

# GPU-parallel Gibbs sampling of a hierarchical model of hybrid vigor in RNA-seq experiments

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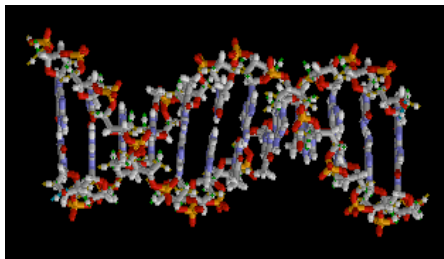
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# DNA



```

... — GTGCATCTGACTCCTGAGGAGAAG ...
... — CACGTAGACTGAGGACTCCTCTTC ...

```

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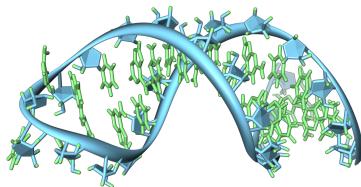
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# RNA



... — GUGCAUCUGACUCCUGAGGAGAAG — ...

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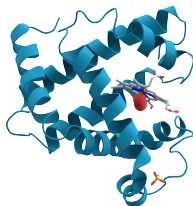
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# Proteins



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# Central dogma: how organisms make proteins

... GTGCATCTGACTCCTGAGGAGAAG ...  
... CACGTAGACTGAGGACTCCTCTTC ... DNA

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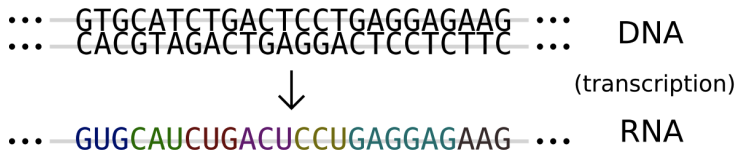
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# Central dogma: how organisms make proteins



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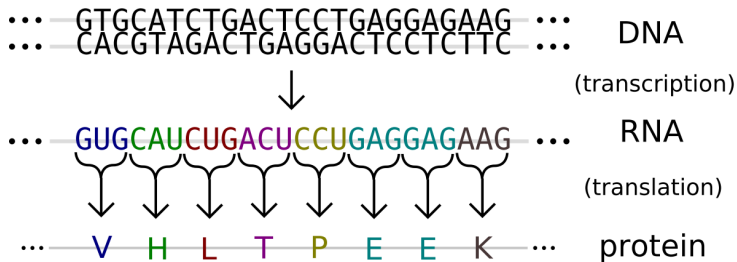
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# Central dogma of genetics



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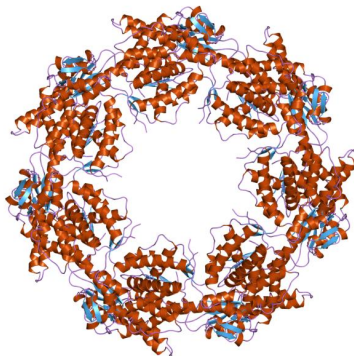
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# HSP60

- ▶ HSP = heat shock protein.
- ▶ Prevent heat damage to other proteins.



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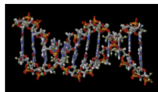
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# Temperature spike triggers HSP60 production.

HSP60 Gene



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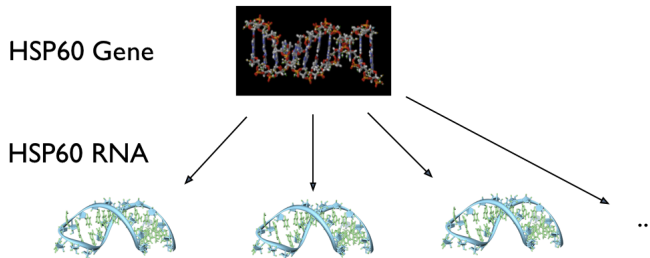
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# Temperature spike causes HSP60 expression.



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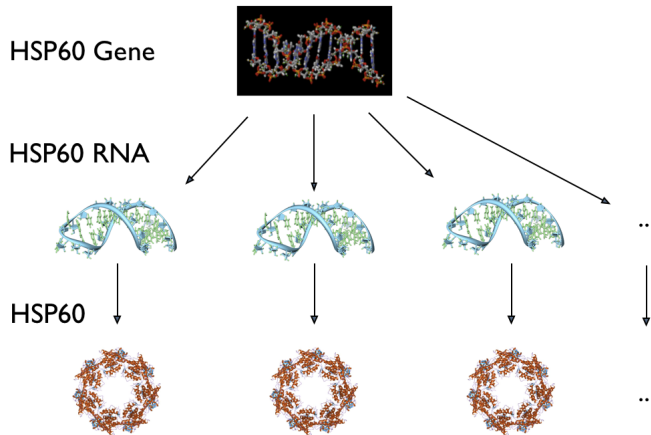
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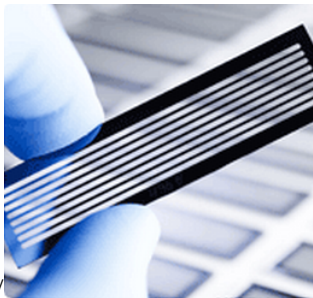
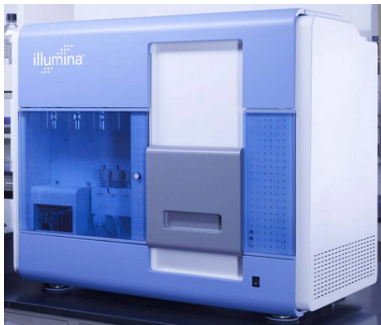
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# RNA-seq

- ▶ RNA sequencing: measure gene expression using relative abundance of RNA.
- ▶ Illumina Genome Analyzer:



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# RNA-seq data: counts of amplified RNA fragments

	Treatment 1			Treatment 2		Treatment 3	
Gene 1	100	225	0	70	279	300	106
Gene 2	0	1	1	50	501	2	7
Gene 3	3	4	2	700	900	0	0
Gene 4	893	400	760	5	5	1000	513
...	...	...	...	...	...	...	...
Gene 34897	10	13	6	819	761	902	912

- Goal: use RNA-seq to study hybrid vigor (heterosis).

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# High-parent heterosis: child's trait surpasses both parents

Parent 1



Parent 2



Child



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# Low-parent heterosis: child's trait is weaker than in each parent

Parent 1



Parent 2



Child



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# Mid-parent heterosis: child's trait is different than average of parents

Parent 1



Parent 2



Child



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# High-parent heterosis in gene expression

	Parent 1			Child		Parent 2	
Gene 1	100	225	0	70	279	300	106
Gene 2	0	1	1	50	501	2	7
Gene 3	3	4	2	700	900	0	0
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$$y_{g,n} \stackrel{\text{ind}}{\sim} \text{Poisson}(y_{g,n} \mid \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g)))$$

$$c_n \stackrel{\text{ind}}{\sim} N(c_n \mid 0, \sigma_c^2)$$

$$\sigma_c \sim U(\sigma_c \mid 0, \sigma_{c0})$$

$$\varepsilon_{g,n} \stackrel{\text{ind}}{\sim} N(\varepsilon_{g,n} \mid 0, \eta_g^2)$$

$$\eta_g^2 \stackrel{\text{ind}}{\sim} \text{Inv-Gamma} \left( \eta_g^2 \mid \text{shape} = \frac{d}{2}, \text{rate} = \frac{d \cdot \tau^2}{2} \right)$$

$$d \sim U(d \mid 0, d_0)$$

$$\tau^2 \sim \text{Gamma}(\tau^2 \mid \text{shape} = a_\tau, \text{rate} = b_\tau)$$

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$$\mu(n, \phi_g, \alpha_g, \delta_g) = \begin{cases} \phi_g - \alpha_g & \text{sample } n \text{ from parent 1} \\ \phi_g + \delta_g & \text{sample } n \text{ from child} \\ \phi_g + \alpha_g & \text{sample } n \text{ from parent 3} \end{cases}$$

$$y_{g,n} \stackrel{\text{ind}}{\sim} \text{Poisson}(y_{g,n} \mid \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g)))$$

$$\phi_g \stackrel{\text{ind}}{\sim} \text{N}(\phi_g \mid \theta_\phi, \sigma_\phi^2)$$

$$\theta_\phi \sim \text{N}(\theta_\phi \mid 0, \gamma_\phi^2)$$

$$\sigma_\phi \sim \text{U}(\sigma_\phi \mid 0, \sigma_{\phi 0})$$

$$\alpha_g \stackrel{\text{ind}}{\sim} \pi_\alpha^{1-I(\alpha_g)} [(1 - \pi_\alpha) \text{N}(\alpha_g \mid \theta_\alpha, \sigma_\alpha^2)]^{I(\alpha_g)}$$

$$\theta_\alpha \sim \text{N}(\theta_\alpha \mid 0, \gamma_\alpha^2)$$

$$\sigma_\alpha \sim \text{U}(\sigma_\alpha \mid 0, \sigma_{\alpha 0})$$

$$\pi_\alpha \sim \text{Beta}(\pi_\alpha \mid a_\alpha, b_\alpha)$$

$$\delta_g \stackrel{\text{ind}}{\sim} \pi_\delta^{1-I(\delta_g)} [(1 - \pi_\delta) \text{N}(\delta_g \mid \theta_\delta, \sigma_\delta^2)]^{I(\delta_g)}$$

$$\theta_\delta \sim \text{N}(\theta_\delta \mid 0, \gamma_\delta^2)$$

$$\sigma_\delta \sim \text{U}(\sigma_\delta \mid 0, \sigma_{\delta 0})$$

$$\pi_\delta \sim \text{Beta}(\pi_\delta \mid a_\delta, b_\delta)$$

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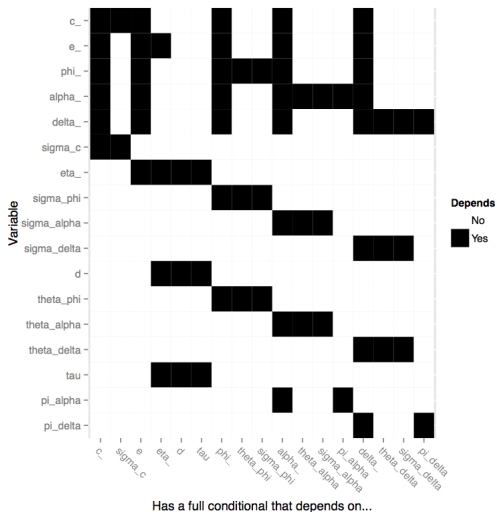
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# Partition parameters by conditional independence.



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# Use these partitions as Gibbs steps.

$$y_{g,n} \stackrel{\text{ind}}{\sim} \text{Poisson}(y_{g,n} \mid \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g)))$$

- From the appropriate full conditional distributions, sample the following:

1.  $c_1, \dots, c_N$
2.  $\tau, \pi_\alpha, \pi_\delta$
3.  $d, \theta_\phi, \theta_\alpha, \theta_\delta$
4.  $\sigma_c, \sigma_\phi, \sigma_\alpha, \sigma_\delta, \eta_1^2, \dots, \eta_G^2$
5.  $\varepsilon_{1,1}, \varepsilon_{1,2}, \dots, \varepsilon_{1,N}, \varepsilon_{2,N}, \dots, \varepsilon_{G,N}$
6.  $\phi_1, \dots, \phi_G$
7.  $\alpha_1, \dots, \alpha_G$
8.  $\delta_1, \dots, \delta_G$

- and then repeat.

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# Estimated heterosis probabilities

$$\mu(n, \phi_g, \alpha_g, \delta_g) = \begin{cases} \phi_g - \alpha_g & \text{sample } n \text{ from parent 1} \\ \phi_g + \delta_g & \text{sample } n \text{ from child} \\ \phi_g + \alpha_g & \text{sample } n \text{ from parent 3} \end{cases}$$

$$y_{g,n} \stackrel{\text{ind}}{\sim} \text{Poisson}(y_{g,n} \mid \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g)))$$

Consider one chain with  $M$  iterations.

$$P(\text{high-parent heterosis in gene } g) \approx \frac{1}{M} \sum_{i=1}^M I(\delta_g^{(i)} > |\alpha_g^{(i)}|)$$

$$P(\text{low-parent heterosis in gene } g) \approx \frac{1}{M} \sum_{i=1}^M I(\delta_g^{(i)} < -|\alpha_g^{(i)}|)$$

$$P(\text{mid-parent heterosis in gene } g) \approx \frac{1}{M} \sum_{i=1}^M I(\delta_g^{(i)} \neq 0)$$

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# Tons of opportunity for GPU parallelism across genes!

$$y_{g,n} \stackrel{\text{ind}}{\sim} \text{Poisson}(y_{g,n} \mid \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g)))$$

► Sample in parallel:

- $\phi_g$ 's
- $\alpha_g$ 's
- $\delta_g$ 's
- $\varepsilon_{g,n}$ 's
- $\eta_g$ 's

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# Example: $\phi_g$ 's

$$y_{g,n} \stackrel{\text{ind}}{\sim} \text{Poisson}(y_{g,n} \mid \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g)))$$

$$\phi_g \stackrel{\text{ind}}{\sim} \text{N}(\phi_g \mid \theta_\phi, \sigma_\phi^2)$$

$$\theta_\phi \sim \text{N}(\theta_\phi \mid 0, \gamma_\phi^2)$$

$$\sigma_\phi \sim \text{U}(\sigma_\phi \mid 0, \sigma_{\phi 0})$$

- Using parallel random walk Metropolis steps, sample the  $\phi_g$ 's from their full conditional distributions,

$$p(\phi_g \mid \dots) \propto \exp \left( \sum_{n=1}^N [y_{g,n} \cdot \mu(n, \phi_g, \alpha_g, \delta_g) - \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g))] - \frac{(\phi_g - \theta_\phi)^2}{2\sigma_\phi^2} \right)$$

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- Use **parallel reductions** to calculate sufficient statistics for:
  - $c_n$ 's
  - $\tau, d$
  - $\theta_\phi, \theta_\alpha, \theta_\delta$
  - $\sigma_\phi, \sigma_\alpha, \sigma_\delta, \sigma_c$
  - $\pi_\alpha, \pi_\delta$

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# Pairwise summation: an example reduction

- ▶ Let's take the pairwise sum of the vector,

$$(5, 2, -3, 1, 1, 8, 2, 6)$$

using 1 block of 4 threads.

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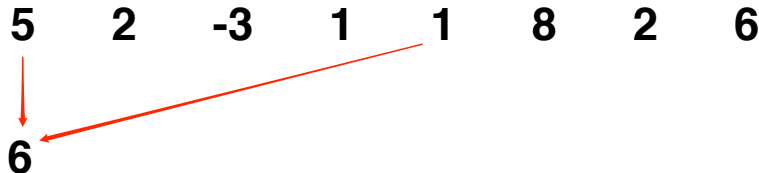
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# Pairwise summation: an example reduction



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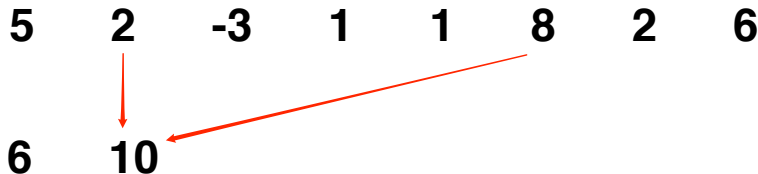
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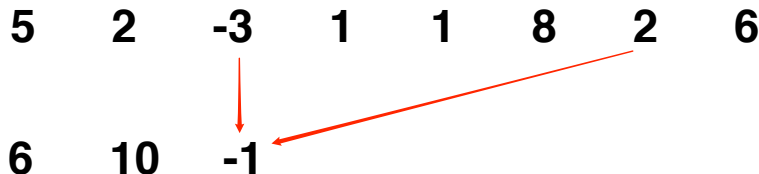
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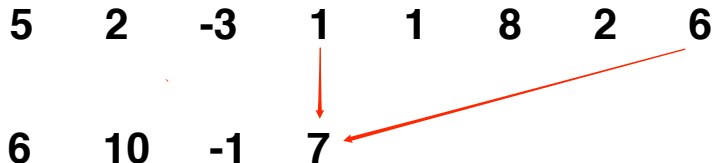
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# Pairwise summation: an example reduction



**Thread 3**

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# Pairwise summation: an example reduction

5	2	-3	1	1	8	2	6
6	10	-1	7				



## Synchronize threads

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# Pairwise summation: an example reduction

5      2      -3      1      1      8      2      6

6      10      -1      7

↓

5



**Thread 0**

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# Pairwise summation: an example reduction

5      2      -3      1      1      8      2      6

6      10      -1      7

5      17

**Thread 1**

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# Pairwise summation: an example reduction

5      2      -3      1      1      8      2      6

6      10      -1      7

5      17

**Synchronize Threads**

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# Pairwise summation: an example reduction

5      2      -3      1      1      8      2      6

6      10      -1      7

5      17

**Thread 0**

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# Example: $\tau^2$

$$y_{g,n} \stackrel{\text{ind}}{\sim} \text{Poisson}(y_{g,n} \mid \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g)))$$

$$\varepsilon_{g,n} \stackrel{\text{ind}}{\sim} \text{N}(\varepsilon_{g,n} \mid 0, \eta_g^2)$$

$$\eta_g^2 \stackrel{\text{ind}}{\sim} \text{Inv-Gamma} \left( \eta_g^2 \mid \text{shape} = \frac{d}{2}, \text{rate} = \frac{d \cdot \tau^2}{2} \right)$$

$$d \sim \text{U}(d \mid 0, d_0)$$

$$\tau^2 \sim \text{Gamma}(\tau^2 \mid \text{shape} = a_\tau, \text{rate} = b_\tau)$$

$$p(\tau^2 \mid \dots)$$

$$= \text{Gamma} \left( \tau^2 \mid \text{shape} = a_\tau + \frac{Gd}{2}, \text{rate} = b_\tau + \frac{d}{2} \sum_{g=1}^G \frac{1}{\eta_g^2} \right)$$

- Using a parallel reduction (NVIDIA's CUDA C/C++ Thrust library), calculate the sufficient statistic:

$$\sum_{g=1}^G \frac{1}{\eta_g^2}$$

- Use an efficient rejection sampler to sample  $\tau^2$ .

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# Example: $d$

$$y_{g,n} \stackrel{\text{ind}}{\sim} \text{Poisson}(y_{g,n} \mid \exp(c_n + \varepsilon_{g,n} + \mu(n, \phi_g, \alpha_g, \delta_g)))$$

$$\varepsilon_{g,n} \stackrel{\text{ind}}{\sim} \text{N}(\varepsilon_{g,n} \mid 0, \eta_g^2)$$

$$\eta_g^2 \stackrel{\text{ind}}{\sim} \text{Inv-Gamma} \left( \eta_g^2 \mid \text{shape} = \frac{d}{2}, \text{rate} = \frac{d \cdot \tau^2}{2} \right)$$

$$d \sim \text{U}(d \mid 0, d_0)$$

$$\tau^2 \sim \text{Gamma}(\tau^2 \mid \text{shape} = a_\tau, \text{rate} = b_\tau)$$

$$p(d \mid \dots) \propto \Gamma(d/2)^{-G} \left( \frac{d \cdot \tau^2}{2} \right)^{Gd/2} \left( \prod_{g=1}^G \eta_g^2 \right)^{-(d/2+1)} \exp \left( -\frac{d \cdot \tau^2}{2} \sum_{g=1}^G \frac{1}{\eta_g^2} \right) I(0 < d < d_0)$$

- Using parallel reductions (NVIDIA's CUDA C/C++ Thrust library), calculate the sufficient statistics:

$$\prod_{g=1}^G \eta_g^2 \qquad \sum_{g=1}^G \frac{1}{\eta_g^2}$$

- Use a random-walk metropolis step to sample  $d$ .

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- ▶ Ordinary C and GPU-accelerated versions, along with an R package wrapper, are available for download at <https://github.com/wlandau/heterosis>.
- ▶ Time for a demo...

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# Thanks for coming.

- ▶ Slides and video will be available at  
<http://will-landau.com/research.html>.

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