## Lecture Summary 3

Change of Notation: We have seen that we need to sample a density function, call it f(x), in order to implement Monte Carlo integration. We have to start somewhere and it is assumed we can sample uniform random variables from (0,1). Computers can do this via clever deterministic sequences.

Some easy distributions can then be sampled via a single uniform U;

$$X = F^{-1}(U).$$

It is then straightforward to show that  $P(X \le x) = F(x)$ , so this method provides a random variable with distribution F.

For harder distributions we need something more sophisticated and the first attempt is known as Rejection Sampling. The basic idea is to sample a density g(x) somehow close to f(x), and to see whether we can use this as a sample from f. Suppose we can write f(x) = h(x) g(x) where  $h(x) \leq M < \infty$ . To this end, consider the joint density function

$$f(x, u) = M \mathbf{1}(0 < u < h(x)/M) \mathbf{1}(0 < u < 1) g(x).$$

The marginal density for x is f(x) so we can sample (x, u) from f(x, u) and keep the x sample as coming from f.

We can sample from f(x, u) by keep sampling from the density

$$\tilde{f}(x, u) = \mathbf{1}(0 < u < 1) g(x)$$

unitil we get a (x, u) satisfying u < h(x)/M. We then keep the x value as coming from f. We can clearly sample from  $\tilde{f}$  by taking u uniform from (0,1) and x from g.

The probability of acceptance is important; too small and this method won't work. Now by the law of total probability,

$$P(U < h(X)/M) = \int h(x) g(x) dx/M = 1/M.$$

Note that  $M \geq 1$  since M is the maximum value of f(x)/g(x) which must be greater than 1 since one density can not sit under another as they both must integrate to 1. The closer g is to f the smaller M becomes and hence the larger the probability of acceptance.