## SCALABLE GRADIENTS OF GAUSSIAN PROCESS MODELS WITH CELERITE

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## ABSTRACT

This research note presents a derivation and implementation of reverse-mode gradients of the *celerite* algorithm for scalable inference with Gaussian Process (GP) models. These gradients can be integrated into existing automatic differentiation frameworks to provide a scalable and computationally efficient method for evaluating the gradients of the GP likelihood with respect to the kernel hyperparameters. The algorithm derived in this note uses less memory and is more efficient than versions using automatic differentiation and the computational cost scales linearly with the number of data points.

## INTRODUCTION

Gaussian Processes (GPs Rasmussen & Williams 2006) are a class of models used extensively in the astrophysical literature to model stochastic processes. The applications are broad-ranging and some example include the time domain variability of astronomical sources (Brewer & Stello 2009; Kelly et al. 2014; Haywood et al. 2014; Rajpaul et al. 2015; Foreman-Mackey et al. 2017), data-driven models of light curves or stellar spectra (Wang et al. 2012; Luger et al. 2016; Czekala et al. 2017), and the cosmic microwave background (Bond & Efstathiou 1987; Wandelt & Hansen 2003). In all of these applications, the calculation and optimization of the GP marginalized likelihood function is of central importance and this can often be of the computational bottleneck. The details of this model are omitted here (more details can be found in Rasmussen & Williams 2006 and Foreman-Mackey et al. 2017), but the key point is that, for a dataset with N data points, every evaluation of a GP model requires computation of the inverse and log-determinant of the  $N \times N$  covariance matrix. The computational cost of these operations scales as  $\mathcal{O}(N^3)$  in the general case, but the celerite method was recently introduced in the astronomical literature to compute the GP likelihood for a class of one-dimensional models with  $\mathcal{O}(N)$  scaling (Foreman-Mackey et al. 2017). The details of these models can be found elsewhere (Foreman-Mackey et al. 2017), but the basic idea is that for some covariance functions, the matrix K can be written as a rank-J semi-separable matrix

$$K = \operatorname{diag}(\boldsymbol{a}) + \operatorname{tril}(U V^{\mathrm{T}}) + \operatorname{triu}(V U^{\mathrm{T}})$$
(1)

where a is a N-vector, U and V are  $N \times J$  matrices, the diag function creates a diagonal matrix from a vector, and the tril and triu extract the strict lower and upper triangular matrices from a dense matrix. For a matrix of this form, Foreman-Mackey et al. (2017) derived an algorithm with  $\mathcal{O}(NJ^2)$  scaling for computing the Cholesky factorization

$$K = L \operatorname{diag}(\boldsymbol{d}) L^{\mathrm{T}} \tag{2}$$

where

$$L = I + \operatorname{tril}(U W^{\mathrm{T}}) \tag{3}$$

for some  $N \times J$  matrix W. The algorithm from Foreman-Mackey et al. (2017) includes an  $N-1 \times J$  "preconditioning" matrix<sup>2</sup> that we include with the notation P. The Cholesky factorization algorithm derived by Foreman-Mackey et al. (2017) is given is reproduced in the following algorithm:

<sup>&</sup>lt;sup>1</sup> Or, more specifically, a linear operator that can left multiply a vector or a matrix by the inverse of this matrix.

<sup>&</sup>lt;sup>2</sup> This matrix is called  $\phi$  by Foreman-Mackey et al. (2017).

```
function celerite_factor(U, P, d, W)

# If at input d = a, W = V, and K is given by Equation (1),

# at output K = LL^{\mathrm{T}} where L is given by Equation (3).

S \leftarrow \mathsf{zeros}(J, J)

w_1 \leftarrow w_1/d_1

for n = 2, \dots, N:

S \leftarrow \mathsf{diag}(p_{n-1}) \left[S + d_{n-1} \, w_{n-1}^{\mathrm{T}} \, w_{n-1}\right] \mathsf{diag}(p_{n-1})

d_n \leftarrow d_n - u_n \, S \, u_n^{\mathrm{T}}

w_n \leftarrow \left[w_n - u_n \, S\right]/d_n

return d, W, S
```

In this algorithm, the zeros(J, K) function creates a  $J \times K$  matrix of zeros, the diag function creates a diagonal matrix from a vector, and  $x_n$  indicates a row vector made from the n-th row of the matrix X.

Similarly, Foreman-Mackey et al. (2017) derived a  $\mathcal{O}(NJ)$  algorithm to apply the inverse of K as shown in the following algorithm:

```
function celerite_solve(U, P, \boldsymbol{d}, W, Z)
# If at input Z = Y, at output Z = K^{-1} Y.

F \leftarrow \operatorname{zeros}(J, N_{\operatorname{rhs}})
G \leftarrow \operatorname{zeros}(J, N_{\operatorname{rhs}})
for n = 2, \dots, N:

F \leftarrow \operatorname{diag}(\boldsymbol{p}_{n-1}) [F + \boldsymbol{w}_{n-1}^{\mathrm{T}} \boldsymbol{z}_{n-1}]
\boldsymbol{z}_n \leftarrow \boldsymbol{z}_n - \boldsymbol{u}_n F
for n = 1, \dots, N:

\boldsymbol{z}_n \leftarrow \boldsymbol{z}_n / d_n
for n = N - 1, \dots, 1:
G \leftarrow \operatorname{diag}(\boldsymbol{p}_n) [G + \boldsymbol{u}_{n+1}^{\mathrm{T}} \boldsymbol{z}_{n+1}]
\boldsymbol{z}_n \leftarrow \boldsymbol{z}_n - \boldsymbol{w}_n G
return Z, F, G

(Murray 2016) and (Giles 2008).
```

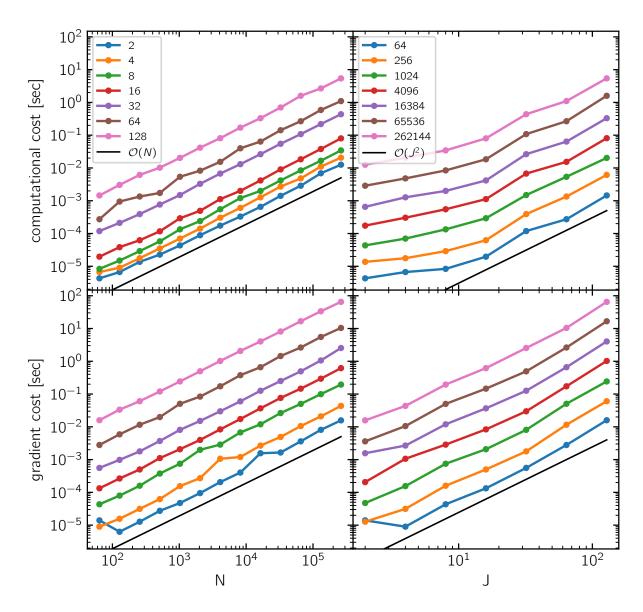


Figure 1. Scaling.

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