Advanced Simulation - Lecture 3

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Rejection Sampling

Recall

Algorithm (Rejection Sampling). Given two densities π , q with $\pi(x) \le M q(x)$ for all x, we can generate a sample from π by

- **1** Draw $X \sim q$, draw $U \sim \mathcal{U}_{[0,1]}$.
- 2 Accept X = x as a sample from π if

$$U \leq \frac{\pi(x)}{M q(x)},$$

otherwise go to step 1.

Proposition

The distribution of the samples accepted by rejection sampling is π .

Rejection Sampling

• Often we only know π and q up to some normalising constants; i.e.

$$\pi = \widetilde{\pi}/Z_{\pi}$$
 and $q = \widetilde{q}/Z_{q}$

where $\widetilde{\pi}$, \widetilde{q} are known but Z_{π} , Z_q are unknown. You still need to be able to sample from $q(\cdot)$.

■ If you can upper bound:

$$\widetilde{\pi}(x)/\widetilde{q}(x) \leq \widetilde{M},$$

then using $\widetilde{\pi}$, \widetilde{q} and \widetilde{M} in the algorithm is correct.

■ Indeed we have

$$\frac{\widetilde{\pi}\left(x\right)}{\widetilde{q}\left(x\right)} \leq \widetilde{M} \Leftrightarrow \frac{\pi\left(x\right)}{q\left(x\right)} \leq \widetilde{M} \frac{Z_{q}}{Z_{\pi}} = M.$$

Rejection Sampling

Let T denote the number of pairs (X, U) that have to be generated until X is accepted for the first time.

Lemma

T is geometrically distributed with parameter 1/M and in particular $\mathbb{E}(T) = M$.

In the unnormalised case, this yields

$$\mathbb{P}\left(X \text{ accepted}\right) = \frac{1}{M} = \frac{Z_{\pi}}{\widetilde{M}Z_{q}},$$

$$\mathbb{E}\left(T\right)=M=\frac{Z_{q}\widetilde{M}}{Z_{\pi}},$$

and it can be used to provide unbiased estimates of Z_{π}/Z_{q} and Z_{q}/Z_{π} .

Examples:Uniform from bounded subset of \mathbb{R}^p

■ Let $B \subset \mathbb{R}^p$, a bounded subset of \mathbb{R}^p :

$$\pi(x) \propto \mathbb{I}_B(x)$$
.

Let *R* be a rectangle containing $B \subset R$ and

$$q(x) \propto \mathbb{I}_R(x)$$
.

■ Then we can use $\widetilde{M} = 1$ and

$$\widetilde{\pi}(x) / (\widetilde{M}'\widetilde{q}(x)) = \mathbb{I}_B(x).$$

■ The probability of accepting a sample is then Z_{π}/Z_{q} .

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Example: Normal density

■ Let $\widetilde{\pi}(x) = \exp\left(-\frac{1}{2}x^2\right)$ and $\widetilde{q}(x) = 1/(1+x^2)$. We have

$$\frac{\widetilde{\pi}\left(x\right)}{\widetilde{q}\left(x\right)} = \left(1 + x^{2}\right) \exp\left(-\frac{1}{2}x^{2}\right) \le 2/\sqrt{e} = \widetilde{M}$$

which is attained at ± 1 .

■ Let $X \sim \tilde{q}$. The acceptance probability is

$$\mathbb{P}\left(U \leq \frac{\widetilde{\pi}\left(X\right)}{\widetilde{M}\widetilde{q}\left(X\right)}\right) = \frac{Z_{\pi}}{\widetilde{M}Z_{q}} = \frac{\sqrt{2\pi}}{\frac{2}{\sqrt{e}}\pi} = \sqrt{\frac{e}{2\pi}} \approx 0.66,$$

and the mean number of trials to success is approximately $1/0.66 \approx 1.52$.

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Examples: Genetic Linkage model

■ We observe

$$(Y_1, Y_2, Y_3, Y_4) \sim \mathcal{M}\left(n; \frac{1}{2} + \frac{\theta}{4}, \frac{1}{4}(1-\theta), \frac{1}{4}(1-\theta), \frac{\theta}{4}\right)$$

where \mathcal{M} is the multinomial distribution and $\theta \in (0,1)$.

■ The likelihood of the observations is thus

$$\begin{split} & p\left(y_{1},...,y_{4};\theta\right) \\ & = \frac{n!}{y_{1}!y_{2}!y_{3}!y_{4}!} \left(\frac{1}{2} + \frac{\theta}{4}\right)^{y_{1}} \left(\frac{1}{4}\left(1 - \theta\right)\right)^{y_{2} + y_{3}} \left(\frac{\theta}{4}\right)^{y_{4}} \\ & \propto (2 + \theta)^{y_{1}} \left(1 - \theta\right)^{y_{2} + y_{3}} \theta^{y_{4}}. \end{split}$$

■ Bayesian approach where we select $p\left(\theta\right) = \mathbb{I}_{\left[0,1\right]}\left(\theta\right)$ and are interested in

$$p(\theta|y_1,...,y_4) \propto (2+\theta)^{y_1} (1-\theta)^{y_2+y_3} \theta^{y_4} \mathbb{I}_{[0,1]}(\theta).$$

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Examples: Genetic linkage model

- Rejection sampling using the prior as proposal $q(\theta) = \widetilde{q}(\theta) = p(\theta)$ to sample from $p(\theta|y_1, ..., y_4)$.
- To use accept-reject, we need to upper bound

$$\frac{\widetilde{\pi}\left(\theta\right)}{\widetilde{q}\left(\theta\right)} = \widetilde{\pi}\left(\theta\right) = (2+\theta)^{y_1} \left(1-\theta\right)^{y_2+y_3} \theta^{y_4}$$

- Maximum of $\widetilde{\pi}$ can be found using standard optimization procedure to perform rejection sampling.
- For a realisation of (Y_1, Y_2, Y_3, Y_4) equal to (69, 9, 11, 11) obtained with n = 100 and $\theta^* = 0.6$, results shown in following figure.

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Examples: Genetic linkage model

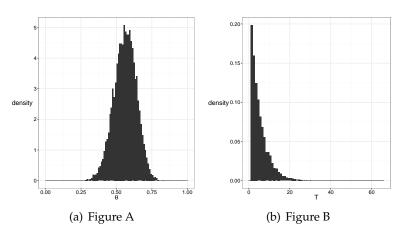


Figure: Histogram of 10,000 samples drawn from posterior obtained by rejection sampling (left); and histogram of waiting time distribution before acceptance (right).

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Rejection Sampling Recap

Rejection sampling requires

1 Samples from some distribution *q*;

2 evaluation of $\pi(\cdot)$ point-wise, or unnormalized $\widetilde{\pi}$;

3 an upper bound M on $\pi(x)/q(x)$, or $\tilde{\pi}/q$ and so on.

Sometimes the upper bound is not feasible.

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Importance Sampling

We want to compute

$$I = \mathbb{E}_{\pi}(\varphi(X)) = \int_{\mathbb{X}} \varphi(x) \, \pi(x) \, dx.$$

- We do not know how to sample from the target π but have access to a proposal distribution of density q.
- We only require that

$$\pi(x) > 0 \Rightarrow q(x) > 0$$
;

i.e. the support of q includes the support of π .

 \blacksquare *q* is called the proposal, or importance distribution.

Importance Sampling

We have the following identity

$$I = \mathbb{E}_{\pi}(\varphi(X)) = \mathbb{E}_{q}(\varphi(X)w(X)),$$

where $w : \mathbb{X} \to \mathbb{R}^+$ is the importance weight function

$$w\left(x\right) = \frac{\pi\left(x\right)}{q\left(x\right)}.$$

■ Hence for $X_1, \ldots, X_n \stackrel{\text{i.i.d.}}{\sim} q$,

$$\widehat{I}_n^{\mathrm{IS}} = \frac{1}{n} \sum_{i=1}^n \varphi(X_i) w(X_i).$$

Importance Sampling Properties

Proposition

- (a) *Unbiased*: $\mathbb{E}_q[\widehat{I}_n^{IS}] = I$;
- (b) *Strongly consistent:* If $\mathbb{E}_q(|\varphi(X)| w(X)) < \infty$ then

$$\lim_{n\to\infty}\widehat{I}_n^{IS}=I,\quad a.s.$$

(c) CLT: $\mathbb{V}_q(\widehat{I}_n^{IS}) = \sigma_{IS}^2/n$ where

$$\sigma_{IS}^{2} := \mathbb{V}_{q} \left(\varphi(X) w \left(X \right) \right)$$

If $\sigma_{IS}^2 < \infty$ *then*

$$\lim_{n\to\infty}\sqrt{n}\left(\widehat{I}_n^{IS}-I\right)\stackrel{D}{\to}\mathcal{N}\left(0,\sigma_{IS}^2\right).$$

Importance Sampling: Practical Advice

■ Consistency does not require $\sigma_{\rm IS}^2 < \infty$ but highly recommended in practice (!).

■ Sufficient condition: If $\mathbb{E}_{\pi}\left(\varphi^{2}(X)\right) < \infty$ and $w\left(x\right) \leq M$ for all x for some $M < \infty$, then $\sigma_{\text{IS}}^{2} < \infty$.

■ In practice ensure $w(x) \le M$ although it is neither necessary nor sufficient, as seen in the following example.

Importance Sampling: Example

 $\pi(x) = \mathcal{N}(x; 0, 1), q(x) = \mathcal{N}(x; 0, \sigma^2)$ $w(x) = \frac{\pi(x)}{q(x)} \propto \exp\left[-x^2\left(1 - \frac{1}{\sigma^2}\right)\right].$

- For $\sigma^2 \ge 1$, $w(x) \le M$ for all x, and for $\sigma^2 < 1$, $w(x) \to \infty$ as $|x| \to \infty$.
- For $\varphi(x) = x^2$, we have $\sigma_{\text{IS}}^2 < \infty$ for all $\sigma^2 > 1/2$.
- For $\varphi(x) = \exp\left(\frac{\beta}{2}x^2\right)$, we have $I < \infty$ for $\beta < 1$ but $\sigma_{\text{IS}}^2 = \infty$ for $\beta > 1 \frac{1}{2\sigma^2}$.

Optimal Importance Distribution I

Question

Is there a best proposal that minimizes the variance σ_{IS}^2 ?

Proposition

The optimal proposal minimising $\mathbb{V}_q\left(\widehat{I}_n^{IS}\right)$ is given by

$$q_{opt}(x) = \frac{|\varphi(x)| \, \pi(x)}{\int_{\mathbb{X}} |\varphi(x)| \, \pi(x) \, dx}.$$

Optimal Importance Distribution II

Proof.

We have indeed

$$\sigma_{\mathrm{IS}}^{2} = \mathbb{V}_{q}\left(\varphi(X)w\left(X\right)\right) = \mathbb{E}_{q}\left(\varphi^{2}(X)w^{2}\left(X\right)\right) - I^{2}.$$

We also have by Jensen's inequality for any *q*

$$\mathbb{E}_{q}\left(\varphi^{2}(X)w^{2}\left(X\right)\right) \geq \left(\int_{\mathbb{X}}\left|\varphi(x)\right|\pi\left(x\right)dx\right)^{2}.$$

For $q = q_{opt}$, we have

$$\mathbb{E}_{q_{\text{opt}}}\left(\varphi^{2}(X)w^{2}(X)\right) = \int_{\mathbb{X}} \frac{\varphi^{2}(x)\pi^{2}(x)}{|\varphi(x)|\pi(x)} dx \times \int_{\mathbb{X}} |\varphi(x)|\pi(x) dx$$
$$= \left(\int_{\mathbb{X}} |\varphi(x)|\pi(x) dx\right)^{2}.$$

Optimal Importance Distribution

- \blacksquare $q_{\text{opt}}(x)$ can never be used in practice!
- For $\varphi(x) > 0$ we have $q_{\text{opt}}(x) = \varphi(x)\pi(x) / I$ and $\mathbb{V}_{q_{\text{opt}}}(\widehat{I}_n^{\text{IS}}) = 0$ but this is because

$$\varphi(x) w(x) = \varphi(x) \frac{\pi(x)}{q_{\text{opt}}(x)} = I,$$

it requires knowing *I*!

- This can be used as a guideline to select q; i.e. select q(x) such that $q(x) \approx q_{\text{opt}}(x)$.
- Particularly interesting in rare event simulation, not quite in statistics.

Normalised Importance Sampling

- Standard IS has limited applications in statistics as it requires knowing $\pi(x)$ and q(x) exactly.
- Assume $\pi(x) = \widetilde{\pi}(x)/Z_{\pi}$ and $q(x) = \widetilde{q}(x)/Z_{q}$, $\pi(x) > 0 \Rightarrow q(x) > 0$ and and define

$$\widetilde{w}(x) = \frac{\widetilde{\pi}(x)}{\widetilde{q}(x)}.$$

An alternative identity is

$$I = \mathbb{E}_{\pi}(\varphi(X)) = \frac{\int_{\mathbb{X}} \varphi(x) \, \widetilde{w}(x) \, q(x) dx}{\int_{\mathbb{X}} \widetilde{w}(x) q(x) dx}.$$

SLLN for NIS

Proposition (SLLN for NIS)

Let $X_1,...,X_n \overset{i.i.d.}{\sim} q$ and assume that $\mathbb{E}_q(|\phi(X)| \, w(X)) < \infty$. Then

$$\widehat{I}_n^{NIS} = \frac{\sum_{i=1}^n \varphi(X_i)\widetilde{w}(X_i)}{\sum_{i=1}^n \widetilde{w}(X_i)}$$

is strongly consistent.

Proof.

Divide numerator and denominator by n. Both converge almost surely by the strong law of large numbers.



CLT for NIS

Proposition

If
$$\mathbb{V}_q(\varphi(X)w(X)) < \infty$$
 and $\mathbb{V}_q(w(X)) < \infty$ then

$$\sqrt{n}(\widehat{I}_n^{NIS}-I)\Rightarrow \mathcal{N}(0,\sigma_{NIS}^2),$$

where

$$\sigma_{NIS}^2 := \mathbb{V}_q \left(\left[\varphi(X) w(X) - I w(X) \right] \right)$$
$$= \int \frac{\pi(x)^2 \left(\varphi(x) - I \right)^2}{q(x)} dx.$$

Proof

Proof.

First notice that with X_1, \ldots, X_n i.i.d. $\sim q$

$$\sqrt{n}(\widehat{I}_n^{\text{NIS}} - I) = \frac{\frac{1}{\sqrt{n}} \sum_{i=1}^n \widetilde{w}(X_i) [\varphi(X_i) - I]}{\frac{1}{n} \sum_{i=1}^n \widetilde{w}(X_i)}$$

where since $\widetilde{w}(x) = \widetilde{\pi}/\widetilde{q}$

$$\mathbb{E}_q\Big[\widetilde{w}(X_n)(\varphi(X_i)-I)\Big]=0.$$

Since $\mathbb{V}_q(\varphi(X)w(X)) < \infty$ by standard CLT

$$\frac{1}{\sqrt{n}}\sum_{i=1}^{n}\widetilde{w}(X_{i})\big[\varphi(X_{i})-I\big]\Rightarrow \mathcal{N}\Big(0,\mathbb{V}_{q}\Big(\widetilde{w}(X_{1})[\varphi(X_{1})-I]\Big)\Big).$$

Proof ctd...

Proof.

The strong law of large numbers applied to the denominator

$$\frac{1}{n}\sum_{i=1}^{n}\widetilde{w}(X_{i})\to\mathbb{E}_{q}[\widetilde{w}(X_{1})]=Z_{\pi}/Z_{q},\quad \text{a.s.}$$

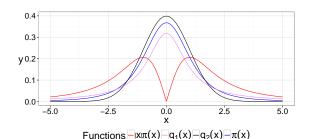
By Slutsky's theorem, combining the two

$$\sqrt{n}(\widehat{I}_n^{\text{NIS}} - I) \Rightarrow \mathcal{N}\left(0, \mathbb{V}_q\left(\widetilde{w}(X_1)[\varphi(X_1) - I]\right) \frac{Z_q^2}{Z_\pi^2}\right)$$
$$\sim \mathcal{N}\left(0, \sigma_{\text{NIS}}^2\right).$$

Alternatively, use Delta method.

Toy Example: t-distribution

- We want to compute $I = \mathbb{E}_{\pi}(|X|)$ where $\pi(x) \propto (1 + x^2/3)^{-2}$ (t₃-distribution).
- 1 Directly sample from π .
- 2 Use $q_1(x) = g_{t_1}(x) \propto (1 + x^2)^{-1}$ (t₁-distribution).
- Use $q_2(x) \propto \exp(-x^2/2)$ (normal).



Toy Example: t-distribution

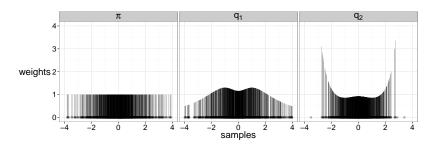


Figure: Sample weights obtained for 1000 realisations of X_i , from the different proposal distributions.

Toy Example: t-distribution

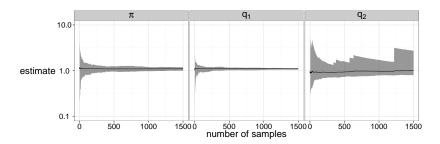


Figure: Estimates \hat{I}_n of I obtained after 1 to 1500 samples. The grey shaded areas correpond to the range of 100 independent replications.

Variance of importance sampling estimators

■ Standard Importance Sampling: $X_1, ..., X_n \stackrel{iid}{\sim} q$,

$$\widehat{I}_n^{\mathrm{IS}} = \frac{1}{n} \sum_{i=1}^n \varphi(X_i) w(X_i).$$

■ Asymptotic Variance:

$$\mathbb{V}_{as}\left(\widehat{I}_{n}^{\text{IS}}\right) = \mathbb{E}_{q}\left[\left(\varphi(X)w(X) - \mathbb{E}_{q}\left(\varphi(X)w(X)\right)\right)^{2}\right]$$
$$\approx \frac{1}{n}\sum_{i=1}^{n}\left(\varphi(X_{i})w(X_{i}) - \widehat{I}_{n}^{\text{IS}}\right)^{2}.$$

 Thus the asymptotic variance can be estimated consistently with

$$\frac{1}{n}\sum_{i=1}^{n}\left(\varphi(X_{i})w(X_{i})-\widehat{I}_{n}^{\mathrm{IS}}\right)^{2}.$$

Variance of importance sampling estimators

■ Normalised Importance Sampling: $X_1, ..., X_n \stackrel{iid}{\sim} q$,

$$\widehat{I}_n^{\text{NIS}} = \frac{\sum_{i=1}^n \varphi(X_i)\widetilde{w}(X_i)}{\sum_{i=1}^n \widetilde{w}(X_i)}.$$

Asymptotic Variance:

$$\mathbb{V}_{as}\left(\widehat{I}_{n}^{\text{NIS}}\right) = \frac{\mathbb{E}_{q}\left[\left(\varphi(X)w(X) - I \times w(X)\right)^{2}\right]}{\mathbb{E}_{q}\left[w(X)\right]^{2}}.$$

■ Thus the asymptotic variance can be estimated consistently with

$$\frac{\frac{1}{n}\sum_{i=1}^{N}\widetilde{w}(X_i)^2\left(\varphi(X_i)-\widehat{I}_n^{\text{NIS}}\right)^2}{\left(\frac{1}{n}\sum_{i=1}^{N}\widetilde{w}(X_i)\right)^2}.$$

Diagnostics

- Importance sampling works well when all weights roughly equal.
- If dominated by one $\widetilde{w}(X_i)$,

$$\widehat{I}_n^{\text{NIS}} = \frac{\sum_{i=1}^n \varphi(X_i) \widetilde{w}(X_i)}{\sum_{i=1}^n \widetilde{w}(X_i)} \approx \widetilde{w}(X_j) \varphi(X_j).$$

The "effective sample size" is one.

■ To how many unweighted samples correspond our weighted samples of size n? Solve for n_e in

$$\frac{1}{n} \mathbb{V}_{as} \left(\widehat{I}_n^{\text{NIS}} \right) = \frac{\sigma^2}{n_e},$$

where σ^2/n_e corresponds to the variance of an unweighted sample of size n_e .

Diagnostics

■ We solve by matching $\varphi(X_i) - \widehat{I}^{\text{NIS}}$ with $\varphi(X_i) - I \approx \sigma$ as if they were i.i.d samples:

$$\frac{1}{n} \frac{\frac{1}{n} \sum_{i=1}^{N} \widetilde{w}(X_i)^2 \left(\varphi(X_i) - \widehat{I}_n^{\text{NIS}}\right)^2}{\left(\frac{1}{n} \sum_{i=1}^{n} \widetilde{w}(X_i)\right)^2} \approx \frac{\sigma^2}{n_e}$$
i.e.
$$\frac{1}{n} \frac{\frac{1}{n} \sum_{i=1}^{N} \widetilde{w}(X_i)^2}{\left(\frac{1}{n} \sum_{i=1}^{n} \widetilde{w}(X_i)\right)^2} = \frac{1}{n_e}.$$

■ The solution is

$$n_e = \frac{\left(\sum_{i=1}^n \widetilde{w}(X_i)\right)^2}{\sum_{i=1}^n \widetilde{w}(X_i)^2},$$

and is called the effective sample size.

Rejection and Importance Sampling in High Dimensions

Toy example: Let $X = \mathbb{R}^d$ and

$$\pi(x) = \frac{1}{(2\pi)^{d/2}} \exp\left(-\frac{\sum_{i=1}^{d} x_i^2}{2}\right)$$

and

$$q(x) = \frac{1}{(2\pi\sigma^2)^{d/2}} \exp\left(-\frac{\sum_{i=1}^{d} x_i^2}{2\sigma^2}\right).$$

■ How do Rejection sampling and Importance sampling scale in this context?

Lecture 3

Performance of Rejection Sampling

We have

$$w\left(x\right) = \frac{\pi\left(x\right)}{q\left(x\right)} = \sigma^{d} \exp\left(-\frac{\sum_{i=1}^{d} x_{i}^{2}}{2} \left(1 - \frac{1}{\sigma^{2}}\right)\right) \le \sigma^{d}$$

for $\sigma > 1$.

Acceptance probability is

$$\mathbb{P}\left(X \text{ accepted}\right) = \frac{1}{\sigma^d} \to 0 \text{ as } d \to \infty,$$

i.e. exponential degradation of performance.

■ For d = 100, $\sigma = 1.2$, we have

$$\mathbb{P}(X \text{ accepted}) \approx 1.2 \times 10^{-8}.$$

Lecture 3

Performance of Importance Sampling

We have

$$w\left(x\right) = \sigma^{d} \exp\left(-\frac{\sum_{i=1}^{d} x_{i}^{2}}{2} \left(1 - \frac{1}{\sigma^{2}}\right)\right).$$

■ Variance of the weights:

$$\mathbb{V}_{q}\left[w\left(X\right)\right] = \left(\frac{\sigma^{4}}{2\sigma^{2} - 1}\right)^{d/2} - 1$$

where $\sigma^4/\left(2\sigma^2-1\right)>1$ for any $\sigma^2>1/2$.

■ For d = 100, $\sigma = 1.2$, we have

$$\mathbb{V}_q\left[w\left(X\right)\right] \approx 1.8 \times 10^4$$
.

Lecture 3

Wait a minute...

Lecture 1:

■ Simpson's rule for approximating integrals: error in $\mathcal{O}(n^{-1/d})$.

Lecture 2:

■ Monte Carlo for approximating integrals: error in $\mathcal{O}(n^{-1/2})$ with rate independent of d.

And now:

■ Importance Sampling standard deviation in the Gaussian example in $\exp(d)n^{-1/2}$.

The rate is indeed independent of d but the "constant" (in n) explodes exponentially (in d).

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Markov chain Monte Carlo

- Revolutionary idea introduced by Metropolis et al., J. Chemical Physics, 1953.
- **Key idea**: Given a target distribution π , build a Markov chain $(X_t)_{t>1}$ such that, as $t \to \infty$, $X_t \sim \pi$ and

$$\frac{1}{n}\sum_{t=1}^{n}\varphi\left(X_{t}\right)\to\int\varphi\left(x\right)\pi\left(x\right)dx$$

when $n \to \infty$ e.g. almost surely.

- Central limit theorems with a rate in $1/\sqrt{n}$.
- In some cases the constant (in *n*) does not explode exponentially with the dimension *d*, but polynomially.

Lecture 3 High dimensions

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Side Dish: Control Variates

- Variance reduction techniques, not always applicable but useful in some cases.
- Suppose that we want to compute

$$I = \int \varphi(x)\pi(x)dx$$

and that we know exactly

$$J = \int \psi(x)\pi(x)dx.$$

■ Sample $X_1, ..., X_n$ from π and compute

$$\widehat{I}_n = \frac{1}{n} \sum_{i=1}^n (\varphi(X_i) - \lambda(\psi(X_i) - J)).$$

■ What is the benefit of \widehat{I}_n over the standard Monte Carlo estimator?

Lecture 3 High dimensions