., 2018, 52, 1787-1796
- Simultaneous Determination of Plastic Particle Identity and Adsorbed Organic Compounds by TD-Pyrolysis GC-MS Molecules 2020, 25, 4985; doi:10.3390/molecules25214985



#### Pyrolysis GC-MS: Good, but....

- Samples are small in mass (0.1 1 mg, ideally)
- Samples small in size (few square mm<sup>2</sup>)
- For filters, several punches must be taken



GERSTEL Pyrolyzer
 Pyrolyzing the whole filter risks overloading the GCMS (pyrolyzers are connected directly to the GCMS)

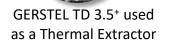
Thus many smaller samples from a single filtrate need to be run together to get a representative sample Or, the sample must be cryomilled to make it homogeneous



#### So what does it all mean?

#### The Short Answers

- Thermal Extraction GC-MS analyzes larger representative samples but is indirect: polymers must have unique additives and samples should be relatively clean (e.g., PET MP's in drinking water); <u>MP analysis range is</u> <u>limited / undetermined</u>
- Pyrolysis GC-MS offers direct MP ID and analysis of a wide range of polymers, can also do additives if needed, and can handle a wider range of matrices; sample sizes are smaller & multiple reps and/ or cryomilling required for representative sampling
- TED GC-MS can do a wide range of polymers, additives runs, larger representative samples without cyromilling, and being off-line to the GC-MS makes it the most robust; more investment (two instruments and training) are needed
- Thermal Extraction/Desorption (TED-GCMS) System





GERSTEL PYRO



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#### The end at the end

- Pyrolysis GC-MS is a great tool for the ID and measurment of Microplastics
- Advances in open source data analysis software make analysis more reliable
- The same advances make data work-up faster and largely automated
- The sample sizes are too small, resulting in more manual sample prep work





- Large filters are great for large samples!
- "PYRO-Sized" samples are 1.2 mm diameter punches
- One sample alone could 'miss' analytes
- Best answer is cryomilling, but it's labor intensive
- Also, TEFLON is a BAD IDEA

~ 1 mm punch visualization



Synthetic Grey Water

Street Runoff

Pond Water

**Bottled Water** 



#### Thanks to:

# BAM

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Ulrike Braun Erik Dümichen Paul Eisentraut et al.

Eike Kleine-Benne Huan-Xiao Zhou, Jackie Whitecavage, John R. Stuff, Nicole Kfoury



#### **Reference Material**

## Instrumentation Comparison



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	Thermal Extraction (~315 °C)	Pyrolysis (600 to 1000 °C)	TED (600 to 1000 °C)
Sample Intro	On-line to GCMS	On-line to GCMS	Off-line from GCMS
Process	Thermally extract to trap, then to column (through valve / transfer line or direct to column)	Pyrolyze direct to GCMS (direct to column, or optional trapping step before column)	Extraction/Pyrolysis in TGA; trap on PDMS and transfer to TD-GCMS (PDMS trap is desorbed in TD; re-trapping before column recommended)
Bake-out	Up to 450 °C, $N_2$ or He	Up to 1000 °C, $N_2$ or He	Up to 1000 °C, in $N_2$ , He, $O_2$
Sample Size	Typically 10-50 mg	0.1 - 10 mg (0.1 – 1 mg typically; all goes to inlet)	Typically 10-50 mg
Representative Sample	One Run	Multiple Runs	One Run
Type of Data	Mass Spectral	Mass Spectral	MS and TGA both



#### Analytical Comparison

	Thermal Extraction (~310 ° <b>C</b> )	Pyrolysis (600 to 1000 °C)	TED (600 to 1000 °C)
Sample Prep (filter media)	Dry, solvent rinse of filter, dry again, analyze	Dry, punch out correct sized samples, analyze (multiple punches very likely needed)	Dry, punch out correct sized samples, analyze (one, larger punch is typically sufficient)
Polymer ID	Yes, Indirect / Inferred (marker additives)	Direct (un-zipped monomer or targeted degradant markers)	Direct (un-zipped monomer or targeted degradant markers)
Range of Polymers	Limited to uniqueness of additives (typ. PET)	Eight or more common MP polymers (PET, PE, PP, PS,)	Eight or more common MP polymers (PET, PE, PP, PS,)
Mass Balance	Yes, Indirect / Inferred quant through additives	Yes, direct quant through monomers / degradants	Yes, direct quant through monomers / degradants
Additives (note: not needed for mass balance)	Yes, Direct, one step	Yes, Direct, one or two steps (300 °C and then 600 to 1000 °C), if two steps needed	Yes, Direct, one or two steps (300 °C and then 600 to 1000 °C), if two steps needed



## Cost and Complexity

	Thermal Extraction (~310 'C)	Pyrolysis (600 to 1000 'C)	TED (600 to 1000 'C)
Instruments	Three (TD, GC, and MS)	Three (PYRO, GC, and MS)	Four (TGA, TD, GC, and MS)
Cost	\$\$	\$\$	\$\$\$
Support	One Provider	One Provider	Two Providers (TGA and TD-GCMS)
Whole Filtrate in One Run	Possible	No	Possible
Carry-over Risk (sludge, tissue,)	Highest (TE only = lowest temp technique)	High (PYRO only less risky) or Higher if TE step used	High (or Moderate with O <sub>2</sub> cleaning step)