Quick slides on RMC

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Basics, sampling from a CDF

Given a non-normalized probability distribution, $p^*(\mathbf{x})$, defined over a composite domain $X = V\Omega$, we can determine a normalized probability distribution, $p(\mathbf{x})$, with

$$p(\mathbf{x}) = \frac{p^*(\mathbf{x})}{\int_X p^*(\mathbf{x}) d\mathbf{x}}.$$
 (1)

We can then determine a CDF from

$$c(\mathbf{x}) = \int_{\mathbf{x}_0}^{\mathbf{x}} p(\mathbf{x}) d\mathbf{x} \tag{2}$$

where \mathbf{x}_0 is some starting domain point.

We can then determine an inverse CDF, $\mathbf{x}_i = c^{-1}(\theta)$, $\theta \in [0,1], \theta \in \mathbb{R}$, from which we can compute the average probability over the sub-domain (or bin), $Y \subset X$,

$$\rho_{Y,avg} = \frac{1}{N} \sum_{i=1}^{N} \begin{cases} 1, & \text{if } \mathbf{x}_i \in Y \\ 0, & \text{if } \mathbf{x}_i \notin Y \end{cases}$$
 (3)

This value integrated over the portion of the composite domain, *A*, can be written as

$$\int_{\Omega} p_{Y,avg} d\Omega = \left(\int_{\Omega} d\Omega \right) \frac{1}{N} \sum_{i=1}^{N} \begin{cases} 1, & \text{if } \mathbf{x}_i \in Y \\ 0, & \text{if } \mathbf{x}_i \notin Y \end{cases}$$
 (4)

Finally:

$$\int_{\Omega} p_{Y,avg}^* d\Omega = \left(\int_{X} p^*(\mathbf{x}) d\mathbf{x} \right) \int_{\Omega} p_{Y,avg} d\Omega \tag{5}$$

Monte Carlo integration

We seek $\int_X p^*(\mathbf{x}) d\mathbf{x}$ via Monte Carlo integration we can be determined simply from a uniform sampling in Y

$$\int_{X} p^{*}(\mathbf{x}) d\mathbf{x} = \sum_{Y} \int_{Y} p^{*}(\mathbf{x}) d\mathbf{x}$$
 (6)

$$\int_{Y} p^{*}(\mathbf{x}) d\mathbf{x} = \left(\int_{V_{Y}} .dV \int_{\Omega} .d\Omega \right) \frac{1}{N_{Y}} \sum_{i=1}^{N_{Y}} p^{*}(\mathbf{x}_{i})$$
 (7)

Residual

Our residual is given by

$$r^* = r_{interior}^* + r_{surface}^* \tag{8}$$

where we lumped surface sources into $r_{surface}^*$. These surface sources present themselves through discontinuities between the present cell flux, ϕ^P , and the neighboring flux, ϕ^N . The latter can be a neighboring cell or a boundary.

Interior residual

The interior residual is then given by:

$$r_{interior}^* = \frac{1}{4\pi} \left(q - \sigma_a \phi - \mathbf{\Omega} \cdot \nabla \phi \right)$$
 (9)

from which we can determine the phase-space integration as

$$\int_{X} r_{interior}^{*}.d\mathbf{x} = \int_{V} \int_{\Omega} \frac{1}{4\pi} \left(q - \sigma_{a}\phi - \mathbf{\Omega} \cdot \nabla \phi \right) . d\mathbf{\Omega} . dV$$

$$= \sum_{Y} \int_{V_{Y}} \int_{\Omega} r_{Y,interior}^{*}. d\mathbf{\Omega} . dV$$

$$= \sum_{Y} \left(\int_{V_{Y}} . dV \int_{\Omega} . d\mathbf{\Omega} \right) r_{Y,interior,avg}^{*} \tag{10}$$

Via Monte Carlo:

$$r_{Y,interior,avg}^* = \frac{1}{N_Y} \sum_{i=1}^{N_Y} r_{Y,interior}^*(\mathbf{x}_i)$$
 (11)

Surface residual

The surface residual can be presented as

$$\int_{S} \int_{2\pi} \int_{-1}^{0} \mu \ r_{surface}^{*}.d\mu.d\varphi.dA = \int_{S} \int_{2\pi} \int_{-1}^{0} \mu \frac{1}{4\pi} (\phi^{N} - \phi^{P}).d\mu.d\varphi.dA \tag{12}$$

$$\therefore \int_{X} r_{surface}^{*}.d\mathbf{x} = \sum_{Y} \left(\int_{S_{Y}} .dS \int_{2\pi} \int_{-1}^{0} \mu.d\mu.d\varphi \right) r_{Y,surface,avg}^{*}$$

$$= \sum_{Y} \sum_{f} \left(A_{f} \pi \right) r_{Y,f,avg}^{*}$$
(13)

Via Monte Carlo:

$$r_{Y,f,avg}^* = \frac{1}{N_f} \sum_{i=1}^{N_f} r_{Y,f}^*(\mathbf{x}_i)$$
 (14)

Sampling strategy

Sample the CDF, first interior vs surface:

▶ If interior, sample cell (\rightarrow cell Y), sample position randomly in cell, sample direction randomly:

$$r_{i,interior}^* = \frac{1}{4\pi} \left(q - \sigma_a \phi - \mathbf{\Omega} \cdot \nabla \phi \right)$$
 (15)

Particle weight: $w = \frac{r_{i,interior}^*}{r_{Y,interior,avg}^*}$

► If surface, sample face (f), sample random position on face, sample direction using cosine-law:

$$r_{i,surface}^* = \frac{1}{4\pi} (\phi^N - \phi^P) \tag{16}$$

Particle weight: $w = \frac{r_{i,surface}^*}{r_{Y,surface,avg}^*}$

Global normalization:

$$\int_{X} r^* . d\mathbf{x} = \int_{X} r_{interior}^* . d\mathbf{x} + \int_{X} r_{surf}^* . d\mathbf{x}$$
 (17)

