

Making SimDist Faster and More Robust

Minimizing the impact of retention time drift

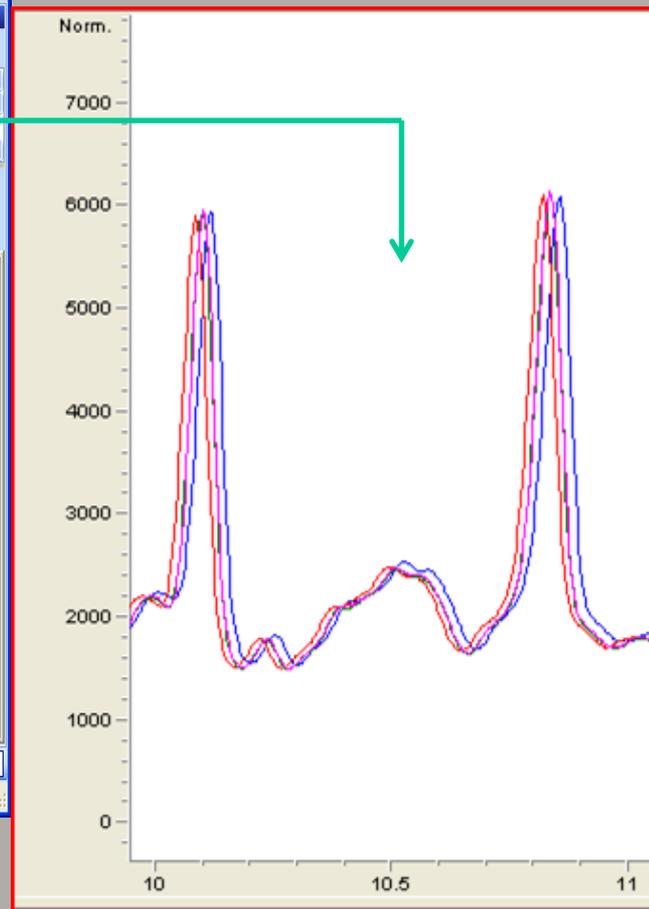
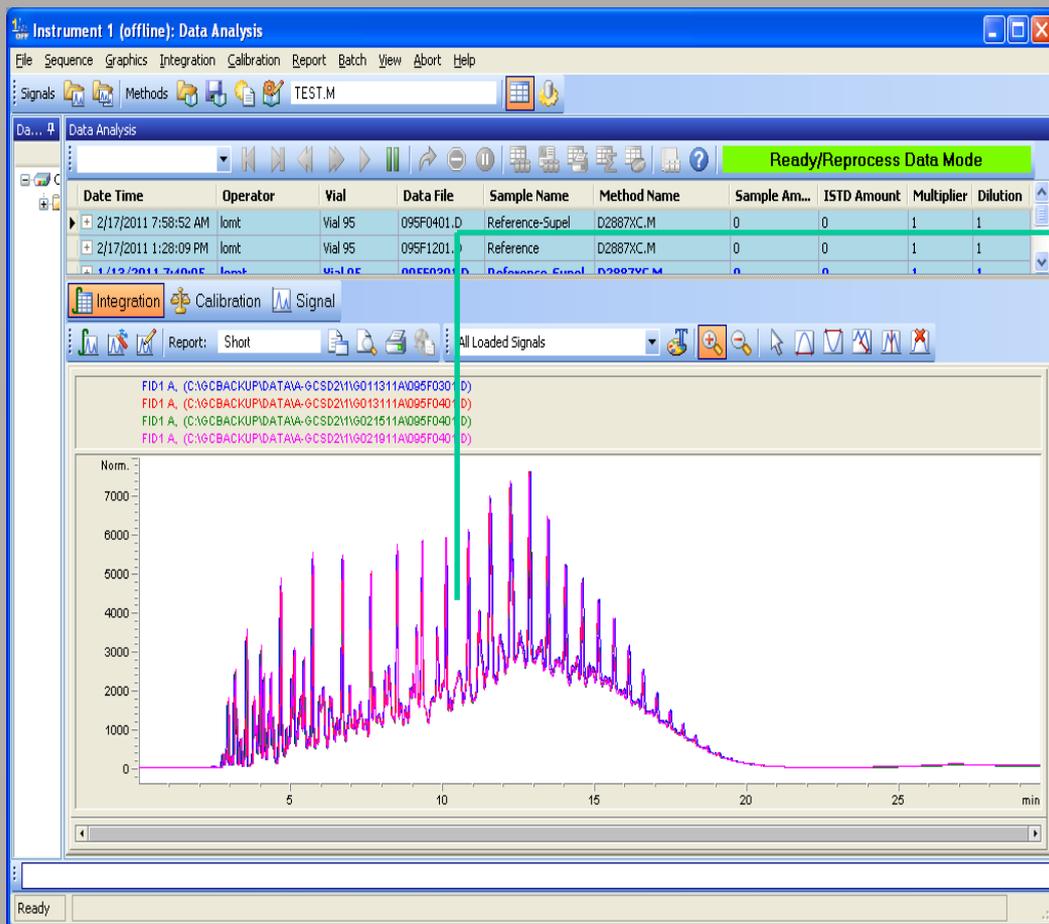
Brian Rohrback, Infometrix, Inc., Bothell, WA

Gulf Coast Conference, October 2016

Processing Whole Chromatograms

- Chromatograms will show an x-axis (retention time) shift for a variety of reasons:
 - Changing columns
 - Aging columns
 - Different instruments
- We need to eliminate retention time variability to improve the precision of our assessments.
- Think about how alignment relates to simulated distillation...
 - We run an n-paraffin standard to correlate temperature to retention time.
 - We use this new axis to map the cumulative percent of total area as we progress along this set of temperatures.

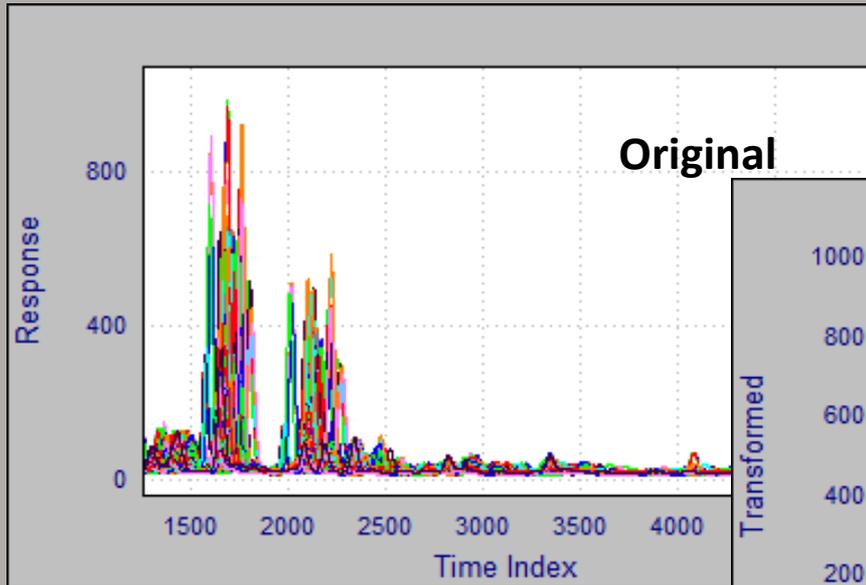
Retention Time Misalignment is a General Chromatography Problem



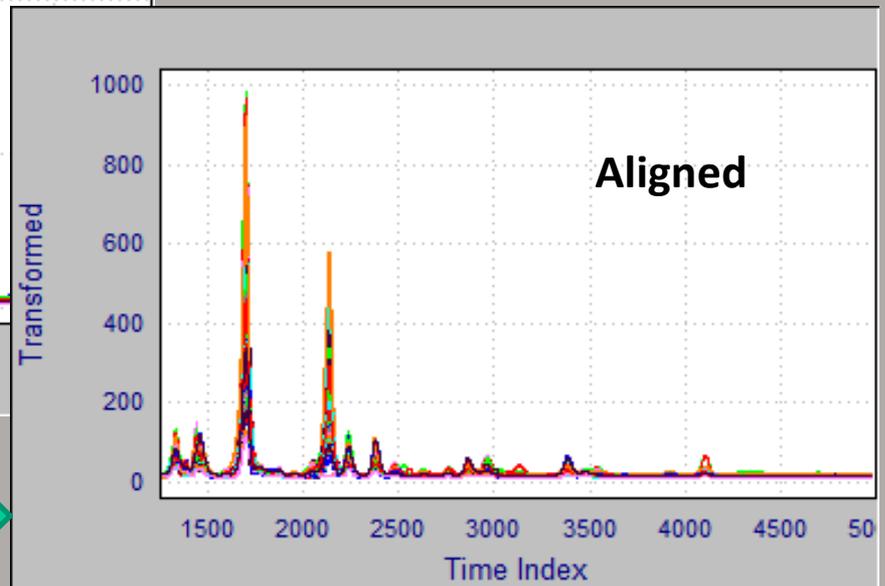
You can spot small changes in retention time even in proximal runs.

Alignment via Software

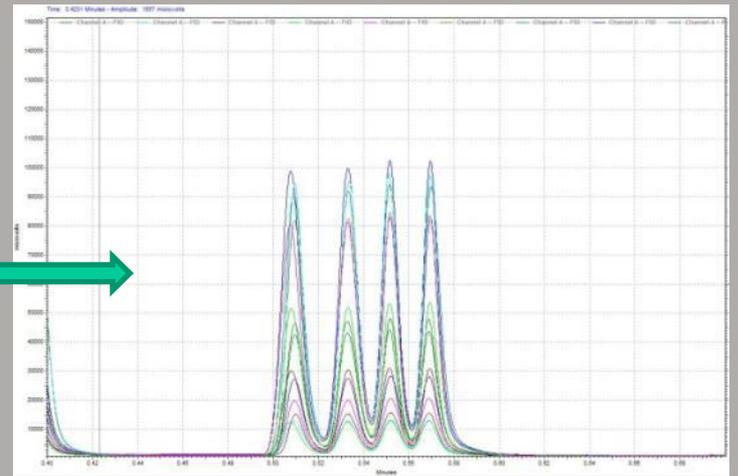
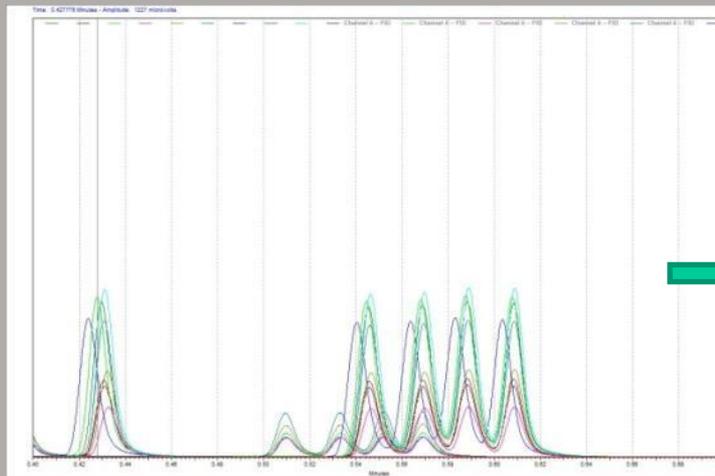
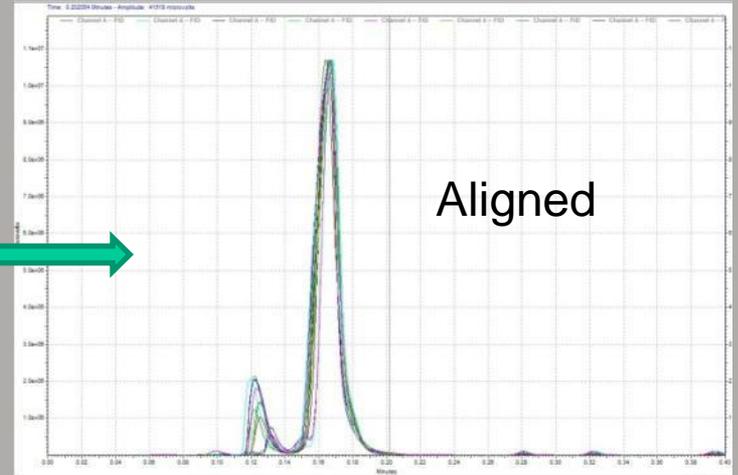
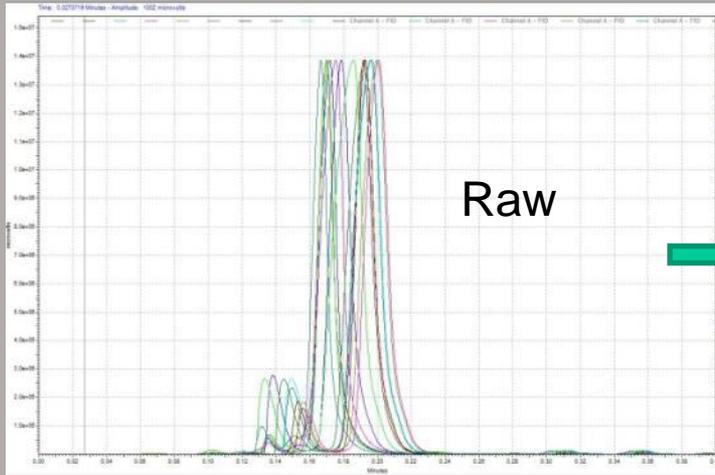
Over time, original chromatograms often show large variation in retention pattern; aligned chromatograms do not



6-year period



Alignment Across Instruments

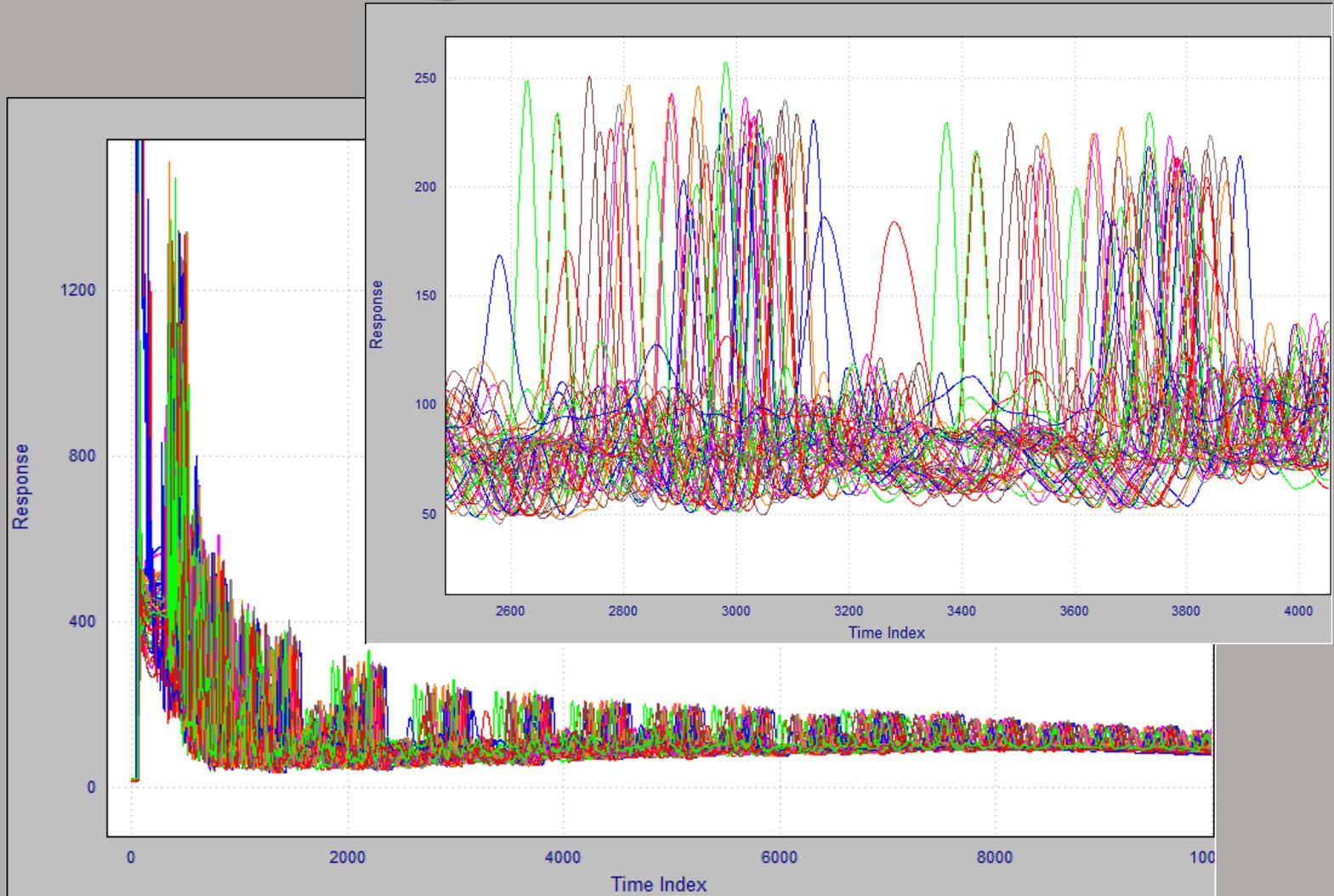


Two instruments with nominally the same method and columns, manual injection

Alaska North Slope Crude Oil: The quintessential column drift problem

- Same container of oil analyzed over 2 ½ years
- 1% crude in CS₂
- The chromatography is challenging
 - Column changes every 3-6 weeks
 - Inlet liner every week
 - Work burden: need to recalibrate every 8-12 hours
 - Some band focusing due to inlet at +30°C and column at -20°C

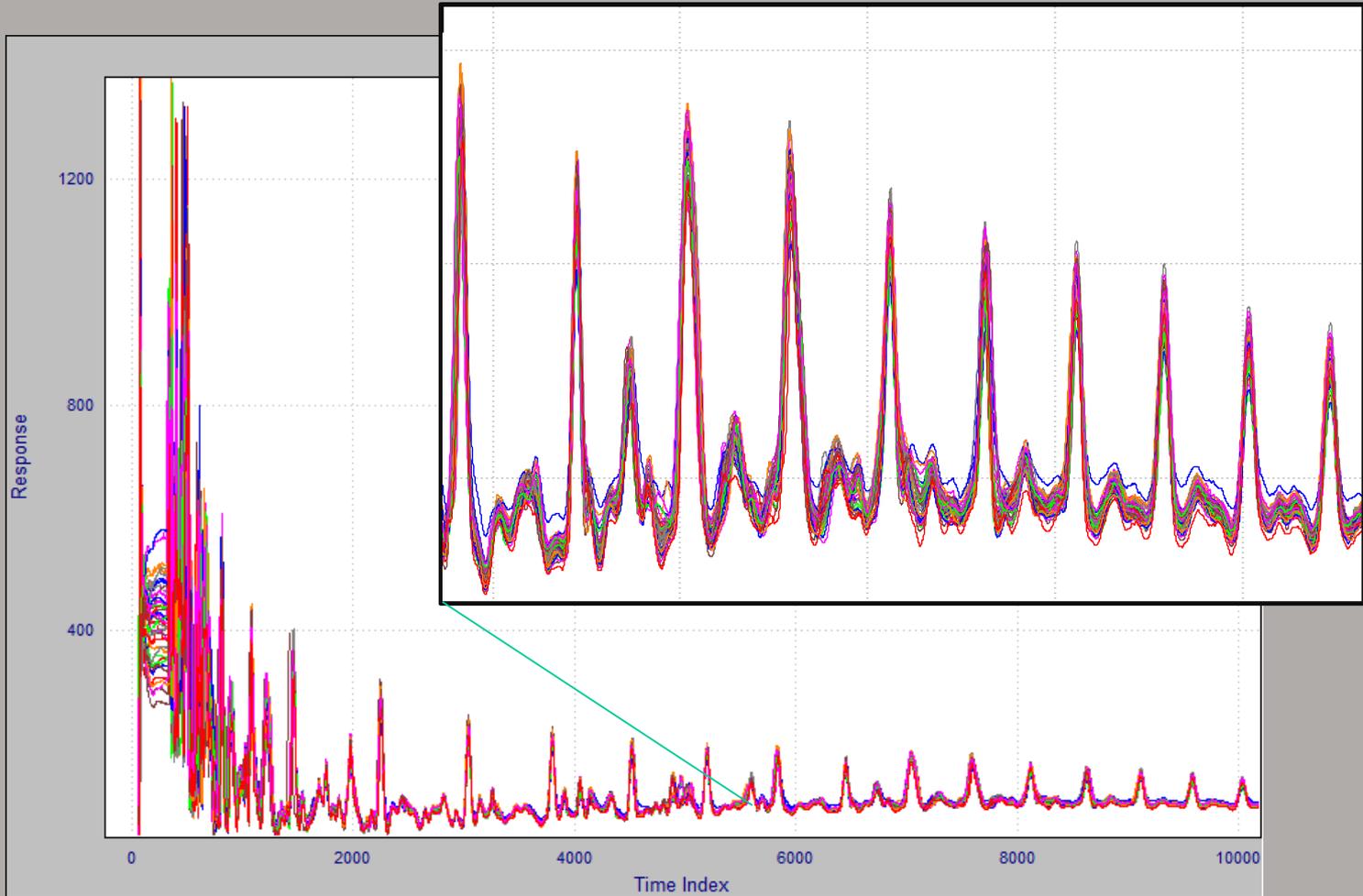
Chromatograms – RT Drift



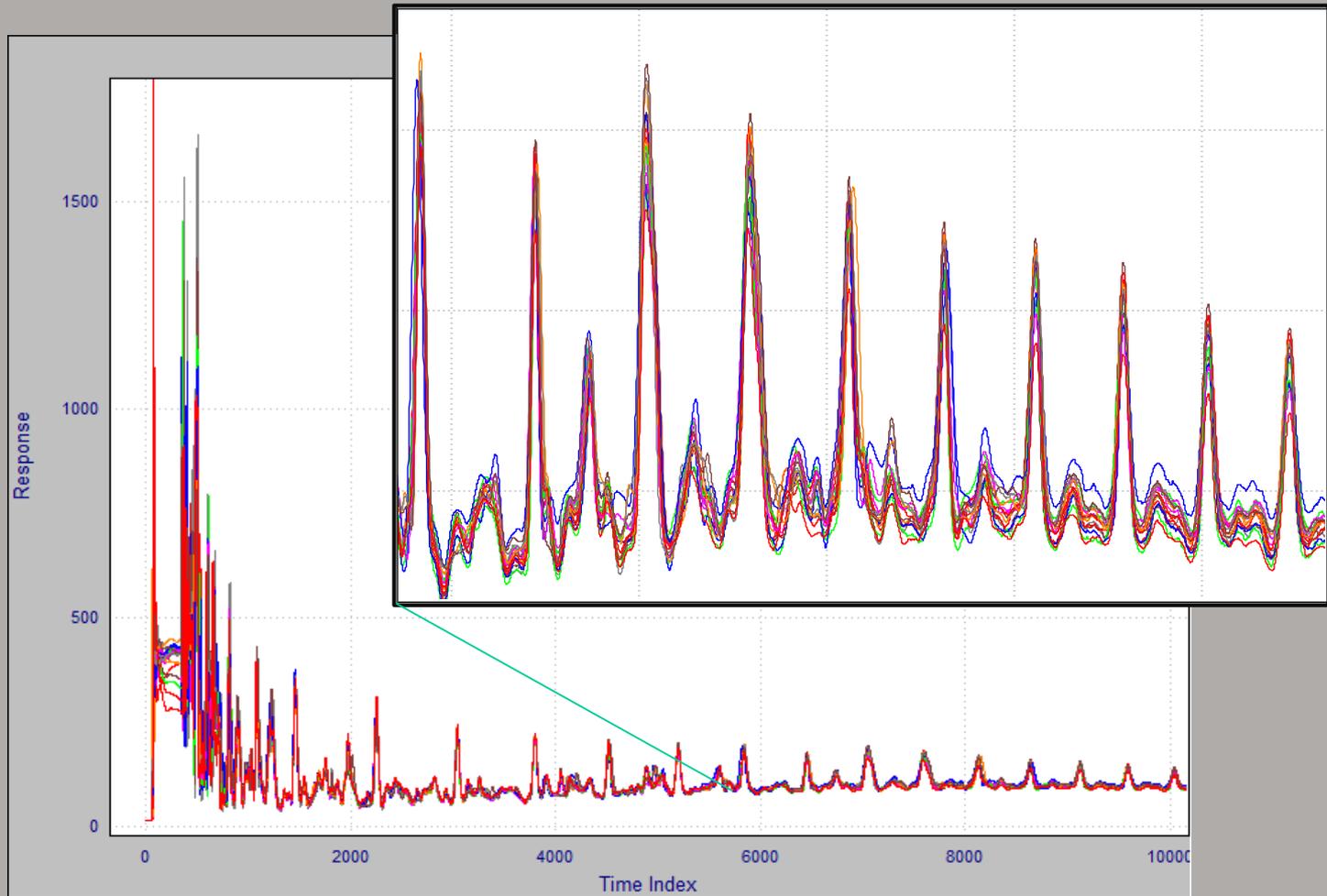
Alignment

- Alignment with COW algorithm is challenged because in these data early-eluting compounds are shifted considerably more than late-eluting compounds, for some samples; drift can be as much as the separation between n-paraffins
- After evaluating several combinations of alignment parameters, these were chosen:
 - Segment = 30; Warp = 6; Start = 201; Stop = 15000
 - Processing requires about 85 sec
- 70 of the 83 samples processed between 2014 and 6/2016 aligned well; the remaining 13 samples were much more challenging

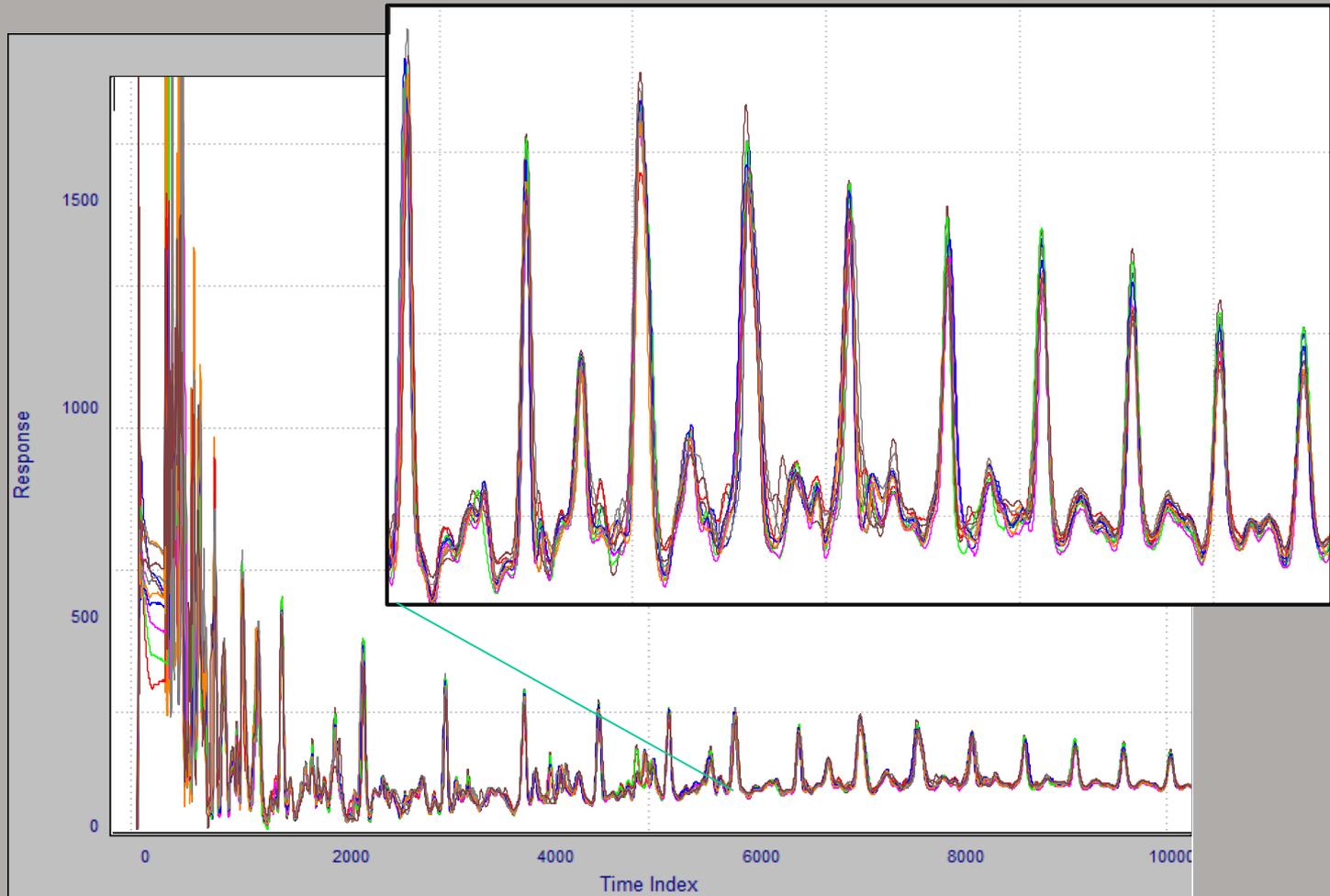
Well Aligned Profiles – 2014



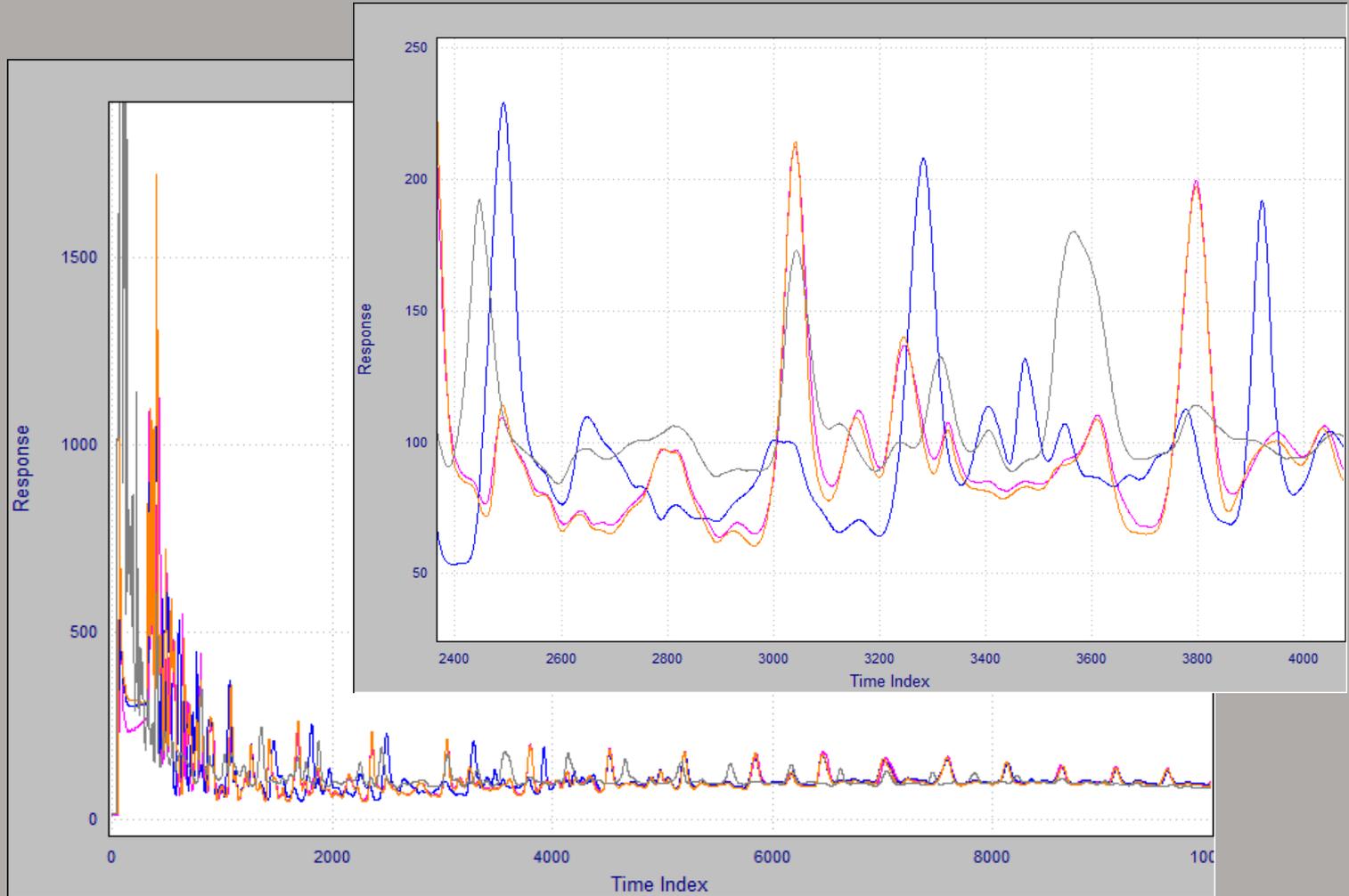
Well Aligned Profiles – 2015



Well Aligned Profiles – 2016



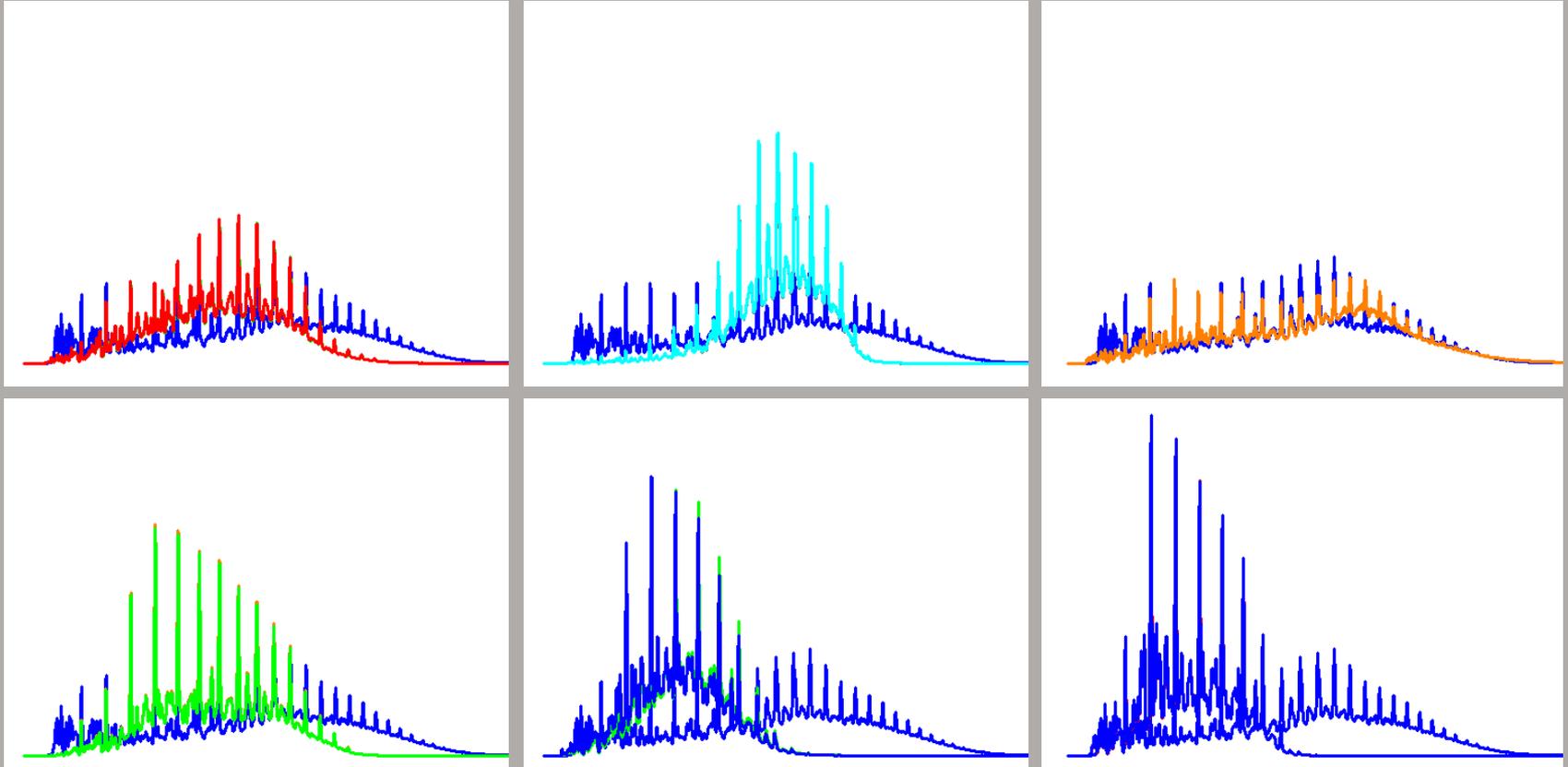
Poorly Aligned Profiles



Takeaway

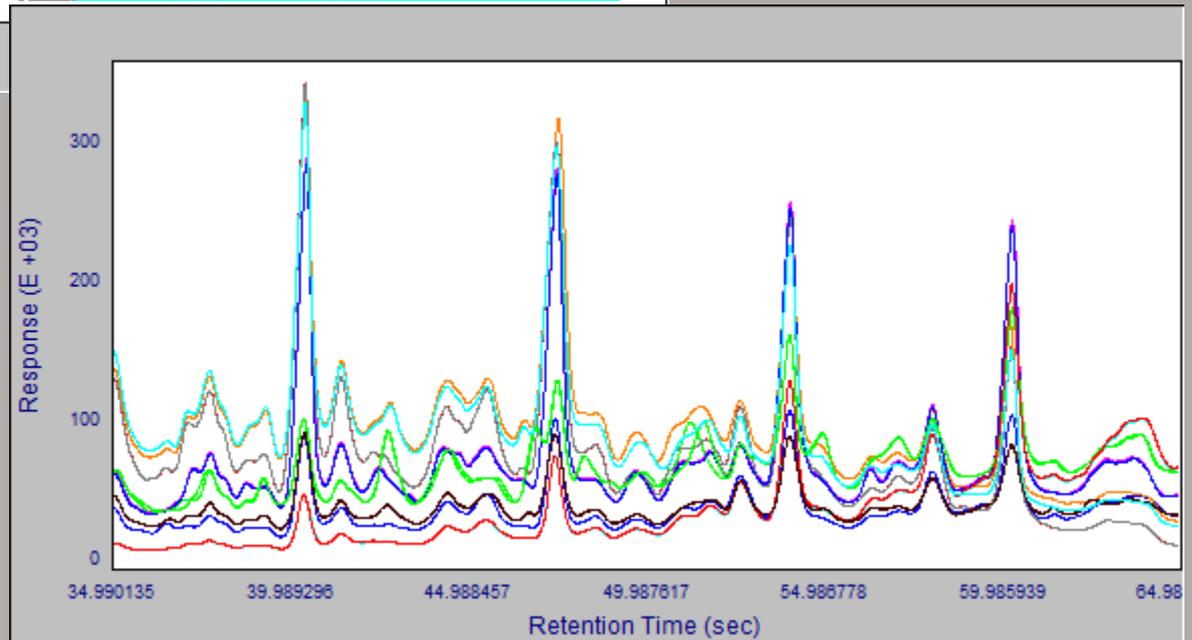
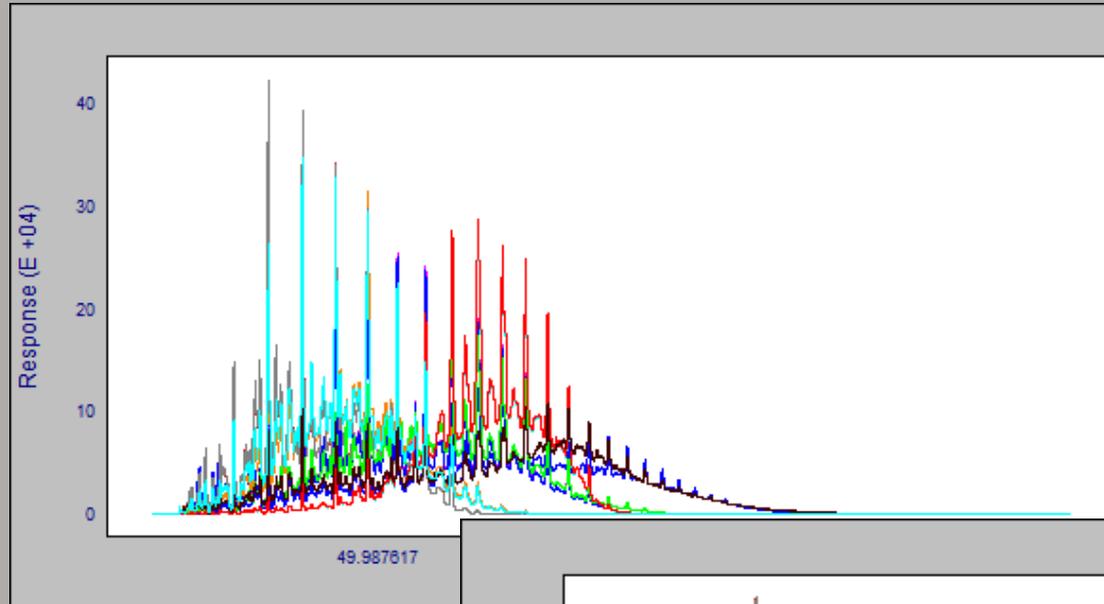
- Alignment (coupled with pattern recognition) provides a hard decision point on when to change the column or recalibrate
- 3-3½% change in Boiling Points was the experience prior to alignment
- 1-1½% variation is the experience for the aligned chromatograms

SimDist Test Samples vs. RGO

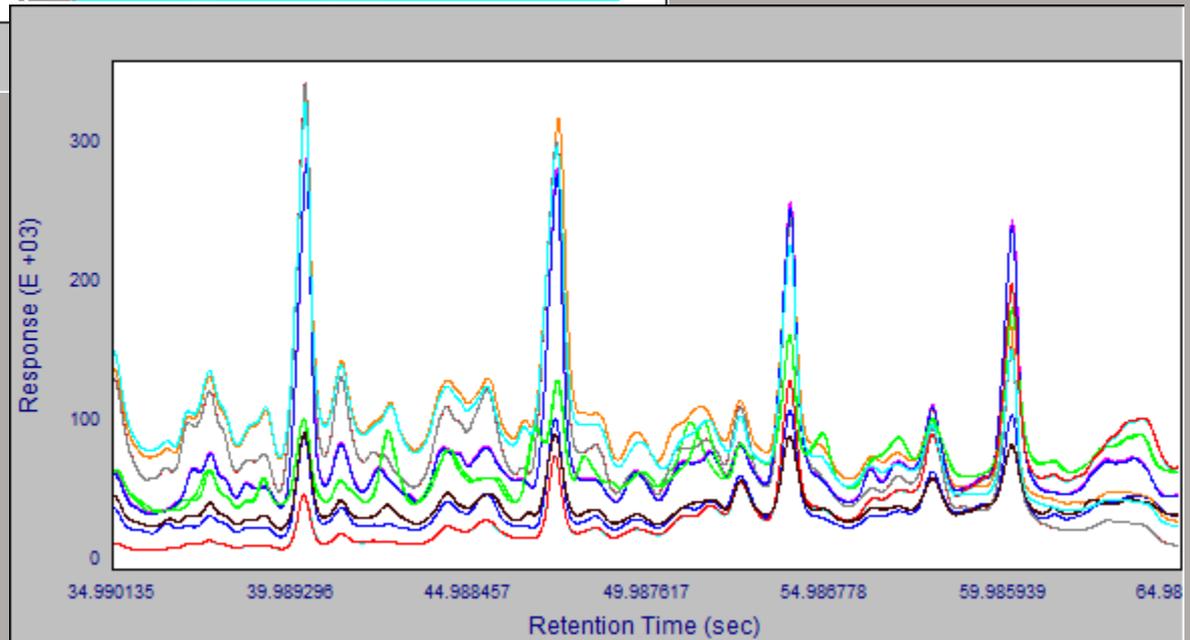
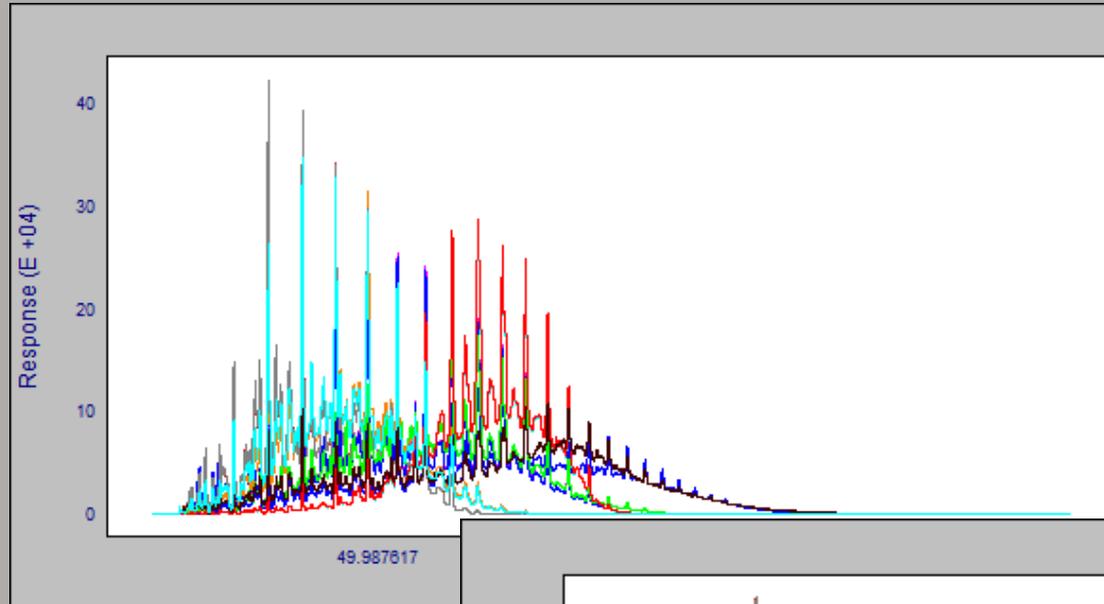


ASTM 7798 ILS

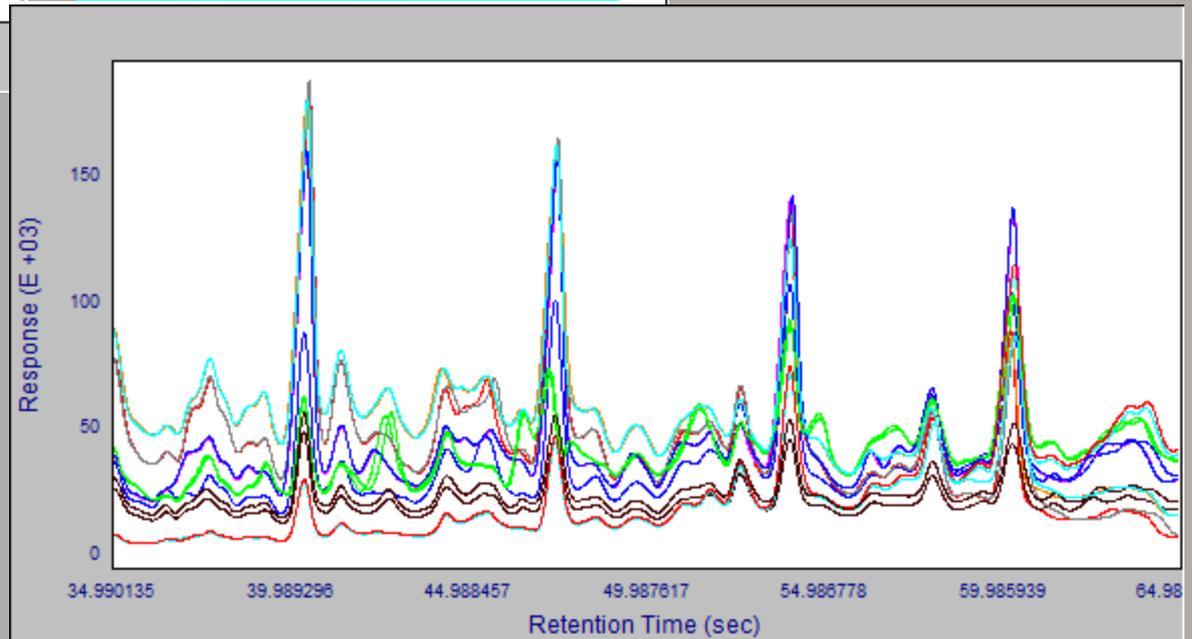
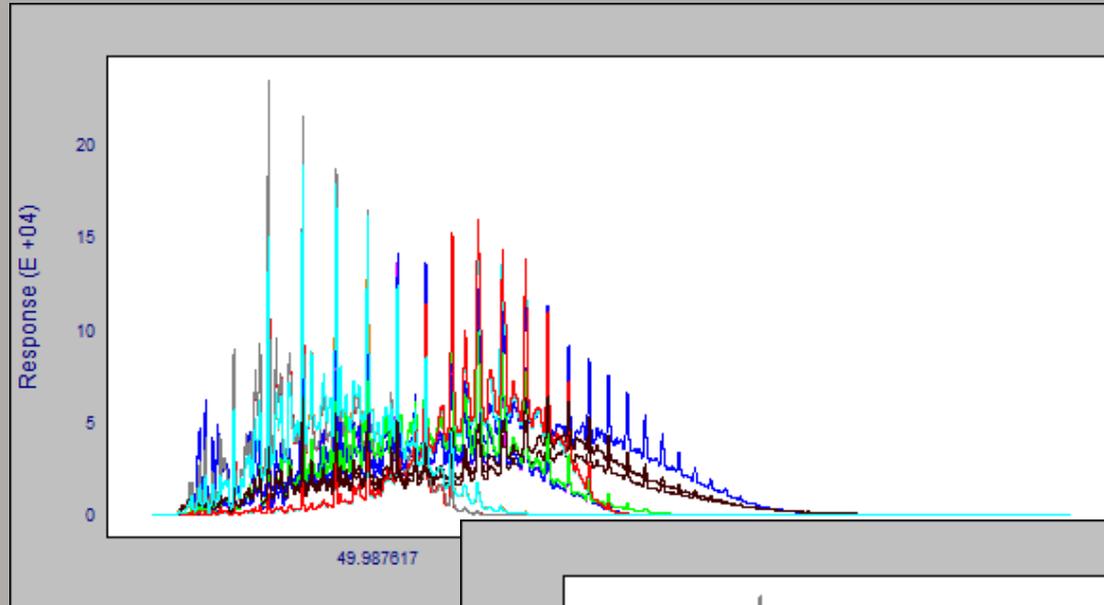
Location I – Aligned to the Location I Standard



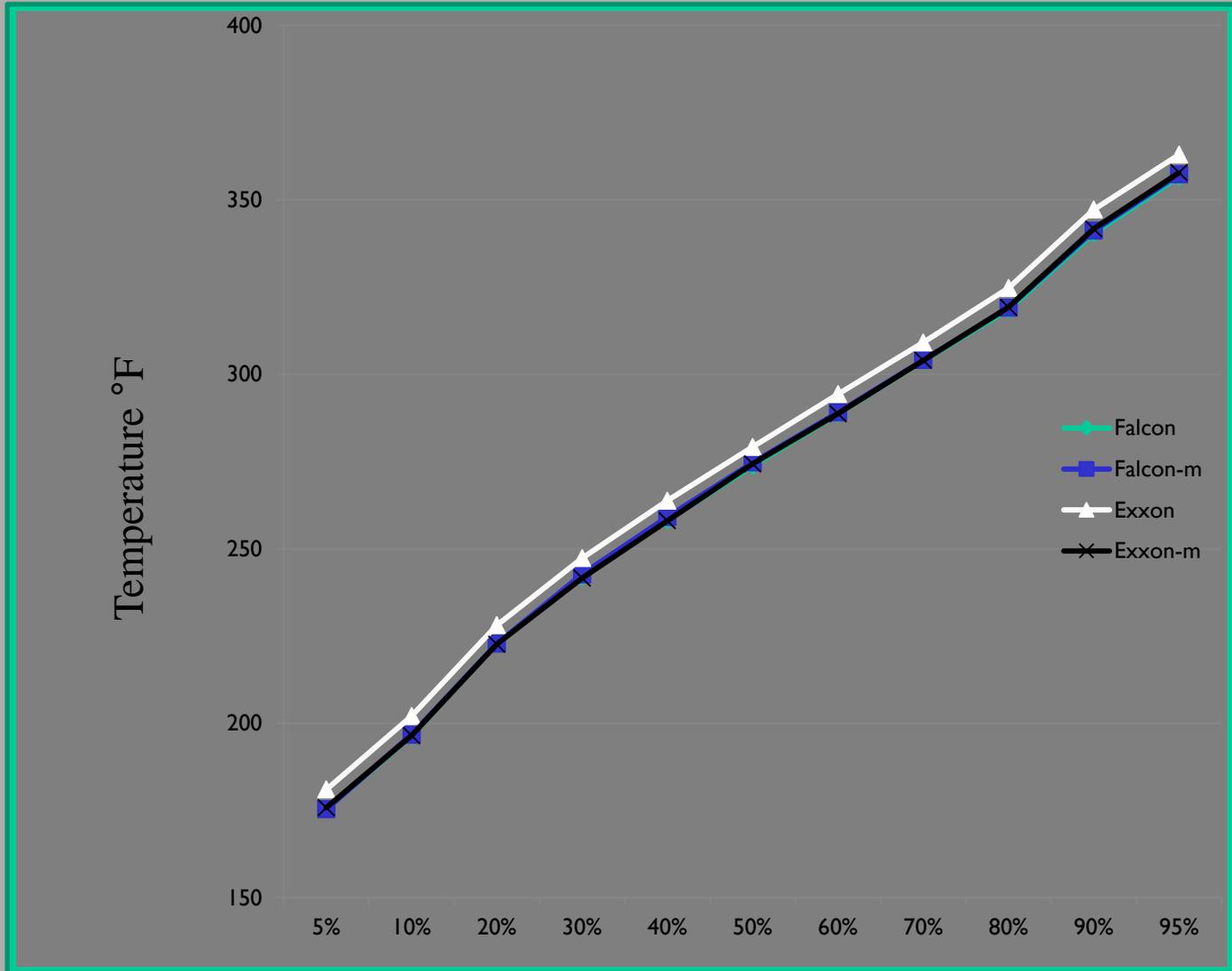
Location 2 – Aligned to the Location 2 Standard



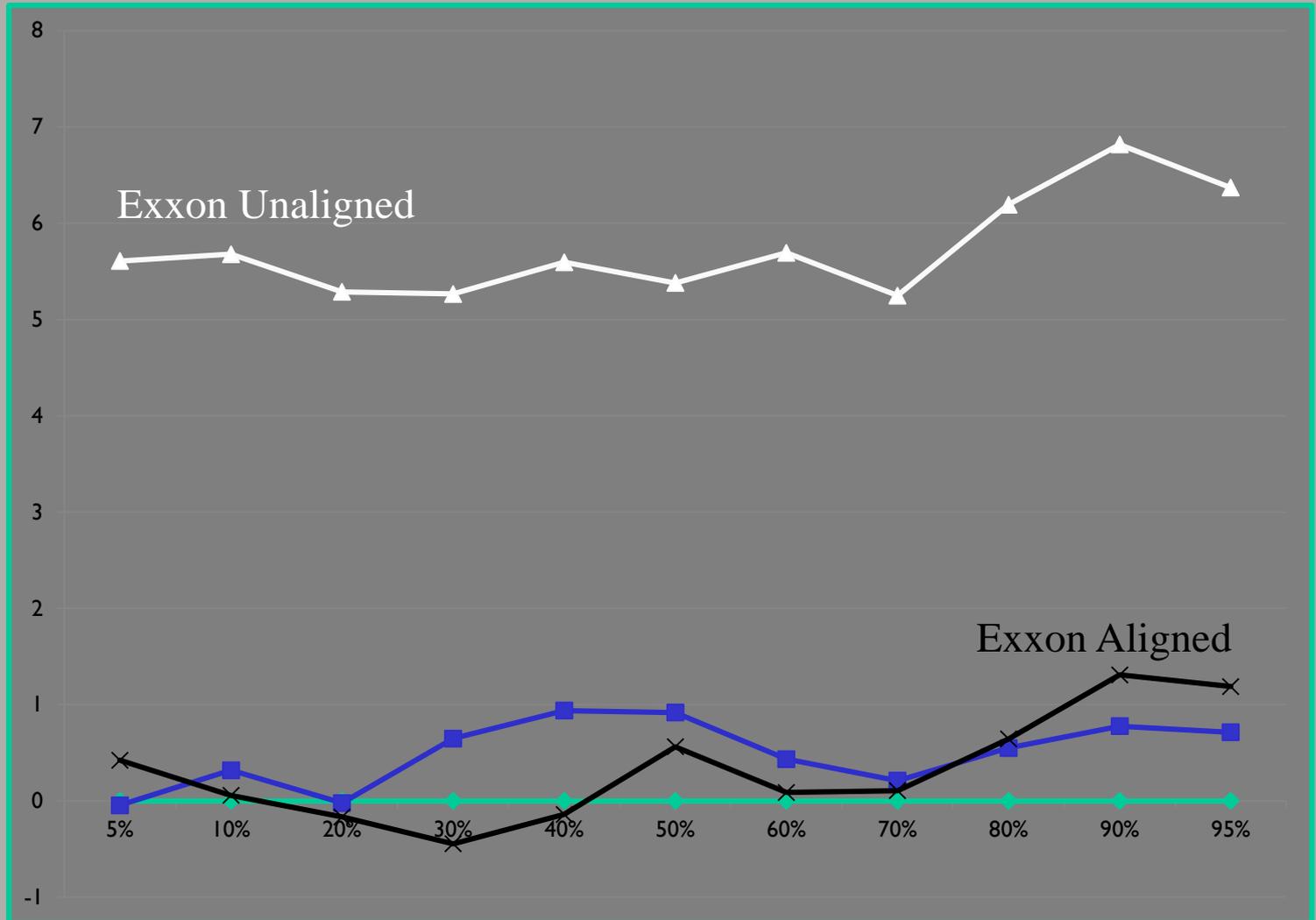
Location 2 – Aligned to the Location 1 Standard



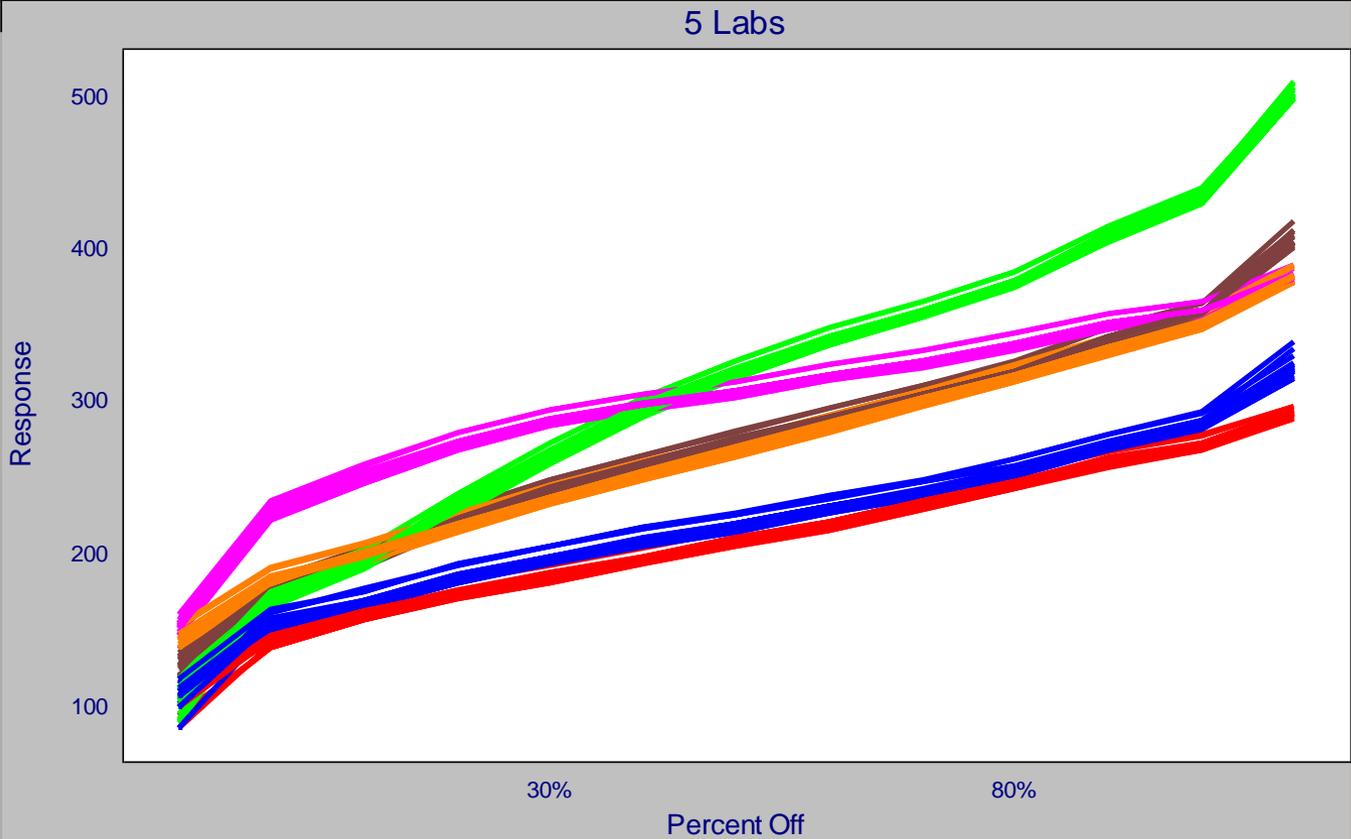
Distillation Profile



Adjusting for Out of Date Calibration

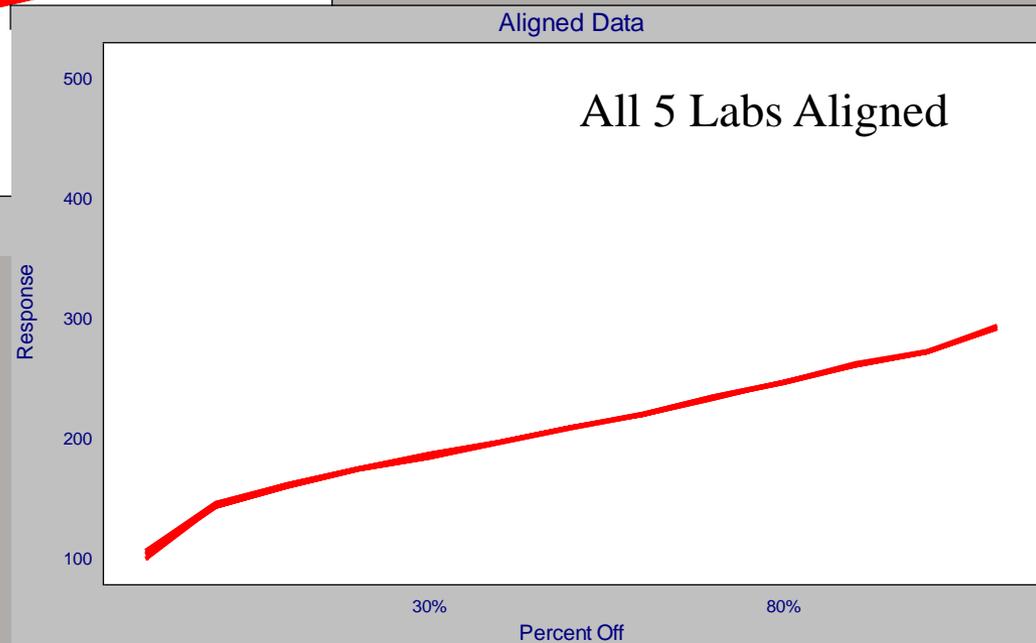
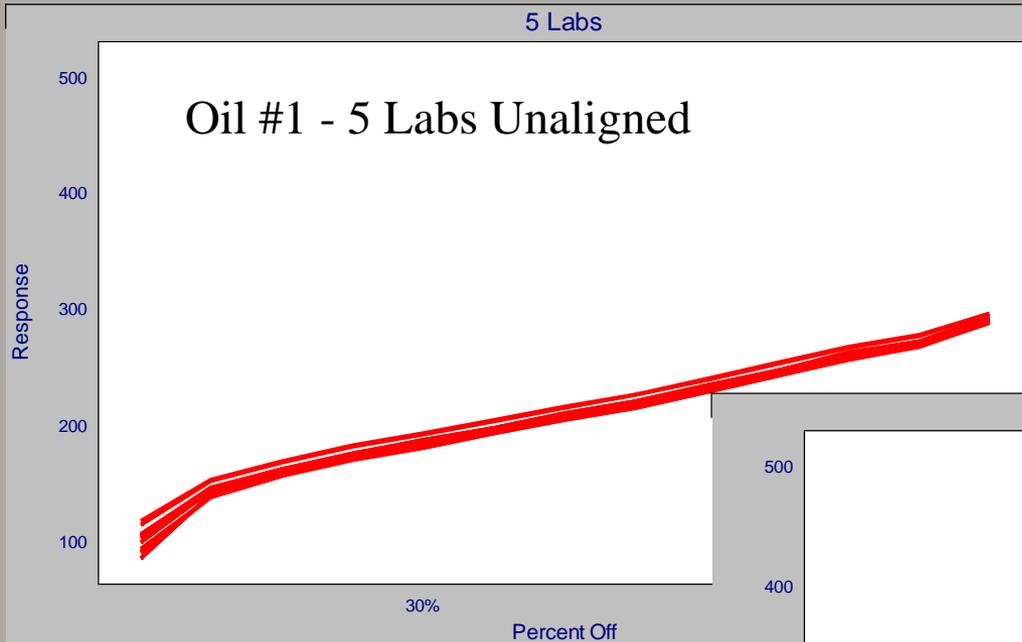


Six Different Oils, Five Labs

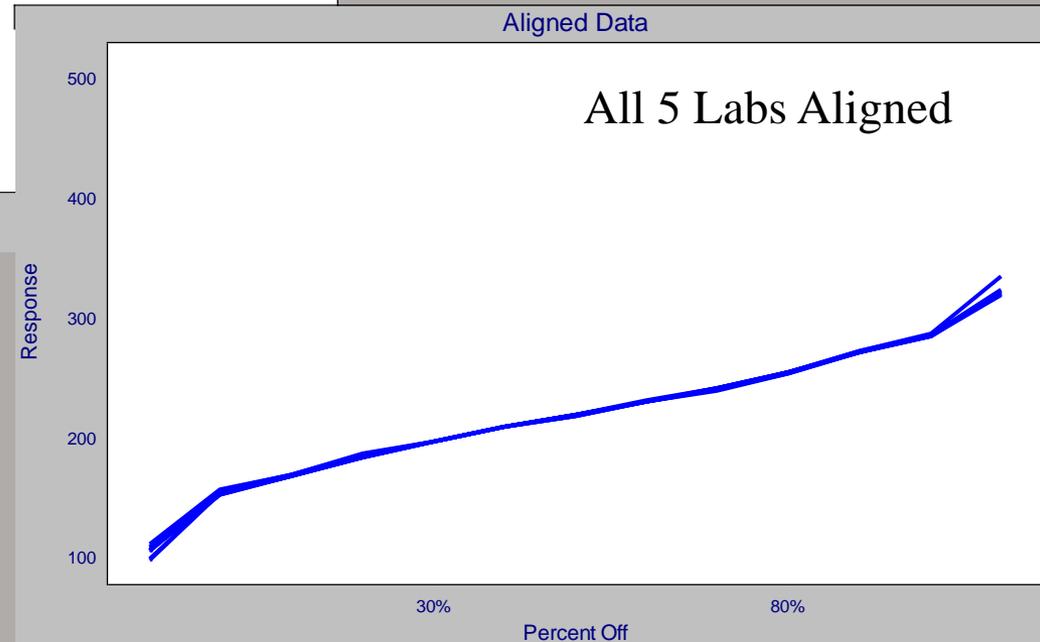
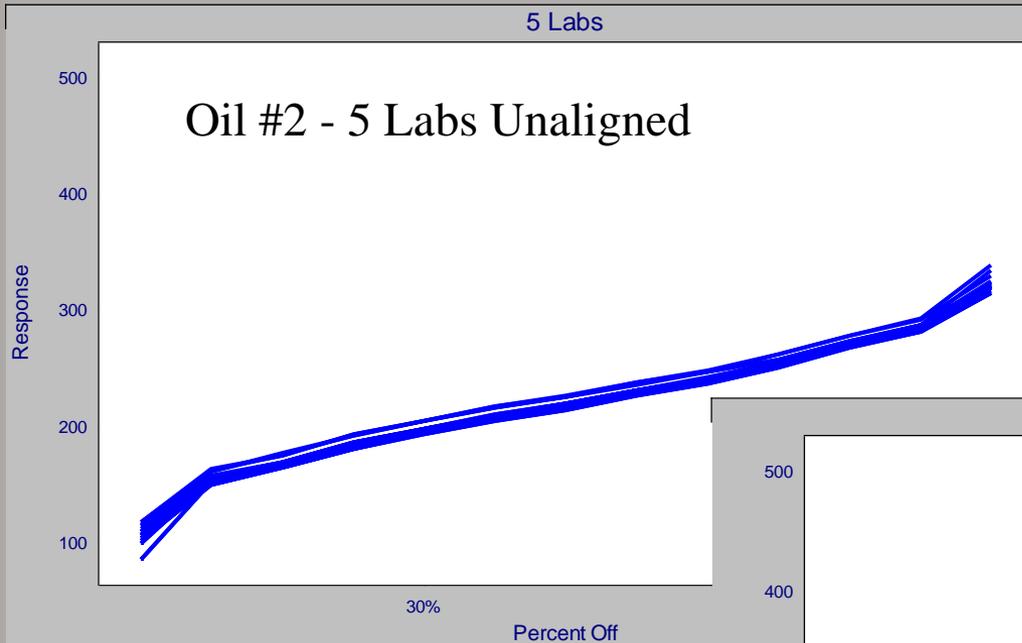


No Alignment

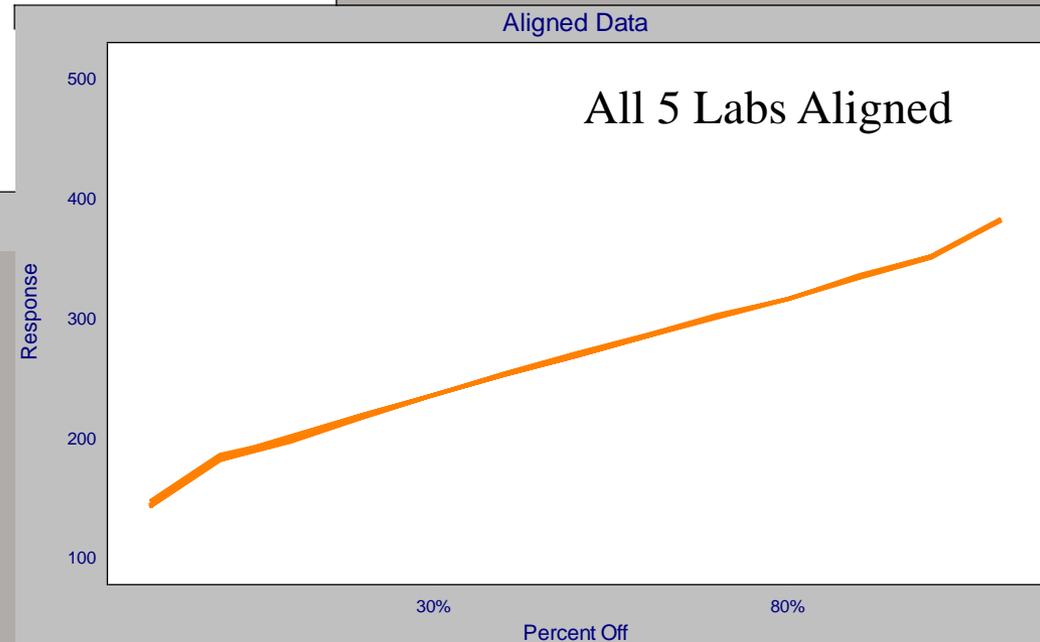
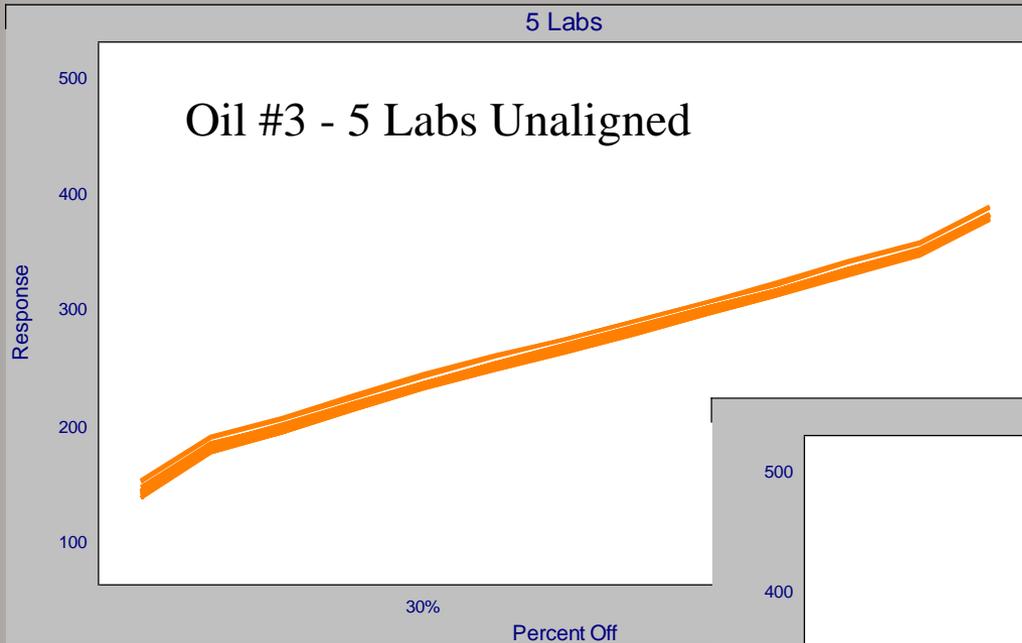
Unaligned and All 5 Labs Aligned to Location #1



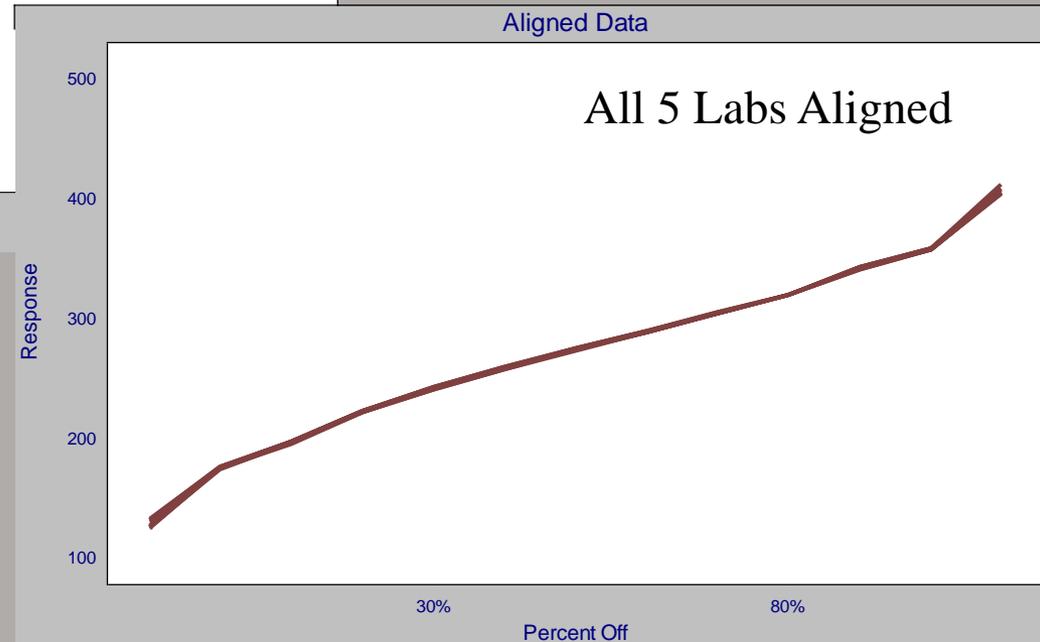
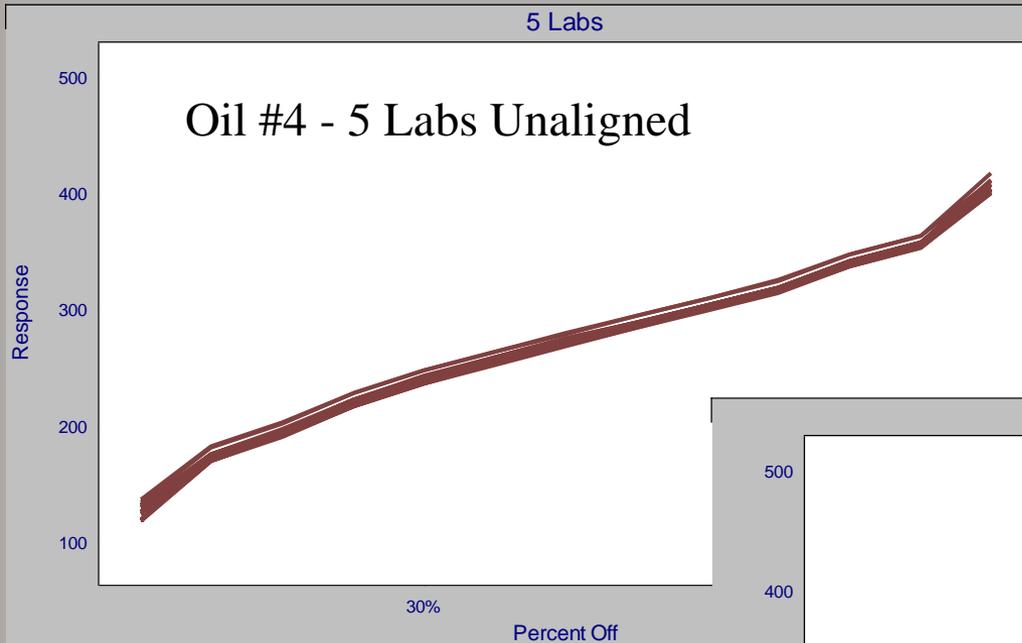
Unaligned and All 5 Labs Aligned to Location #1



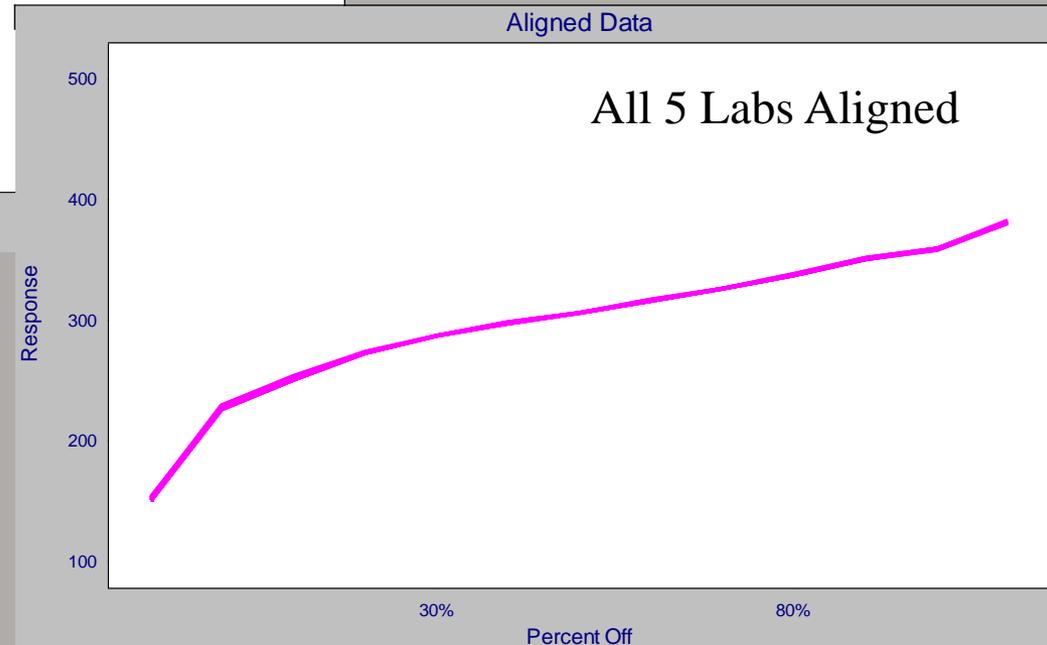
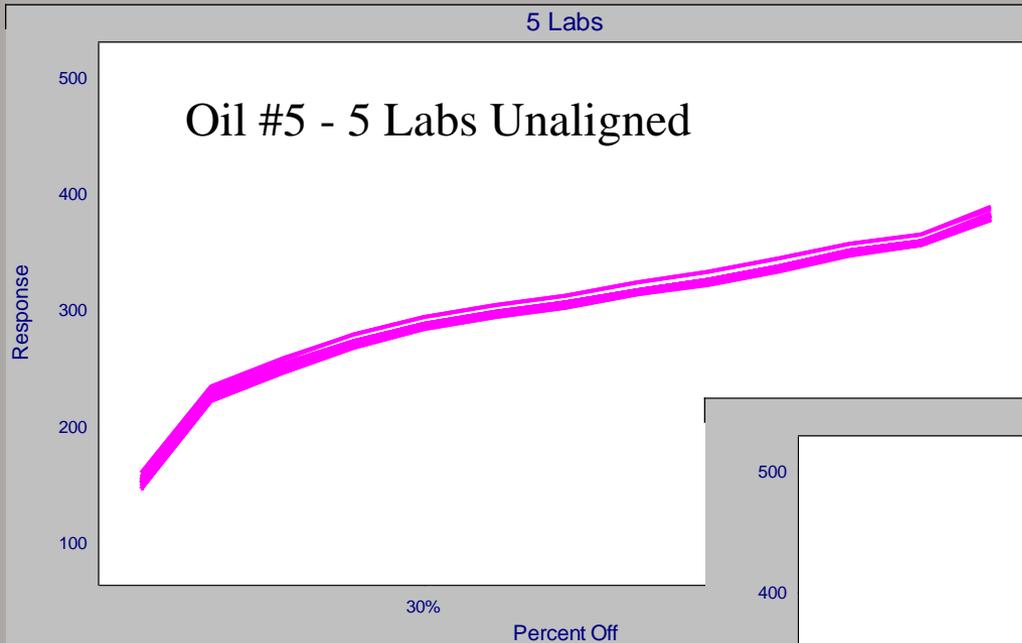
Unaligned and All 5 Labs Aligned to Location #1



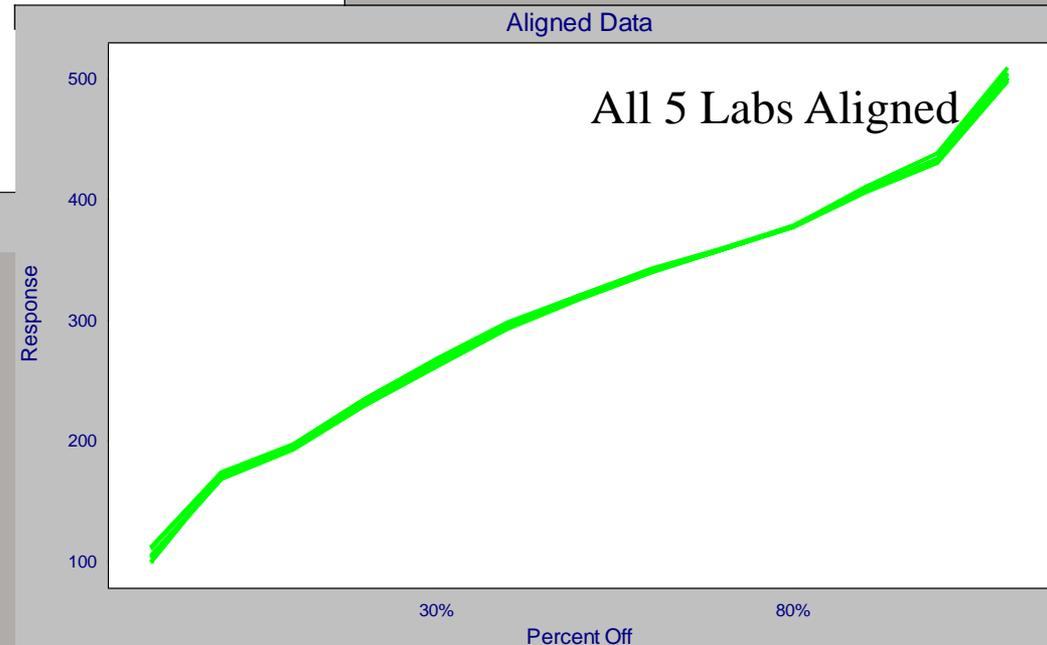
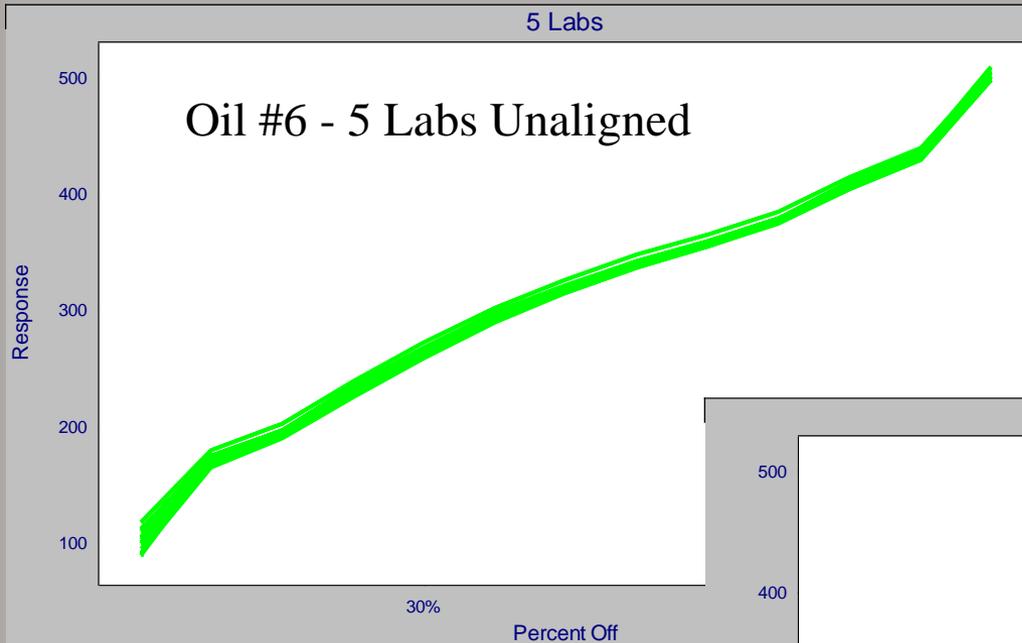
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Unaligned and All 5 Labs Aligned to Location #1



Continuous data interpretation *PLUS* validation of a multivariate instrument

We can correct retention times to match an application-specific relevant sample

You can use this to make all instruments performing a similar task to look identical (Plug and Play)

This raises the possibility of having a universal calibration

At the least, the frequency with which we really need to run calibration standards is significantly lower than what is currently being done.

Acknowledgements

- Robert Lorenz, Chevron
- John Crandall, Falcon
- ASTM 7798 ILS Laboratories