Section 6. Fast Stan

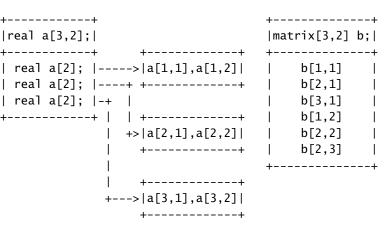
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Part I

Vectorization

Matrices vs Arrays



Vectorizing Models

Stan can calculate parts of the model more efficiently if contributions to the log likelihood are vectorized

· Multilevel model

```
for (n in 1:N)
 y[n] ~ bernoulli_logit(x[n,] * beta + alpha[g[n]]);
   VS
 vector[N] eta:
 for (n in 1:N)
    eta[n] = x[n,] * beta + alpha[g[n]];
 y ~ bernoulli_logit(eta);
   VS
 ~ bernoulli_logit(x * beta + alpha[q]);
```

Part II

Integration

Dimension Reduction

- If a parameter can be integrated from the posterior, doing so can increase performance
- · Often makes model code messier
- Parameter can be recoved by simulating from its conditional distribution in generated quantities

Marginalization Example

Varying intercept, linear model

$$y_i \mid \alpha \sim \text{Normal}(x_i^{\top} \beta + \alpha_{g[i]}, \sigma_y^2),$$

 $\alpha_j \sim \text{Normal}(0, \sigma_{\alpha}^2).$

Integrate out α to obtain:

$$y_i \sim \mathsf{Normal}(x_i^{\mathsf{T}} \beta, \sigma_y^2 + \sigma_\alpha^2),$$

$$\mathsf{Cov}(y_i, y_j) = \begin{cases} \sigma_\alpha^2 & \text{if } g[i] = g[j] \\ 0 & \text{otherwise} \end{cases} \text{ for } i \neq j$$

Stan jumps on K + 2 parameters vs K + J + 2

Marginalization Example pt. 2

```
data {
  int<lower = 0> N:
  int<lower = 0> K;
  int<lower = 0> N_group;
  vector[N] y;
  int<lower = 1, upper = J> group[N];
  matrix[N,K] x;
parameters {
  real<lower = 0> sigma_y;
  real<lower = 0> sigma alpha:
  vector[K] beta;
```

Marginalization Example pt. 3

```
transformed parmeters {
  cov_matrix[N] Sigma_y;
  Sigma_y = diag_matrix(rep_vector(sigma_y * sigma_y, N));
  for (row in 1:N) {
    for (col in 1:N) {
      if (group[row] == group[col]) Sigma_y[row,col] =
        Sigma_v[row,col] + sigma_alpha * sigma_alpha;
model {
  y ~ multi_normal(x * beta, Sigma_y);
  beta \sim cauchy(0, 10);
  sigma_y \sim cauchy(0, 2.5);
  sigma_alpha \sim cauchy(0, 2.5);
```

Marginalization Example pt. 4

Add a generated quantities block to simulate:

$$\alpha_j \mid y \sim \mathsf{Normal}\left(\frac{n_j/\sigma_y^2}{n_j/\sigma_y^2 + 1/\sigma_\alpha^2} \bar{y}_j, \frac{1}{n_j/\sigma_y^2 + 1/\sigma_\alpha^2}\right),$$

where
$$\bar{y}_j = \frac{1}{n_i} \sum_{i \in g_j} (y_i - x_i^{\mathsf{T}} \boldsymbol{\beta})$$

When to Marginalize

- When a model is slow, examine math and see if a parameter can be integrated
- Required for finite mixture models (cluster)
- Required for latent discrete parameters (change point models, noisy categorical measurements)
- · Censored values
- (Not recommended in MLM)

Exploiting Independence

```
data {
  int<lower = 0> N_in_group[N_group];
model {
  for (ng in 1:N_group) {
    cov_matrix[N_in_group[ng]] Sigma_y_ng;
    Sigma_y_ng = rep_matrix(sigma_gamma * sigma_gamma,
                            N_in_group[ng], N_in_group[ng]);
    for (i in 1:N_in_group[ng])
      Sigma_y_ng[i,i] = Sigma_y_ng[i,i] + sigma_y * sigma_y;
    y[group[ng]] ~
      multi_normal(x[group[ng],] * beta, Sigma_y_ng);
```

Part III

Decomposition

```
data {
  int<lower = 0> N:
  int<lower = 0> K;
  vector[N] y;
  matrix[N,K] x;
  cov_matrix[N] Sigma_y;
parameters {
  vector[K] beta;
model {
  y ~ multi_normal(x * beta, Sigma_y);
```

• Each sample requires computing Multi-Normal($x\beta, \Sigma_{\nu}$)

$$p(y) = (2\pi)^{-n/2} |\Sigma_y|^{-1/2} \exp\left\{-\frac{1}{2}(y - x\beta)^{\top} \Sigma_y^{-1}(y - x\beta)\right\}.$$

 Matrix determinants and inverses are slow and potentially unstable

- Internally, Multi-Normal $(x\beta,\Sigma_y)$ is computed by decomposing Σ_y
- Symmetric positive definite matrices behave like positive numbers ($a^{T}Aa > 0$), which have square roots
- · Cholesky decomposition $\Sigma_{\nu} = L_{\nu}L_{\nu}^{\mathsf{T}}$, L_{ν} is lower-triangular

$$\begin{bmatrix} 4 & 12 \\ 12 & 37 \end{bmatrix} = \begin{bmatrix} 2 & 0 \\ 6 & 1 \end{bmatrix} \begin{bmatrix} 2 & 6 \\ 0 & 1 \end{bmatrix}.$$

Determinant of L is product of diagonal, inverse straightforward to compute

```
v \sim \text{Multi-Normal}(x\beta, \Sigma_v),
                    y_0 = y - x\beta.
                    y_0 \sim \text{Multi-Normal}(0, \Sigma_v),
                     z = L^{-1}(\nu - x\beta).
                      z \sim \text{Multi-Normal}(0, I_n).
model {
  vector[N] z;
  z = L.inv * (y - x * beta);
  z \sim normal(0, 1);
```

```
data {
  cov_matrix[N] Sigma_y;
transformed data {
  cholesky_factor_cov[N] L_y;
  L_y = cholesky_decompose(Sigma_y);
model {
  v ~ multi_normal_cholesky(x * beta, L_y);
```

- · Avoid for loops if possible, use vectorized constructions
- · Reuse decompositions if possible
- If changing on every iteration, gain nothing by storing decomp
- Further work on efficient matrix construction email to ask/request

```
https://groups.google.com/forum/#!forum/
stan-users
```