VE414 Lecture 8

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• We have considered both deterministic and distributional approximations for

$$\int_{\infty}^{\infty} f_{Y\mid X}\left(y\mid x\right) \, dy$$

now we turn to the 3rd approach, which is what people use often nowadays.

- Q: Given a needle, a ruler, a pen and a piece of paper, how would you estimate
 - $\pi = 3.14159265358979323846264338327950288419716939937510?$
- Q: Given a needle of length l dropped on a piece of paper with parallel lines t units apart, what is the probability that the needle will land across a line?
 - ullet Let y be the distance from the centre of the needle to the closest line, and let heta be the acute angle between the needle and the lines, then

$$f_Y(y) = \begin{cases} \frac{2}{t}, & \text{for } 0 \leq y \leq \frac{t}{2} \text{ ,} \\ 0, & \text{otherwise.} \end{cases} \quad \text{and} \quad f_\theta(\theta) = \begin{cases} \frac{2}{\pi}, & \text{for } 0 \leq \theta \leq \frac{\pi}{2} \text{ ,} \\ 0, & \text{otherwise.} \end{cases}$$

ullet The two random variables, Y and heta, are independent, so the joint pdf is

$$f_{Y, heta}(y, heta) = egin{cases} rac{4}{t\pi}, & ext{for } 0 \leq y \leq rac{t}{2}, \, 0 \leq heta \leq rac{\pi}{2} \ , \ 0, & ext{otherwise}. \end{cases}$$

• The needle crosses a line if

$$y \leq \frac{l}{2} \sin \theta$$

• Assume l < t, then the probability is given by

$$p = \int_0^{\pi/2} \int_0^{l/2 \sin \theta} \frac{4}{t\pi} \, dy \, d\theta = \frac{2l}{t\pi}$$

• Back to our question of estimating π , how can we make use of this?

$$X_k \implies \pi = \lim_{k \to \infty} \frac{2l}{t} \frac{k}{x_k}$$

• Using needle dropping to estimate π illustrates the essence of

Monte Carlo simulation

• By dropping the needle repeatedly, we have obtained the information on

$$y \le \frac{l}{2}\sin\theta$$

in terms of x_k number of success out of k number of trials, which leads to

$$\frac{x_k}{k} \approx p = \int_0^{\pi/2} \int_0^{l/2\sin\theta} \frac{4}{t\pi} \, dy \, d\theta = \frac{2l}{t\pi}$$

• In general, the essence of the above idea is applied to the following integral

$$A = \int_{\mathcal{D}} f_{\mathbf{X}|\mathbf{Y}} \left(\mathbf{x} \mid \mathbf{y} \right) f_{\mathbf{Y}} \left(\mathbf{y} \right) d\mathbf{y}$$

where \mathcal{D} is region in high-dimensional space.

• If we can draw independent and identically distributed samples

$$\mathbf{y}^{(1)}, \dots, \mathbf{y}^{(m)}$$

from the target distribution, e.g. the posterior distribution, then we have

$$\hat{A}_{m} = \frac{1}{m} \left[f_{\mathbf{X}|\mathbf{Y}} \left(\mathbf{x} \mid \mathbf{y}^{(1)} \right) f_{\mathbf{Y}} \left(\mathbf{y}^{(1)} \right) + \dots + f_{\mathbf{X}|\mathbf{Y}} \left(\mathbf{x} \mid \mathbf{y}^{(m)} \right) f_{\mathbf{Y}} \left(\mathbf{y}^{(m)} \right) \right]$$

 The law of large numbers states the average of many independent random variables with common mean and finite variances tends to stabilise at their common mean; that is,

$$\lim_{m \to \infty} \hat{A}_m = A, \quad \text{with probability 1.}$$

• Its convergence rate can be assessed by the central limit theorem,

$$\sqrt{m}\left(\hat{A}_m - A\right) \to \operatorname{Normal}\left(0, \sigma^2\right), \quad \text{in distribution}$$

where $\sigma^2 = \text{Var}\left(f_{\mathbf{X}|\mathbf{Y}}\left(\mathbf{x} \mid \mathbf{y}\right) f_{\mathbf{Y}}\left(\mathbf{y}\right)\right)$ in terms of \mathbf{Y} .

• Like normal approximation, the "error term" of Monte Carlo approximation

$$\hat{A}_{m}$$

depends on the number of points in your sample, i.e.

m

regardless of the dimensionality of \mathcal{D} , that is, a high-dimension integral can be approximated just as efficient as a one-dimensional integral.

- This makes Monte Carlo very appealing, because Gauss quadrature is highly efficient in 1-dimension, but its counterpart in high-dimensional space is not, i.e. the error depends on the dimension of \mathcal{D} .
- Another aspect of Monte Carlo is its simplicity or convenience when comes to various inference that rooted on posterior distribution, e.g. consider

$$\mathbb{E}\left[\mathbf{Y} \mid \mathbf{X}\right] = \int_{\mathcal{D}} \mathbf{Y} f_{\mathbf{X} \mid \mathbf{Y}} \left(\mathbf{x} \mid \mathbf{y}\right) f_{\mathbf{Y}} \left(\mathbf{y}\right) d\mathbf{y}$$

• In fact, the focus of using Monte Carlo is not about the normalising constant

$$A = \int_{\mathcal{D}} f_{\mathbf{X}|\mathbf{Y}} \left(\mathbf{x} \mid \mathbf{y} \right) f_{\mathbf{Y}} \left(\mathbf{y} \right) d\mathbf{y}$$

It provides inference directly, to illustrate that, consider the following again

$$\begin{array}{rcl} P & \sim & \mathrm{Uniform}\left(0,1\right) \\ X_k \mid P & \sim & \mathrm{Binomial}\left(k,p\right) \\ \Longrightarrow & P \mid X_3 = 2 & \sim & \mathrm{Beta}\left(3,2\right) \end{array}$$

• The easiest sampling method is based discretising on a grid approximation,

```
> # Given
> k = 3; x = 2; a = 0; b = 1;
> # No. nodes
> n = 10;
> # Define grid
> p_grid = seq(from=a, to=b, length.out=n)
```

```
> # define prior
> prior = rep(1, n)

> # compute likelihood at each value in grid
> likelihood = dbinom(x , size=k, prob=p_grid)

> # compute product of likelihood and prior
> unstd.posterior = likelihood * prior
```

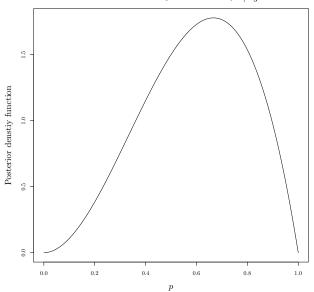
- Notice the following step is different from deterministic approximation
 - > # discretising the distribution, so it sums to 1
 - > posterior = unstd.posterior / sum(unstd.posterior)
- ullet Once we have the discretisation, we sample m values from it
 - > # sample size
 - > m = 10

- > # Take a sample from this discrete distribution
 > samples = sample(p_grid, prob=posterior,
- + size=m, replace=TRUE)
- > samples
 - [1] 0.5555556 0.6666667 0.5555556 0.5555556
 - [5] 0.5555556 0.6666667 0.6666667 0.6666667
 - [9] 0.222222 0.7777778

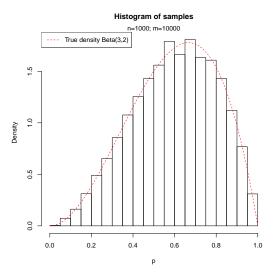
> p_grid

- [1] 0.0000000 0.1111111 0.222222 0.3333333
- [5] 0.4444444 0.5555556 0.6666667 0.7777778
- [9] 0.8888889 1.0000000
- Note some values occurring more often than other values which reflects the posterior knowledge regarding p in a very rough sense.

Posterior density function $f_{P|X_3=2}$



ullet Of course, we should use more nodes n=1e3 and a larger sample m=1e4.



- A sample form a Monte Carlo simulation offers more than just a crude plot of the posterior density, which we already have before the simulation, i.e.
 - > # normalising constant
 - > A = (b-a) * sum(unstd.posterior) / n
- Having a sample means we can easily find an expectation of some function,

$$\mathbb{E}[g(y)] = \int_{\infty}^{\infty} g(y) f_{Y|X}(y \mid x) \, dy$$

> mean(samples)

[1] 0.5989582

- > mean(log(samples/(1-samples))) # log odds/logit
- [1] 0.4952244

• We could also obtain an estimate of the probability of p in certain interval

```
> sum(samples < 0.5) / m
```

```
[1] 0.3168
```

> sum(samples > 0.25 & samples < 0.75) / m

```
[1] 0.6899
```

• Of course, we could also reverse it

```
> median(samples); quantile(samples, 0.5)
```

```
[1] 0.6136136
50%
```

0.6136136

0.1941942 0.9309309

- Since we have discrete values, optimisation is easy
 - > p_grid[which.max(posterior)]

[1] 0.6666667

Q: How about the posterior predictive distribution?

$$\begin{split} f_{X_k^*|X_3=2} &= \int_0^1 f_{X_k^*|P} \cdot f_{P|X_3=2} \, dp \\ &= \frac{k^*!}{x_k^*!(k^*-x_k^*)!} \cdot \frac{\Gamma\left(\alpha^*+\beta^*\right)}{\Gamma\left(\alpha^*\right)\Gamma\left(\beta^*\right)} \cdot \frac{\Gamma\left(x_k^*+\alpha^*\right)\Gamma\left(k^*-x_k^*+\beta^*\right)}{\Gamma\left(k^*+\alpha^*+\beta^*\right)} \end{split}$$

```
> ppsamples = double(m)
> for (i in 1:m){
+    p = samples[i]
+    ppsamples[i] = rbinom(1, size=3, p)
+ }
> head(ppsamples)
```

[1] 1 1 3 3 2 1

ullet Note the above example has only a single scalar unknown y,

$$f_{Y|X} \propto f_{X|Y} f_Y$$

it illustrates what Monte Carlo simulation can provide. It is chosen since it is simple/familiar, and knowing the exact posterior makes comparisons easy.

In practice, Monte Carlo simulation is used for complicated Bayesian models

$$f_{\{\mathbf{P},\alpha,\beta\}|\mathbf{Y}} \propto f_{\mathbf{Y}|\{\mathbf{P},\alpha,\beta\}} \cdot f_{\mathbf{P}|\{\alpha,\beta\}} \cdot f_{\alpha,\beta}$$

$$= f_{\alpha,\beta} \prod_{j=1}^{71} \binom{k_j}{y_j} \frac{\Gamma(\alpha+\beta)}{\Gamma(\alpha)\Gamma(\beta)} p_j^{y_j+\alpha-1} (1-p_j)^{k_j-y_j+\beta-1}$$

with a large number of unknowns, however, what Monte Carlo can provide and how it can provide them remain largely the same.

- Q: But do you notice any flaw with this simple sampling scheme?
 - The above sampling scheme is based on a grid approximation, which means it will fail miserably in high dimensions, we must replace this direct sampling scheme before we can apply Monte Carlo in high dimensions.