Advanced Analytical Methods for Fuels Characterization

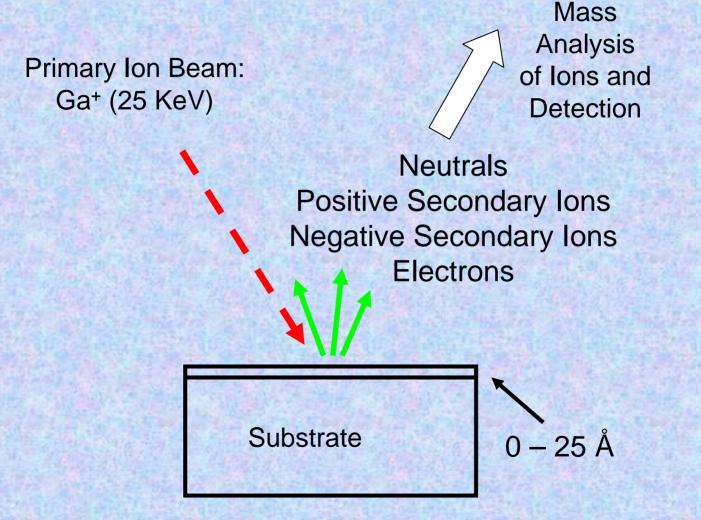
A Time-of-Flight Secondary Ion Mass Spectrometry (ToF-SIMS)/ Chemometrics Analysis of Coal

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> > Presentation given at ACERC February 28, 2007

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Static ToF-SIMS: A Chemical Microscope



Note: 4 eV corresponds to ~ 92 kcal/mol

BYU Time-of-Flight Secondary Ion Mass Spectrometer (ToF-SIMS)

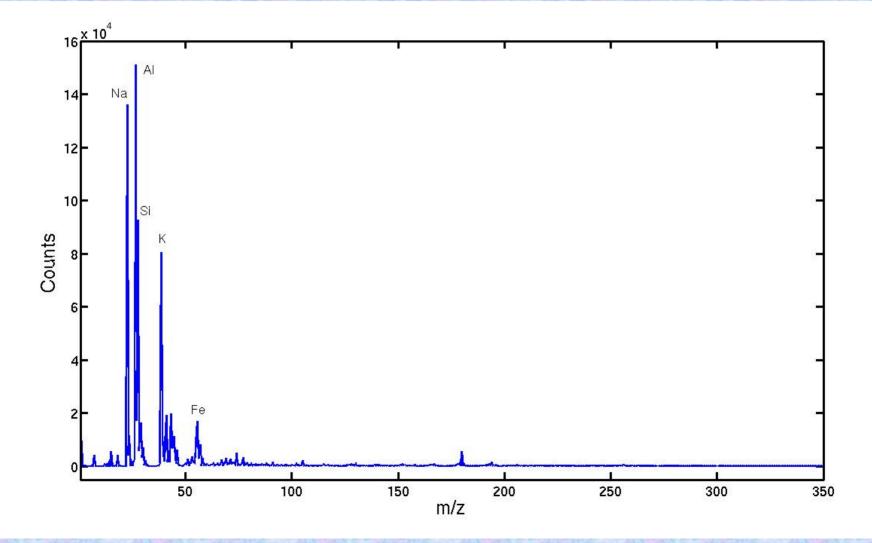


ToF-SIMS Analysis of Coal

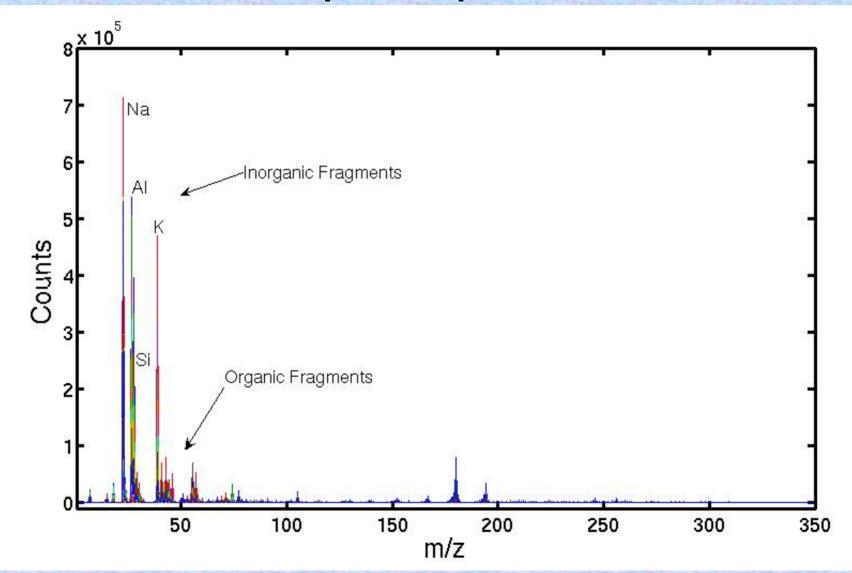
Sample Prep:

- Coal samples were ground with a mortar and pestle
- They were immediately mounted on double sticky tape
- 500μm x 500μm regions were then analyzed by ToF-SIMS
- ca. 100 spectra were taken of ca. 30 different coal samples

Average ToF-SIMS Spectrum of Coal Samples

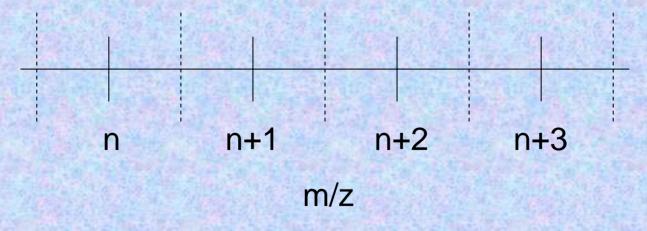


All of the ToF-SIMS Spectra Superimposed

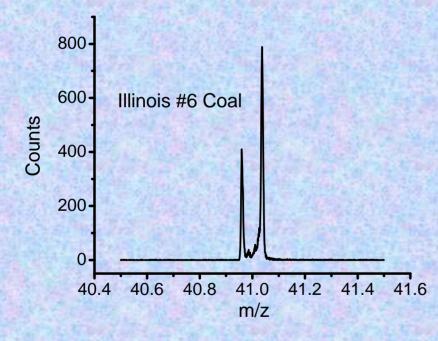


Data Preprocessing

- To simplify spectral analyses of complicated spectra, mass spectral data are commonly binned, typically to unit mass resolution.
- For our spectra we binned 1099 regions.



Binning to Half Unit Mass Resolution

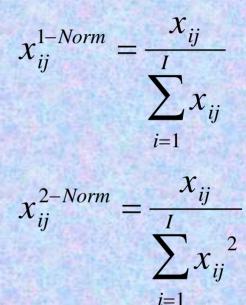


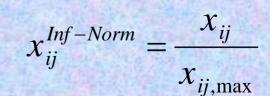
Mass excess vs. mass deficit

Isotope	mass	abundance
¹ H	1.0079	99.99%
⁴ He	4.0026	100.00%
⁷ Li	7.0160	92.48%
⁹ Be	9.0123	100.00%
¹¹ B	11.0093	81.02%
12 C	12.0000	98.89%
¹⁴ N	14.0031	99.64%
¹⁶ O	15.9949	95.77%
¹⁹ F	18.9984	100.00%
²⁰ Ne	19.9924	90.92%

Data Preprocessing, Cont.

- All mass spectra were normalized with the 1-Norm. The 1-Norm is useful for correcting spectra that are identical, except to some constant factor.
- Other possible normalization schemes are the 2-Norm, which creates a vector of unit length, or the Inf-Norm, which gives the maximum peak a value of one.





Normalization, Mean Centering

	Var 1	Var 2	Var 3		Var m	
Spectrum 1	S ₁₁	S ₁₂	S ₁₃		S _{1m}	
Spectrum 2	S ₂₁	S ₂₂	S ₂₃		S _{2m}	
Spectrum 3	S ₃₁	S ₃₂	S ₃₃	:	S _{3m}	
Spectrum n	S _{n1}	S _{n2}	S _{n3}	:	S _{nm}	

mean centering -- a column operation $cen x_{ij} = x_{ij} - \overline{x}_j; \quad \overline{x}_j = \overline{\overline{x}_{ij}}$ (not essential for cluster analysis but critical for PCA)

normalization -- a row operation

 $\sum x_{ij}$

n

Each Spectrum Can be Viewed as a Vector and/or a Point in a Hyperspace

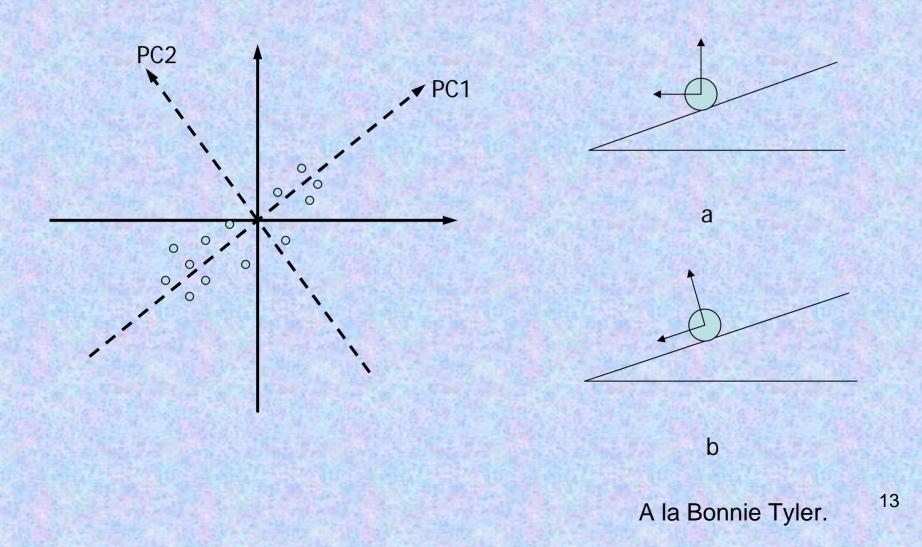
Spectrum 1 = $(S_{11}, S_{12}, S_{13}, ..., S_{1m})$ Spectrum 2 = $(S_{21}, S_{22}, S_{23}, ..., S_{2m})$ Spectrum 3 = $(S_{31}, S_{32}, S_{33}, ..., S_{3m})$

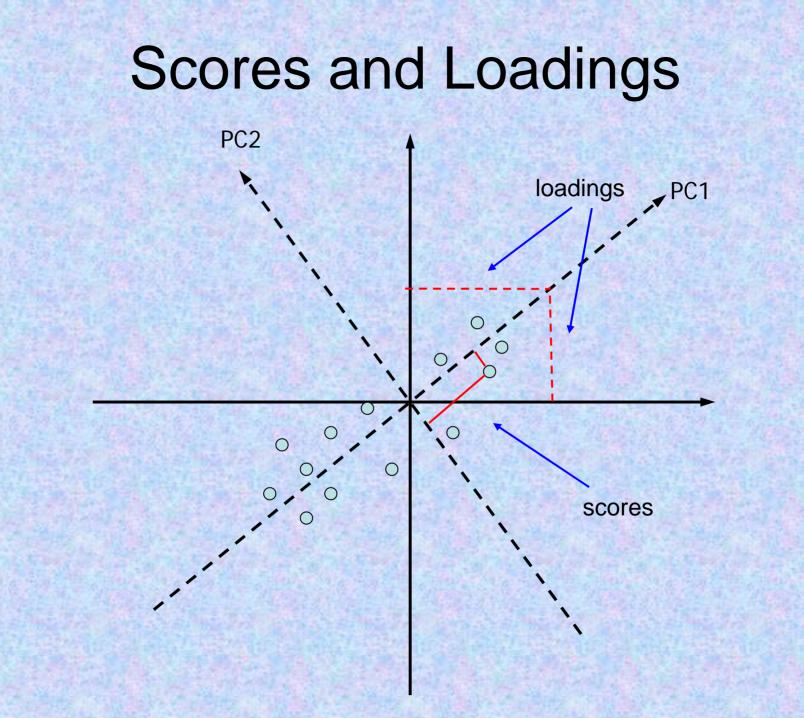
Spectrum n = $(S_{n1}, S_{n2}, S_{n3}, ..., S_{nm})$

Principal Components Analysis (PCA) of Mass Spec Data

- Each peak or spectral region is considered to be a unique variable. Complete spectra are represented as vectors and/or single points in a hyperspace.
- The data are binned, normalized, and mean centered.
- Each spectrum is then "plotted" as a single point in a hyperspace of these variables
- The coordinate system is rotated to capture the maximum variation in the data along the axes.
- This analysis is called Principal Components Analysis (PCA).
- The projections of the original data points (spectra) on the new axes (principal components) are called scores.
- The contributions of the original axes (variables) to the new axes (principal components) are called loadings.

A Simple Way to Explain PCA





Cluster Analysis

- Calculate the distance between points in a hyperspace.
- The simplest measure of distance between two points x_i and x_j in a hyperspace is the Euclidean distance:

$$d_{ij} = \left[\left(\vec{x}_i - \vec{x}_j \right)^T \left(\vec{x}_i - \vec{x}_j \right)^T \right]^{1/2}$$

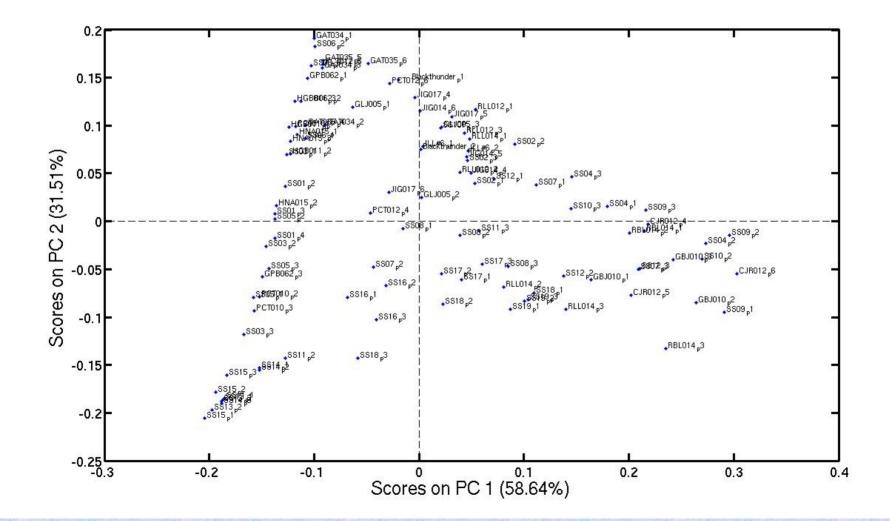
- Create a plot (a dendrogram) that reflects the distances between the points.
- This is an established method that has been used for many years, especially in the biological sciences as a classification tool.

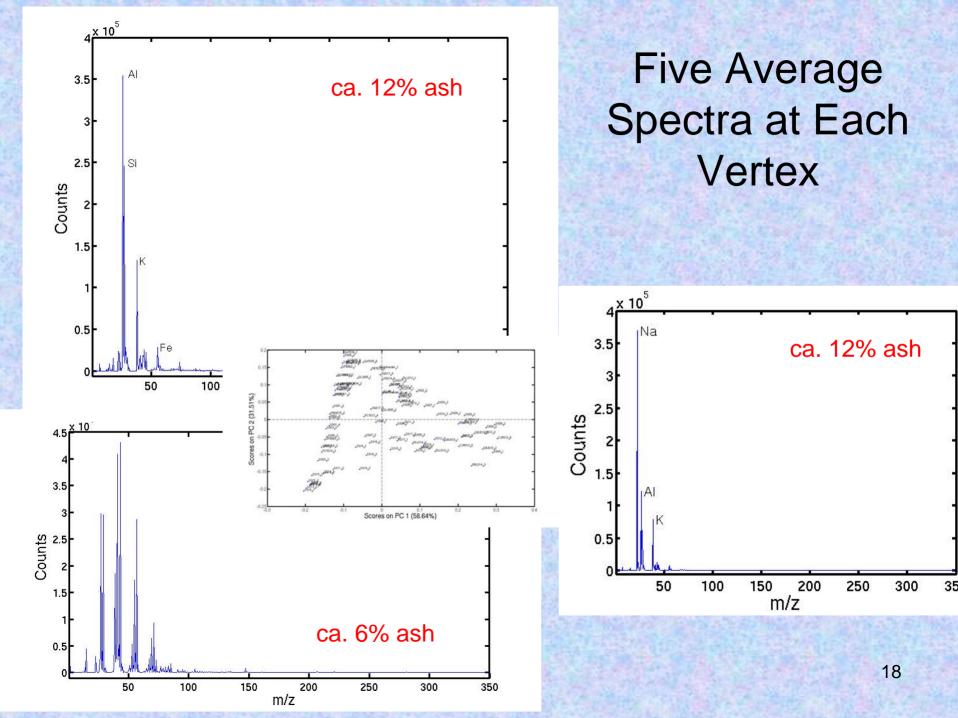
Use the K-Means Method of Clustering

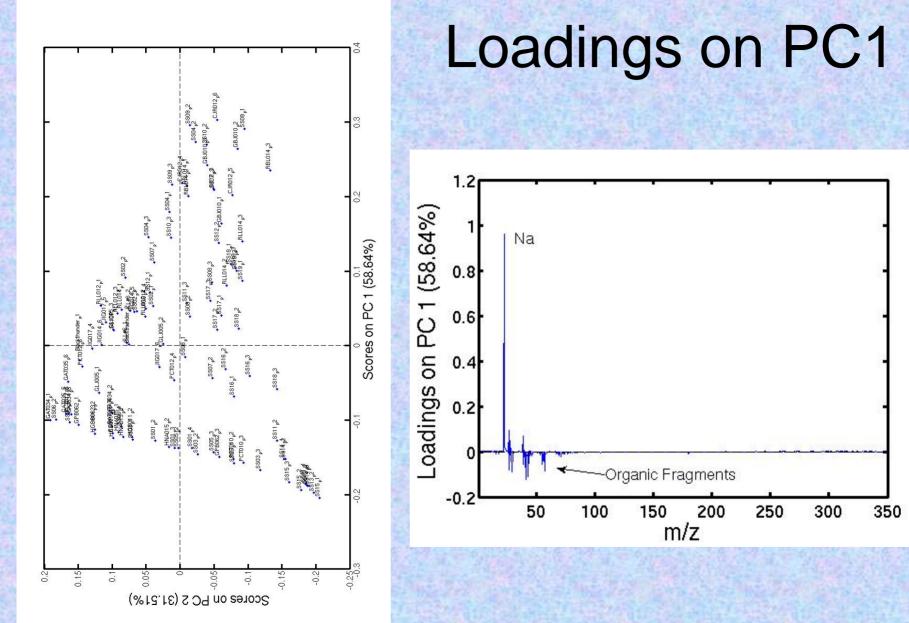
points in a hyperspace

Scores Plot of PC1 vs. PC2

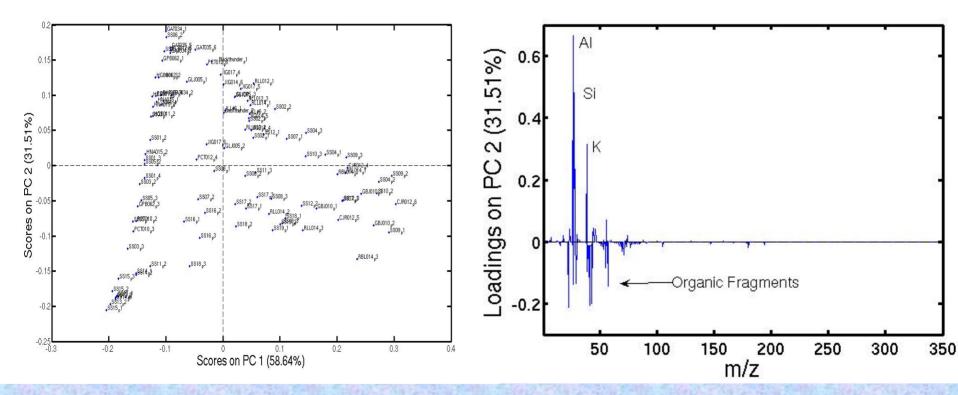
(PC1 and PC2 account for just over 90% of the variance in the data.)



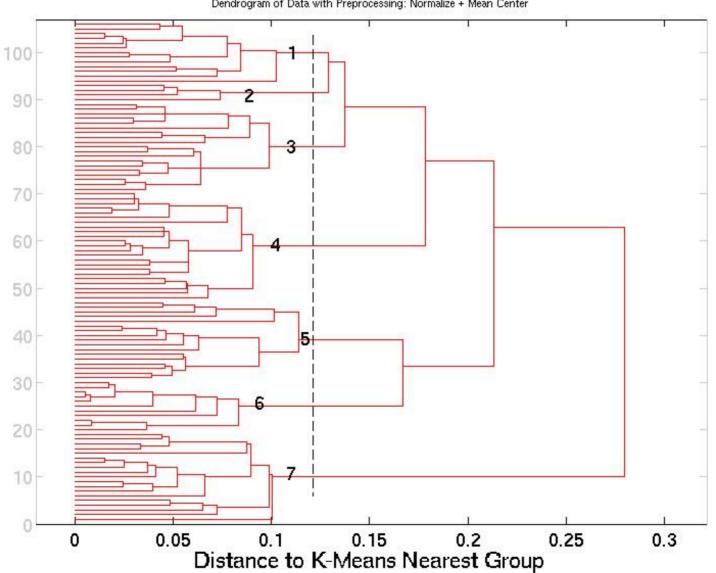




Loadings on PC2

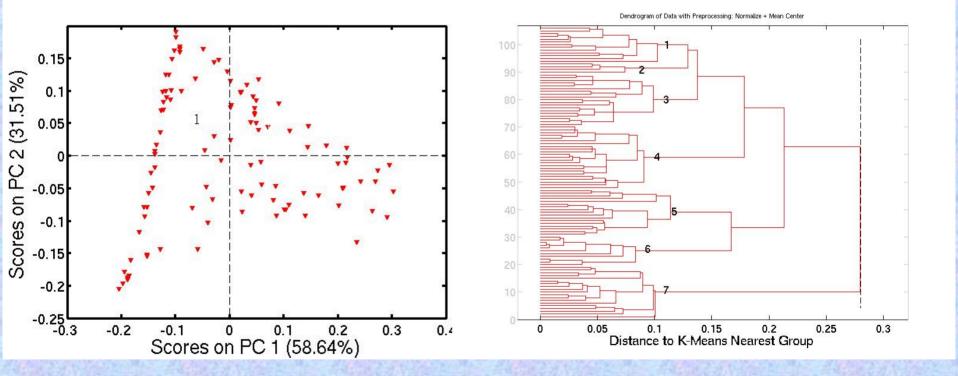


Dendrogram from a Cluster Analysis

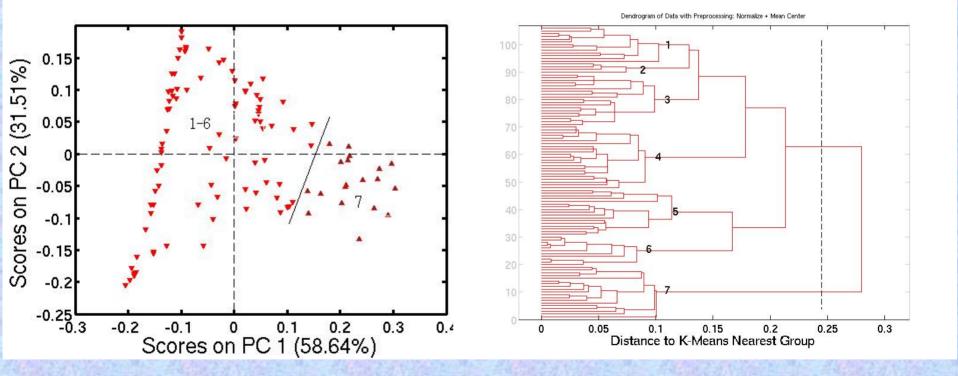


Dendrogram of Data with Preprocessing: Normalize + Mean Center

Integrating the PCA and Cluster Analyses: One Cluster

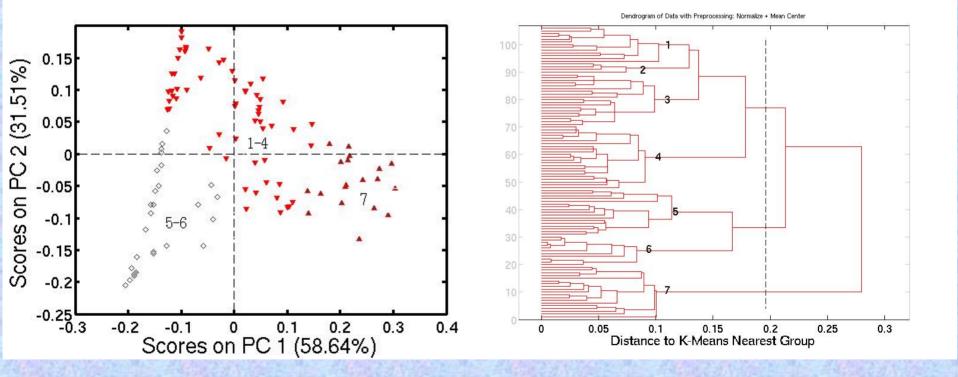


Integrating the PCA and Cluster Analyses: Two Clusters

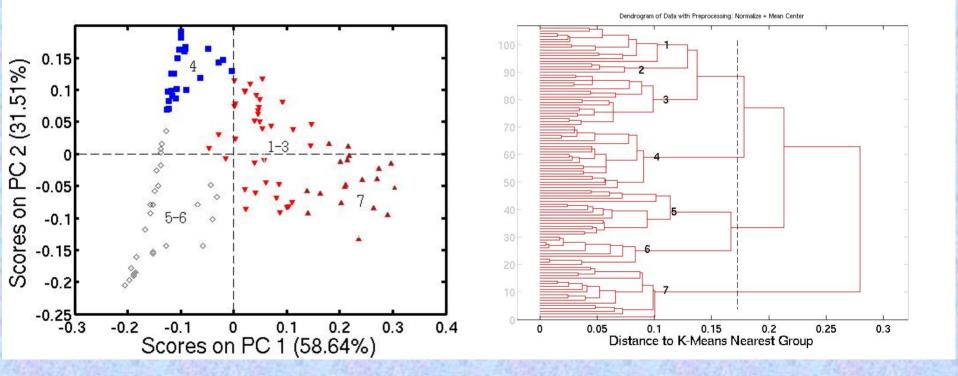


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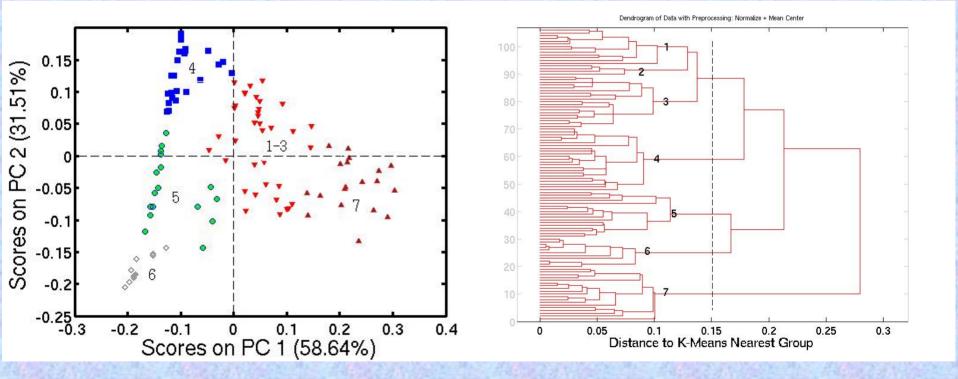
Integrating the PCA and Cluster Analyses: Three Clusters



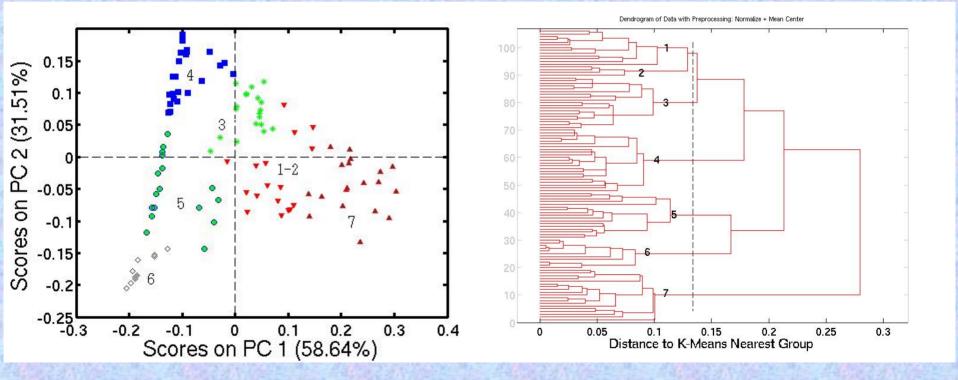
Integrating the PCA and Cluster Analyses: Four Clusters



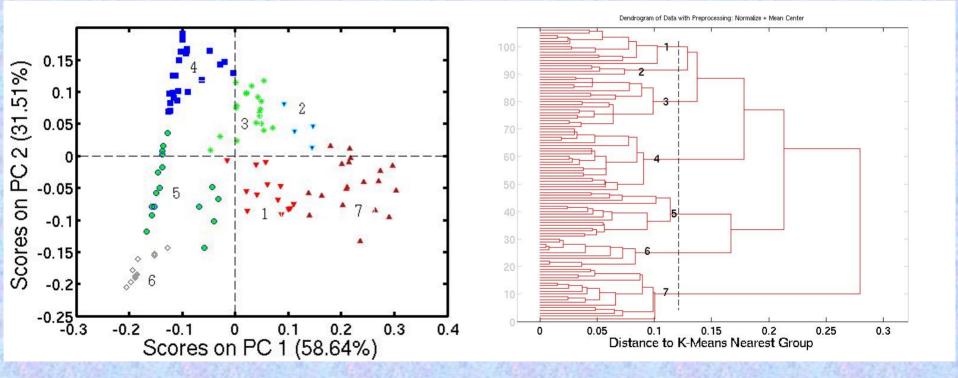
Integrating the PCA and Cluster Analyses: Five Clusters



Integrating the PCA and Cluster Analyses: Six Clusters

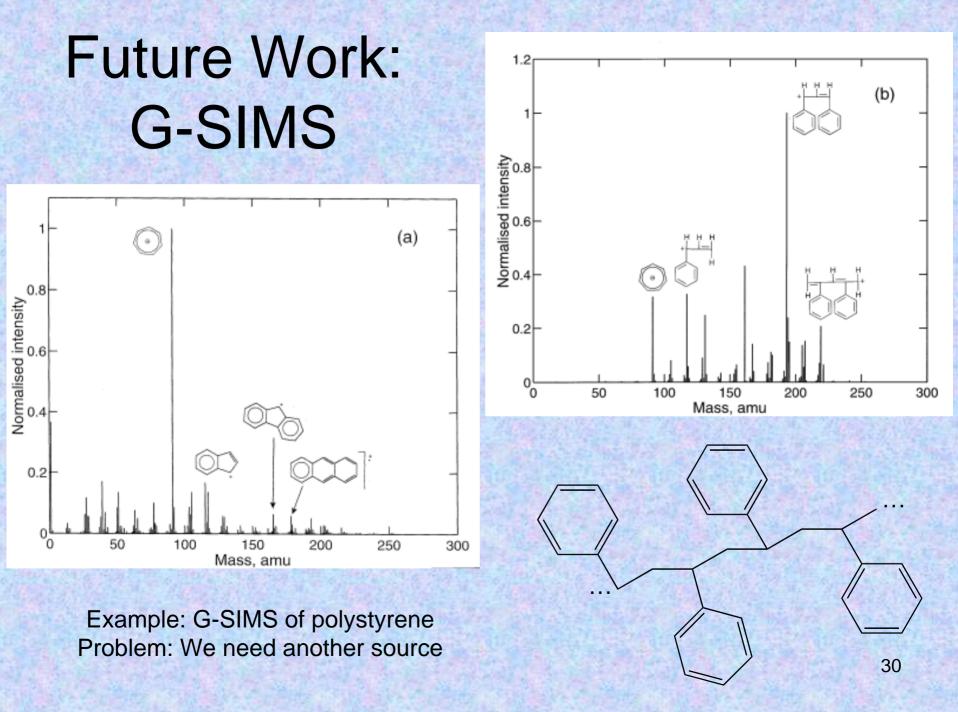


Integrating the PCA and Cluster Analyses: Seven Clusters



Future Work: New Sources

- New sources, such as gold and bismuth clusters, or C₆₀ ions can produce significantly more high mass, organic ions than monatomic sources, such as Ga⁺.
- High mass ions are highly characteristic of the sample they come from.
- These ion sources would allow some speciation of the coal.
- They should also increase the number of organic, compared to inorganic, ions that are produced by ToF-SIMS.



Conclusions

- Good spectra can be obtained when coal is analyzed by ToF-SIMS.
- The spectra contain both an organic and an inorganic component.
- Principal components analysis and cluster analysis group the spectra in a similar way.
- There seems to be some correlation between the ToF-SIMS spectra and the physical/chemical properties of the coal samples.
- Other sources would allow us to obtain more high-mass ions, which are highly characteristic of the materials they come from.
- Other sources would also allow us to do G-SIMS.

Thank You