

More model fitting

Mark Blyth

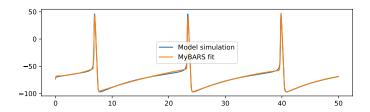


Presentation points

- Models for wind tunnel data
- A BARS redesign
- ₭ BARS potential pitfalls



An exciting aside

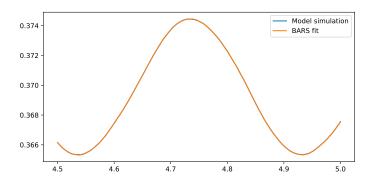


My BARS implementation works!

- Allows arbitrary interpolation
- Slower than the C implementation



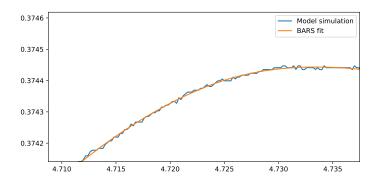
BARS - wind tunnel



2500 datapoints, 41s run-time



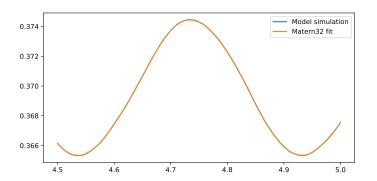
BARS - wind tunnel



2500 datapoints, 40s run-time



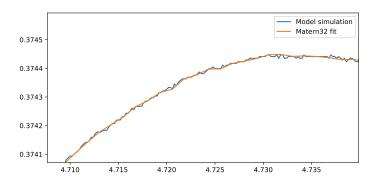
Matern32 (GPR) - wind tunnel



2500 datapoints, 119s run-time



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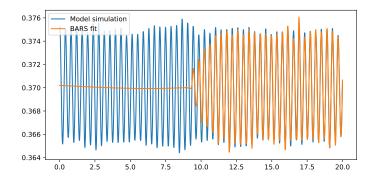


Timings

- Unoptimized C code is still faster than optimized python / matlab
 - BARS is using optimized C implementation, GPRs is using my unoptimized python implementation
 - GPFlow, GPyTorch would be much faster for GPR
 - Probabilistic programming might speed up BARS
- **W** Both methods are $\mathcal{O}(n^3)$; require inverting $n \times n$ matrix
 - Both perform poorly on lots of datapoints!



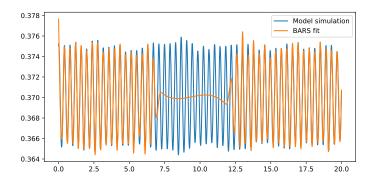
BARS



10,000 datapoints [crashes with full dataset], 142s run-time



BARS



10,000 datapoints [crashes with full dataset], max. number of knots [80], more

burn-in; 590s (9 mins 50s) run-time

bristol.ac.uk



BARS issues

- C implementation doesn't allow validation
 - Can't interpolate at arbitrary points
 - Python implementation does! But it's slower
- C implementation caps the max. number of knots
 - Python implementation doesn't! But it's slower
 - Doesn't work for long time series
- $\operatorname{\mathscr{U}}(n^3)$ means data need downsampling to be used
 - Took about 10 minutes to fit a model to 10% of the data



Experimental data

- If experiments use small numbers of periods (1, 2, 3), current methods work
 - ► Similar to how I tested models on small numbers of neuron spikes
 - Few periods = less data = fewer knots needed, and quicker fit obtained
- More datapoints mean we need to get creative...
 - Sparse variational BARS
 - Periodic BARS
 - Semiperiodic methods

Or... speed up BARS by not using BARS



IDEA: Bayesian adaptive evolutionary splines; genetic algorithm

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 - Each MCMC acceptance/rejection becomes an evolutionary scoring step



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- 5. Can maximise posterior, instead of likelihood, by including a prior term



BARS vs evolution

- Would be interesting to compare this to the MCMC method
 - ► MCMC sets up a Markov chain whose stationary distribution is the posterior
 - ► This aims to find a Markov chain whose stationary distribution is the argmax
 - Grounds for a rigorous justification / proof of convergence
- Evolution will likely be faster
 - Could leverage existing genetic optimization packages
 - Easily parallelised for more speed-up
 - No RJ-MCMC makes it easier target for probabilistic programming



SV-BARS

Another idea: remodel BARS to work similarly to sparse GPR

- - **Each MCMC** step requires inverting an $n \times n$ matrix SLOW
- $m{k}$ Choose a [small] set of maximally informative surrogate datapoints (x_i^*, y_i^*)
- Run MCMC step on surrogate datapoints
 - Much faster to invert the smaller matrix
- Well-chosen surrogate points means we get the same result as running on real data
- Fewer datapoints means it runs a lot faster



SV-BARS implementation

- - Find inducing points by minimising $D_{\mathrm{KL}}\left[p_{y}\|p_{y^{*}}\right]$
 - ► Use variational Bayes to approximate this
- \checkmark Find posterior knots $p(k, \xi|y^*)$ [instead of $p(k, \xi|y)$]
 - $ightharpoonup \mathcal{O}(m^3), m \ll n$
 - ▶ Sparse GPR is $\mathcal{O}(nm^3)$, so if my complexity is correct, we get a bigger speed-up / outperform SVGPR!

Would require learning RJ-MCMC, variational Bayes, sparse GPR in-depth



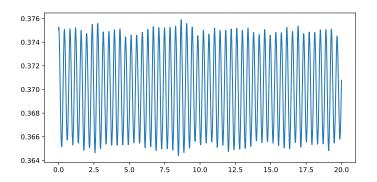
Periodic BARS

An approach for if we need to model more than a few spikes:

- **K** Assume data are given by $y_i = f(x_i) + \varepsilon$
 - f(t) = f(t+T)
- Find either
 - ► T-periodic knot-set
 - ▶ T-periodic basis splines
 - Nicer approach
- ...in such a way that we...
 - k minimise fitting time
 - k balance fit against number of knots



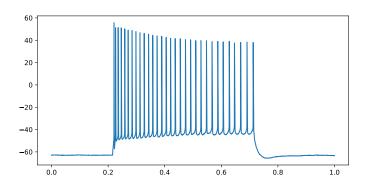
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Experimental data aren't perfectly periodic [could Ca²⁺, or experiment setup!]



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- ▶ BARS, GPR, NARMAX all interesting model options
- Would likely give similar results to sparse BARS



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- (Semi)periodic BARS would be less generally applicable, but potentially faster when applicable



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- 4. Make a periodic BARS setup, then use that in CBC
 - Periodic BARS is a less general method



My proposal

- Validate models
 - ► Also try simple data transformations for GPR
- Write up notes on everything so far

Then...

- 1. Adapt BARS for sparsity, evolution
 - Fast, SOTA, scalable
 - Needs variational Bayes learned
- 2. Demonstrate sparse BARS on other problems (NDC, comp-synth-bio, ML)
 - Not sure which problems would benefit from splines?
- 3. Apply the shiny new splines method to CBC