# Lessons Learned from NIST's Two Recent Polymer MS Interlaboratory Comparisons

Charles M. Guttman<sup>1</sup>, <u>William E. Wallace<sup>1</sup></u>, Stephanie J. Wetzel<sup>2</sup>, Kathleen M. Flynn<sup>1</sup>, David L. VanderHart<sup>1</sup> <sup>1</sup>National Institute of Standards and Technology, Gaithersburg, MD <sup>2</sup>Duguesne University, Pittsburgh, PA

### Purpose for Interlaboratory Comparison

- Compare data among laboratories as a check on technique consistency.
- Obtain some measure of current experimental practice.
- Compare aggregate MS data from many labs with measures of the same properties by classical methods, i.e. "benchmarking".
- Begin a dialog in the synthetic polymer MS community on issues related to the quantitative measurement of singlechain properties by MS.
- NIST has a national mandate to produce measurement standards and to monitor the US Measurement System.

#### Participation

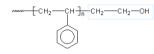
- First Interlaboratory Comparison
  - 20 Laboratories returned data
    - 10 Industrial Labs
    - 4 Government Labs
      6 Academic Labs
  - 0 Countries Door
- 8 Countries Represented
- Second Interlaboratory Comparison
  - 14 Laboratories returned data
    - 5 Industrial Labs
    - 5 Government Labs
  - 4 Academic Labs
  - 7 Countries Represented

#### Results from First Interlaboratory Comparison of Polystyrene 7000 u to obtain Molecular Mass Distribution

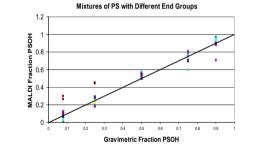
- Comparison of MALDI with Classical Methods
- Light Scattering gives M<sub>w</sub> = 7300 u uncertainty including type A & type B ±600 u
- NMR gives M<sub>n</sub> = 7050 u uncertainty including type A & type B ±400 u
- MALDI Interlaboratory Data Average
   M<sub>w</sub> = 6700 u type A uncertainty ±85 u
   M<sub>a</sub> = 6600 u type A uncertainty ±125 u
- Data Analysis
- Three outliers: All due to data analysis errors
- Weakness of analysis software
   Background/Baseline Subtraction
- Peak Integration
- Smoothing
- $M_n$  computed by participants at times did not agree with the  $M_n$  we computed from their data
- <u>These errors are easily preventable by more careful</u> use of data analysis software programs.
- For All Moments
- The variance within each lab is less than the variance among labs.
- There is no statistically significant difference between retinoic acid and dithranol as matrices.
- There is no statistically significant difference between moments for data run in linear or reflectron mode.
- There is a statistically significant difference between instruments.
- Conclusions
- MALDI seems to be consistently lower than classical methods by about 3% to 5%.
- There seems to be differences that are easily corrected by improved calibration and analysis methods.
- The overall agreement between labs is very good for this polymer in this mass range.
- Differences between MALDI procedures, (matrix, instrument, etc.) can often be seen statistically but were generally small.

#### Results from Second Interlaboratory Comparison on Mixtures of Polystyrene with Different End Groups – A Study of Mass Fraction

One PS had  $-CH_2CH_2OH$  as an end group (shown below outlined in light blue). The second PS had -H as an end group.



Comparison of MALDI-TOF-MS Results with Gravimetric Results



- Some Further Results
- Lab-to-Lab variation is significant
- Calibration Problems with about 30% of Labs
   Identification of end groups
- Instrument variation is significant
  - S/N variation
  - · Some Instruments had problems with the recipe
- Conclusions
- Overall MALDI is in Good Agreement with Gravimetric Methods
- Strong Variation amongst Labs
- Calibration again an important issue
- Optimization of Instrument for S/N

## **Lessons Learned**

1) The instrument must be optimized for best signal to noise in order to identify the minor components in a mixture. At this time there seems to be no generally-accepted, systematic procedure or set of necessary and sufficient criteria to optimize MALDI-TOF mass spectrometers.

2) There is a need for generally-accepted practices for data analysis. This includes, but is not necessarily limited to, baseline subtraction, and peak integration. These procedures need to be supported by statistical theory in order that they may also provide meaningful uncertainties.

3) Instrument mass calibration must be carefully performed. This is best done under measurement conditions as close to the analyte measurement conditions as is feasible. Biomacromolecules may not be the best choice for calibration when the analyte to be measured is a synthetic polymer due to the large differences in operating parameters required for these classes of sample.

#### References

Guttman, C.M.; Wetzel, S.J.; Blair, W.R.; Fanconi, B.M.; Girard, J.E.; Goldschmidt, R.J.; Wallace, W.E.; VanderHart, D.L. *Anal. Chem.*, 2001, **73**, 1252-1262.

Guttman, C.M.; Wetzel, S.J.; Flynn, K.R.; Fanconi, B.M.; J.; Wallace, W.E.; VanderHart, D.L. *Anal. Chem.*, 2005, in press.

## **Other Outputs**

A. See our web site:

www.nist.gov/maldi

For our Polymer MALDI Recipes page, *MassSpectator* our online peak integration tool, and announcements about our Fall 2005 Workshop (Nov. 9-10).

B. ASTM Standard Test Method to Determine the Molecular Mass Distribution of Polystyrene using MALDI Approved as ASTM D7134 This standard uses the *Lessons Learned* from the interlaboratory comparison studies.