

# Efficient Importance Sampling

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Our goal is to calculate integrals of the form

$$G(Y) = \int_{\Theta} \phi(\theta; Y) d\theta.$$

Special case (e.g., posterior moment):

$$G(Y) = \int_{\Theta} \phi(\theta; Y) p(\theta|Y) d\theta.$$

Scenario: analytical solutions to these integrals are unavailable. We will remedy this problem using **numerical approximation methods**.

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In the context of achieving likelihood evaluation and filtering in state space representations, recall that we face the challenge of constructing the filtering density

$$f(s_t|Y_t) = \frac{f(y_t, s_t|Y_{t-1})}{f(y_t|Y_{t-1})} = \frac{f(y_t|s_t, Y_{t-1}) f(s_t|Y_{t-1})}{f(y_t|Y_{t-1})},$$

where

$$f(s_t|Y_{t-1}) = \int f(s_t|s_{t-1}, Y_{t-1}) f(s_{t-1}|Y_{t-1}) ds_{t-1},$$

and

$$f(y_t|Y_{t-1}) = \int f(y_t|s_t, Y_{t-1}) f(s_t|Y_{t-1}) ds_t.$$

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To gain intuition, suppose the integral we face is of the form

$$G(Y) = \int_{\Theta} \phi(\theta; Y) p(\theta|Y) d\theta,$$

and it is possible to obtain pseudo-random drawings  $\theta_i$  from  $p(\theta|Y)$ . Then by the law of large numbers,

$$\overline{G(Y)}_N = \frac{1}{N} \sum_{i=1}^N \phi(\theta_i; Y)$$

converges in probability to  $G(Y)$ . We refer to  $\overline{G(Y)}_N$  as the **Monte Carlo** estimate of  $G(Y)$ . As we shall see, from the standpoint of **numerical efficiency**, this represents a **best-case scenario**.

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The MC estimate of the standard deviation of  $G(Y)$  is given by

$$\bar{\sigma}_N(G(Y)) = \left[ \left( \frac{1}{N} \sum_i^N \phi(\theta_i; Y)^2 \right) - \overline{G(Y)}_N^2 \right]^{1/2}.$$

The numerical standard error associated with  $\overline{G(Y)}_N$  is given by

$$s.e. \left( \overline{G(Y)}_N \right) = \frac{\bar{\sigma}_N(G(Y))}{\sqrt{N}}.$$

Thus for  $N = 10,000$ ,  $s.e. \left( \overline{G(Y)}_N \right)$  is 1% of the size of  $\bar{\sigma}_N(G(Y))$ .

# Importance Sampling (Geweke, 1989 Econometrica)

If  $p(\theta|Y)$  is unavailable as a sampler, one remedy is to augment the targeted integrand with an **importance sampling** distribution  $g(\theta|a)$ :

$$\begin{aligned} G(Y) &= \int_{\Theta} \frac{\varphi(\theta; Y)}{g(\theta|a)} g(\theta|a) d\theta \\ &= \int_{\Theta} \frac{\phi(\theta; Y) p(\theta|Y)}{g(\theta|a)} g(\theta|a) d\theta. \end{aligned}$$

## Key requirements:

- ▶ Support of  $g(\theta|a)$  must span that of  $\varphi(\theta|Y)$
- ▶  $E[G(Y)]$  must exist and be finite.
- ▶  $g(\theta|a)$  must be implementable as a sampler

# Importance Sampling, cont.

- ▶ MC estimate of  $G(Y)$ :

$$\overline{G(Y)}_N = \frac{1}{N} \sum_{i=1}^N \omega_i ; \quad \omega_i = \frac{\varphi(\theta_i|Y)}{g(\theta_i|a)}$$

- ▶ MC estimate of the standard deviation of  $\omega$  w.r.t.  $g(\theta|a)$ :

$$\bar{\sigma}_N(\omega(\theta, Y)) = \left[ \left( \frac{1}{N} \sum \omega_i^2 \right) - \overline{G(Y)}_N^2 \right]^{1/2}.$$

- ▶ Numerical standard error associated with  $\overline{G(\theta, Y)}_N$ :

$$s.e. \left( \overline{G(Y)}_N \right)_I = \frac{\bar{\sigma}_N(\omega(\theta, Y))}{\sqrt{N}}.$$

- ▶ **Note:** Variability in  $\omega$  translates into increased n.s.e. (numerical inefficiency)

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For the special case in which the integrand factorizes as

$$\varphi(\theta_i|Y) = \phi(\theta; Y) p(\theta|Y),$$

$$\overline{G(Y)}_N = \frac{1}{N} \sum_{i=1}^N \phi(\theta_i; Y) w_i, \quad w_i = \frac{p(\theta_i|Y)}{g(\theta_i|a)}$$

$$\bar{\sigma}_N(G(Y)) = \left[ \left( \frac{1}{N} \sum_{i=1}^N \phi(\theta_i; Y)^2 w_i \right) - \overline{G(Y)}_N^2 \right]^{1/2}.$$



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Continuing with the special case, if we lack an integrating constant for either  $p(\theta|Y)$  or  $g(\theta|a)$ , we can work instead with

$$\overline{G(Y)}_N = \left( \frac{1}{\sum w_i} \right) \sum_{i=1}^N \phi(\theta_i; Y) w_i$$

$$\bar{\sigma}_N(G(Y)) = \left[ \left( \left( \frac{1}{\sum w_i} \right) \sum_{i=1}^N \phi(\theta_i; Y)^2 w_i \right) - \overline{G(Y)}_N^2 \right]^{1/2}.$$

The impact of ignoring integrating constants is eliminated by the inclusion of the accumulated weights in the denominators of these expressions.

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Regardless of whether  $p(\theta|Y)$  and  $g(\theta|a)$  are proper p.d.f.s, notice that when  $p(\theta|Y)$  itself can be used as a sampling density (i.e.,  $p(\theta|Y) = g(\theta|a)$ ), then  $w_i = 1 \forall i$ , and we revert to the best-case scenario.

In general, the accuracy of our approximation will fall short of the best-case scenario. To judge the degree of the shortfall, various metrics are available.

## Two metrics for judging numerical accuracy:

- ▶  $\frac{w_{\max}^2}{\sum w_i^2}$  (good practical benchmark: 1%)
- ▶ relative numerical efficiency (**RNE**)

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**Motivation for RNE.** Issue: how close are we to the best-case scenario?

- Under the best-case scenario, recall that n.s.e. is given by

$$\text{s.e.} \left( \overline{G(Y)}_N \right) = \frac{\bar{\sigma}_N(G(Y))}{\sqrt{N}}.$$

- Actual n.s.e. is given by

$$\text{s.e.} \left( \overline{G(Y)}_N \right)_I = \frac{\bar{\sigma}_N(\omega(\theta, Y))}{\sqrt{N}},$$

where recall

$$\bar{\sigma}_N(\omega(\theta, Y)) = \left[ \left( \frac{1}{N} \sum \omega_i^2 \right) - \overline{G(Y)}_N^2 \right]^{1/2}, \quad \omega_i = \frac{\varphi(\theta_i|Y)}{g(\theta_i|a)}$$

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The idea behind RNE is to compare the actual n.s.e. to an estimate of the optimal (best-case) n.s.e.:

$$\begin{aligned} RNE &= \frac{(\text{ideal n.s.e.})^2}{(\text{actual n.s.e.})^2} \\ &= \frac{\left( \frac{\bar{\sigma}_N(G(Y))}{\sqrt{N}} \right)^2}{\left( \text{s.e.} \left( \overline{G(Y)}_N \right)_I \right)^2}. \end{aligned}$$

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Rearranging yields

$$s.e. \left( \overline{G(Y)}_N \right)_I = \frac{\bar{\sigma}_N(G(Y))}{\sqrt{N \cdot RNE}}.$$

Note: relative to the best-case scenario,  $\sqrt{N \cdot RNE}$  replaces  $\sqrt{N}$  in the denominator. Thus the further is RNE from 1, the more draws are required to achieve a given level of accuracy relative to the best-case scenario.

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Suppose  $p(\theta|Y) \sim N(\mu, \Sigma)$ ,

$$\mu = \begin{bmatrix} 1 \\ 2 \\ 3 \\ 4 \\ 5 \end{bmatrix}, \quad \text{corr}(\Sigma) = \begin{bmatrix} 1 & 0.6 & 0 & 0 & 0 \\ 0.6 & 1 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 1 & -0.8 \\ 0 & 0 & 0 & -0.8 & 1 \end{bmatrix},$$

$$\text{sqrt}(\text{diag}(\Sigma))' = [2 \quad 0.2 \quad 5 \quad 1 \quad 0.1]$$

Statistics of interest:

$$E(\text{sumc}(\mu)), \quad E(\text{prodc}(\mu))$$

## Example, cont.

Using `rndseed 123456789`, and  $N = 10,000$ , MC estimates  $(\hat{\mu}, \widehat{\text{sqrt}(\text{diag}(\Sigma))}, \text{n.s.e.}(\hat{\mu}))$ :

1.0077	1.9664	0.0197
2.0031	0.2007	0.0020
2.9157	4.9899	0.0499
4.0041	0.9793	0.0098
4.9993	0.0986	0.0010

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MC estimate of  $\text{corr}(\Sigma)$  :

1.0000	0.6060	0.0044	-0.0139	0.0107
0.6060	1.0000	0.0049	-0.0048	0.0016
0.0044	0.0049	1.0000	-0.0013	-0.0110
-0.0139	-0.0048	-0.0013	1.0000	-0.7969
0.0107	0.0016	-0.0110	-0.7969	1.0000

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MC estimates of statistics (mean, std. dev., n.s.e.(mean)):

$E(\text{sumc}(\mu))$	14.9299	5.4883	0.0549
$E(\text{prodc}(\mu))$	130.7018	533.3792	5.3338

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### Exercise: replicate

#### Hint for exercise.

To obtain draws from  $N(\mu, \Sigma)$  distribution:

- ▶ `swish = chol(sig)'`; `swish` is lower-diagonal Cholesky decomposition of `sig` (i.e., `sig=swish*swish'`).
- ▶ `draw = mu + swish*rndn(n,1);`

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Suppose instead we seek to obtain estimates using an Importance Sampling density

$$g(\theta|a) \sim N(\mu_I, \Sigma_I),$$

with

$$\begin{aligned}\mu_I &= \mu + 1.5 \cdot \text{sqrt}(\text{diag}(\Sigma)), \\ \Sigma_I &= \text{diagrv}(\Sigma, \text{sqrt}(\text{diag}(\Sigma)))\end{aligned}$$

## Example, cont.

Using `rndseed 123456789`, and  $N = 10,000$ , MC versus IS estimates  $(\hat{\mu}, \sqrt{\widehat{\text{diag}}(\Sigma)})$ , n.s.e. $(\hat{\mu})$ :

1.0077	1.9664	0.0197
2.0031	0.2007	0.0020
2.9157	4.9899	0.0499
4.0041	0.9793	0.0098
4.9993	0.0986	0.0010
1.1646	1.5813	0.4140
1.9884	0.1721	0.0572
3.1161	3.5692	0.6526
4.2545	0.8731	0.2691
4.9947	0.0803	0.0173

## Example, cont.

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MC versus IS estimate of  $\text{corr}(\Sigma)$  :

1.0000	0.6060	0.0044	-0.0139	0.0107
0.6060	1.0000	0.0049	-0.0048	0.0016
0.0044	0.0049	1.0000	-0.0013	-0.0110
-0.0139	-0.0048	-0.0013	1.0000	-0.7969
0.0107	0.0016	-0.0110	-0.7969	1.0000
1.0000	0.6712	0.0623	-0.2506	-0.2307
0.6712	1.0000	0.2031	-0.2387	0.0065
0.0623	0.2031	1.0000	-0.1987	0.2975
-0.2506	-0.2387	-0.1987	1.0000	-0.6440
-0.2307	0.0065	0.2975	-0.6440	1.0000

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MC versus estimates of statistics (mean, std. dev.,  
n.s.e.(mean)):

$E(\text{sumc}(\mu))$	14.9299	5.4883	0.0549
$E(\text{prodc}(\mu))$	130.7018	533.3792	5.3338
$E(\text{sumc}(\mu))$	15.5184	3.9212	0.8116
$E(\text{prodc}(\mu))$	168.4666	471.0707	64.8759

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### Accuracy Diagnostics

Summary statistics on weights:

avg,	stdev,	min,	max,	maxsq/totsq:
1.0120	36.3643	0.0000	3341.8561	0.8440

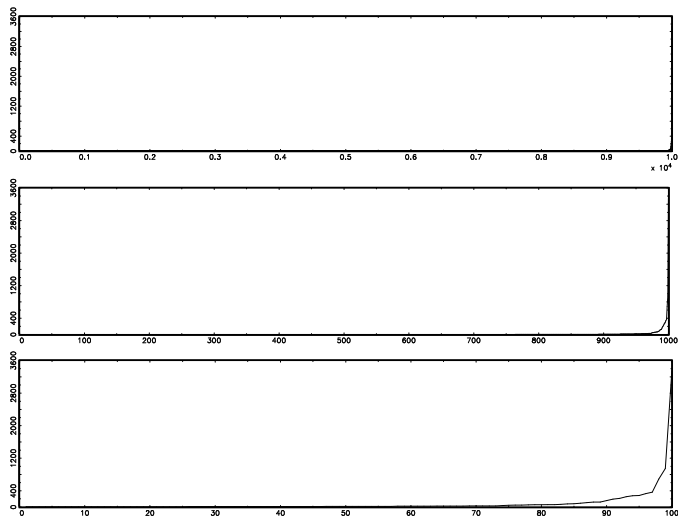
RNEs and 1/RNEs:

0.0023	428.4143
0.0053	189.6681



## Example, cont.

Plot of weights (all, top 1,000, top 100):



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**Exercise: Replicate**

From a programming perspective, two simple approaches to improving efficiency are:

- ▶ Increase  $N$  (RNEs indicate good rules of thumb for necessary increases). This brute-force method is often computationally prohibitive.
- ▶ **Sequential updating.** (Can still be expensive, but less brutish.)

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Sequential updating:

- ▶ Begin with an initial parameterization  $a_0$  for  $g(\theta|a)$  (e.g.,  $(\mu_0, \Sigma_0)$ ).
- ▶ Calculate  $\hat{\theta}_0$ , map into  $a_1$ .
- ▶ Repeat until  $a_i$  yields an acceptable level of numerical accuracy.

## Improving Efficiency, cont.

Returning to the example, RNEs and  $1/\text{RNEs}$  evolve as follows:

Iteration 0:

0.0023	428.4143
0.0053	189.6681

Iteration 1:

0.0575	17.4021
0.0331	30.1702

Iteration 2:

0.8018	1.2472
0.7302	1.3694

Iteration 3:

0.9790	1.0215
0.9535	1.0488

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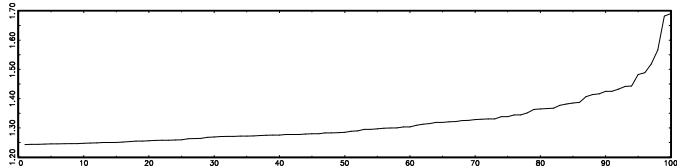
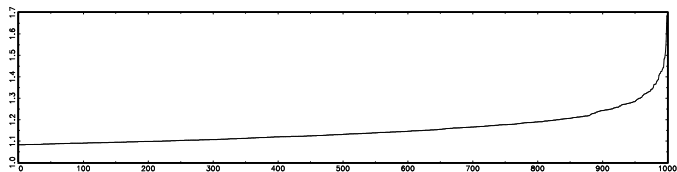
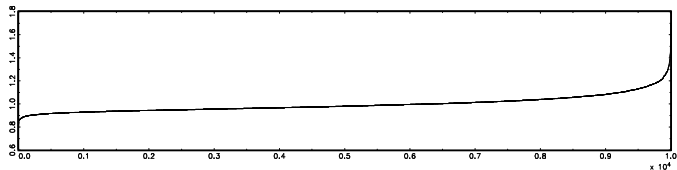
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Weight plots, Iteration 3:



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## Caveats regarding sequential updating:

- ▶ Convergence to target is not guaranteed
- ▶ Performance can be sensitive to starting values
- ▶ Initial sampler should be sufficiently diffuse to ensure coverage of appropriate range for targeted integrand

## Aside regarding coverage:

The multivariate-t density is an attractive sampler relative to the normal density: it has similar location and shape parameters, but has **thicker tails**. Tail thickness controlled by degrees-of-freedom parameter  $\nu$  (smaller  $\nu$ , thicker tails).

- ▶ Parameters of the multivariate-t:  $(\gamma, V, \nu)$ .
- ▶ Mean and second-order moments:  $\gamma, \left(\frac{\nu}{\nu-2}\right) V^{-1}$



## Improving Efficiency, cont.

Algorithm for obtaining drawings  $\mu_i$  from multivariate-t  $(\gamma, V, v)$  :

- ▶ Obtain  $s_i$  from a  $\chi^2(v)$  distribution:

$$s_i = \sum_{j=1}^v x_j^2, \quad x_j \sim N(0, 1)$$

- ▶ Use  $s_i$  to construct the scaling factor

$$\sigma_i = (s_i/v)^{-1/2}$$

- ▶ Obtain  $\mu_i$  as

$$\begin{aligned} \mu_i &= \gamma \pm \sigma_i V^{-1/2} w_i, & w_{i,j} &\sim N(0, 1), & j &= 1, \dots, k \\ V^{-1/2} &= \text{chol}(V^{-1})' \end{aligned}$$

- ▶ Note: use of  $\pm$  yields *antithetic acceleration* (Geweke, 1988, *JoE*)

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**Exercise:** Replace the normal densities used as Importance Samplers in the exercise above with multi-t densities, with  $\gamma = \mu$  and  $V^{-1} = \Sigma$ . Experiment with alternative  $v$ 's to assess the impact on numerical efficiency.

**Goal:** Tailor  $g(\theta|a)$  (via the specification of  $a$ ) to minimize the n.s.e. associated with the approximation of

$$G(Y) = \int_{\Theta} \varphi(\theta|Y) d\theta.$$

Write  $g(\theta|a)$  as

$$\begin{aligned} g(\theta|a) &= \frac{k(\theta; a)}{\chi(a)}, \\ \chi(a) &= \int_{\Theta} k(\theta; a) d\theta. \end{aligned}$$

Details regarding the tailoring of  $g(\theta|a)$  are distinct for two special cases:

- ▶  $g(\theta|a)$  is parametric (i.e., a normal distribution)
- ▶  $g(\theta|a)$  is piecewise-linear

When  $g(\theta|a)$  is fully parametric, n.s.e. is (approximately) minimized via iterations on

$$\begin{aligned}(\hat{a}_{l+1}, \hat{c}_{l+1}) &= \arg \min_{a, c} \overline{Q}_N(a, c; Y|\hat{a}_l), \\ \overline{Q}_N(a, c; Y|\hat{a}_l) &= \frac{1}{N} \sum_{i=1}^N d^2(\theta_i^l, a, c, Y) \omega(\theta_i^l; Y, \hat{a}_l), \\ d(\theta_i^l, a, c, Y) &= \ln \varphi(\theta_i^l | Y) - c - \ln k(\theta_i^l; a), \\ \omega(\theta_i^l; Y, \hat{a}_l) &= \frac{\varphi(\theta_i | Y)}{g(\theta_i | \hat{a}_l)}.\end{aligned}$$

The term  $c$  is a normalizing constant that controls for factors in  $\varphi$  and  $g$  that do not depend upon  $\theta$ . Typically, it suffices to set  $\omega(\theta_i^l; Y, \hat{a}_l) = 1 \forall i$ .

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When  $g(\theta|a)$  is piecewise-linear, the parameters  $a$  are grid points:

$$a' = (a_0, \dots, a_R), \quad a_0 < a_1 < \dots < a_R.$$

In this case, the kernel  $k(\theta; a)$  is given by

$$\begin{aligned} \ln k_j(\theta; a) &= \alpha_j + \beta_j \theta \quad \forall \theta \in [a_{j-1}, a_j], \\ \beta_j &= \frac{\ln \varphi(a_j) - \ln \varphi(a_{j-1})}{a_j - a_{j-1}}, \quad \alpha_j = \ln \varphi(a_j) - \beta_j a_j. \end{aligned}$$

Optimization is achieved by selecting  $\hat{a}$  as an equal-probability division of the support of  $\varphi(\theta|Y)$ .

Given final estimates  $(\hat{a}, \hat{c})$ , the EIS estimate of  $G(Y)$  is given by

$$\overline{G(Y)}_N = \frac{1}{N} \sum_{i=1}^N \omega(\theta_i; Y, \hat{a}).$$

N.S.E. is computed as indicated above.

# Implementation, Gaussian Sampler

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To simplify notation, denote the targeted integrand as  $\varphi(\theta|Y) \equiv \varphi(\theta)$ . We'll take  $\theta$  as  $k$ -dimensional, with elements  $(x_1, x_2, \dots, x_k)$ .

With  $g(\theta|a)$  Gaussian,  $a$  consists of the  $k \times 1$  vector of means  $\mu$  and the  $k \times k$  covariance matrix  $\Sigma$ . Since the covariance matrix is symmetric, the number of auxiliary parameters reduces to  $k + k(k+1)/2$ .

The precision matrix  $H = \Sigma^{-1}$ .



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Our goal is to choose  $(\mu, H)$  to approximate optimally  $\ln \varphi(s)$  by a Gaussian kernel:

$$\begin{aligned}\ln \varphi(\theta) &\propto -\frac{1}{2}(\theta - \mu)' H (\theta - \mu) \\ &\propto -\frac{1}{2} (\theta' H \theta - 2\theta' H \mu) .\end{aligned}$$

Recall that by ‘optimally’, we refer to the weighted-squared-error minimization introduced above.

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The term  $\theta' H \theta$  can be written as

$$\begin{pmatrix} x_1 & x_2 & . & . & x_k \end{pmatrix} \begin{pmatrix} h_{11} & h_{21} & . & . & h_{1k} \\ h_{21} & h_{22} & . & . & h_{2k} \\ . & . & . & . & . \\ . & . & . & . & . \\ h_{k1} & h_{k2} & . & . & h_{kk} \end{pmatrix} \begin{pmatrix} x_1 \\ x_2 \\ . \\ . \\ x_k \end{pmatrix} .$$

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Expanding  $\theta' H \theta$ , we obtain

$$\begin{aligned}\theta' H \theta &= h_{11} (x_1^2) + h_{22} (x_2^2) + \dots + h_{jj} (x_{kj}^2) \\ &\quad + 2h_{21} (x_2 x_1) + 2h_{31} (x_3 x_1) + \dots + 2h_{k1} (x_k x_1) \\ &\quad + 2h_{32} (x_3 x_2) + 2h_{42} (x_4 x_2) + \dots + 2h_{k2} (x_k x_2) \\ &\quad \dots \\ &\quad + 2h_{k(k-1)} (x_k x_{k-1}) .\end{aligned}$$

This expression indicates that the coefficients of the squares, pairwise products and the individual components of  $\theta$  are in one-to-one correspondence with the  $\mu$  and  $H$ .

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Given this correspondence, the EIS optimization problem amounts to a weighted-least-squares problem involving the regression of  $\ln \varphi(s)$  on

$$\left[ 1, x_1, \dots, x_k, x_1 x_2, x_1 x_3, \dots, x_{k-1} x_k, x_1^2, x_2^2, \dots, x_k^2 \right].$$

The number of regressors is  $K = \left( 1 + k + \frac{k(k+1)}{2} \right)$ .

# Implementation, Gaussian Sampler, cont.

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## Implementation Algorithm

- Specify  $a_0$ , generate

$$y = \begin{pmatrix} \ln \varphi(\theta_1) \\ \dots \\ \ln \varphi(\theta_M) \end{pmatrix}, \quad w = \begin{pmatrix} \omega_1 \\ \dots \\ \omega_M \end{pmatrix}, \quad \frac{\varphi(\theta_i|Y)}{g(\theta_i|a_0)},$$

$$X = \begin{pmatrix} \kappa_1 \\ \dots \\ \kappa_M \end{pmatrix},$$

$$\kappa_i = [1 \sim \theta_i' \sim \text{vech}(\theta_i \cdot \theta_i')]'$$

(Note:  $M \ll N$ )

## Implementation Algorithm, cont.

- Construct

$$\tilde{y} = y \cdot (w.^2), \quad \tilde{X} = X \cdot (w.^2).$$

(Caution: set  $w = 1$  when using a poor initial sampling density.)

- Estimate

$$\hat{\beta} = (\tilde{X}'\tilde{X})^{-1} (\tilde{X}'\tilde{y}).$$

- Map  $\hat{\beta}$  into  $(\hat{\mu}, \hat{\Sigma})$ . Jointly, these constitute  $a_1$ .
- Replacing  $a_0$  above with  $a_1$ , repeat until convergence.

# Implementation, Gaussian Sampler, cont.

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## Implementation Algorithm, cont.

To map  $\hat{\beta}$  into  $(\hat{\mu}, \hat{\Sigma})$  :

- ▶ Map the  $k + 2$  through  $K$  elements of  $\hat{\beta}$  into a symmetric matrix  $\tilde{H}$ , with  $j^{th}$  diagonal element corresponding to the coefficient associated with the squared value of the  $j^{th}$  element of  $\theta$ , and  $(j, k)^{th}$  element corresponding to the product of the  $j$  and  $k^{th}$  element of  $\theta$ . I.E.,

$$\tilde{H} = \text{xpnd}(\hat{\beta}[k+2:\text{rows}(\text{beta})]);$$

- ▶ Construct  $\tilde{\tilde{H}}$  by multiplying all elements of  $\tilde{H}$  by  $-1$ , then multiplying the diagonal elements by 2.
- ▶  $\hat{\Sigma} = \tilde{\tilde{H}}^{-1}$
- ▶  $\hat{\mu} = \hat{\Sigma} \cdot \hat{\beta}[2 : k + 1]$

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# Implementation, Gaussian Sampler, cont.

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**Exercise:** Return to the example outlined above. Using an initial sampler specified with

$$\begin{aligned}\mu_0 &= \mu + 3 \cdot \text{sqrt}(\text{diag}(\Sigma)), \\ \Sigma_0 &= 10 \cdot \text{diagrv}(\Sigma, \text{sqrt}(\text{diag}(\Sigma))),\end{aligned}$$

show that the EIS algorithm yields

$$\hat{\mu} = \mu, \quad \hat{\Sigma} = \Sigma$$

in one step using  $M = 50$ ,  $w = 1$ . Experiment with alternative initial samplers.



# Implementation, Piecewise-Linear Approximation

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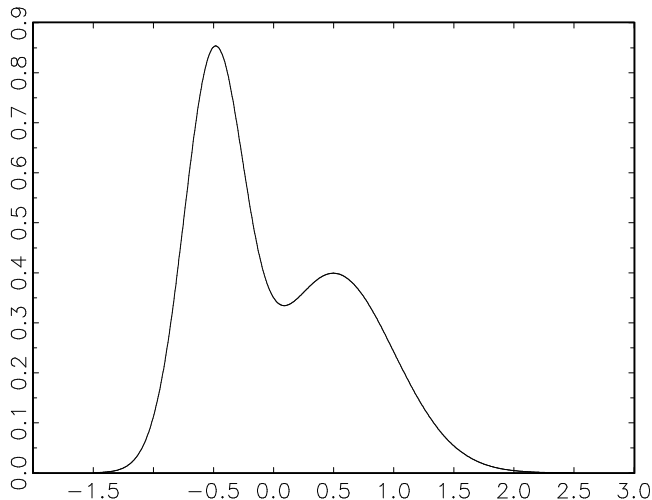
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**Context:** Effective for use in univariate cases featuring obvious deviations from normality.  
(Curse of dimensionality renders implementation problematic in high-dimensional cases.)

## Implementation, PW-L Approx., cont.

Motivation 1. Mixture of Normals,  $(\mu, \sigma) = (-0.5, 0.25)$ ,  $(0.5, 0.5)$ , equal weights



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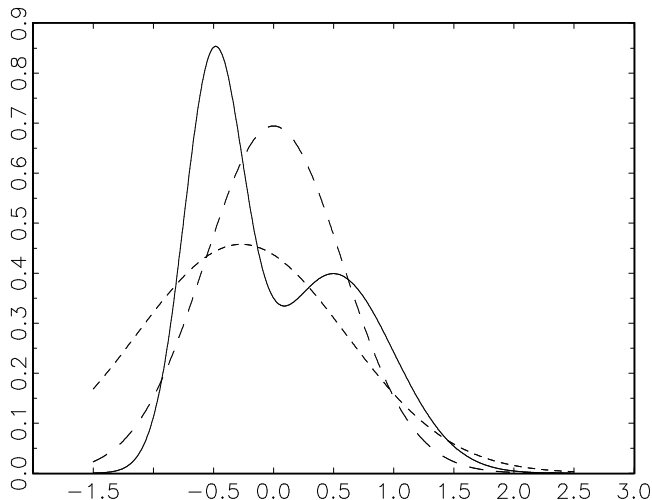
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# Implementation, PW-L Approx., cont.

Approximation via Gaussian samplers (handmade and EIS):



RNEs for calculating  $E(x)$ : 0.61, 0.78.

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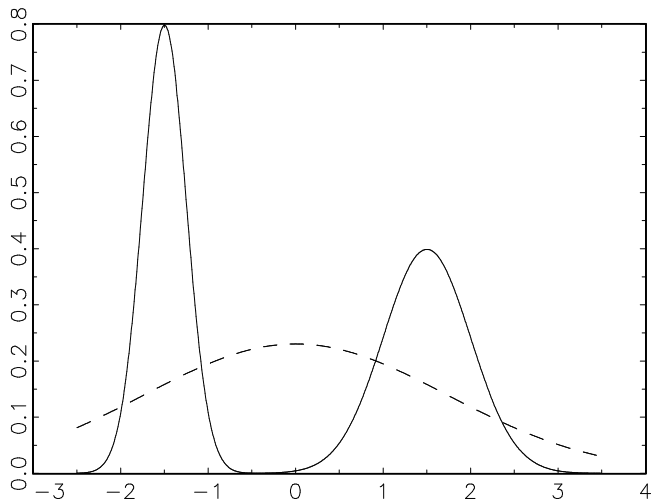
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## Implementation, PW-L Approx., cont.

Motivation 2.  $(\mu, \sigma) = (-1.5, 0.25), (1.5, 0.5)$ , equal weights



RNE: 0.37

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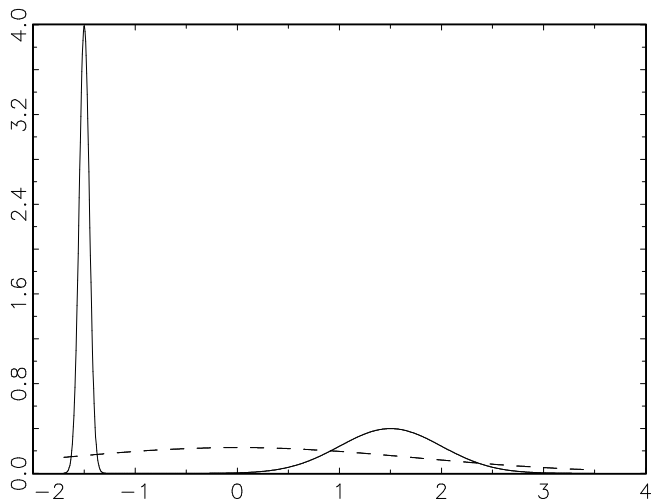
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## Implementation, PW-L Approx., cont.

Motivation 3.  $(\mu, \sigma) = (-1.5, 0.05), (1.5, 0.5)$ , equal weights



RNE: 0.11

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# Implementation, PW-L Approx., cont.

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Reboot: Recall that when  $g(\theta|a)$  is piecewise-linear, the parameters  $a$  are grid points:

$$a' = (a_0, \dots, a_R), \quad a_0 < a_1 < \dots < a_R.$$

In this case, the kernel  $k(\theta; a)$  is given by

$$\begin{aligned} \ln k_j(\theta; a) &= \alpha_j + \beta_j \theta \quad \forall \theta \in [a_{j-1}, a_j], \\ \beta_j &= \frac{\ln \varphi(a_j) - \ln \varphi(a_{j-1})}{a_j - a_{j-1}}, \quad \alpha_j = \ln \varphi(a_j) - \beta_j a_j. \end{aligned}$$

Optimization is achieved by selecting  $\hat{a}$  as an equal-probability division of the support of  $\varphi(\theta|Y)$ .

# Implementation, PW-L Approx., cont.

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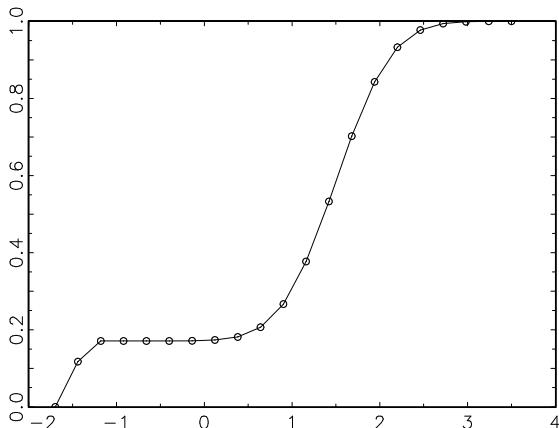
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In order to achieve equal-probability division, and to implement the distribution as a sampler, we must

- ▶ construct its associated CDF
- ▶ invert the CDF.

## Implementation, PW-L Approx., cont.

To gain intuition behind implementation and inversion, consider the CDF associated with Case 3:



Inversion involves inducing a mapping from the  $y$  to the  $x$  axis. Implementation involves mapping drawings obtained from a  $U[0, 1]$  distribution onto the  $x$  axis.



# Implementation, PW-L Approx., cont.

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Letting  $s$  denote the  $x$ -axis variable, the **CDF** of  $k$  can be written as

$$\begin{aligned}K_j(s; a) &= \frac{\chi_j(s; a)}{\chi_n(a)}, \quad \forall s \in [a_{j-1}, a_j], \\ \chi_j(s; a) &= \chi_{j-1}(a) + \frac{1}{\beta_j} [k_j(s; a) - k_j(a_{j-1}; a)], \\ \chi_0(a) &= 0, \quad \chi_j(a) = \chi_j(a_j; a).\end{aligned}$$

Note:  $\chi_n(a)$  is the integrating constant associated with the pdf.

# Implementation, PW-L Approx., cont.

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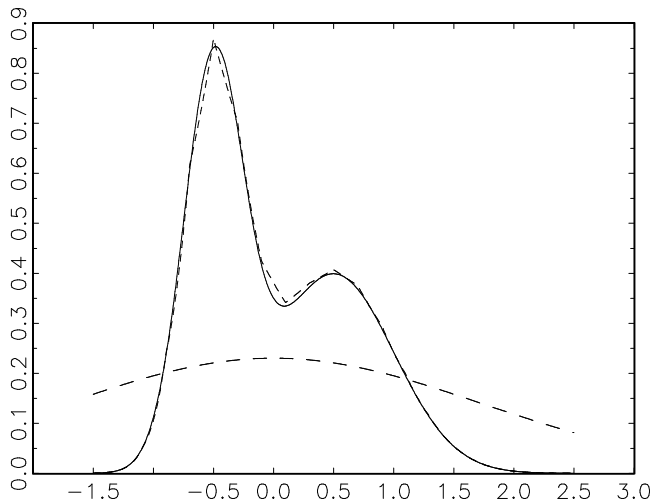
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Inversion/implementation:

$$s = \frac{1}{\beta_j} \left\{ \ln \left[ k_j(a_{j-1}; a) + \beta_j (u \cdot \chi_R(a) - \chi_{j-1}(a)) \right] - \alpha_j \right\},$$
$$u \in ]0, 1[ \quad \text{and} \quad \chi_{j-1}(a) < u \cdot \chi_R(a) < \chi_j(a).$$

## Implementation, PW-L Approx., cont.

Unrefined (equal spacing over x range) piecewise approximation for case 1:



RNE: 0.99

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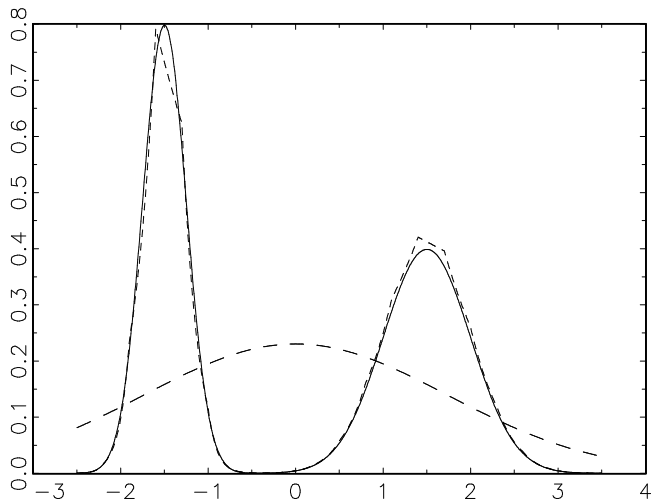
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# Implementation, PW-L Approx., cont.

Unrefined (equal spacing over x range) piecewise approximation for case 2:



RNE: 0.99

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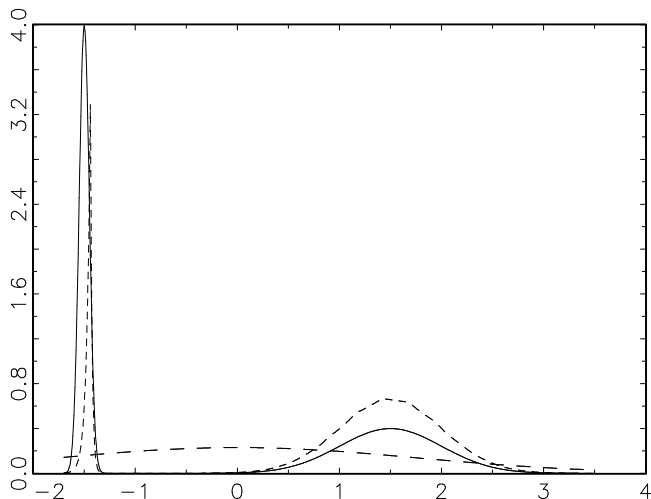
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## Implementation, PW-L Approx., cont.

Unrefined (equal spacing over x range) piecewise approximation for case 3:



RNE: 0.24

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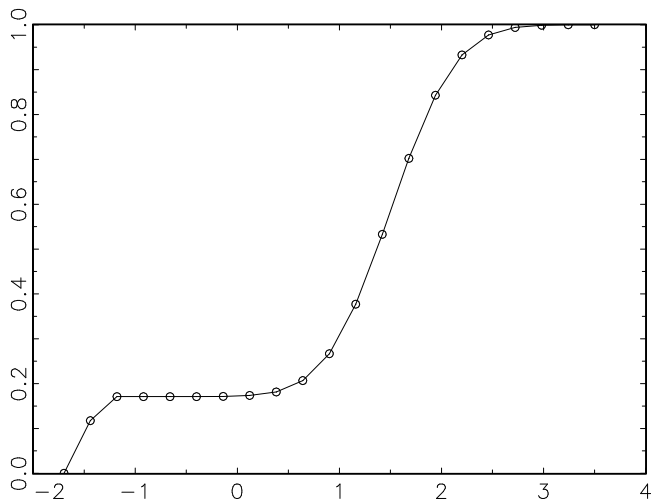
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## Implementation, PW-L Approx., cont.

To understand the source of the problem in Case 3, consider again the CDF of the unrefined sampler:



Note: relatively few points along steep portions of the CDF,

# Implementation, PW-L Approx., cont.

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To address the problem: **equal probability division** of the range for  $s$ . I.e., divide the vertical axis of the CDF into equal portions, then map into  $s$  :

$$u_i = \varepsilon + (2 - \varepsilon) \frac{i}{R}, \quad i = 1, \dots, R - 1,$$

with  $\varepsilon$  sufficiently small (typically  $\varepsilon = 10^{-4}$ ) to avoid tail intervals of excessive length.

## Iterative construction:

Given the step- $l$  grid  $\hat{a}^l$ , construct the density kernel  $k$  and its CDF  $K$  as described above. The step- $l + 1$  grid is then computed as

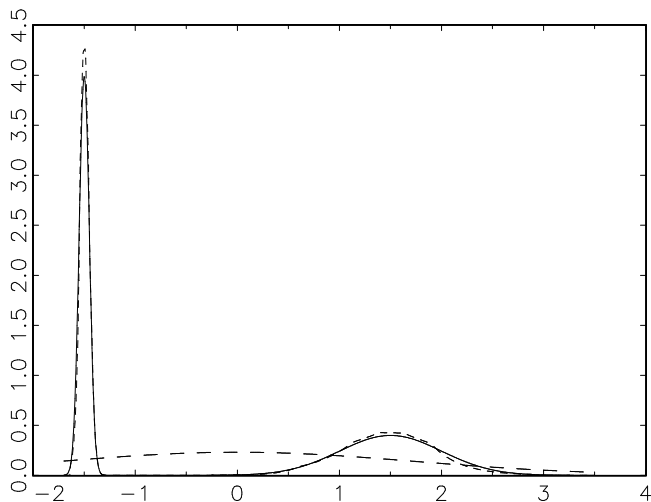
$$\hat{a}_i^{l+1} = K^{-1}(u_i), \quad i = 1, \dots, R - 1.$$

Iterate until (approximate) convergence.



# Implementation, PW-L Approx., cont.

Refined approximation for Case 3:



RNE: 0.95

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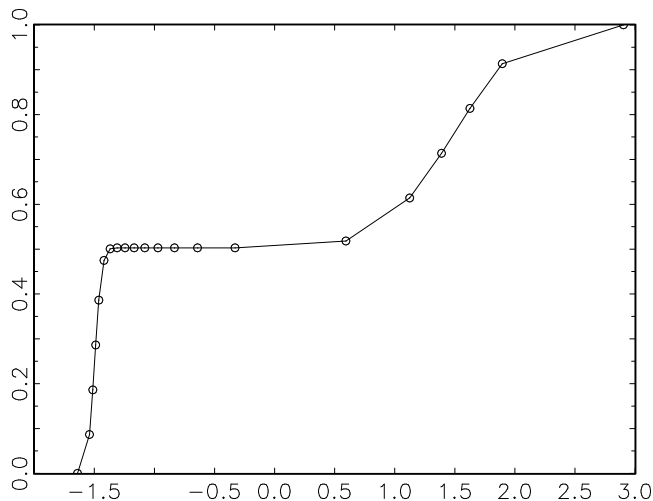
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Refined CDF:



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