



Reference data for peak integration in capillary electrophoresis: another successful CASSS project

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Integration of Electropherograms in GMP Labs Under Increasing Scrutiny Due to Data Integrity Intensive Inspections

Tim Blanc¹, Hermann Wätzig², Cari Sänger-van de Griend^{2, 3, 4}

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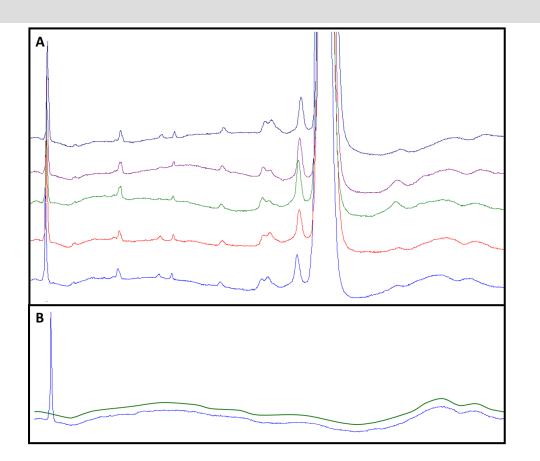
¹Eli Lilly and Company

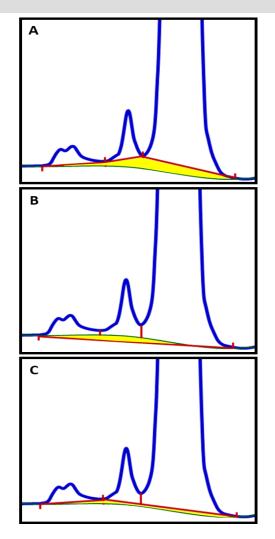
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Peak integration: critical examples









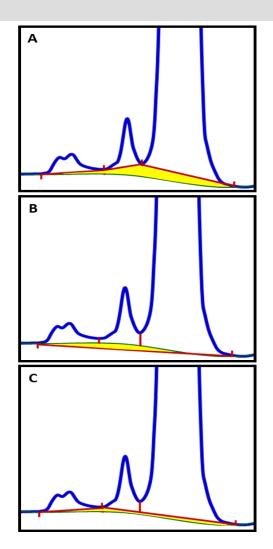
Peak integration: critical examples



Tim Blanc



Cari Sänger





Peak integration

- Integration in GMP laboratories is critically important
 - has become focus of data integrity-centric regulatory inspections
- Data systems developed for chromatograms rather than electropherograms, and the increased regulatory scrutiny call for a resolution.
- This extends to R&D, clinical, and academic labs

Tim Blanc, Hermann Wätzig and Cari Sänger – van de Griend,

Peak Integration of Electropherograms in GMP and Research Labs:

Navigating Increased Scrutiny Amid Data Integrity Audits and Inspections

Electrophoresis, 2025; 46:653–668

http://dx.doi.org/10.1002/elps.70002



Peak integration: Conclusions from this review

- We need APs and SOPs
- We need detailed method descriptions, defaults, parameter lists, illustrations, training
- We need to keep manual integration at the present time
- Collaboration within the industry for generally accepted standards
- Benchmark data sets could be an important step
- Based on this, considerable improvements in integration algorithms are possible
- So probably less manual integration in the future, but at present it is still a must





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- We need APs and SOPs
- We need detailed method descriptions, defaults, parameter lists, illustrations, training
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- Benchmark/Reference data sets could be an important step
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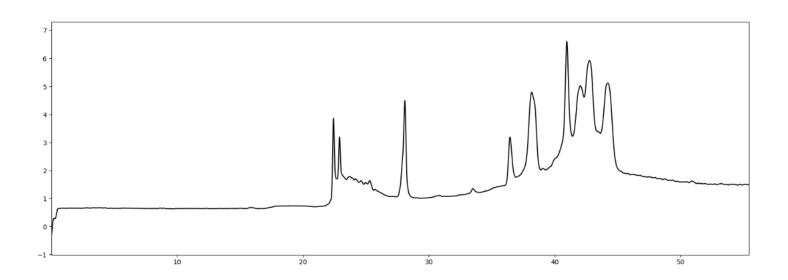
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Interesting Reference Data for Peak Integration



- Broad, asymmetrical peaks
- Fronting/Tailing
- Baseline irregularities

→ Existing softwares can not properly face these challenges





[3]

Intercompany Collaboration on Reference Data



Marlon Krompholz

- Multinational collaboration of CE experts
- Data collection
- Experts set peak limits
- Comparison and Discussion
- Refined integration rules
- Pre-integration by Marlon
- Approval/Confirmation by experts
- Approval by the community
- Reference Data Set CE Integration #1

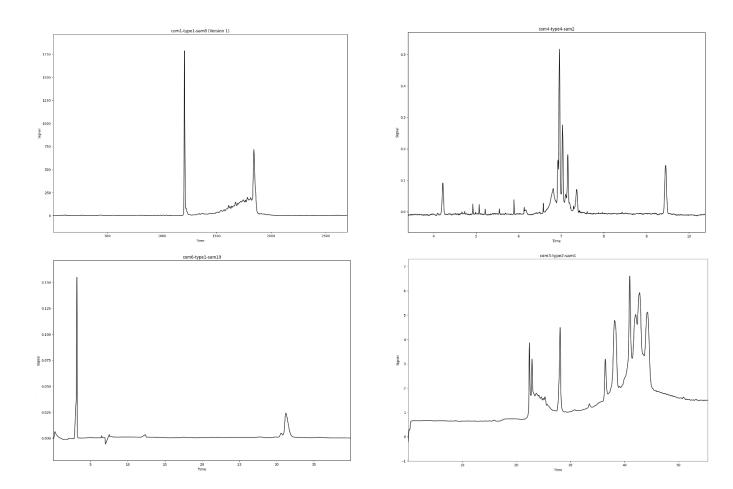


Intercompany Collaboration: a CASSS initiative

Mostafa Abuzeid, **Timothy Blanc, Tao Bo**, M Chan, **Patricia** Christensen, Adriana Coll de Peña, Miyuru de Silva, Hans Dewald, Gangadhar Dhulipala, Tara Enda, Joachim Ermer, Jarvas Gabor, James Geiger, Frederic Ginot, Mervin Gutierrez, Jakob Haglöf, Mark Haverick, Philip Hoang, Robert Hong, Christopher Hood, Philipp Huber, **Andrei** Hutanu, Ryan Hylands, Yun Hwang, Pilsoo Kang, Steffen Kiessig, Marlon Krompholz, Andreas Krupke, David Lasar, Rick Linkous, Huixin Lu, Jane Luo, Will McElroy, Chris Heger, Paul S Michael-Butler, Jana Molitor, Cristina Montealegre Dondarza, **Bernd Moritz**, Matthew Myers, Ryan Nai, **Trang D Nguyen**, John Orlet, **Rebecca Osborne**, Jeremy Osko, David O'Sullivan, **Ashley Prout**, **Timothy Riehlman**, David Ripley, Jacob Rostkowski, Shauna Salem, Cari Sänger-van de Griend, Emineguel Sakka, Carlie M Schaeffer, Johannes Schlecht, Kristin Schultz-Kuszak, Jana Steflova, Christian Streng, Adam Sutton, Lalin Theverapperuma, Amanda Torelli, Ewoud van Tricht, Carolina G. Vega, Yun Wang, Denise Warzak, Brian Wei, Katharina Yandrofski, Tao Zao, Yuling Zhang



Intercompany Collaboration: Data Collection







Intercompany Collaboration: Experts set peak limits

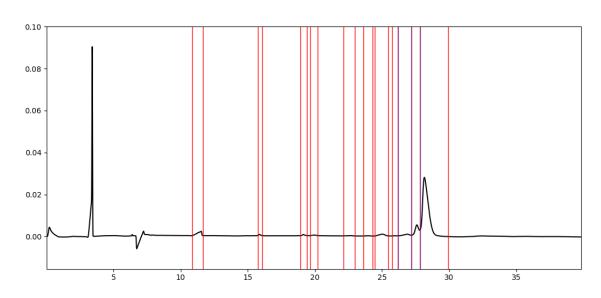


Software tool:

Developed with

Python using the

OpenAl o3 model



(10.85, 0.0003722) (11.65, 0.0004326) (15.74, 0.0004146) (16.09, 0.0003845) (18.93, 0.000397) (19.38, 0.0004314) (19.64, 0.0003967) (20.21, 0.0004049) (22.13, 0.000267) (22.238, 0.0002462)







Intercompany Collaboration: Experts set peak limits

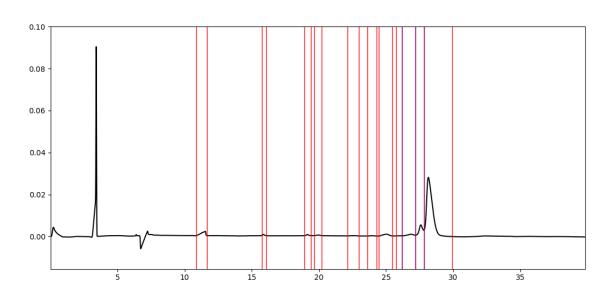


Software tool:

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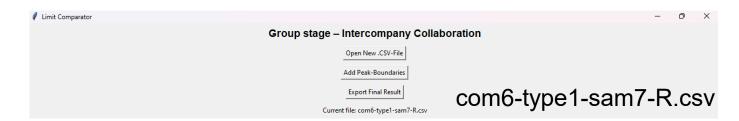
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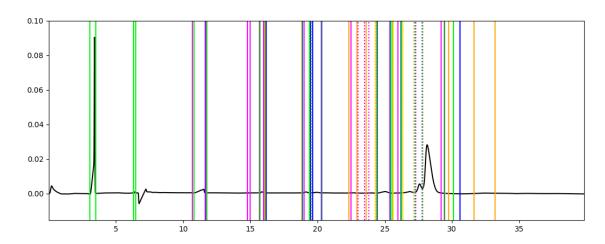




Intercompany Collaboration: Comparison and Discussion



Software tool: Developed with Python using the OpenAl o3







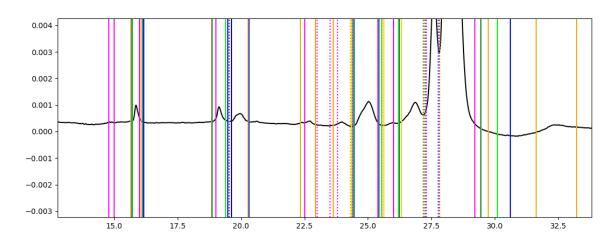
model



Intercompany Collaboration: Comparison and Discussion



Software tool: Developed with Python using the OpenAl o3 model

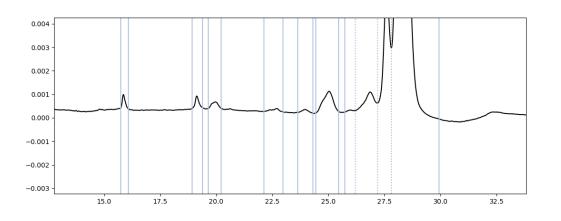




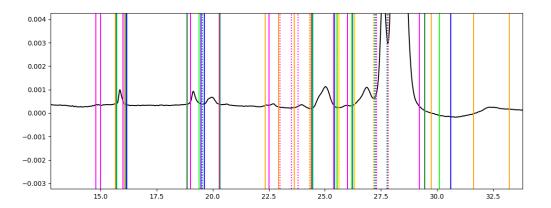




Intercompany Collaboration: Comparison and Discussion



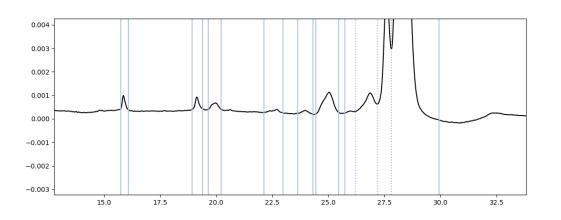
Re-set boundaries



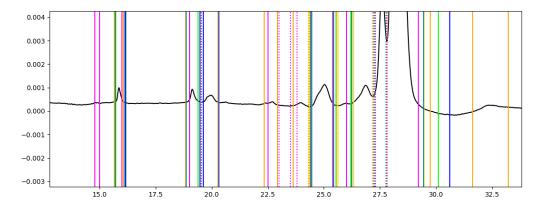
Experts' boundaries



Refining Integration Rules



Re-set boundaries







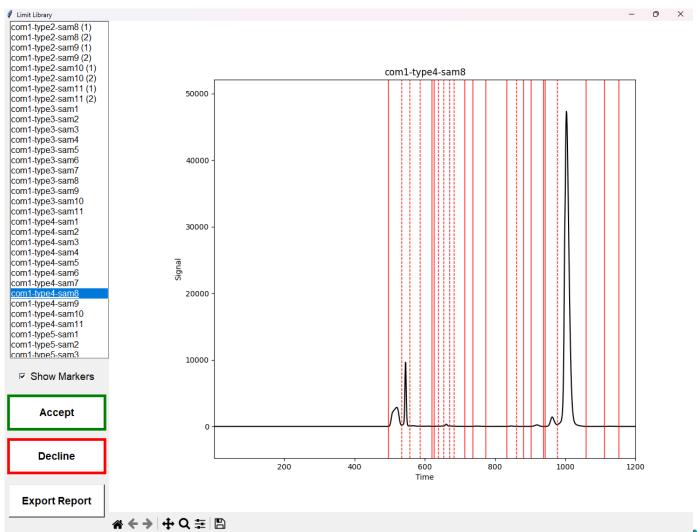
Pre-Integration by Marlon

Software tool:

Developed with

Python using the

OpenAl o3





model

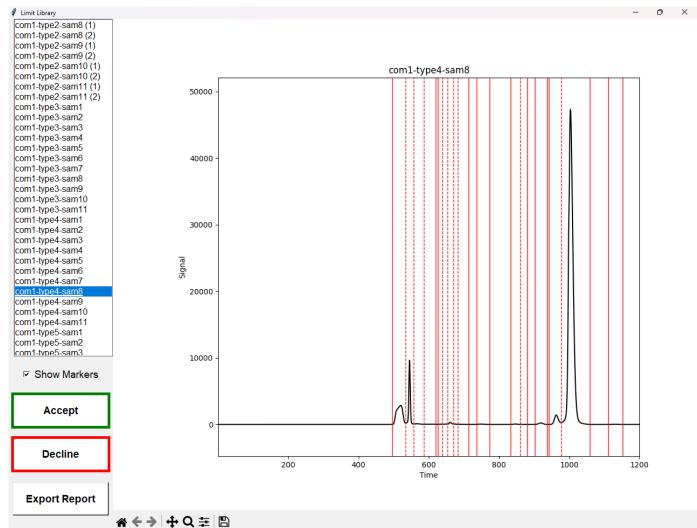


Intercompany Collaboration: Confirmation by the Experts

Software tool:

Developed with Python using the

OpenAl o3 model







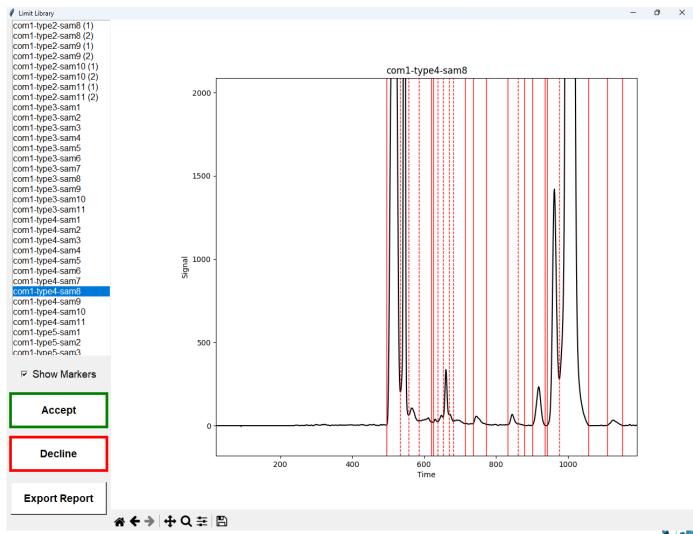
Intercompany Collaboration: Confirmation by the Experts

Software tool:

Developed with

Python using the

OpenAl o3

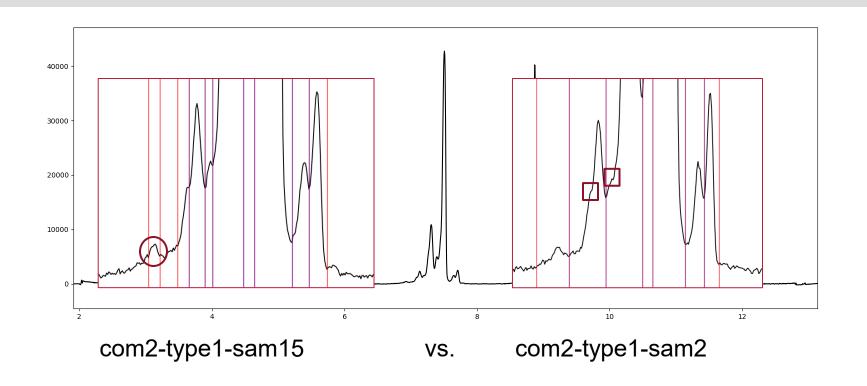




model



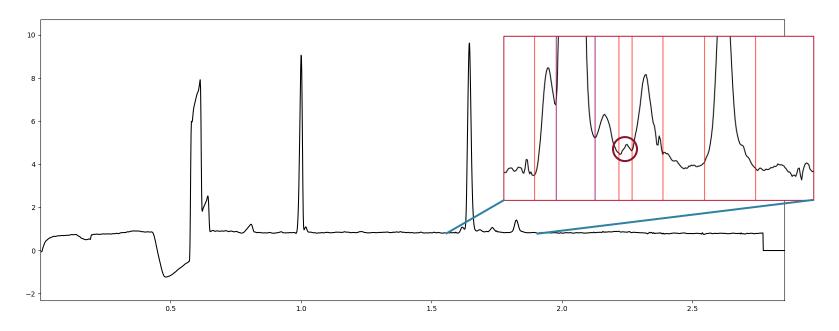
Debatable and context-dependant cases



- Is circle marked area a peak or noise?
- If peak: Which approach is better?
- Should square marked points be integration limits due to the comparison to the left image section?





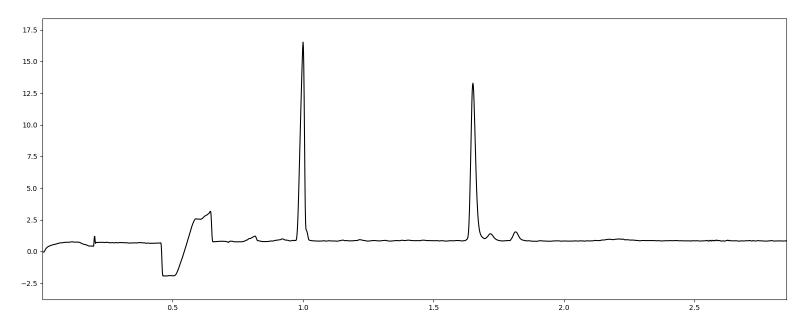


com2-type2-sam2

 Does the marked area belong to the baseline or does it lay above and therefore form a peak?





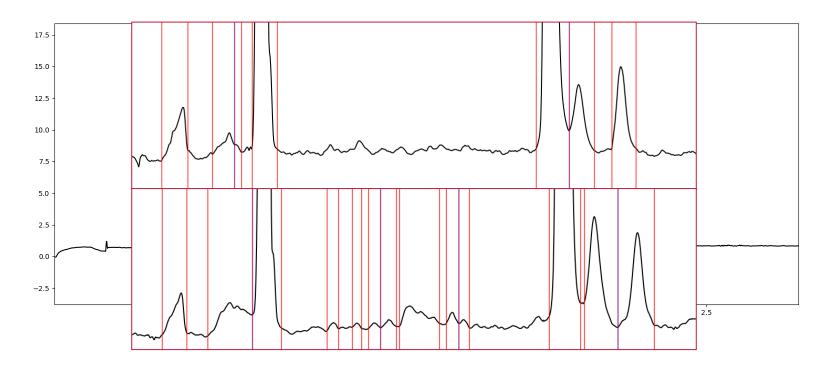


com2-type2-sam25 and com2-type2-sam5

 Are there any peaks to be considered between the main peaks or is it just noise?





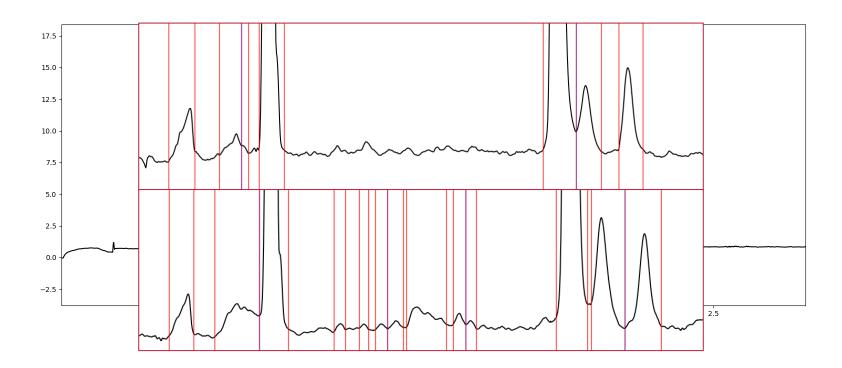


com2-type2-sam25 (top) vs. com2-type2-sam5 (bottom)

 Are there any peaks to be considered between the main peaks or is it just noise?





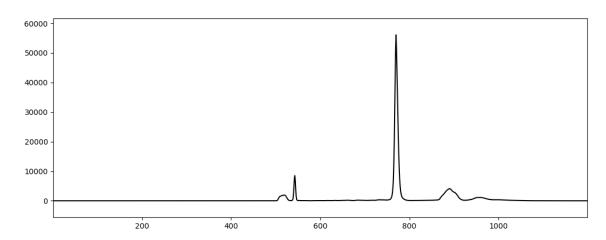


Suggestion, related to the European Pharmacopoeia 2.2.47, Capillary Electrophoresis: Find baseline as close as possible to the possible peaks of interest, 20 times as long as a major peak's $h_{0.5}$, estimate peak-to-peak noise. All signal S/N > 3 considered as peaks



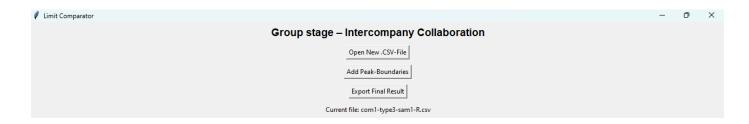




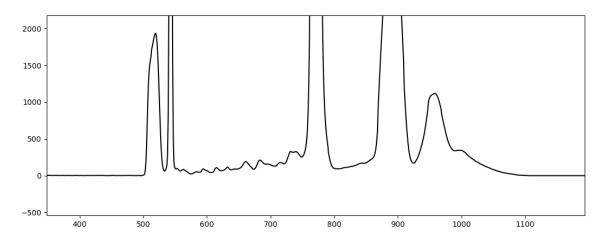








Zoom level is crucial

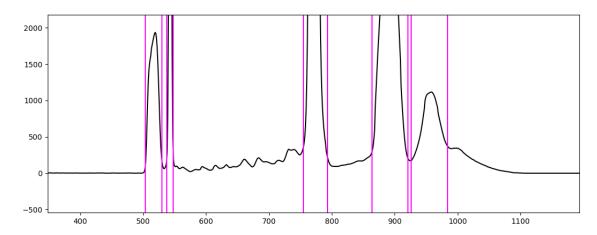


·←→ □ □		(x, y) = (770.9, 1597.)
	Show All	
com1-type3-sam1-P10.csv	Show Delete	<u> </u>
com1-type3-sam1-P14.csv	Show Delete	
com1-type3-sam1-P18.csv	Show Delete	
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com1-type3-sam1-P26.csv	Show Delete	
com1-type3-sam1-P4.csv	Show Delete	







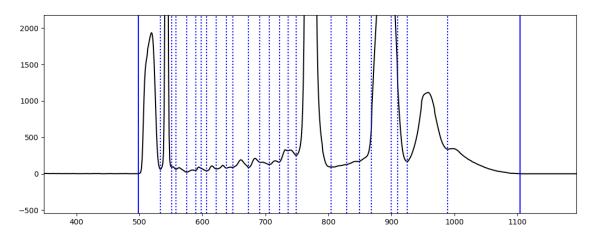












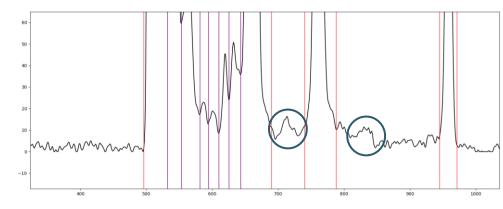




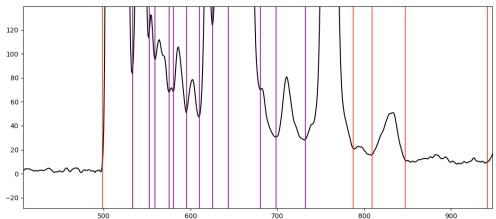


Integration depends on the context:

Peak or baseline?



→ Clear peaks after comparing to other injection

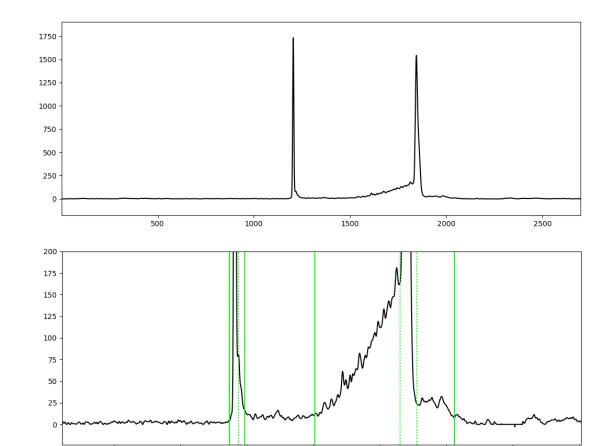






Integration depends on the context:

RNA degradation measurement



1500

1750

2000





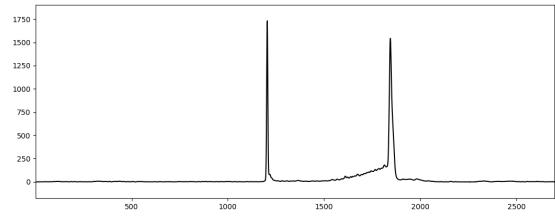
1000

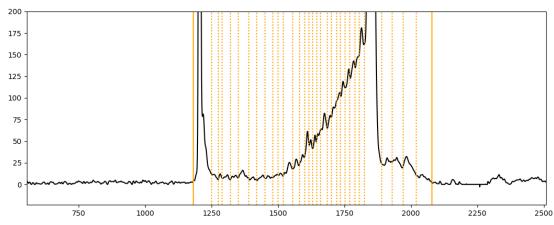
1250

750

Integration depends on the context:

RNA degradation measurement



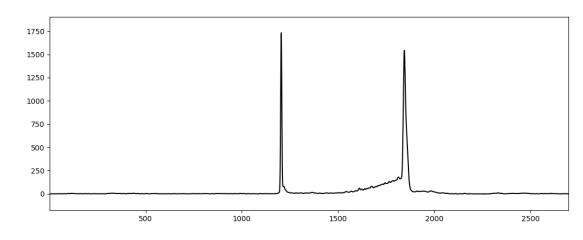




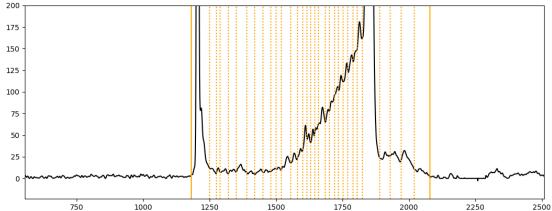


Integration depends on the context:

RNA degradation measurement



→ Cutting the front might not be necessary for the analysis

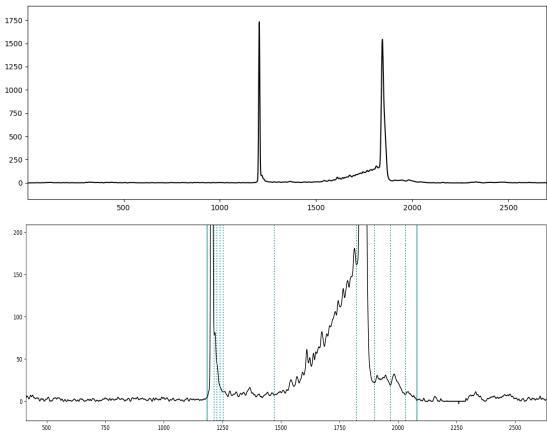






Integration depends on the context:

RNA degradation measurement



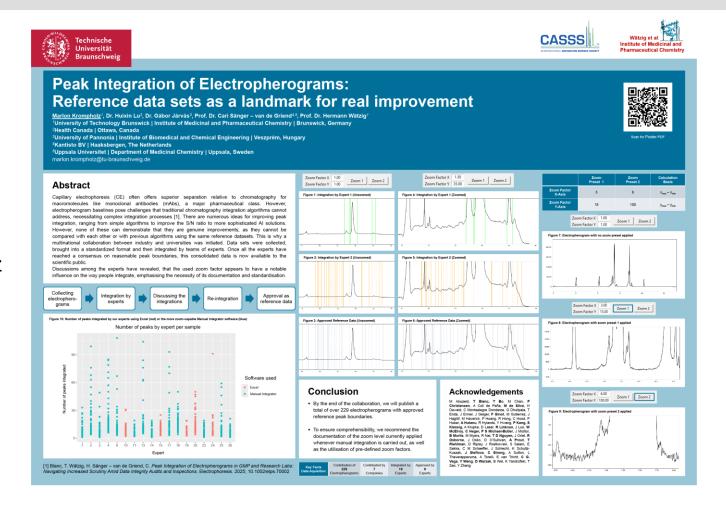




More Examples => Marlon's Poster

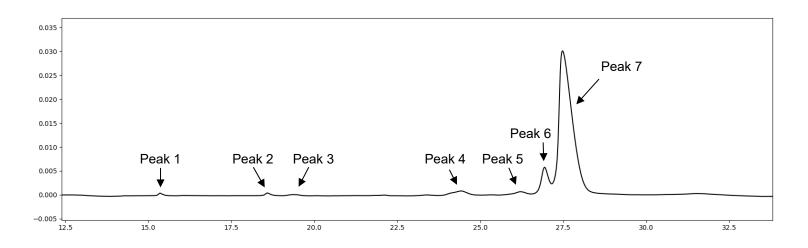


Marlon Krompholz





Potential



Series of n = 21 injections

	Peak Area RSD%							
	Peak 1	Peak 2	Peak 3	Peak 4	Peak 5	Peak 6	Peak 7	
Empower	15.43	8.13	25.56	11.96	23.04	3.20	1.36	
Manual Integration*	12.40	5.75	10.08	9.94	19.16	3.19	1.01	





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The next steps

- Approval by the community
- Reference Data Set CE Integration #1





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Software development and qualification

- Performance tests of existing softwares: precision & accuracy
- Development of algorithmic and Al strategies

 Investigations on Human errors in integration

Extending the concept to LC (HPLC 2025)





Thank you very much!



Jana Haegner, Nelly Luong, Marlon Krompholz, Hermann Wätzig













