

# Organic Speciation for Source Apportionment in DEARS

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## DEARS Objective #1:



(1) To determine the associations between concentrations measured at central site monitors and outdoor residential, indoor residential and personal exposures for selected air toxics, PM constituents, *and PM from specific sources.*

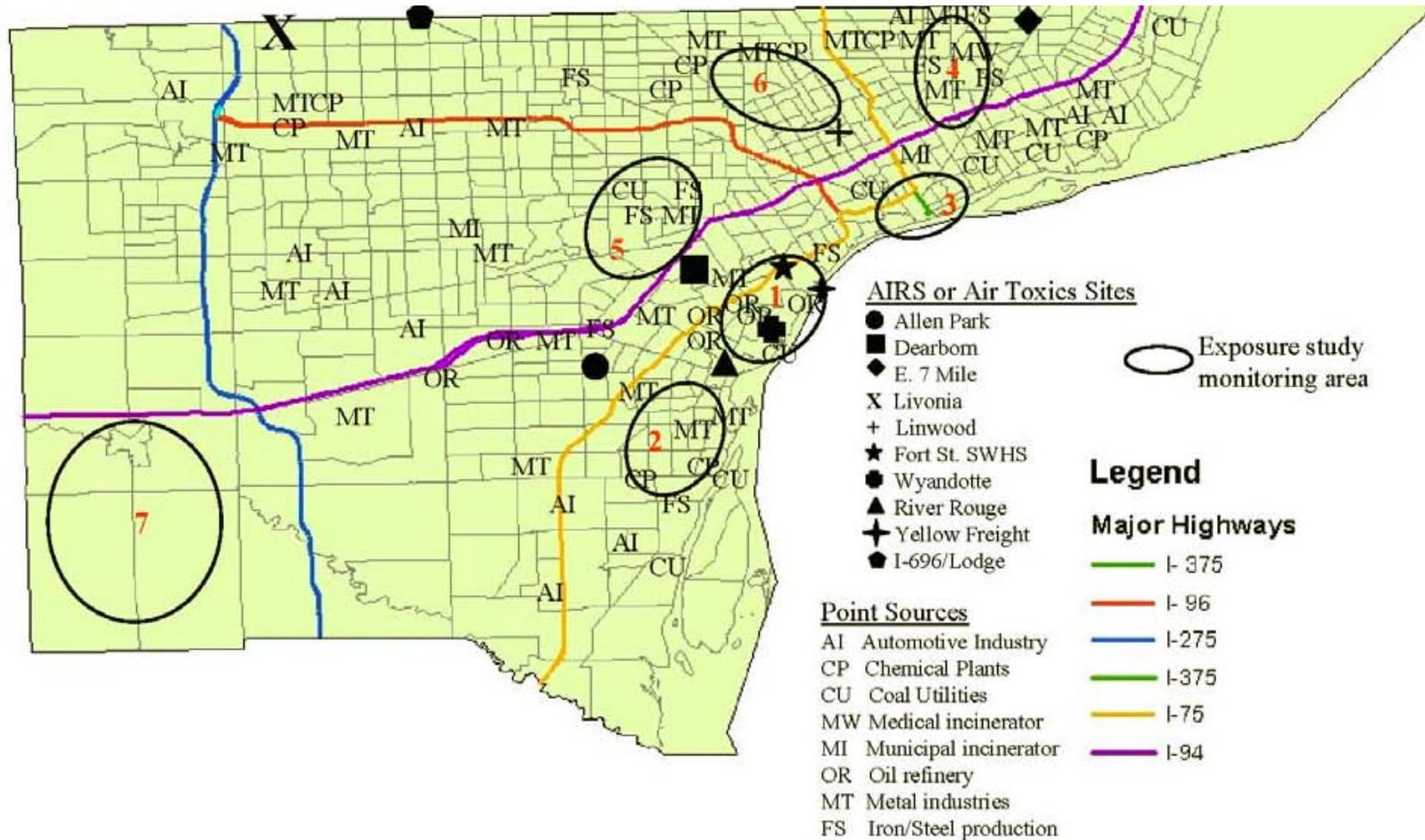
Source: [www.epa.gov/dears](http://www.epa.gov/dears)



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# DEARS Study Design



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# Sampling

## Central Site



## Residential Monitor



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# *Sampling & Analysis Methods*

- Allen Park community air monitoring station in Detroit, MI
- From July 13, 2004 to August 28, 2004
- 34 high-volume (113 liters/minute) PM<sub>2.5</sub> samples



## What's unique to DEARS Organic Analysis:

- Largest dataset planned for organic source apportionment
- Low collection volumes (10 liters/minute)
- Daily samples



## Challenges:

- 1) Can we detect source markers with sufficient sensitivity in 24 hour, 10 liter/minute samples?
- 2) Can we adequately estimate source contributions with a limited set of organic markers?
- 3) Can we achieve low enough uncertainties to adequately resolve sources?



- 1) Can we detect source markers with sufficient sensitivity in 24 hour, 10 liter/minute samples?



## Methods Considered:

- Thermal Desorption
- Large Volume Injection
- Splitless/Selective Ion Monitoring



## Large Volume Injection Problems

- Contamination (solved)
- High Blanks (solved)
- Rapid column deterioration (probably solvent related)\*

\*Recalibration usually required after column maintenance



# Sampling & Analysis Methods



## ORGANIC AEROSOL LAB SAMPLING AND ANALYSIS SCHEME

**Tisch TE-1202  
Quartz Filters  
113 liters/minute**

**Dionex Accelerated Solvent Extractor 200  
Hexane/Dichloromethane/Methanol**

**Zymark Turbovap  
Concentration to 0.30 ml**

**Gas Chromatography/Mass Spectrometry  
Selective Ion Monitoring**



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## Determining Method Detection Limits

- 1) Prepare standard 1-5 x MDL
- 2) Analyze at least 7 aliquots
- 3) Calculate variance ( $s^2$ )
- 4) Calculate MDL for  $t = n-1$ ,  $\alpha=0.99$  ( $=3.143s$ )
- 5) Verify average is 1 to 5 x MDL
- 6) Adjust standard concentration and repeat if necessary

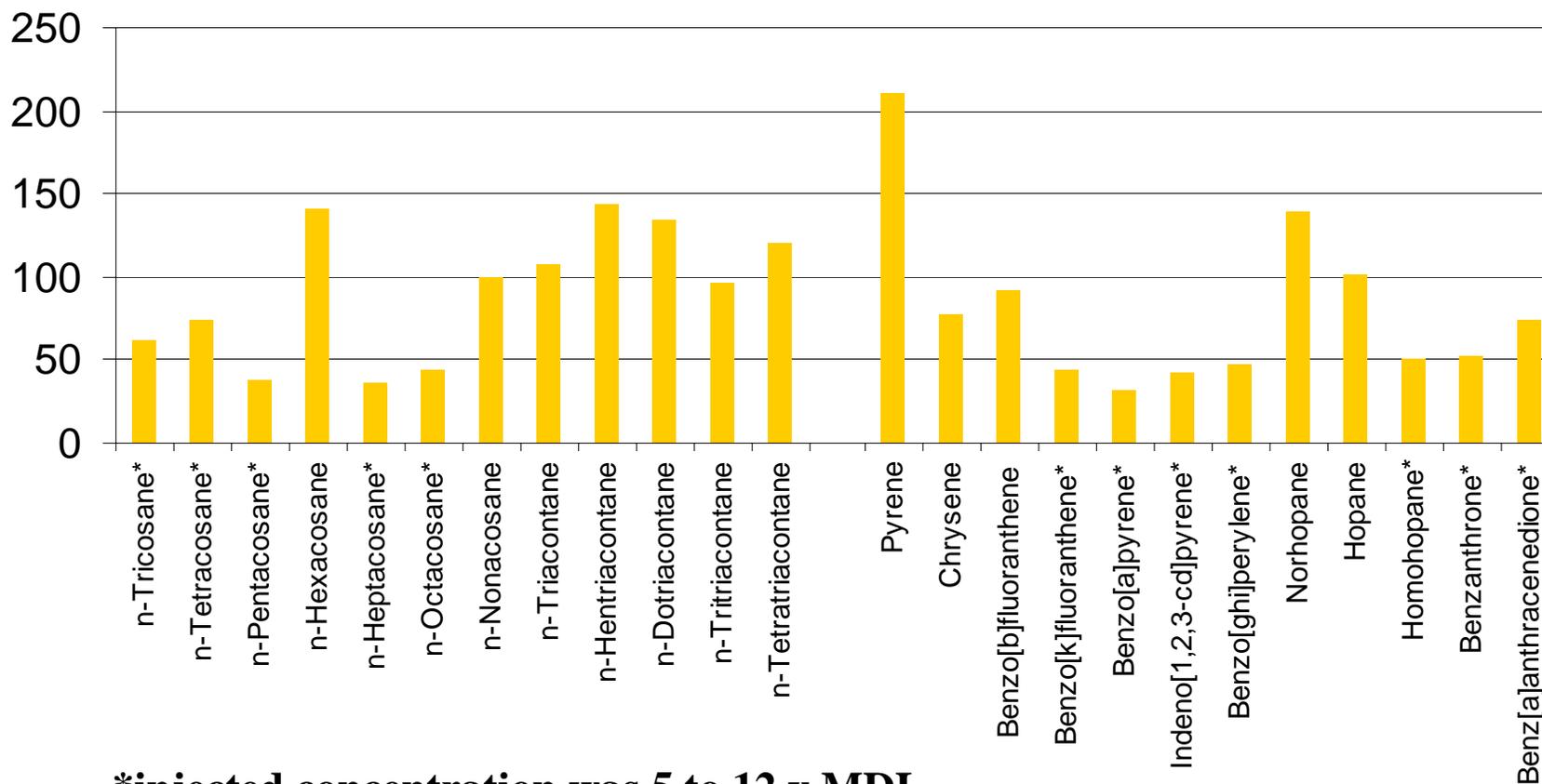
Source: 40CFR136, Appendix B



# GCMS Limit of Quantitation (pg/m<sup>3</sup> equivalent)

## GCMS Limits of Quantitation

LOQ = 10s (from 40CFR136B)



\*injected concentration was 5 to 12 x MDL



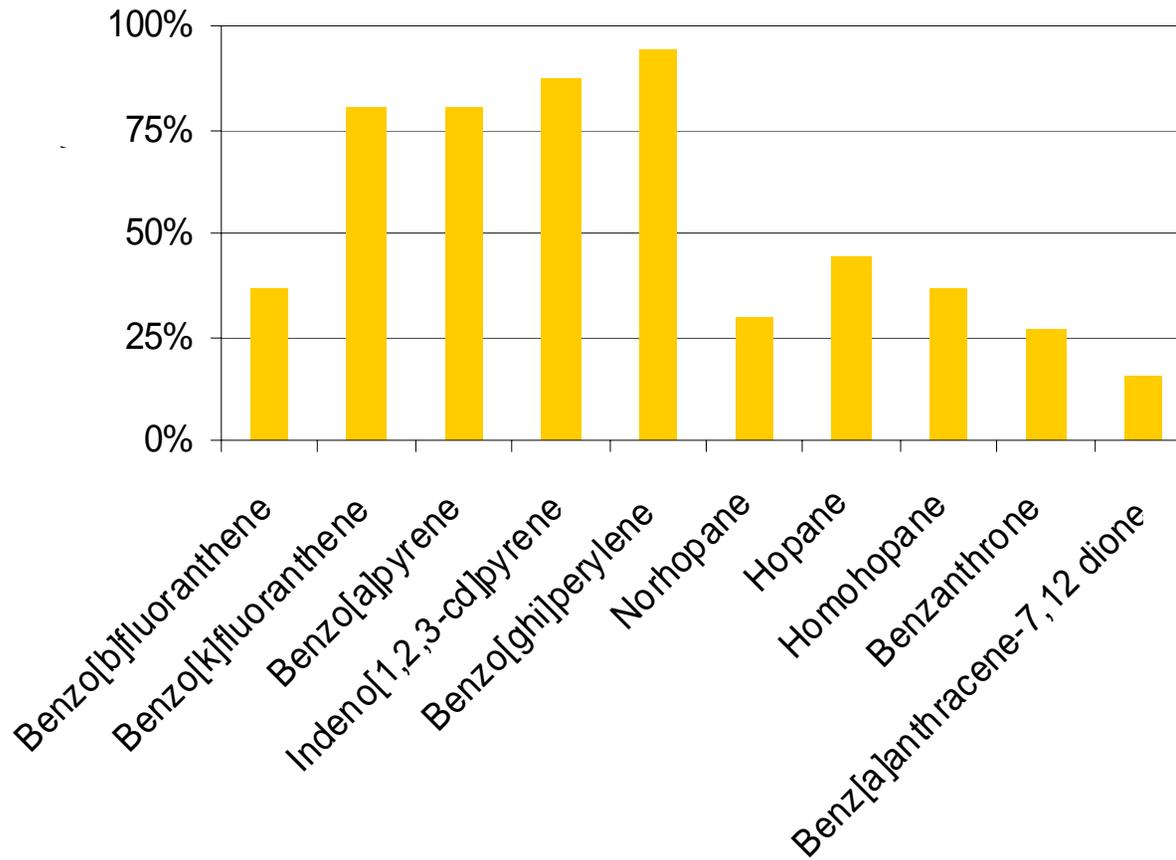
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# Fraction of TACS Samples > GCMS LOQ

n = 84

Fraction > GCMS LOQ

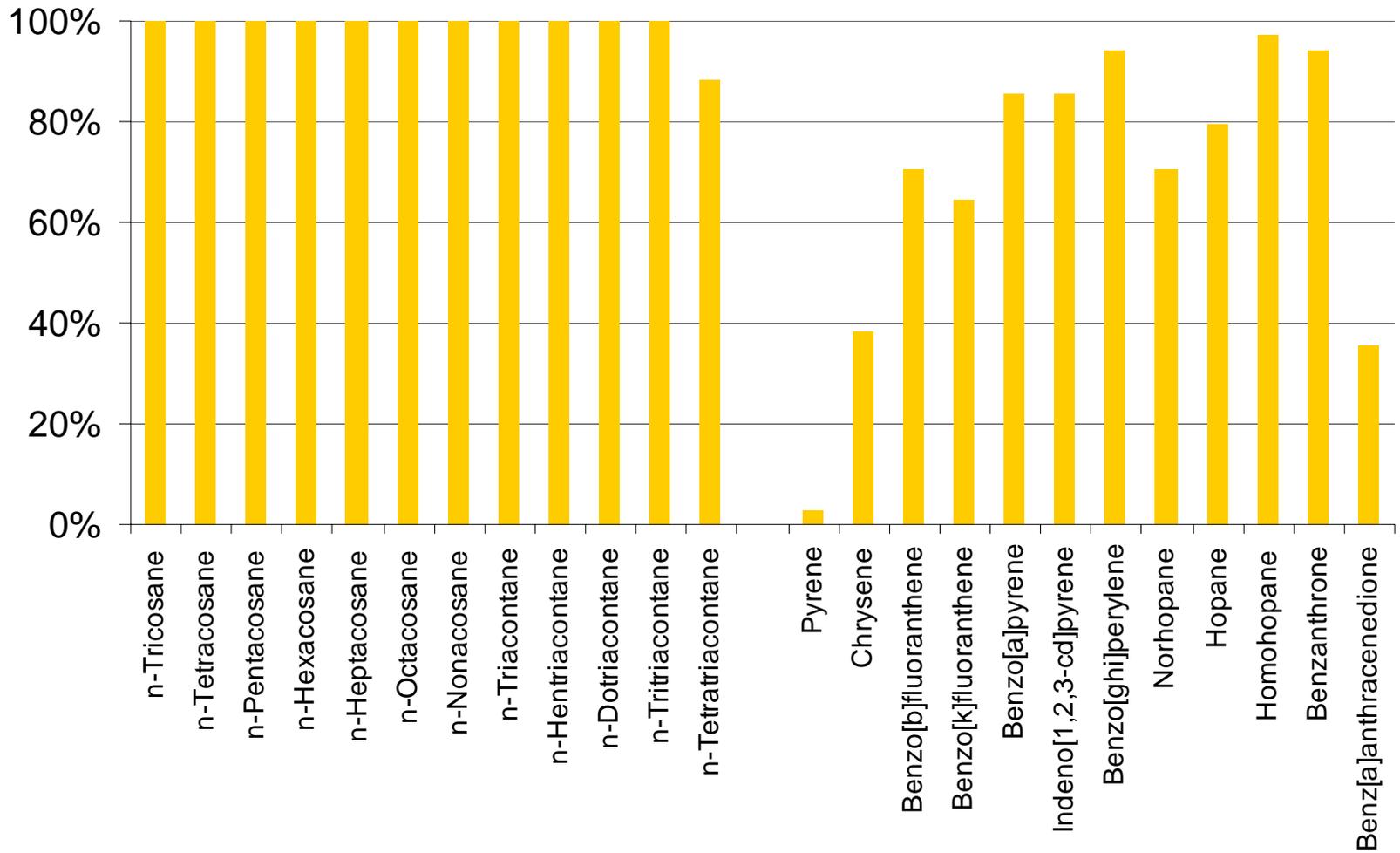


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## Fraction of 113 lpm Samples from DEARS that *would be* > LOQ at 10 lpm (n=34)

**Fraction > GCMS LOQ**



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2) Can we adequately estimate source contributions with a limited set of organic markers?



## DEARS Organic Source Markers (limited for higher throughput + SIM)

Tricosane		
Tetracosane		
Pentacosane		
Hexacosane	Pyrene	Norhopane
Heptacosane	Chrysene/Triphenylene	Hopane
Octacosane	Benzo[b]fluoranthene	Homohopane
Nonacosane	Benzo[k]fluoranthene	
Triacontane	Benzo[a]pyrene	
Hentriacontane	Indeno[1,2,3-cd]pyrene	Hexadecanoic acid
Dotriacontane	Benzo[ghi]perylene	Cis-Hexadecenoic acid
Tritriacontane	Benzanthrone	Octadecanoic acid
Tetratriacontane	Benzanthracenedione	Cis-Octadecenoic acid



## **Fundamental Questions:**

**How many fitting species are needed?**

Friedlander 1973 – Fitting Species: Al, Na, K, Mg, Ca, V, Pb  
Sources: auto, diesel, aircraft, oil, industrial

**How much are CMB fits improved by reducing measurement uncertainties?**

Kowalczyk et al. 1978 – Improved measurements gave better fits

**What is the practical optimum balance between adding fitting species and increasing measurement uncertainty?**





## **Which species influence source contribution?**

The MPIN matrix identifies which fitting species have the largest influence on the source contribution estimates from each profile.

Species with MPIN absolute values of 0.5 to 1.0 are considered influential species.

**WARNING:** binoculars required for next slide...

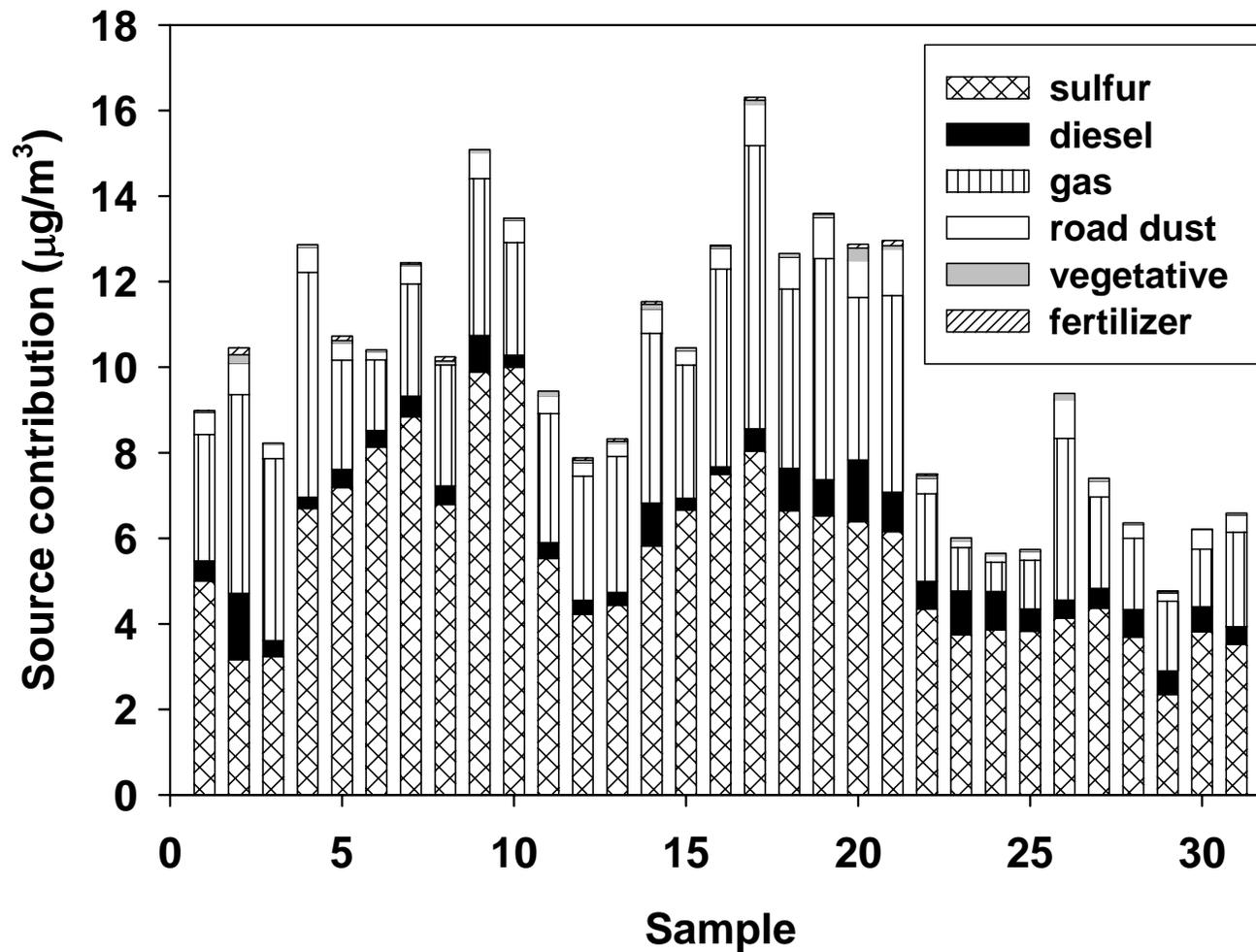


MPIN Matrix:

SPECIES	CLRTON	NFRDSL	PTLEAF	EWHTPS	EHMLKS	GASLIN	FRY90
ECC	0.01	<b>1.00</b>	-0.06	0.02	-0.01	-0.66	-0.42
<b>NAL03C</b>	<b>0.01</b>	<b>0.03</b>	<b>0.01</b>	<b>0.00</b>	<b>0.01</b>	<b>0.03</b>	<b>0.00</b>
<b>NAL04C</b>	<b>0.05</b>	<b>0.15</b>	<b>0.01</b>	<b>-0.01</b>	<b>0.01</b>	<b>-0.06</b>	<b>0.13</b>
<b>NAL05C</b>	<b>0.00</b>	<b>-0.05</b>	<b>0.03</b>	<b>0.00</b>	<b>0.00</b>	<b>0.21</b>	<b>0.15</b>
<b>NAL06C</b>	<b>0.09</b>	<b>0.39</b>	<b>-0.01</b>	<b>-0.02</b>	<b>0.01</b>	<b>-0.18</b>	<b>-0.08</b>
<b>NAL07C</b>	<b>0.03</b>	<b>0.11</b>	<b>0.30</b>	<b>-0.01</b>	<b>-0.01</b>	<b>-0.05</b>	<b>0.00</b>
<b>NAL08C</b>	<b>0.09</b>	<b>0.34</b>	<b>0.03</b>	<b>-0.01</b>	<b>-0.01</b>	<b>-0.32</b>	<b>0.00</b>
NAL09C	0.00	-0.02	<b>0.79</b>	0.02	-0.08	0.00	0.03
NAL11C	-0.01	-0.05	<b>1.00</b>	0.02	-0.10	0.03	-0.03
<b>NAL12C</b>	<b>0.00</b>	<b>0.00</b>	<b>0.08</b>	<b>0.00</b>	<b>-0.01</b>	<b>0.00</b>	<b>0.00</b>
<b>NAL13C</b>	<b>-0.01</b>	<b>-0.02</b>	<b>0.40</b>	<b>0.01</b>	<b>-0.04</b>	<b>0.01</b>	<b>-0.01</b>
NACD6C	-0.01	-0.22	-0.02	0.01	0.02	0.14	<b>0.81</b>
NACD8C	-0.03	0.11	-0.05	0.00	0.00	-0.09	<b>1.00</b>
SNDPMC	-0.03	-0.03	0.00	<b>0.59</b>	-0.13	0.02	0.01
<b>OXDROC</b>	<b>0.00</b>	<b>0.00</b>	<b>0.00</b>	<b>0.07</b>	<b>0.01</b>	<b>0.00</b>	<b>0.00</b>
SYALDC	-0.04	0.00	0.11	-0.19	<b>0.98</b>	0.01	-0.04
ACSYNC	-0.03	0.00	-0.07	-0.22	<b>1.00</b>	0.00	-0.02
<b>SYNCDC</b>	<b>0.00</b>						
LEVOGC	-0.03	0.00	-0.05	-0.16	<b>0.73</b>	0.00	-0.02
CHLTLC	-0.01	-0.15	-0.02	0.01	-0.02	0.09	<b>0.53</b>
HOP2C	-0.14	-0.14	0.00	0.00	0.00	<b>0.90</b>	0.00
HOP3C	-0.15	-0.17	0.00	0.00	0.00	<b>1.00</b>	0.01
<b>STRN1C</b>	<b>0.00</b>	<b>0.05</b>	<b>0.00</b>	<b>0.00</b>	<b>0.00</b>	<b>-0.04</b>	<b>-0.02</b>
<b>STR1BC</b>	<b>0.00</b>						
BNZBF	<b>0.53</b>	0.00	0.00	0.00	0.00	-0.06	0.00
BNZKF	<b>0.82</b>	0.00	-0.01	0.00	0.00	-0.08	0.00
BNZEP	<b>0.69</b>	0.00	-0.01	-0.01	0.00	-0.04	-0.01
INDCDP	<b>0.62</b>	-0.01	0.00	-0.01	0.00	-0.04	0.00
BNZGHI	<b>0.67</b>	-0.05	0.00	-0.01	-0.01	0.09	0.01
CORON	<b>1.00</b>	-0.10	-0.01	0.02	0.01	0.16	0.02
RSNSMC	-0.05	-0.05	0.00	<b>1.00</b>	-0.33	0.03	0.02
SRHHPN	-0.10	-0.06	0.00	0.00	0.00	<b>0.62</b>	-0.02
SRBHPN	-0.09	-0.01	-0.01	0.00	0.00	<b>0.51</b>	-0.03



# TACS CMB Results with limited “DEARS” source markers



3) Can we achieve low enough uncertainties to adequately resolve sources?\*

\*How well are we estimating uncertainty?



## QA Based on EPA Method 8000 Series with modifications

-Initial Demonstration of Proficiency includes:

- Initial calibration
- Multi-level replicate recoveries

- Each batch of 12 samples includes:

- 2 replicate injections
- 1 method blank
- 1 matrix control
- 1 calibration verification standard
  - level?
  - acceptance criteria? (next slide)

- Each sample includes:

- > internal standards by class, volatility (return to later)
- > Surrogate standards by class, volatility



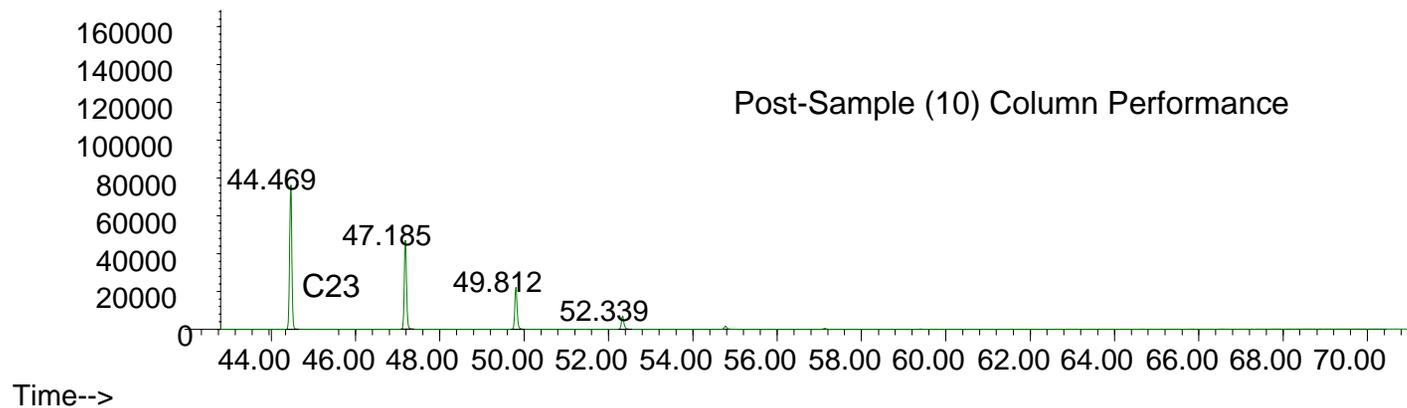
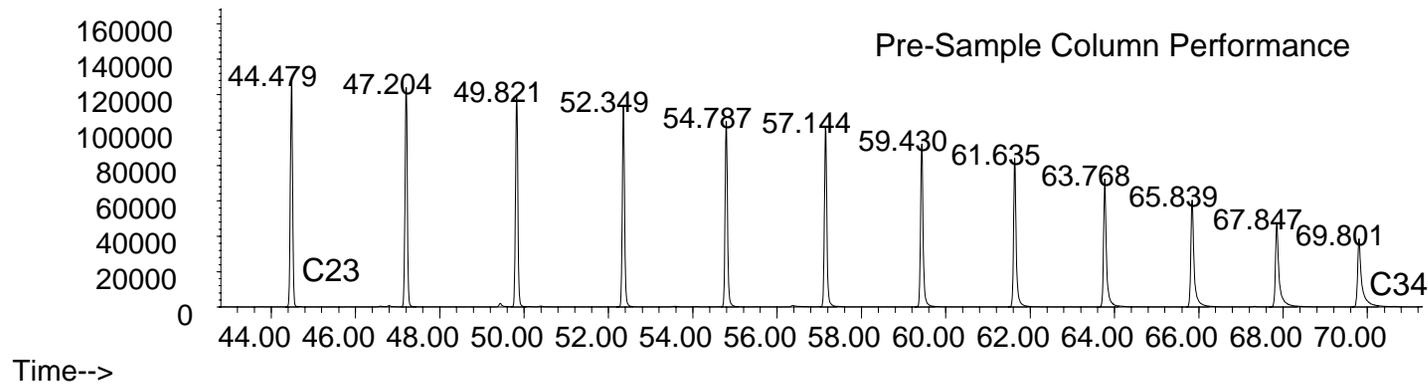
What if we accept calibration if 80% of species are within 20%?

- Important Source Markers:

- > C29 and C31 alkanes
- > Indenopyrene + Benzoperylene
- > Hopane + Norhopane



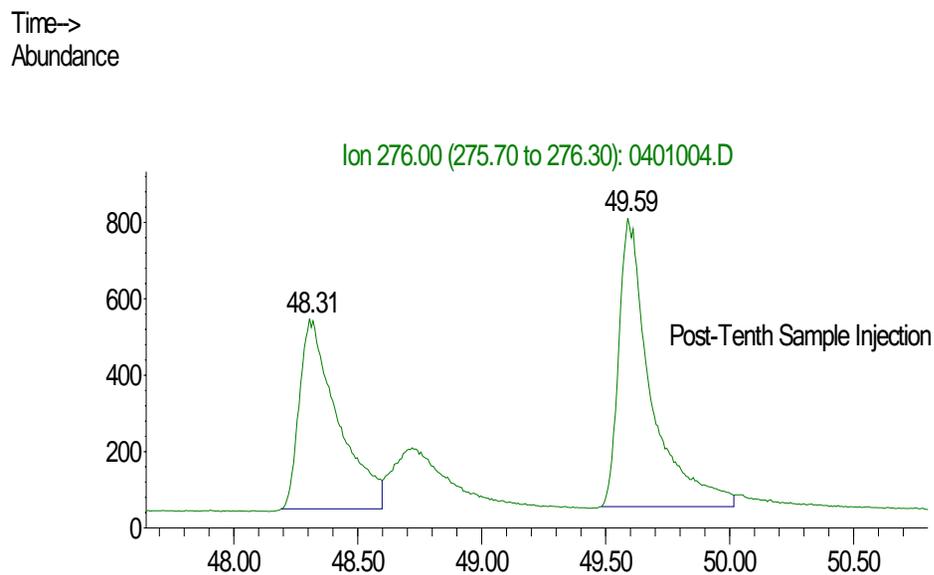
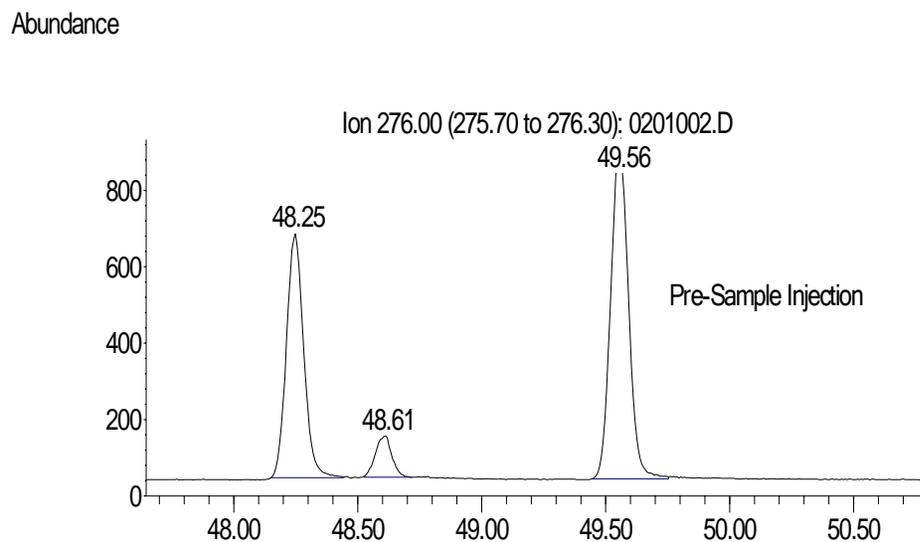
# Pre-Sample vs. Post-Sample Column Performance (TACS)



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# Effect of Column Deterioration on indenopyrene + benzoperylene



Time-->



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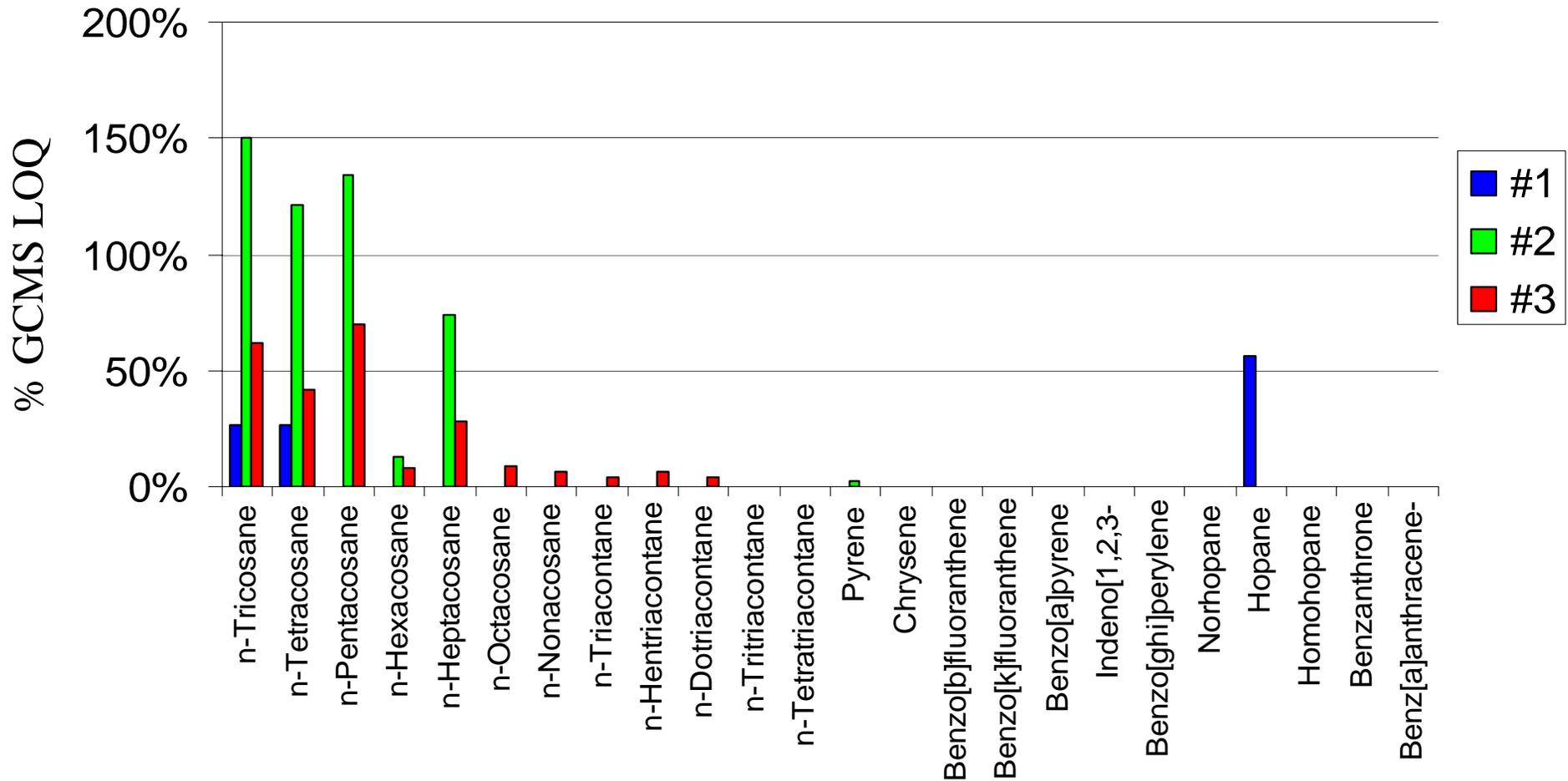
Each week of sampling includes:

- Samples for Tuesday through Sunday
- Duplicate sample each Wednesday (2 seasons)
- Field Blank (placed in sampler for 1 day with no flow)\*
- Travel Blank (not removed from shipping container)\*

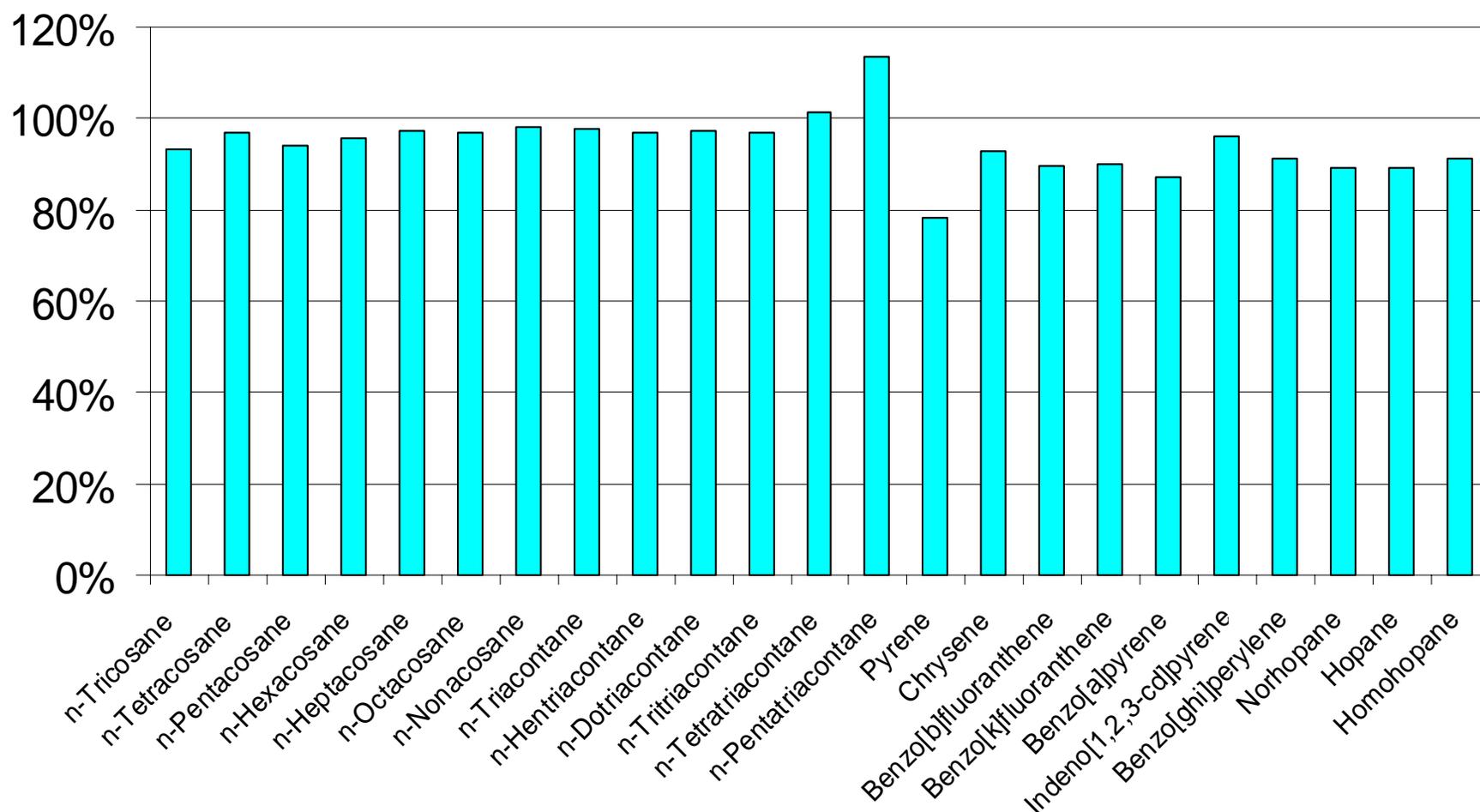
\*Shipping & storage described in detail AS&T, in revision



# DEARS Matrix Blanks



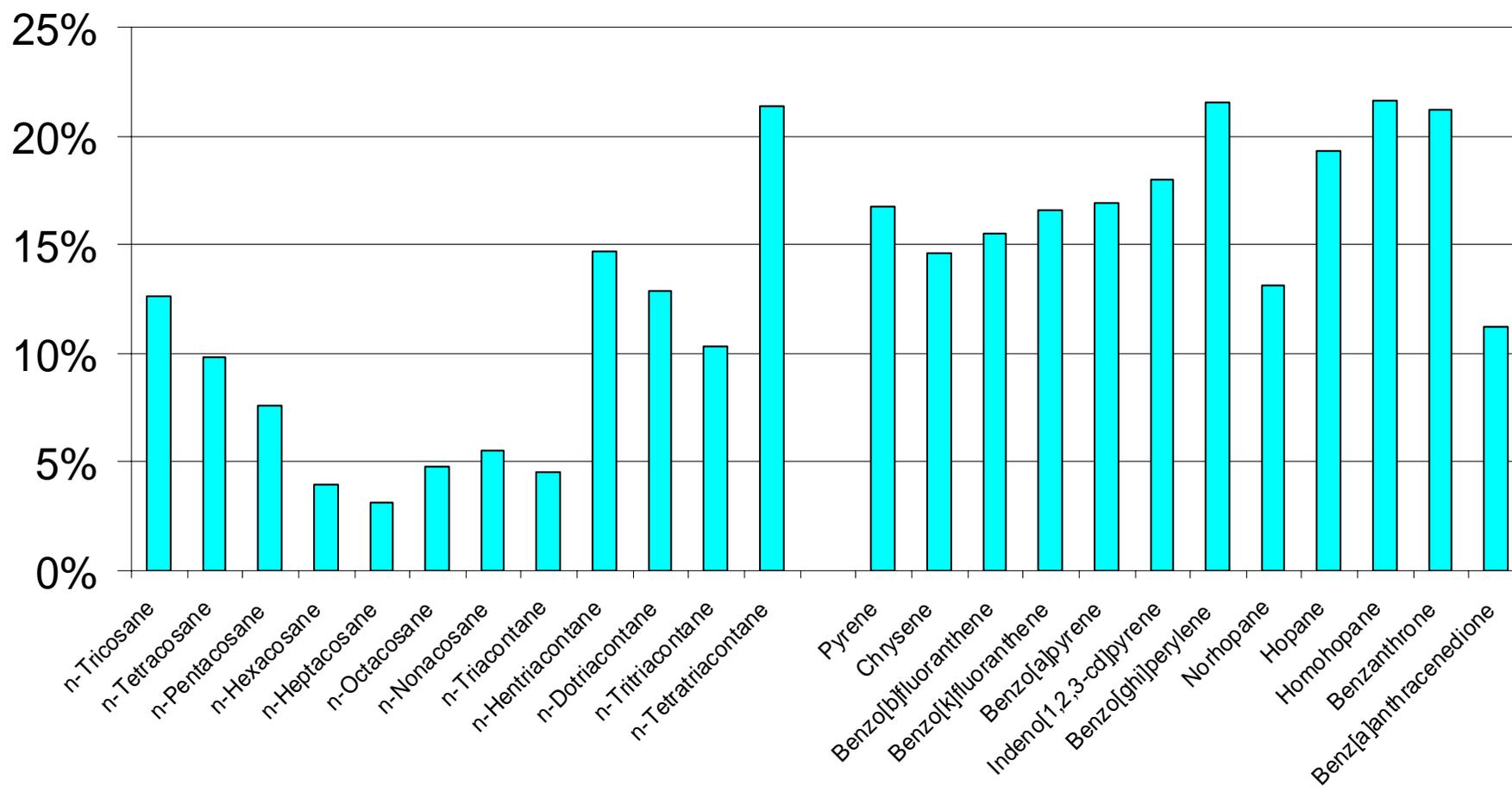
# Preliminary Recovery



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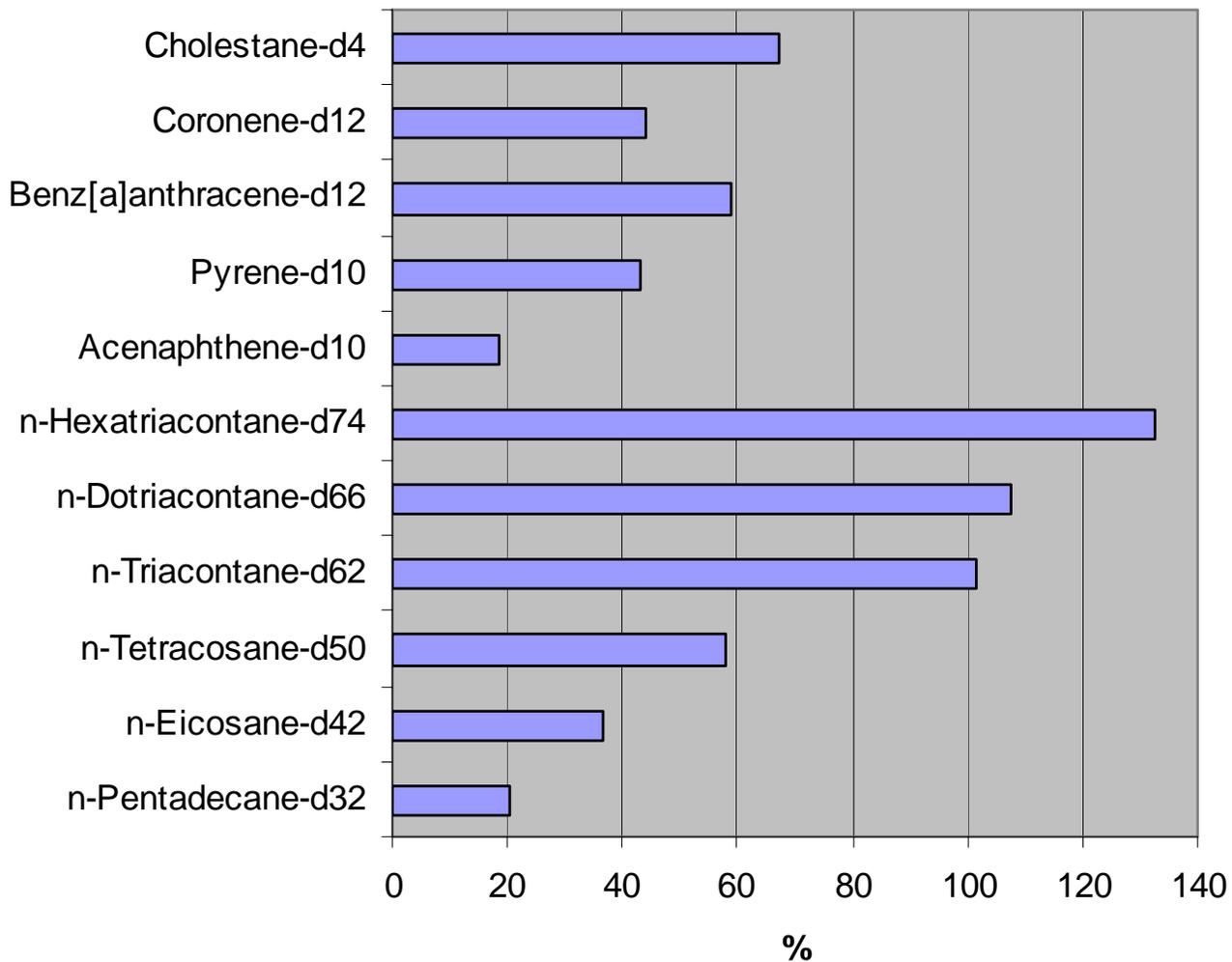
# Duplicate Analysis



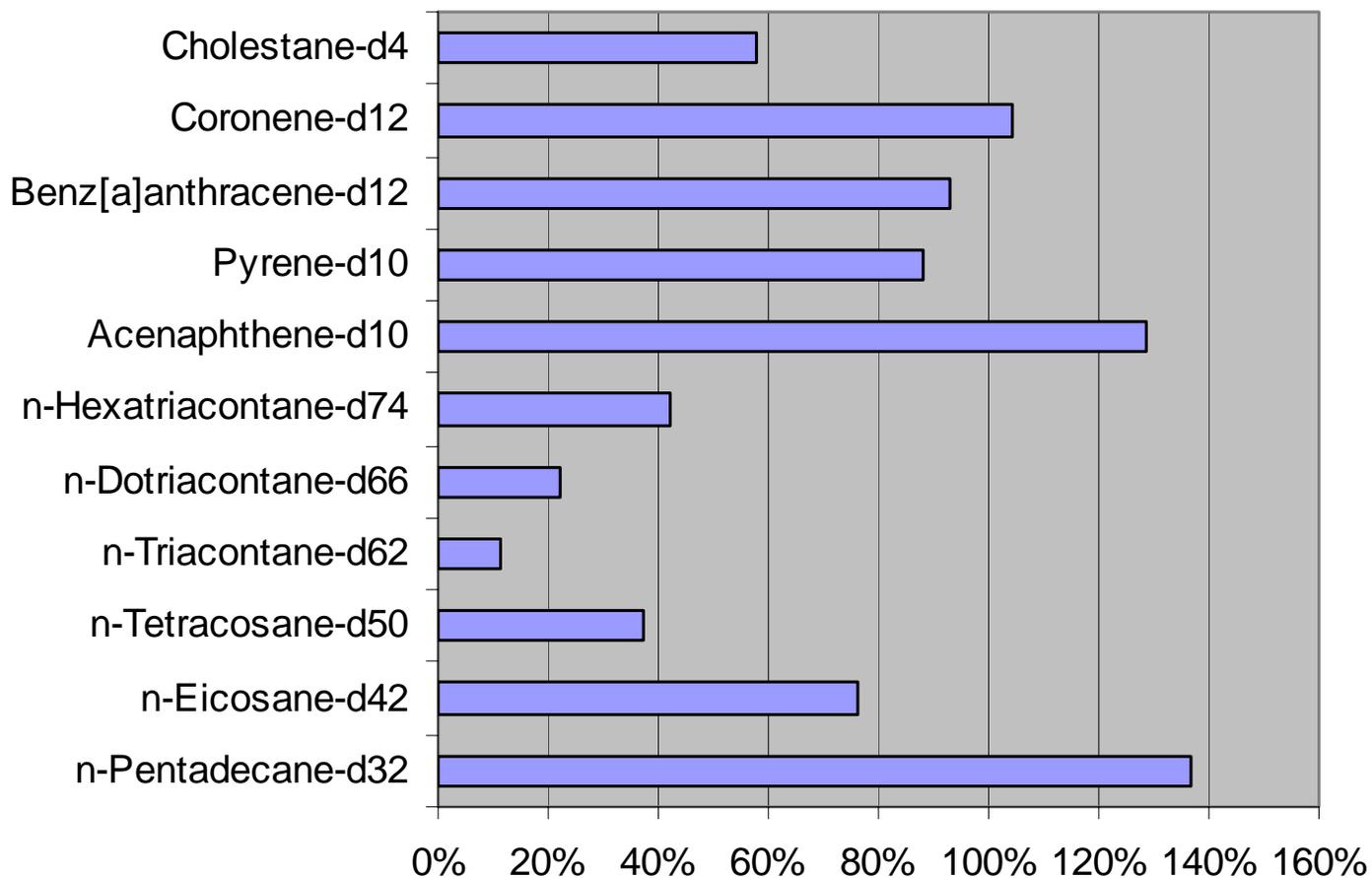
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### Sample Surrogate Recovery (n=34)



## Sample Surrogate Variability (n=34)



## Summary of Season 1 High Volume Samples

Analytical blanks – acceptable

Analytical precision – acceptable

Sampling blanks & precision – not evaluated

Calibration frequency – **not acceptable**

- > solid phase extract step required
- > (extremely low concentrations!)

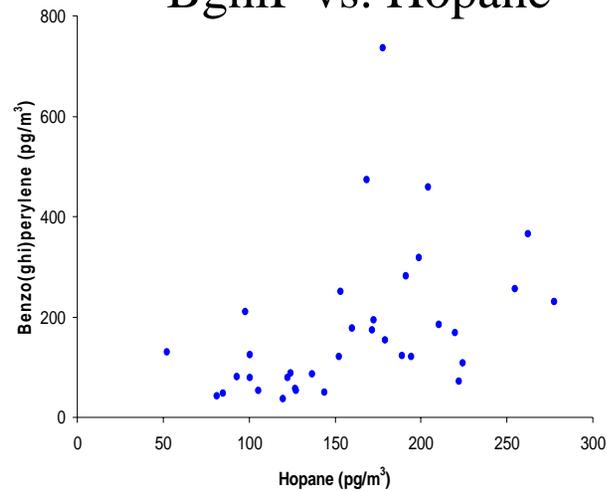
Surrogate recoveries – **not acceptable**

- > collected PM strongly influences surrogate recovery
- > internal standards for a range of volatility are necessary
- > surrogate standards need to better align with targets

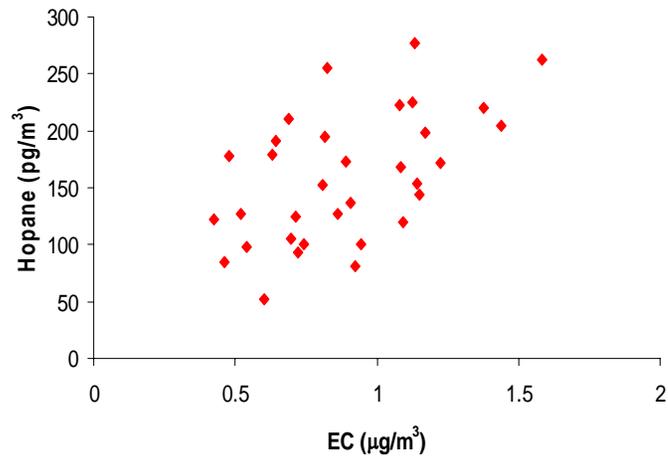


# Associations Between Key Species

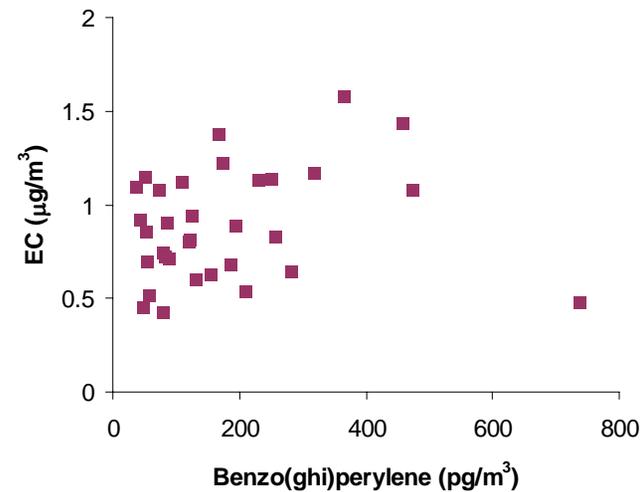
## BghiP vs. Hopane



## Hopane vs. EC



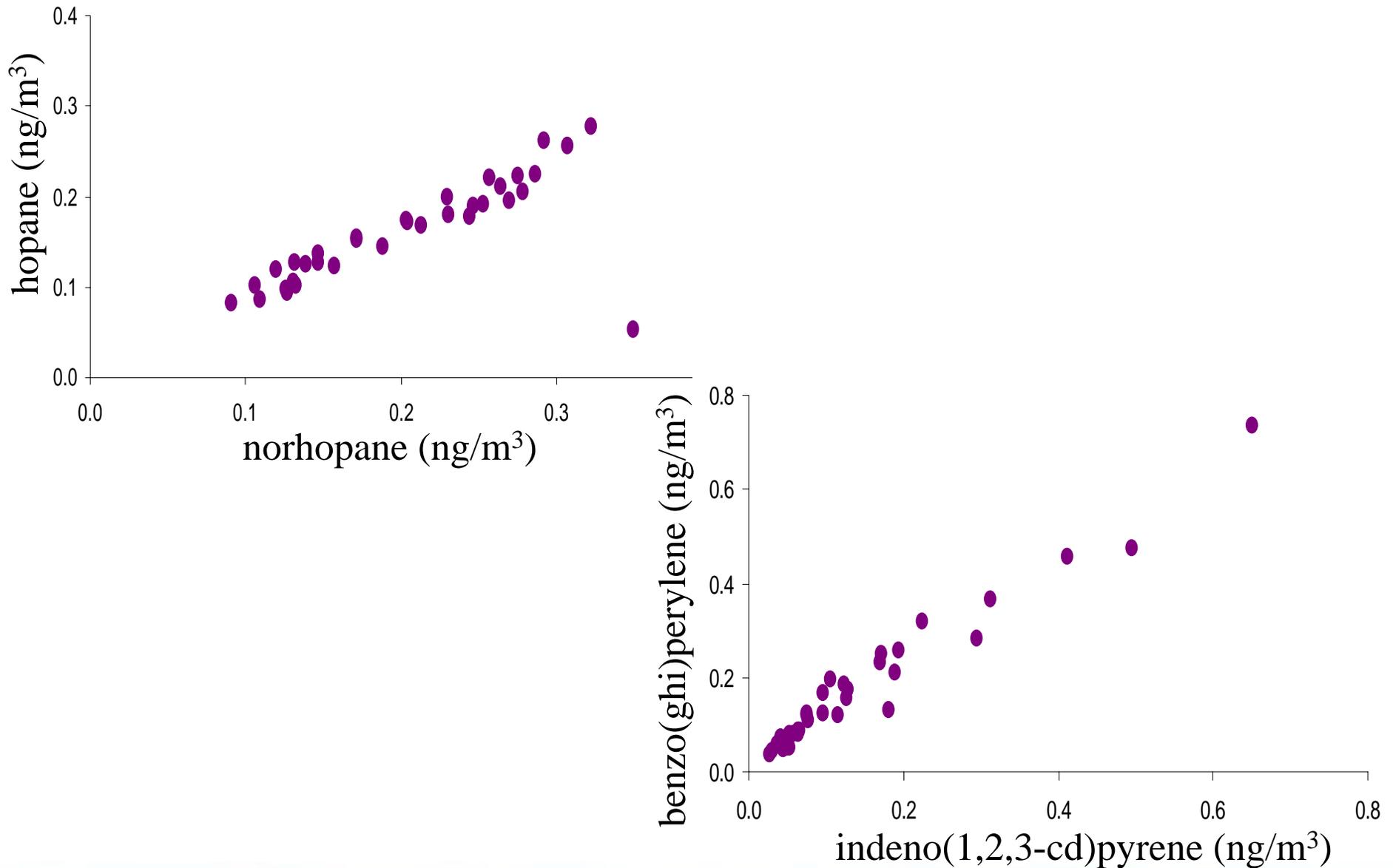
## BghiP vs. EC



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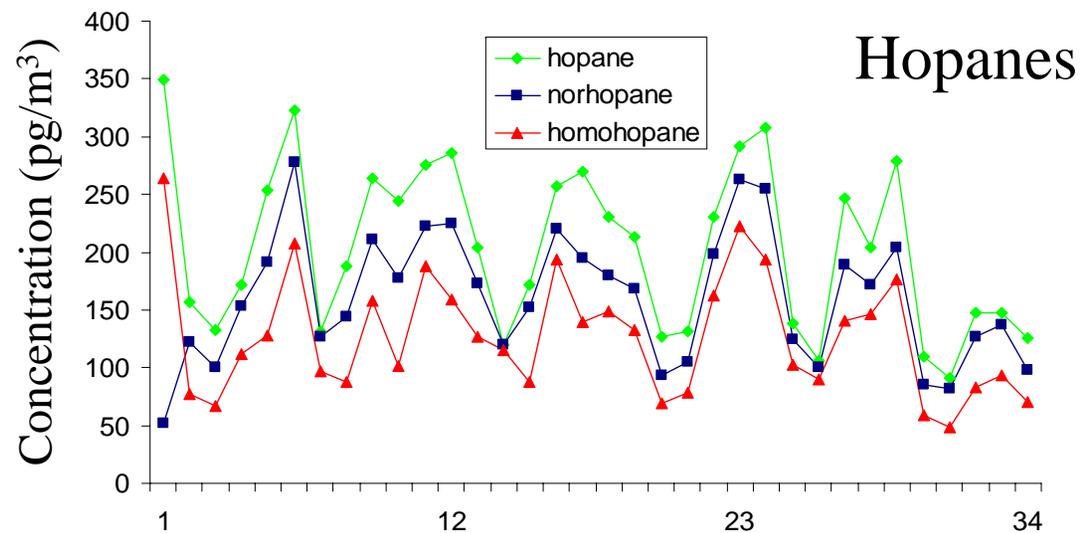
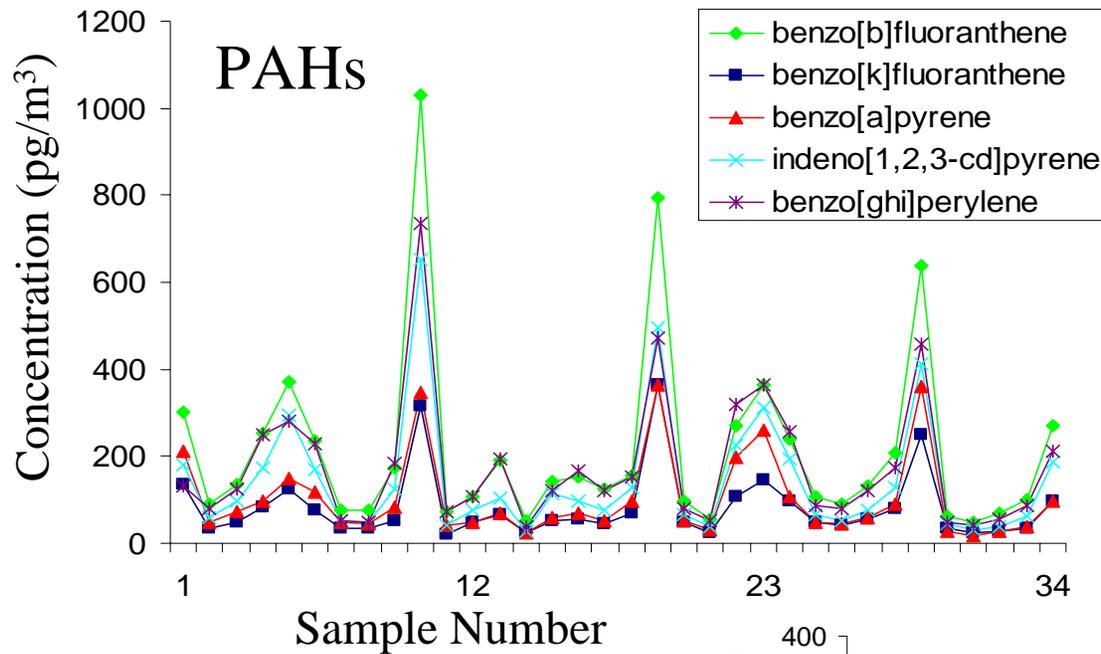
# Associations Between Key Species



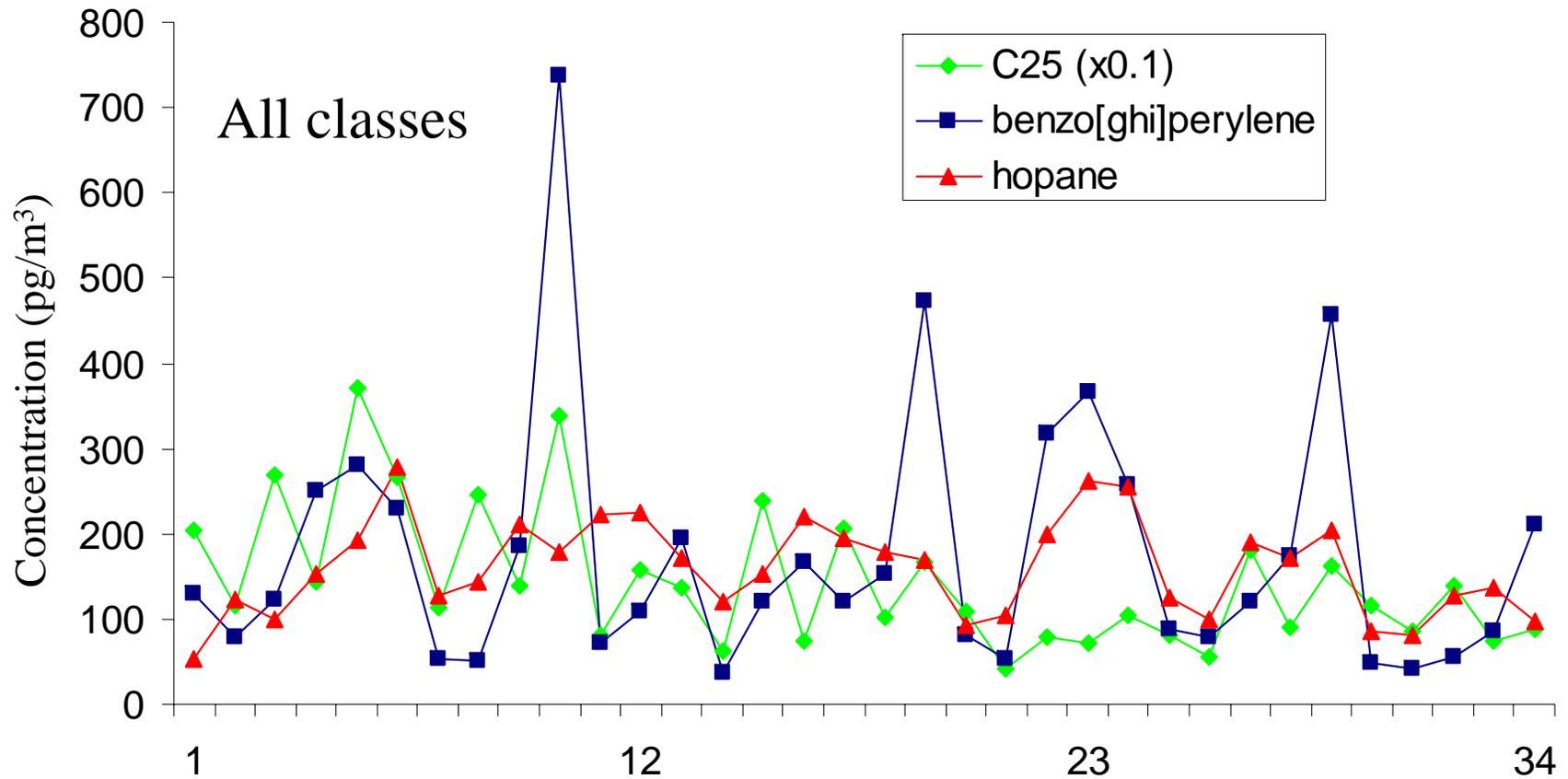
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# Variation in Daily Concentration



# Variation in Daily Concentration



# Conclusions

- Average concentrations fairly low (~100 to 200 pg/m<sup>3</sup>)
- Strong associations observed *between* different PAHs and between different hopanes
- Associations of PAHs *with* hopanes (e.g. hopane vs. benzo[ghi]perylene) generally not as strong
- PAH concentrations exhibited considerably more day to day variability (CV ~ 100%) than hopane concentrations (CV<50%)



## *Next Steps*

- *Add Solid Phase Extraction Step*
- *Determine Method Detection Limits*
- *Demonstrate Method Proficiency (recoveries)*
- *Proceed with Seasons 2-6 and low volume samples*

