Sequential Monte Carlo for Bayesian Analysis of Raman Spectroscopy

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Raman Spectroscopy



Illustration courtesy Jake Carson (U. Warwick)

Sequential Monte Carlo

Observed Spectrum





Sequential Monte Carlo

Raman scattering



Illustration courtesy Jake Carson (U. Warwick)

Surface-enhanced Raman scattering (SERS)

- Raman signal enhanced by proximity to nanoparticles
- Functionalisation using antibodies



Illustration courtesy Kirsten Gracie (U. Strathclyde)

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SuperCam

• 532nm Raman spectroscopy at up to 12 m distance



Functional Model

Separate the hyperspectral signal into 3 components:

$$\mathbf{y}_i = \xi_i(\tilde{\nu}) + s_i(\tilde{\nu}) + \boldsymbol{\epsilon}_i \tag{1}$$

where:

- \mathbf{y}_i is a an observed spectrum, discretised at multiple wavenumbers $\nu_j \in \widetilde{\mathcal{V}}$ (cm⁻¹)
- $\xi_i(\tilde{\nu})$ is a smooth baseline function
- $s_i(\tilde{\nu})$ is the spectral signature of the molecule
 - $\epsilon_{i,j} \sim \mathcal{N}\left(0,\sigma_{\epsilon}^{2}
 ight)$ is additive, zero mean white noise

Baseline

Penalised spline:

$$\xi_i(\tilde{\nu}) = \sum_{m=1}^M B_m(\tilde{\nu}) \alpha_{i,m}$$
(2)

$$\pi(\alpha_{i,\cdot}) \sim \mathcal{N}_{M}(0, \Sigma_{\lambda})$$
(3)

where $B_m(\tilde{\nu})$ are Demmler-Reinsch or B-spline basis functions



Spectral Signature

An additive mixture of radial basis functions:

$$s_{i}(\tilde{\nu}) = \sum_{p=1}^{P} A_{i,p} f\left(\tilde{\nu} \mid \ell_{p}, \varphi_{p}\right)$$
(4)

where:

 ℓ_p is the location of peak p $A_{i,p}$ is the amplitude φ_p is the scale (broadening) Sequential Monte Carlo

Squared exponential

Peak broadening function is an unnormalised Gaussian density:

$$f(\nu_{j} | \ell_{p}, \varphi_{p}) = \exp\left\{\frac{(\nu_{j} - \ell_{p})^{2}}{2\varphi_{p}^{2}}\right\}$$
(5)

$$FWHM = 2\sqrt{2 \ln 2}\varphi_{p}$$
(6)



Lorentzian Peaks

Long-range dependence between peaks can be modelled using an unnormalised Cauchy density:

$$f(\nu_{j} | \ell_{p}, \varphi_{p}) = \frac{\varphi_{p}^{2}}{(\nu_{j} - \ell_{p})^{2} + \varphi_{p}^{2}}$$
(7)

$$FWHM = 2\varphi_{p}$$
(8)



Informative priors

Obtained from manual peak fitting of independent data:



RRUFF Project

Anorthite R040059

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Name: Anorthite RRUFF ID: R040059 Ideal Chemistry: Ca(Al₂Si₂O₈) Locality: Miyakejima, Japan Source: University of Arizona Mineral Museum 4079 [view label] Owner: RRUFF Description: Colorless fragments of crystals Status: The identification of this mineral has been confirmed by X-ray diffraction and chemical analysis

Mineral Group: [feldspar (58)] Quick search: [All Anorthite samples (12)]

Lafuente, Downs, Yang & Stone (2015) In: *Highlights in Mineralogical Crystallography*, pp 1–30.

Raman Spectrum



Sequential Monte Carlo

Peak Locations



Sharma, Simons & Yoder Jr. (1983) American Mineralogist, 68: 1113-1125.

Time-dependent density functional theory

Locations of Raman peaks can be predicted from chemical structure using a quantum mechanical model (TD-DFT):



Watanabe, Hayazawa, Inouye & Kawata (2005) J. Phys. Chem. B 109(11)

TD-DFT, continued

Off-the-shelf implementations:

- Gaussian 09, Amsterdam Density Functional (ADF), Quantum Espresso (QE), etc.
- B3LYP functional with the basis set of 6-311++G(d, p)

Rhodamine B: C₂₈H₃₁CIN₂O₃ (PubChem ID 6694)

- Use the crystal structure of the molecule as an initial geometry
- Optimise to obtain the resting state (energy minimum)
- Calculate potential energy distributions for the vibrational modes (C–H out-of-plane bend, C=C symmetric stretch, etc.)
- Apply selection rules to determine Raman scattering frequencies and infrared absorption frequencies

Becke (1993) *J Chem. Phys.* **98**: 5648. Lee, Yang & Parr (1988) *Phys. Rev. B* **37**: 785.

Markov chain Monte Carlo

MCMC targeting the joint posterior π (**A**, φ , $\ell \mid y_i(\tilde{\nu})$)

Marginal likelihood is available in closed form:

$$p(y_i(\tilde{\nu}) \mid \mathbf{A}, \varphi, \ell) = \int \int p(y_i(\tilde{\nu}) \mid \mathbf{\Theta}) \pi(\alpha) \pi(\sigma_{\epsilon}^2) \, d\alpha \, d\sigma_{\epsilon}$$
$$= \frac{p(y_i(\tilde{\nu}) \mid \mathbf{\Theta}) \pi(\alpha) \pi(\sigma_{\epsilon}^2)}{p(\alpha, \sigma_{\epsilon}^2 \mid y_i(\tilde{\nu}), \mathbf{A}, \varphi, \ell)}$$

Given random walk proposals for $\mathbf{A}', \varphi', \ell'$, accept with probability min $(1, \rho_t)$ where:

$$\rho_t = \frac{p(y_i(\tilde{\nu}) \mid \mathbf{A}', \varphi', \ell') \pi(\mathbf{A}') \pi(\varphi') \pi(\ell')}{p\left(y_i(\tilde{\nu}) \mid \mathbf{A}^{(t-1)}, \varphi^{(t-1)}, \ell^{(t-1)}\right) \pi(\mathbf{A}^{(t-1)}) \pi(\varphi^{(t-1)}) \pi(\ell^{(t-1)})}$$

Chib (JASA, 1995) "Marginal Likelihood from the Gibbs Output."

Rao-Blackwellized Particle Filter

Particle-based method targeting a sequence of partial posteriors $\pi_t (\mathbf{A}, \varphi, \ell \mid y_i(\tilde{\nu}))$

Algorithm 1 SMC

- 1: Initialise $arphi^{(q)}$, $\mathbf{A}^{(q)}$, $\boldsymbol{\ell}^{(q)}$ $orall \, q \in \{1,\dots,Q\}$
- 2: Initialise importance weights, $w_0^{(q)} = \frac{1}{Q}$
- 3: for all iterations $t = 1, \ldots, T$ do
- 4: Update importance weights:

$$w_t^{(q)} \propto w_{t-1}^{(q)} \frac{p\left(y_i(\tilde{\nu}) \mid \boldsymbol{\ell}, \mathbf{A}, \boldsymbol{\varphi}\right)^{\kappa_t}}{p\left(y_i(\tilde{\nu}) \mid \boldsymbol{\ell}, \mathbf{A}, \boldsymbol{\varphi}\right)^{\kappa_{t-1}}}$$
(9)

- 5: Resample particles if ESS_t is below threshold
- 6: for all particles $q \in \{1, \ldots, Q\}$ do
- 7: Update $\varphi^{(q)}$, $\mathbf{A}^{(q)}$, $\ell^{(q)}$ using MCMC steps
- 8: end for
- 9: end for

Del Moral, Doucet & Jasra (2006)

Likelihood Tempering



ESS







- Multinomial (bootstrap particle filter)
- Systematic
- Stratified
- Residual
- Can parallelize by ordering the ancestry vector

Douc, Cappé & Moulines (*Proc. 4th IEEE ISPA*, 2005) "Comparison of resampling schemes for particle filtering." Murray, Lee & Jacob (*JCGS*, 2016) "Parallel resampling in the particle filter."

SMC collapse



SMC iteration

serrsBayes

An R package for Bayesian modelling and quantification of Raman spectroscopy using sequential Monte Carlo (SMC) algorithms:

- RcppEigen for fast linear algebra in C++
- OpenMP for parallelism

```
library(serrsBayes)
library(hyperSpec)
spec ~ read.spc("spectrum.spc")
lPriors ~ list(..)
result ~ fitSpectraSMC(spec$wl,spec$spc,lPriors)
```

Bates & Eddelbuettel (2013) Fast and Elegant Numerical Linear Algebra Using the RcppEigen Package. *J. Stat. Soft.* **52**(5): 1–24. Beleites & Sergo (2014) hyperSpec: a package to handle hyperspectral data sets in R.

Posterior distribution



Summary

serrsBayes provides an open-source approach to analysis of spectroscopy:

- Joint estimation of baseline and peaks
- 95% CI for peak locations, amplitudes, and FWHM

Ongoing and future work:

- Scalable computation using a divide-and-conquer algorithm
- T-optimum experimental design for multiplex Raman
- Spatial and temporal modelling of Raman maps
- Other types of spectroscopy (RF, X-ray, LIBS)

For Further Reading I

Moores, Gracie, Carson, Faulds, Graham & Girolami Bayesian modelling and quantification of Raman spectroscopy. arXiv preprint arXiv:1604.07299 [stat.AP]

Noonan, Asiala, Grassia, MacRitchie, Gracie, Carson, Moores, et al. In vivo multiplex molecular imaging of vascular inflammation using surface-enhanced Raman spectroscopy.

To appear in Theranostics.



Gracie, Moores, Smith, Harding, Girolami, Graham, & Faulds Preferential attachment of specific fluorescent dyes and dye labelled DNA sequences in a SERS multiplex. *Anal. Chem.*, 88(2): 1147–1153, 2016.

Zhong, Girolami, Faulds & Graham

Bayesian methods to detect dye-labelled DNA oligonucleotides in multiplexed Raman spectra.

J. R. Stat. Soc. Ser. C, 60(2): 187–206, 2011.

For Further Reading II

📔 Särkkä, Vehtari & Lampinen

Rao-Blackwellized particle filter for multiple target tracking *Information Fusion* 8(1): 2–15, 2007.

- R. Douc, O. Cappé & E. Moulines
 Comparison of resampling schemes for particle filtering
 In Proc. 4th IEEE Int. Symp. Image and Signal Processing and Analysis, 2005.

L.M. Murray, A. Lee & P.E. Jacob Parallel resampling in the particle filter *J. Comput. Graph. Stat.* **25**(3): 789–805, 2016.

F. Lindsten, A.M. Johansen, C.A. Naesseth, B. Kirkpatrick, T.B. Schön, J.A.D. Aston & A. Bouchard-Côté Divide-and-conquer with sequential Monte Carlo J. Comput. Graph. Stat. 26(2): 445–458, 2017.

Dilution study for FAM

315 spectra at 21 different concentrations, from 0.13 to 24.7 nM



Results: Baseline correction



(a) Posterior means of the baselines

(b) Baseline-corrected spectra

Results: Quantification

SERS peak intensities at 650cm⁻¹: 95% CI [257.7; 262.5] $\times c_i$

