

# A statistical framework for joint pre-processing and deconvolution of solid-state NMR mixture spectra

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Piotr Prostko

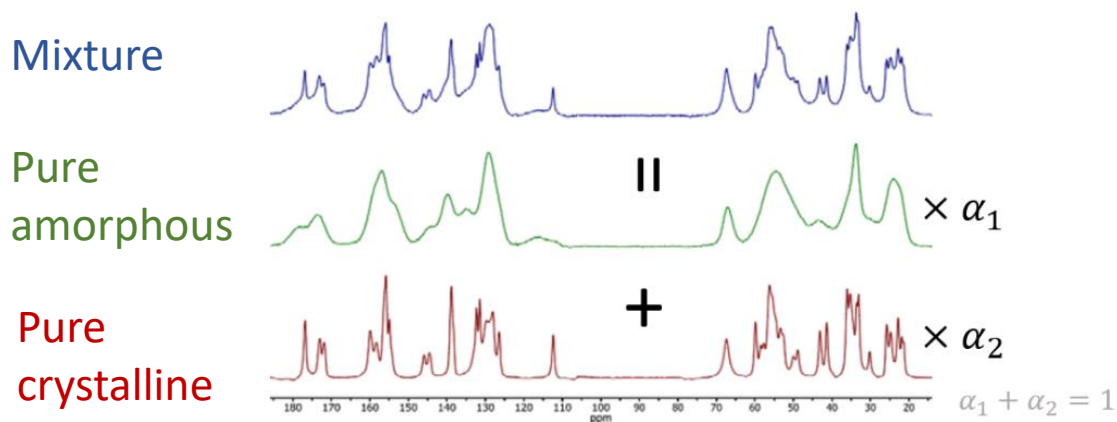
[piotr.prostko@uhasselt.be](mailto:piotr.prostko@uhasselt.be)



PHARMACEUTICAL COMPANIES OF  
*Johnson & Johnson*

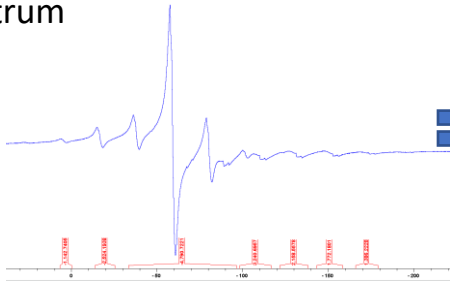
- Nuclear Magnetic Resonance (NMR) spectroscopy useful in studying various drug product properties, e.g. bioavailability
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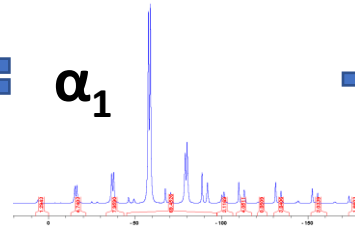


**Goal:** given ssNMR spectra of pure crystalline and amorphous reference samples, estimate  $\alpha_1$  and  $\alpha_2$  proportions in analytical, mixed samples

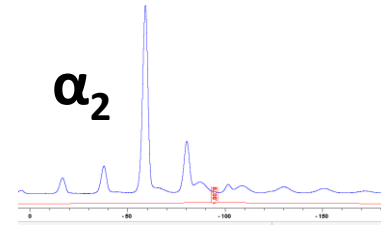
Raw analytical mixture spectrum



Pure solid-state form1 template



Pure solid-state form2 template



Mixing proportions  
 $\alpha_1, \alpha_2$

Non-linear preprocessing  
(phase correction and  
horizontal shifting)

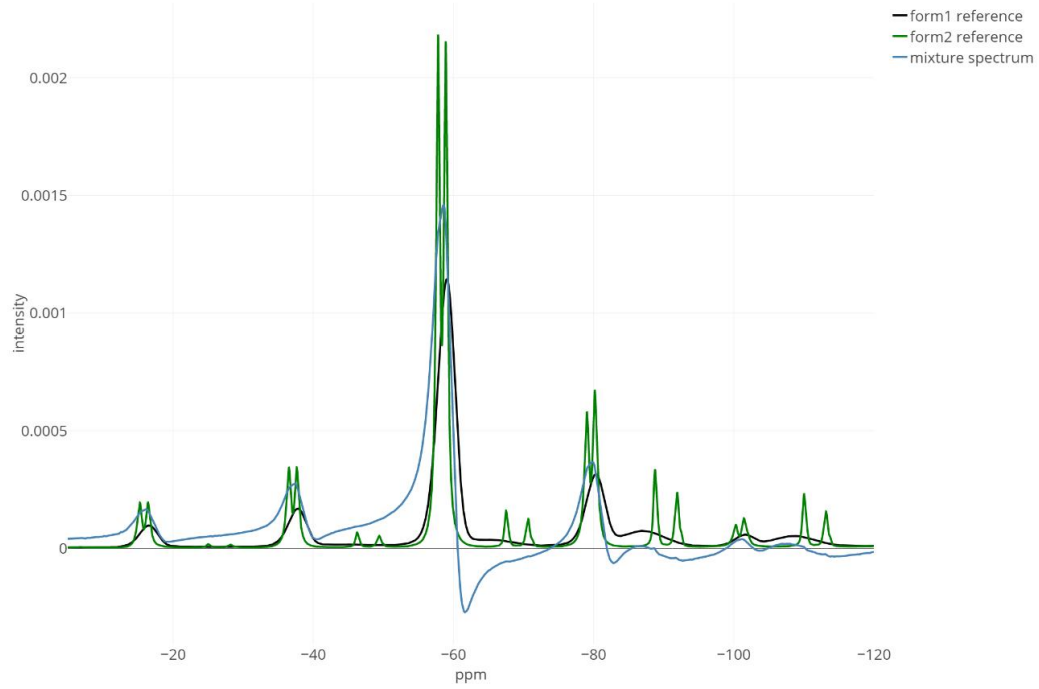


Constrained, non-linear optimization

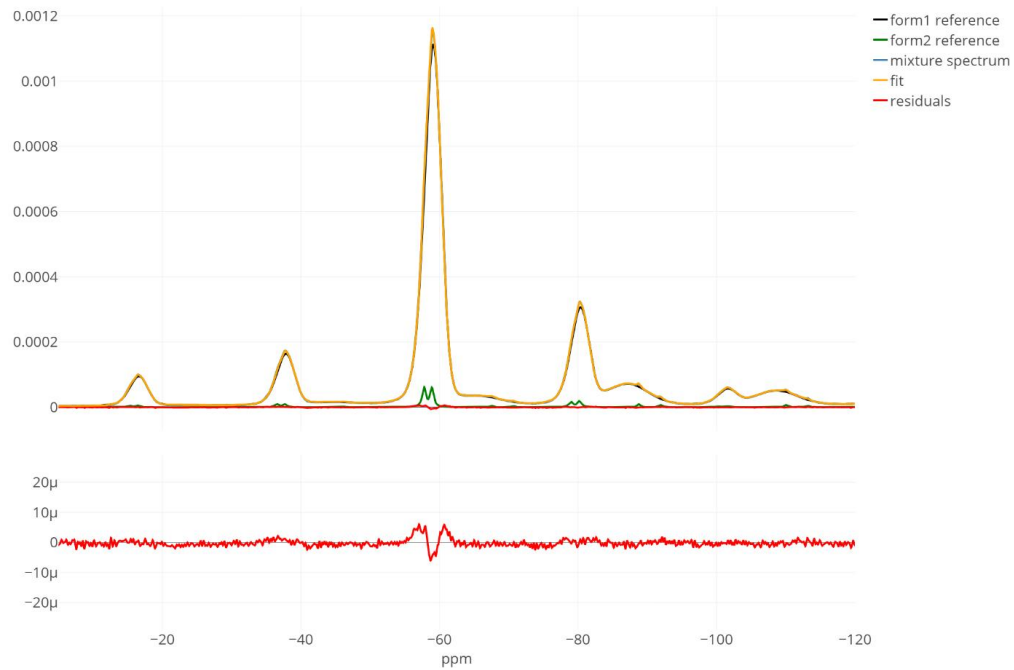
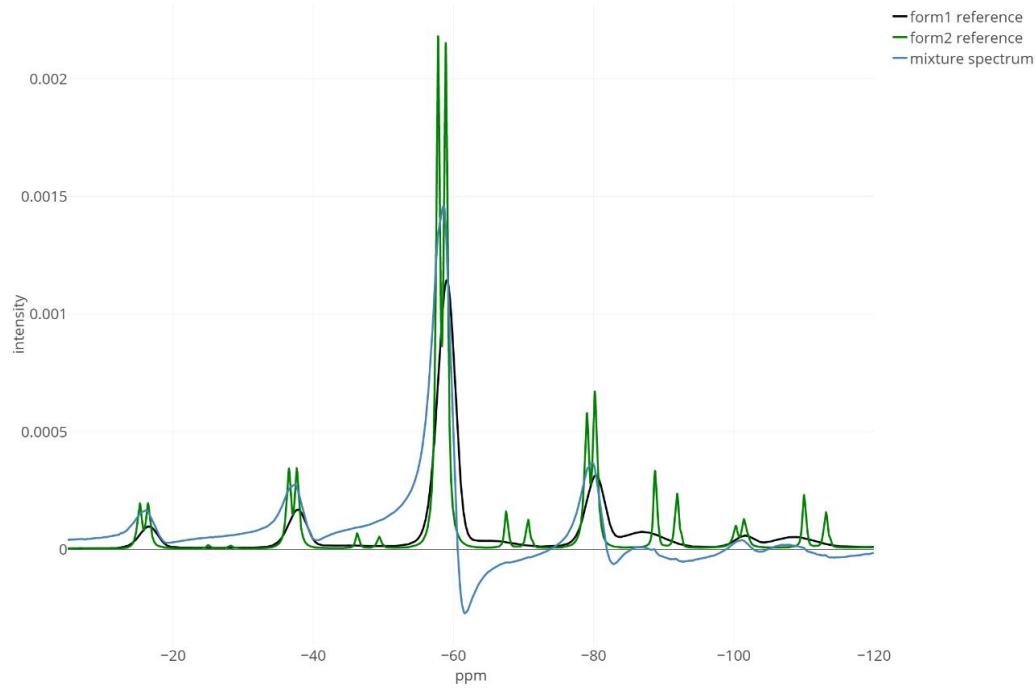
Objective: Residual Sum of Squares

Software: R nloptr()

The method applied to 3% crystalline mixture and reference spectra ...



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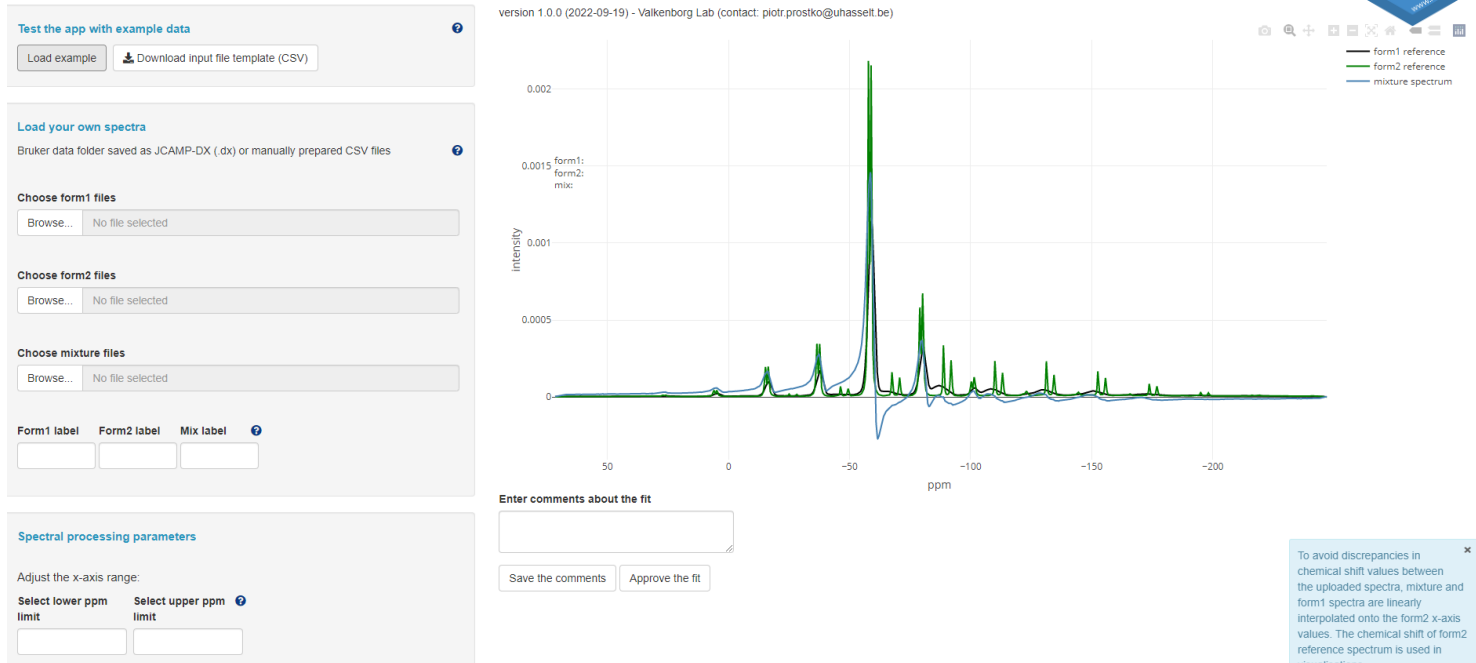


... returned 2.84% estimate and a good residual trace

# Try it for yourself!



## Automated deconvolution of solid-state NMR mixture spectra



- Safe to use (data processed on-the-fly)
- Flexible and interactive
- Promotes reproducibility
- Nucleus agnostic ( $^{19}\text{F}$ ,  $^{13}\text{C}$ , ...)

# Try it for yourself!



<https://valkenborg-lab.shinyapps.io/ssNMRdeconvolution/>

