

HIGHER ORDER DISCONTINUOUS FINITE ELEMENT METHODS FOR
DISCRETE ORDINATES THERMAL RADIATIVE TRANSFER

A Dissertation

by

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ABSTRACT

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DEDICATION

To my wife, Kelli, for traveling with me along all the unforeseen curves of life.
To my mom and dad, for providing my foundation and inspiration.

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NOMENCLATURE

DFEM	Discontinuous finite element method
DSA	Diffusion synthetic acceleration
LDFEM	Linear discontinuous finite element method
S2SA	S_2 synthetic acceleration
SL	Self-lumping
S_N	Discrete ordinates method
TL	Traditional lumping
TRT	Thermal radiative transfer
$\vec{\Omega}$	Particle direction
c	Speed of light
I	Photon radiation intensity
ψ	Neutron transport angular flux
T	Temperature
σ_t	Total interaction opacity
σ_s	Scattering interaction opacity
σ_a	Absorption opacity
B	Planck function
B_g	Planck function integrated across photon energy group g
\underline{i}	Basis function i
Σ_t	Macroscopic neutron total interaction cross section
Σ_s	Macroscopic neutron scattering interaction cross section
Σ_a	Macroscopic neutron absorption interaction cross section
Σ_f	Macroscopic neutron fission interaction cross section

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1. INTRODUCTION

This dissertation is dedicated to the solution of thermal radiative transfer (TRT) equations. The TRT equations:

$$\frac{1}{c} \frac{dI}{dt} + \vec{\Omega} \cdot \vec{\nabla} I + \sigma_t I = \int_0^\infty \int_{4\pi} \sigma_s(\vec{\Omega}' \rightarrow \vec{\Omega}, E' \rightarrow E) I, d\vec{\Omega}' dE' + \sigma_a B \quad (1.1a)$$

$$C_v \frac{dT}{dt} = \int_0^\infty \sigma_a (\phi - 4\pi B) dE, \quad (1.1b)$$

are a nonlinear system of equations that describe the exchange of energy between a photon radiation field and a non-moving material. The radiation intensity, I , is a seven dimensional field dependent upon spatial location, \vec{x} ; photon energy, E ; photon direction of travel, $\vec{\Omega}$; and time t . c is the speed of light. Material opacities for all interactions, σ_t ; absorption, σ_a ; and scattering, σ_s are functions of photon energy and material temperature, T . Material heat capacity, C_v , is also a function of material temperature. The angle integrated radiation intensity is an integral over all photon directions of the the photon intensity and is a function of space and photon energy. Finally, the Planck function, B , is a function of photon energy and material temperature. While materials at all temperatures emit photon radiation, the radiation emission is proportional to T^4 . Thus, solution of the radiative transfer equations is most important in situations where materials are very hot. Solving the thermal radiative transfer equations is an important component of the simulation of different scientific and engineering problems including astrophysics supernova explosions and high energy density laboratory physics experiments like those conducted at the National Ignition Facility.

1.1 Simplifications of the Thermal Radiative Transfer Equations

In this dissertation, we make a number of simplifying assumptions to make solution of Eqs. (1.1) more tractable. First, we limit our focus to 1-D Cartesian (slab) geometry. The assumption of slab geometry is not required, but slab geometry radiation transport simulations require significantly less computational time. Further, any methods that have a possibility of being viable for radiation transport in multiple spatial dimensions must also work well in slab geometry.

Second, we approximate the continuous angle dependence of the intensity using the discrete ordinates (S_N) method. The S_N method approximates the true definition of the angle integrated intensity,

$$\phi(\vec{x}, E, t) = \int_{4\pi} I(\vec{x}, \vec{\Omega}, E, t) d\vec{\Omega},$$

using quadrature integration,

$$\phi(\vec{x}, E, t) \approx \sum_{d=1}^{N_{dir}} w_d I(\vec{x}, \vec{\Omega}_d, E, t). \quad (1.2)$$

In Eq. (1.2), $\{w_d, \vec{\Omega}_d\}_{d=1, \dots, N_{dir}}$ is the set of N_{dir} quadrature weights w_d and discrete directions, $\vec{\Omega}_d$ and corresponding intensities I_d .

Finally, we treat the photon energy dependence using the multi-frequency method. The multi-frequency method approximates photon energy dependence by discretizing the continuous photon energy dependence with G discrete groups such that:

$$\int_0^\infty I(\vec{x}, \vec{\Omega}, t, E) dE = \sum_{g=1}^G I_g, \quad (1.3)$$

where

$$I(\vec{x}, \vec{\Omega}, t)_g = \int_{E_{g+1/2}}^{E_{g-1/2}} I(\vec{x}, \vec{\Omega}, t, E) dE, \quad (1.4)$$

$E_{g+1/2}$ is the lower photon energy bound of group g , $E_{g-1/2}$ is the upper photon energy bound of group g , and we have maintained the traditional neutron transport number of higher energy particles belonging to lower number energy groups.

1.2 Spatial and Temporal Discretization

To complete a description of the approach we will take to solve Eqs. (1.1), we now describe how we will discretize the spatial and temporal variables.

1.2.1 Time Integration

The appearance of the speed of light in Eq. (1.1) results in the TRT equations being very stiff. To solve the such a stiff system of equations would require either an impractically small time step, or the use of implicit methods. We elect to use Diagonally Implicit Runge-Kutta (DIRK) methods to advance our TRT solution in time. The simplest of DIRK scheme is the first order implicit Euler scheme, but more advanced DIRK higher order methods in time [1].

1.2.2 Spatial Discretization with Discontinuous Finite Elements

The linear discontinuous finite element method (LDFEM) has long been used to solve the discrete ordinates neutron transport equation [2]. Through manipulation, the thermal radiative transfer equations can be transformed into a form that is equivalent to the neutron transport equation with pseudo- scattering, fission, and fixed sources. This makes it possible to use the same methods and techniques developed for neutron transport to assist in solving the thermal radiative transfer equations. LDFEM has achieved wide spread acceptance in the neutron transport community because it is accurate [3] and highly damped. Because it possesses the thick diffusion

limit [4], LDFEM has also been applied to the S_N TRT equations. Morel, Wareing, and Smith first considered the application of LDFEM to the S_N TRT equations in [5]. Mass matrix lumped LDFEM was shown to preserve the thick equilibrium diffusion limit [5]. This suggests that discontinuous finite element (DFEM) schemes can be used to accurately solve the TRT equations in both diffusive and transport effects dominated regions.

1.3 Progression Towards Higher Order DFEM Thermal Radiative Transfer

For higher order (quadratic and higher polynomial degree trial spaces) DFEM to be accurate and practical for solving Eqs. (1.1) we must demonstrate that higher order DFEM:

1. are “robust”,
2. account for within cell spatial variation of opacity accurately, and
3. can be accelerated using appropriate iterative acceleration techniques.

By “robust”, we mean that that calculated radiation outflow from a spatial cell is strictly positive for all cell widths and optical thicknesses.

In Section 2 we use a steady-state, mono-energetic, source-free pure absorber neutron transport problem with a cross section that is constant in space to examine the robustness of different radiation transport DFEM matrix lumping techniques. Next, we extend the techniques developed by Adams [6, 7], for a spatial discretization scheme related to LDFEM to address the within cell spatial variation of opacity, for higher order DFEM in Section 3. Then, we examine iterative acceleration techniques compatible with higher order DFEM spatial discretizations that account for the spatial variation of interaction cross (in neutron transport problems) or opacity (TRT simulations) in Section 4.

In preparation for solving the coupled, non-linear TRT equations, in Section 5 we combine all of the strategies we have developed in Sections 2-4 and apply them to a coupled system of linear equations. Section 5, is devoted to the development and solution of a two-group fuel depletion problem that uses explicit Euler time differencing. Finally, in Section 6 we solve the energy integrated, or grey, thermal radiative transfer equations using higher order DFEM. We then provides solutions to analytic benchmarks to verify our methods [8], and use the method of manufactured solutions [9] to demonstrate the increased accuracy of applying higher order DFEM and DIRK time integration techniques to the grey TRT equations.

2. DISCONTINUOUS FINITE ELEMENTS FOR RADIATION TRANSPORT

In Section 1, we briefly mentioned that through manipulation, the thermal radiative transfer equations can be put into a form equivalent to the neutron transport equation with pseudo-scattering, fission, and fixed sources. We will (repeatedly) go through this process in Section 6, but for now we take for granted that solving for the neutron transport equation's angular flux, ψ , is related to solving Eq. (1.1) for I . Additionally, we will assume that a steady-state, source-free, pure absorber neutron transport problem taxes DFEM schemes in a manner similar to the way DFEM schemes are tested in time-dependent thermal radiative transfer simulations, in particular Marshak wave type problems [10]. This chapter draws primarily from our previously published work in [11] and [12].

2.1 History of DFEM for Neutron Transport

The linear discontinuous finite element method (LDFEM) for discrete ordinates neutron transport is widely used and has been extensively studied [13, 14, 15, 16]. However, the DFEM technique is not limited to linear trial spaces. Reed et. al [2] used arbitrary order DFEM S_N neutron transport in TRIPLET but, due to data storage limitations at the time, only LDFEM was computationally practical. As a result of these historical computing limitations, the accuracy of LDFEM [3], and LDFEM possessing the thick diffusion limit [4], the majority of reported DFEM radiation transport literature has focused on the LDFEM approximation. Higher order DFEM methods have received periodic attention; some older examples include the work of Walters [17] and Hennart and del Valle [18, 19]. More recent investigations of higher order DFEM trial spaces include those of Warsa and Prinja [20] and Wang and Ragusa [21, 22]. The primary focus of the work in [18, 19, 20, 21, 22] was the

convergence rate of arbitrary order DFEM schemes.

Negative angular flux solutions of the neutron transport equation obtained with LDFEM have been well documented in [14, 15, 16]. While these negativities do not affect the order of convergence and can be tolerated for certain applications [23], some nonlinear problems, particularly radiative transfer calculations, can diverge if the angular intensities are negative. As a result, several methods to eliminate or inhibit negative solutions have been developed and can be categorized into one of three categories: ad-hoc fix-ups [14], strictly non-negative solution representations [15], and matrix lumping [16]. The first two methods result in nonlinear systems of equations, while matrix lumping yields linear systems of equations. By definition, ad-hoc fix-ups and strictly non-negative solution representations yield non-negative outflows in 1-D, 2-D, and 3-D geometries, regardless of material properties. Mass matrix lumping (applied to LDFEM) yields strictly positive outflows only in 1-D geometries, though it does otherwise inhibit negativities [16].

Solution positivity of even degree unlumped DFEM methods for 1-D problems has been noted previously [17, 18, 19]. In comparing DFEM methods to nodal transport methods, Walters derived the quadratic DFEM scheme from the nodal transport equations using the Padé(2,3) approximation to the exponential term and noted that this approximation would result in a strictly positive outflow, regardless of cell optical thickness [17]. Hennart and del Valle then showed for slab geometry that all even P degree polynomial DFEM schemes approximate the cell outflow angular flux as a Padé($P, P + 1$) function, which is a strictly positive approximation of the exponential [18, 19]. The positivity of even degree unlumped DFEM for 1-D problems was also shown in [20].

2.2 Mass Matrix Lumping Techniques

In this Section, we examine the idea of mass matrix lumping and its ability to ensure positive angular flux solutions of the neutron transport equation for arbitrary degree DFEM trial spaces in non-scattering 1-D slab geometries. We consider traditional lumping (TL), that constructs a diagonal mass matrix by collapsing all off-diagonal entries onto the main diagonal [16], and quadrature-based self-lumping (SL) methods [24], that yield a diagonal mass matrix by numerically integrating the DFEM equations using the DFEM interpolatory points as quadrature points. Restricting ourselves to equally-spaced interpolation points, self-lumping numerical integration with the greatest degree of accuracy is achieved through the use of closed Newton-Cotes formulae [25]. However, Newton-Cotes formulas with a large number of integration points are known to be oscillatory and are of relatively low-order accuracy, integrating polynomials at most of degree equal to the number of integration points. By considering solution representations that employ quadrature points as the interpolatory points, for example Gauss-Legendre (hereafter Gauss) or Lobatto-Gauss-Legendre (hereafter Lobatto) quadrature points [25], we wish to find methods that are self-lumping with a significantly higher accuracy. We analyze the combinations of Lagrange interpolatory points and numerical integration strategies given in Table 2.1 for positivity of the angular flux solution, local truncation error order, and spatial convergence order as a function of trial space polynomial degree.

We limit the consideration of exact numerical integration schemes to those with equally-spaced interpolatory points, due to the fact that exact integration with any particular set of interpolatory points will always yield the same DFEM solution.

It has long been noted that traditional lumping (TL) with equally-spaced interpolatory points for 1-D LDFEM is equivalent to using the trapezoidal quadrature

Table 2.1: Nomenclature of numerical schemes.

Interpolation Point Type	Integration Strategy	Method Short Hand Name
Equally- Spaced	Exact spatial integration, with collapsing of mass matrix entries to the main diagonal	TL
Equally- Spaced	Numerical integration via Newton-Cotes quadrature restricted to interpolation points	SL Newton-Cotes
Gauss Quadrature	Numerical integration via Gauss quadrature restricted to interpolation points	SL Gauss
Lobatto Quadrature	Numerical integration via Lobatto quadrature restricted to interpolation points	SL Lobatto
Equally- Spaced	Exact spatial integration	Exact DFEM

rule to approximately integrate the mass matrix [26] while exactly integrating the gradient operator. Since the trapezoidal rule is identical to the closed Newton-Cotes formula with two points, we hypothesize that, for finite elements of arbitrary order using equally-spaced interpolatory points, traditional lumping is equivalent to using a closed Newton-Cotes formula to compute the mass matrix while exactly integrating the gradient operator. We demonstrate the equivalence between traditional lumping and closed Newton-Cotes formulae in the computation of the mass matrix.

Self-lumping (SL) based on Newton-Cotes formulae differs from traditional lumping in that SL Newton-Cotes generally does not exactly integrate the gradient operator. Coincidentally, the gradient operator is exactly integrated for linear/quadratic trial spaces using a 2-point/3-point Newton-Cotes formula, respectively. However, for higher degree polynomial trial spaces, the corresponding Newton-Cotes formula does not exactly integrate the gradient operator.

Self-lumping based on either Gauss or Lobatto quadratures exactly integrates the gradient operator in 1-D slab geometry for all degree of polynomial trial spaces; thus, there is no need to distinguish between exact integration and quadrature integration of the gradient operator for the SL Gauss and SL Lobatto schemes.

2.3 Lumping Techniques for the 1-D S_N Neutron Transport Equation with Arbitrary Order DFEM

We now derive the weak form of the 1-D S_N neutron transport equations discretized with DFEM and define the different mass matrix lumping techniques.

2.3.1 Weak Form Derivation

Consider the 1-D slab geometry S_N neutron transport equation:

$$\mu_d \frac{d\psi_d}{dx} + \Sigma_t \psi_d = Q_d, \quad (2.1)$$

where ψ_d is the angular flux [$1/[cm^2 - sec - ster]$] in the μ_d direction, μ_d is the d 'th directional cosine relative to the x -axis, Σ_t is the total interaction cross section [cm^{-1}], and Q_d is a total source (fixed+scattering+fission) angular source in the direction of μ_d [$1/[cm^3 - sec - ster]$]. In all that follows, we consider only non-scattering, non-fissioning media (pure absorbers), thus Q_d will only be non-trivial if a fixed source is present in the problem. The scalar flux, ϕ [$n/cm^2 - sec$], is defined as

$$\phi(x) = 2\pi \int_{-1}^1 \psi(x, \mu_d) d\mu_d. \quad (2.2)$$

For simplicity, we derive the DFEM equations for a single-cell domain, with $x \in [x_L, x_R]$. A known angular flux, $\psi_{in,d}$, is defined on the incoming face of the domain for all μ_d . For $\mu_d > 0$, $\psi_{in,d}$ is defined only at x_L and for $\mu_d < 0$, $\psi_{in,d}$ is defined

at x_R . We begin our derivation by first transforming the physical geometry to a reference element, $s \in [-1, 1]$. This affine transformation is such that:

$$x = \bar{x} + \frac{\Delta x}{2} s, \quad (2.3a)$$

$$dx = \frac{\Delta x}{2} ds, \quad (2.3b)$$

with $\bar{x} = \frac{x_L + x_R}{2}$, and $\Delta x = x_R - x_L$. We seek a numerical approximation to the true angular flux ψ_d using Lagrange polynomials of degree P :

$$\psi_d(s) \approx \tilde{\psi}_d(s) = \sum_{j=1}^{N_P} \psi_{j,d} b_j(s), \quad (2.4)$$

where the $\tilde{\psi}_d(s)$ denotes the numerical approximation. The basis functions b_j are the canonical Lagrange polynomials with interpolatory points s_j ,

$$b_j(s) = \prod_{\substack{k=1 \\ k \neq j}}^{N_P} \frac{s - s_k}{s_j - s_k}, \quad (2.5)$$

and $N_P = P + 1$. To determine the N_P unknown coefficients of Eq. (2.4), we follow a standard discontinuous Galerkin procedure, successively multiplying Eq. (2.1) by weight function b_i and integrating by parts, hence generating N_P moment equations ($1 \leq i \leq N_P$). We assume that the cross sections are constant per cell. Inserting our solution representation $\tilde{\psi}_d$, the i -th moment equation is given by:

$$\begin{aligned} \mu_d \left[b_i(1) \tilde{\psi}_d(1) - b_i(-1) \tilde{\psi}_d(-1) - \int_{-1}^1 \tilde{\psi}_d(s) \frac{db_i}{ds} ds \right] + \frac{\Delta x \Sigma_t}{2} \int_{-1}^1 b_i(s) \tilde{\psi}_d(s) ds \\ = \frac{\Delta x}{2} \int_{-1}^1 b_i(s) Q_d(s) ds. \end{aligned} \quad (2.6)$$

We now introduce the upwind approximation to define the angular flux at the cell edges. For $\mu_d > 0$ the angular flux at the cell interfaces is

$$\tilde{\psi}_d(-1) = \psi_{in,d} \text{ and} \quad (2.7a)$$

$$\tilde{\psi}_d(1) = \sum_{j=1}^{N_P} \psi_{j,d} b_j(1). \quad (2.7b)$$

Similarly for $\mu_d < 0$:

$$\tilde{\psi}_d(-1) = \sum_{j=1}^{N_P} \psi_{j,d} b_j(-1) \text{ and} \quad (2.8a)$$

$$\tilde{\psi}_d(1) = \psi_{in,d}. \quad (2.8b)$$

In Eq. (2.7a) and Eq. (2.8b), $\psi_{in,d}$ is either the known angular flux outflow from the upwind cell or a boundary condition. Inserting the definition of $\tilde{\psi}_d(s)$, Eq. (2.6) becomes, for $\mu_d > 0$,

$$\begin{aligned} \mu_d \left[b_i(1) \left(\sum_{j=1}^{N_P} \psi_{j,d} b_j(1) \right) - b_i(-1) \psi_{in,d} - \int_{-1}^1 \left(\sum_{j=1}^{N_P} \psi_{j,d} b_j(s) \right) \frac{db_i}{ds} ds \right] + \\ \frac{\Delta x \Sigma_t}{2} \int_{-1}^1 b_i(s) \left(\sum_{j=1}^{N_P} \psi_{j,d} b_j(s) \right) ds = \frac{\Delta x}{2} \int_{-1}^1 b_i(s) Q_d(s) ds, \quad (2.9) \end{aligned}$$

and, for $\mu_d < 0$,

$$\begin{aligned} \mu_d \left[b_i(1) \psi_{in,d} - b_i(-1) \left(\sum_{j=1}^{N_P} \psi_{j,d} b_j(-1) \right) - \int_{-1}^1 \left(\sum_{j=1}^{N_P} \psi_{j,d} b_j(s) \right) \frac{db_i}{ds} ds \right] \\ + \frac{\Delta x \Sigma_t}{2} \int_{-1}^1 b_i(s) \left(\sum_{j=1}^{N_P} \psi_{j,d} b_j(s) \right) ds = \frac{\Delta x}{2} \int_{-1}^1 b_i(s) Q_d(s) ds. \quad (2.10) \end{aligned}$$

Considering all of the N_P moment equations at once we can write both Eq. (2.9) and Eq. (2.10) in a single matrix form:

$$\left(\mu_d \mathbf{G} + \frac{\Sigma_t \Delta x}{2} \mathbf{M} \right) \vec{\psi}_d = \frac{\Delta x}{2} \vec{Q}_d + \mu_d \psi_{in,d} \vec{f}. \quad (2.11)$$

In Eq. (2.11) we have made use of the following definitions: the vector of unknowns is given by

$$\vec{\psi}_d = [\psi_{1,d} \ \dots \ \psi_{N_P,d}]^T, \quad (2.12)$$

the mass matrix \mathbf{M} is:

$$\mathbf{M}_{ij} = \int_{-1}^1 b_i(s) b_j(s) ds, \quad (2.13)$$

the fixed source moment vector, \vec{Q}_d , is a column vector of length N_P :

$$\vec{Q}_{d,i} = \int_{-1}^1 b_i(s) Q_d(s) ds, \quad (2.14)$$

and \vec{f} is a column vector of length N_P :

$$\vec{f}_i = \begin{cases} b_i(-1) & \text{for } \mu_d > 0 \\ -b_i(1) & \text{for } \mu_d < 0 \end{cases}. \quad (2.15)$$

\mathbf{G} is a $N_P \times N_P$ matrix which we refer to as the gradient operator. When $\mu_d > 0$, \mathbf{G} is given by:

$$\mathbf{G}_{ij} = b_i(1) b_j(1) - \int_{-1}^1 \frac{db_i}{ds} b_j(s) ds. \quad (2.16a)$$

For $\mu_d < 0$, \mathbf{G} is:

$$\mathbf{G}_{ij} = -b_i(-1) b_j(-1) - \int_{-1}^1 \frac{db_i}{ds} b_j(s) ds. \quad (2.16b)$$

When interpolatory points are not located at the cell interfaces (i.e., at $s = \pm 1$), it can be noted that

1. \vec{f} has N_P non-zero entries and
2. $b_i(\pm 1)b_j(\pm 1) \neq 0$ for all $i, j = 1, \dots, N_P$.

When a Lagrange interpolatory point exists on the cell edges, then \vec{f} has only one non-zero entry and the product $b_i(\pm 1)b_j(\pm 1) \neq 0$ only when $i = j = N_P$ for $\mu_d > 0$ or when $i = j = 1$ for $\mu_d < 0$, as is the case when equally-spaced points or a Lobatto quadrature are used as interpolation points.

We evaluate the integrals of Eq. (2.13) and Eq. (2.16) using a numerical quadrature. A method exactly integrates a quantity when the quadrature rule used to evaluate the integral is accurate for polynomials of degree equal to or greater than the polynomial degree of the integrand. In general, the matrices are dense and their entries are computed as:

$$\mathbf{M}_{ij} \approx \sum_{q=1}^{N_q} w_q b_i(s_q) b_j(s_q), \quad (2.17)$$

$$\mathbf{G}_{ij} \approx sg(\mu_d) b_i(sg(\mu_d)) b_j(sg(\mu_d)) - \sum_{q=1}^{N_q} w_q \left. \frac{db_i}{ds} \right|_{s=s_q} b_j(s_q), \quad (2.18)$$

where N_q is the number of quadrature points to be used, w_q are the weights associated with quadrature points s_q , and $sg(a)$ is the sign function defined as

$$sg(a) = \begin{cases} +1 & \text{if } a > 0 \\ -1 & \text{if } a < 0 \end{cases}. \quad (2.19)$$

2.3.2 Traditional Lumping

The traditional lumping (TL) scheme replaces \mathbf{M} with $\widehat{\mathbf{M}}$, the latter being formed by collapsing row entries onto the main diagonal via the following formula [16]:

$$\widehat{\mathbf{M}}_{ij} = \begin{cases} \sum_{j=1}^{N_P} \mathbf{M}_{ij} & \text{for } i = j \\ 0 & \text{otherwise} \end{cases}. \quad (2.20)$$

2.3.3 Quadrature-Based Lumping

An alternative method of mass matrix lumping restricts the quadrature points to the interpolatory points where:

$$b_i(s_j) = \begin{cases} 1 & \text{if } s_i = s_j \\ 0 & \text{otherwise} \end{cases}, \quad i = 1, \dots, N_P, \quad (2.21)$$

and the quadrature integration of Eq. (2.17) reduces to:

$$\mathbf{M}_{ij} = \begin{cases} w_i & i = j \\ 0 & \text{otherwise} \end{cases}. \quad (2.22)$$

As mentioned previously, we refer to the implicit lumping of Eq. (2.22) as self-lumping (SL). Self-lumping is a method to automatically generate a diagonal mass matrix. We note that self-lumping does not imply that the quadrature formula inexactly integrates the mass matrix.

2.3.4 Source Moment Evaluation

Historically, when discussing lumping techniques, the focus has been on matrix lumping [16] and little attention was paid to lumping source terms. For instance,

consider a δ -shaped volumetric sources (i.e., equal to 0 everywhere except at one given point). In such a case, the evaluation of \vec{Q}_d using quadrature-based self-lumping schemes is an open question. Obviously, quadrature-based schemes cannot evaluate Eq. (2.14) for δ -sources. To address this, we expand the source on a Legendre polynomial basis:

$$\hat{S}_d(s) = \sum_{n=0}^P S_n P_n(s) \quad (2.23a)$$

$$\text{with } S_n = \frac{2n+1}{2} \int_{-1}^1 Q_d(s) P_n(s) ds, \quad (2.23b)$$

and evaluate \vec{Q}_d as follows

$$\vec{Q}_{d,i} = \int_{-1}^1 b_i(s) \hat{S}_d(s) ds. \quad (2.24)$$

Note that if the right-hand-side of Eq. (2.24) is exactly integrated, this is equivalent to exactly integrating Eq. (2.14).

2.4 Quadrature Point Selection

We now discuss the properties of different numerical quadratures as applied to the 1-D DFEM S_N neutron transport equations. We consider three different types of interpolatory points: equally-spaced, Gauss quadrature, and Lobatto quadrature. On the $[-1, 1]$ interval, the $N_P = P + 1$ equally spaced interpolation points for a degree P polynomial trial space are:

$$s_j = -1 + (j-1) \frac{2}{P}, \quad j = 1, \dots, N_P. \quad (2.25)$$

Self-lumping using equally-spaced interpolation points requires numerical integration with closed Newton-Cotes quadrature formulae. The N_P weights, w_j , used for

Newton-Cotes numerical integration at the interpolation points do not follow a concise pattern, so we refer the reader to [25]. The Gauss quadrature points are the N_P roots of the Legendre polynomial, $P_{N_P}(s)$ [25]. The corresponding weights are:

$$w_j = \frac{2}{(1 - s_j^2)} [P'_{N_P}(s_j)]^2 . \quad (2.26)$$

Lobatto quadrature points have fixed endpoints, $s_1 = -1$, $s_{N_P} = 1$. The remaining $N_P - 2$ points are the roots of $P'_{N_P-1}(s)$ [25], with corresponding weights:

$$w_j = \begin{cases} \frac{2}{N_P(N_P-1)} & j = 1, j = N_P \\ \frac{2}{N_P(N_P-1)[P'_{N_P-1}(s_j)]^2} & \text{otherwise} \end{cases} . \quad (2.27)$$

The highest polynomial degree a particular self-lumping quadrature formula exactly integrates is given in Table 2.2 for Newton-Cotes, in Table 2.3 for Gauss, and Table 2.4 for Lobatto quadratures. Also listed in Table 2.2 - Table 2.4 is the maximum polynomial degree of the integrands present in the gradient and mass matrices.

Table 2.2: Accuracy of self-lumping closed Newton-Cotes quadratures for DFEM trial spaces of polynomial degree P .

Polynomial Degree of $\tilde{\psi}$	$N_P =$ $P + 1$	Degree of M integrand	Degree of G integrand	Quadrature Accuracy
1	2	2	1	1
2	3	4	3	3
3	4	6	5	3
4	5	8	7	5
5	6	10	9	5
P	$P + 1$	$2P$	$2P - 1$	Odd $\tilde{\psi}$: P Even $\tilde{\psi}$: $P + 1$

Table 2.3: Accuracy of self-lumping Gauss quadratures for DFEM trial spaces of polynomial degree P .

Polynomial Degree of $\tilde{\psi}$	$N_P =$ $P + 1$	Degree of \mathbf{M} integrand	Degree of \mathbf{G} integrand	Quadrature Accuracy
1	2	2	1	3
2	3	4	3	5
3	4	6	5	7
4	5	8	7	9
5	6	10	9	11
P	$P + 1$	$2P$	$2P - 1$	$2P + 1$

Table 2.4: Accuracy of self-lumping Lobatto quadratures for DFEM trial spaces of polynomial degree P .

Polynomial Degree of $\tilde{\psi}$	$N_P =$ $P + 1$	Degree of \mathbf{M} integrand	Degree of \mathbf{G} integrand	Quadrature Accuracy
1	2	2	1	1
2	3	4	3	3
3	4	6	5	5
4	5	8	7	7
5	6	10	9	9
P	$P + 1$	$2P$	$2P - 1$	$2P - 1$

Since the accuracy of an $N_P = P + 1$ point Gauss quadrature integration exceeds the polynomial degree of the \mathbf{M} and \mathbf{G} integrands for a trial space of degree P , using the SL Gauss scheme will strictly yield the same numerical solution as any DFEM scheme that exactly integrates \mathbf{M} and \mathbf{G} . Thus, the SL Gauss scheme yields the same numerical solution as the Exact DFEM scheme.

For linear and quadratic trial spaces, self-lumping methods using either Lobatto or equally-spaced interpolation points will yield identical solutions. This is a direct result of the two- and three-point Lobatto quadrature formulae being identical to the two- and three-point closed Newton-Cotes quadratures. This equivalence does

not hold for higher degree polynomial trial spaces because the Lobatto quadrature points will no longer correspond to the equally-spaced quadrature points.

By definition, TL uses equally-spaced interpolation points and exactly integrates the gradient operator. For cell-wise constant cross sections, TL is equivalent to a numerical integration scheme that:

1. uses equally-spaced interpolation points,
2. integrates the gradient operator exactly, and
3. uses a Newton-Cotes quadrature restricted to the DFEM interpolation points to compute the mass matrix.

To prove the third point, consider the following. With traditional lumping, \mathbf{M}_{ij} is exactly computed and then a row-sum operation is performed on the rows of \mathbf{M} ; thus the entries of the diagonal mass matrix computed for TL are

$$\begin{aligned}\widehat{\mathbf{M}}_{ii} &= \sum_{j=1}^{N_P} \int_{-1}^1 b_i(s) b_j(s) ds = \int_{-1}^1 b_i(s) \left[\sum_{j=1}^{N_P} b_j(s) \right] ds \\ &= \int_{-1}^1 ds b_i(s) \quad \forall i = 1, \dots, N_P,\end{aligned}\tag{2.28}$$

because $\sum_j^{N_P} b_j(s) = 1 \ \forall s \in [-1, +1]$ by definition. The integral $\int_{-1}^1 ds b_i(s)$ is exactly integrated using a closed Newton-Cotes formula with $N_P = P + 1$ points since $b_i(s)$ is a polynomial of degree P . Finally, when the b_i functions are defined using equally-spaced points, the use of a closed Newton-Cotes formula with N_P points yields

$$\widehat{\mathbf{M}}_{ii} = \int_{-1}^1 ds b_i(s) = \sum_{q=1}^{N_P} w_q b_i(s_q) = w_i,\tag{2.29}$$

because $b_i(s_q) = \delta_{iq}$. Thus, the diagonal mass matrix computed using TL contains the closed Newton-Cotes weights as diagonal entries and is equivalent to approximating

\mathbf{M} using closed Newton-Cotes quadrature in Eq. (2.22). We also numerically verify this in Table 2.5 for polynomial degrees up to 4. For linear and quadratic trial

Table 2.5: Equivalence of traditional lumping and closed Newton-Cotes quadrature approximation of the mass matrix.

P	Exact Integration of \mathbf{M}	Row Sums of \mathbf{M}	Newton-Cotes w with $P + 1$ points
1	$\begin{bmatrix} \frac{2}{3} & \frac{1}{3} \\ \frac{1}{3} & \frac{2}{3} \end{bmatrix}$	$\begin{bmatrix} 1 \\ 1 \end{bmatrix}$	$\begin{bmatrix} 1 \\ 1 \end{bmatrix}$
2	$\begin{bmatrix} \frac{4}{15} & \frac{2}{15} & -\frac{1}{15} \\ \frac{2}{15} & \frac{16}{15} & \frac{1}{15} \\ -\frac{1}{15} & \frac{2}{15} & \frac{4}{15} \end{bmatrix}$	$\begin{bmatrix} \frac{1}{3} \\ \frac{4}{3} \\ \frac{1}{3} \end{bmatrix}$	$\begin{bmatrix} \frac{1}{3} \\ \frac{4}{3} \\ \frac{1}{3} \end{bmatrix}$
3	$\begin{bmatrix} \frac{16}{105} & \frac{38}{280} & -\frac{3}{70} & \frac{19}{840} \\ \frac{33}{280} & \frac{27}{35} & -\frac{27}{280} & -\frac{3}{70} \\ -\frac{3}{70} & -\frac{27}{280} & \frac{27}{35} & \frac{33}{280} \\ \frac{19}{840} & -\frac{3}{70} & \frac{38}{280} & \frac{16}{105} \end{bmatrix}$	$\begin{bmatrix} \frac{1}{4} \\ \frac{3}{4} \\ \frac{3}{4} \\ \frac{1}{4} \end{bmatrix}$	$\begin{bmatrix} \frac{1}{4} \\ \frac{3}{4} \\ \frac{3}{4} \\ \frac{1}{4} \end{bmatrix}$
4	$\begin{bmatrix} \frac{292}{2835} & \frac{296}{2835} & -\frac{58}{945} & \frac{8}{405} & -\frac{29}{2835} \\ \frac{296}{2835} & \frac{256}{405} & -\frac{128}{945} & \frac{256}{2835} & \frac{8}{405} \\ -\frac{58}{945} & -\frac{128}{945} & \frac{208}{315} & -\frac{128}{945} & -\frac{58}{945} \\ \frac{8}{405} & \frac{256}{2835} & -\frac{128}{945} & \frac{256}{405} & \frac{296}{2835} \\ -\frac{29}{2835} & \frac{8}{405} & -\frac{58}{945} & \frac{296}{2835} & \frac{292}{2835} \end{bmatrix}$	$\begin{bmatrix} \frac{7}{45} \\ \frac{32}{45} \\ \frac{4}{15} \\ \frac{32}{45} \\ \frac{7}{45} \end{bmatrix}$	$\begin{bmatrix} \frac{7}{45} \\ \frac{32}{45} \\ \frac{4}{15} \\ \frac{32}{45} \\ \frac{7}{45} \end{bmatrix}$

spaces, the 2-point and 3-point Newton-Cotes quadrature formulae exactly integrate the gradient operator, as shown in Table 2.2. Thus, for linear and quadratic trial spaces, schemes that use

1. equally-spaced interpolation points and traditional lumping,
2. equally-spaced interpolation points and self-lumping numerical integration, or
3. Lobatto quadrature interpolation points and self-lumping numerical integration,

will yield identical solutions.

2.5 Numerical Results

In this Section, we present numerical results for two 1-D slab problems. For the first problem, we consider a source-free pure absorber with vacuum boundary conditions on the right, a known angular flux $\psi_{in,d}$ incident on the left face, and a spatially constant total cross section Σ_t . The second problem consists of a slab with vacuum boundary conditions on both sides, no scattering, constant Σ_t , and a fixed δ -source.

For $\mu_d > 0$, the numerical approximations to the angular flux near the cell inflow and outflow are as follows:

$$\tilde{\psi}_{in,d} = \sum_{j=1}^{N_P} \psi_{j,d} b_j(-1) \text{ and} \quad (2.30)$$

$$\tilde{\psi}_{out,d} = \sum_{j=1}^{N_P} \psi_{j,d} b_j(1). \quad (2.31)$$

Regardless of the sign of μ_d , the numerical approximation to the cell average angular flux is defined as:

$$\tilde{\psi}_{A,d} = \frac{1}{2} \sum_{j=1}^{N_P} w_j \psi_{j,d}. \quad (2.32)$$

We used the following quadrature weight normalization: $\sum_{j=1}^{N_P} w_j = 2$. In Eq. (2.30), Eq. (2.31), and Eq. (2.32), $\psi_{j,d}$ are the components of $\vec{\psi}_d$, the numerical solution obtained by solving Eq. (2.11). Hence, the numerical angular flux solution of any of the previously discussed DFEM schemes can be obtained as a function of h , the number of mean free paths divided by μ_d ,

$$h = \frac{\Sigma_t \Delta x}{\mu_d}. \quad (2.33)$$

2.5.1 Incident Flux Single-Cell Outflow Comparison

For the incident-flux problem, the analytical solution of Eq. (2.1) is:

$$\psi(x, \mu_d) = \begin{cases} \psi_{in,d} \exp \left[-\frac{\Sigma_t(x-x_L)}{\mu_d} \right] & \text{for } \mu_d > 0 \\ 0 & \text{for } \mu_d < 0 \end{cases} . \quad (2.34)$$

The analytic angular flux outflow, $\psi_{out,d} = \psi(x_R, \mu_d)$, is:

$$\psi_{out,d} = \psi_{in,d} \exp[-h] . \quad (2.35)$$

Similarly, the analytic average angular flux within the cell, $\psi_{A,d}$, is:

$$\psi_{A,d} = \frac{1}{\Delta x} \int_{x_L}^{x_R} \psi(x, \mu_d) dx = \frac{\psi_{in,d}}{h} (1 - \exp[-h]) . \quad (2.36)$$

The solution components are given by

$$\vec{\psi}_d = \psi_{in,d} \left(\mathbf{G} + \frac{h}{2} \mathbf{M} \right)^{-1} \vec{f} , \quad (2.37)$$

which allows us to compare the various choices of interpolatory points and numerical integration strategies solely as a function of h .

Figures 2.1-2.4 show the numerically calculated cell outflow, $\tilde{\psi}_{out,d}$, as a function of h for all methods considered. All methods converge to the analytical solution as $h \rightarrow 0$, thus we have zoomed in the range where the methods visually differ (i.e., $h \geq 1$). We observe that:

- SL Gauss yields strictly positive outflows for even degree polynomial trial spaces,

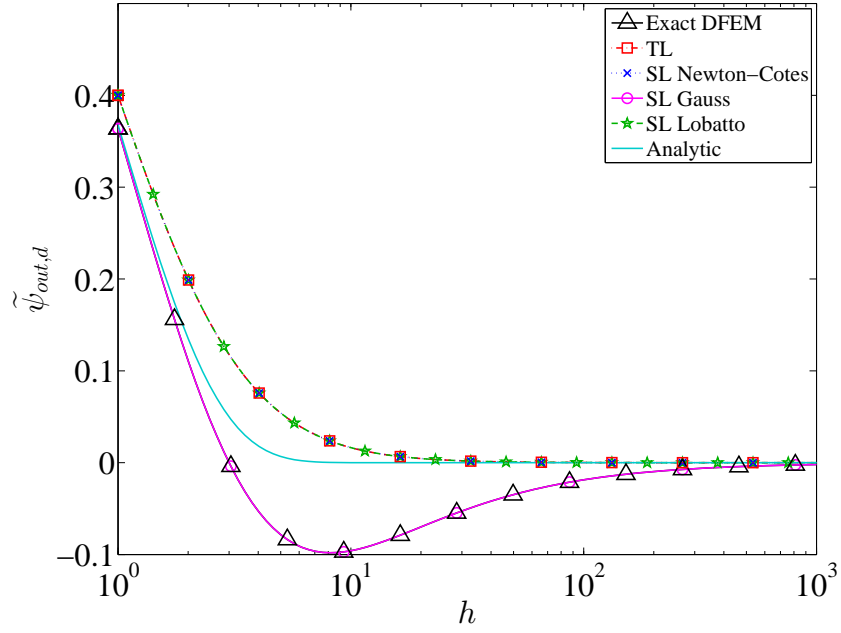


Figure 2.1: Numerical outflow values as a function of h , for a single cell homogeneous absorber problem with a linear DFEM trial space.

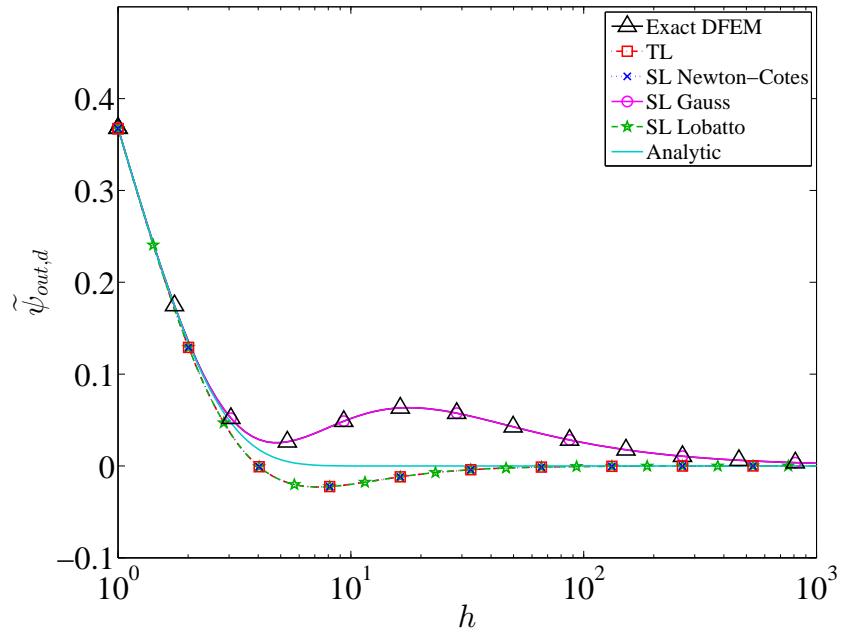


Figure 2.2: Numerical outflow values as a function of h , for a single cell homogeneous absorber problem with a quadratic DFEM trial space.

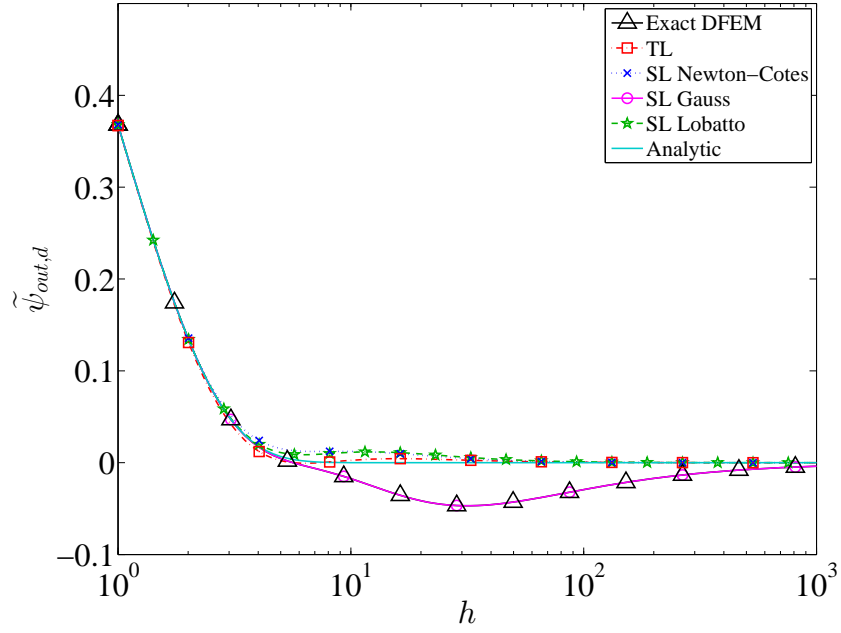


Figure 2.3: Numerical outflow values as a function of h , for a single cell homogeneous absorber problem with a cubic DFEM trial space.

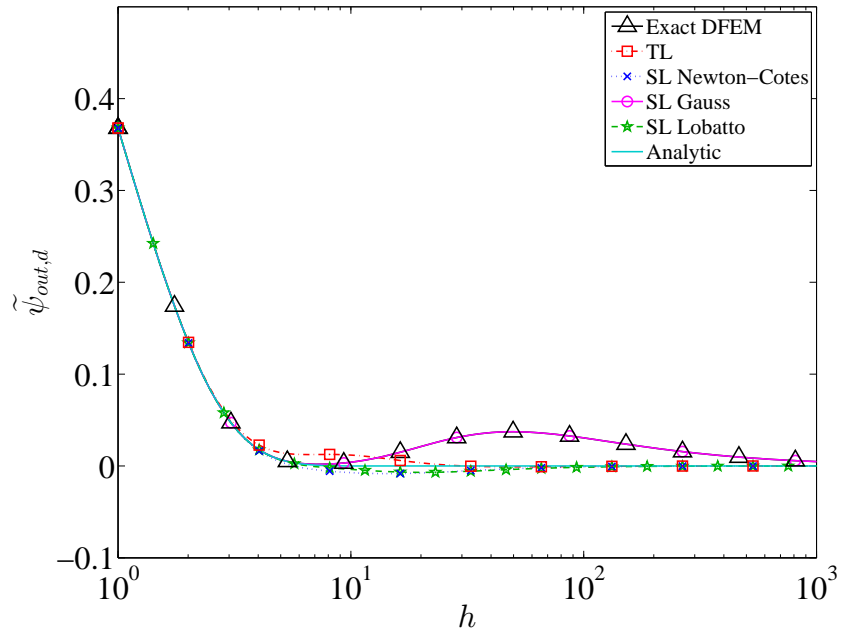


Figure 2.4: Numerical outflow values as a function of h , for a single cell homogeneous absorber problem with a quartic DFEM trial space.

- SL Lobatto and SL Newton-Cotes yield strictly positive outflows for odd degree polynomial trial spaces, and
- TL yields strictly positive outflows only for a linear trial space.

We also numerically verify the remarks made of Section 2.4, that is:

- SL Gauss is equivalent to Exact DFEM,
- SL Lobatto, SL Newton-Cotes, and TL are equivalent for linear and quadratic trial spaces, and
- for even degree trial spaces, the outflow value computed by SL Gauss is not monotonically decreasing as a function of h for cells of intermediate optical thickness (the same was noted in [20] for Exact DFEM).

2.5.2 Fixed Source Single-Cell Inflow Comparison

As noted in [15], it is possible for LDFEM to yield negative solutions near cell inflows for source driven problems. In this second problem, we use a δ -source:

$$Q_d(x) = \begin{cases} \delta(x - x_o) & \text{for } \mu_d > 0 \\ 0 & \text{for } \mu_d < 0 \end{cases}, \quad (2.38)$$

$x \in [-1, 1]$, and $-1 \leq x_o \leq 1$. The analytic solution to this problem for $\mu_d > 0$ is:

$$\psi(x, \mu_d) = \begin{cases} \exp \left[-\frac{\Sigma_t(x-x_o)}{\mu_d} \right] & x \geq x_o \\ 0 & x < x_o \end{cases}. \quad (2.39)$$

(For $\mu_d < 0$, $\psi(x, \mu_d) = 0$.) We now examine the numerical approximation to the angular flux near the cell inflow, $\tilde{\psi}_{in,d}$, for various integration schemes, trial space degrees, and as a function of the ratio of the first Legendre moment of the source, S_1 ,

to the zero-th Legendre moment of the source, S_0 . Note that the physical range of that ratio, $\frac{S_1}{S_0}$, is $[-3, 3]$, corresponding to a δ -source at the left cell edge ($\frac{S_1}{S_0} = -3$) or at the right edge ($\frac{S_1}{S_0} = 3$).

We first consider the case of a vacuum ($\Sigma_t = 0$), thus only testing the effect of quadrature accuracy in evaluating \vec{Q}_d and \mathbf{G} . In Figs. 2.5-2.8, we plot $\tilde{\psi}_{in,d}$ for three schemes:

1. Lobatto quadrature, which is exact for \mathbf{G} and approximate for the source moments, Eq. (2.24) ,
2. Gauss quadrature: which is exact for both \mathbf{G} and the source moments, and
3. Newton-Cotes quadrature: which is approximate for both \mathbf{G} and the source moments.

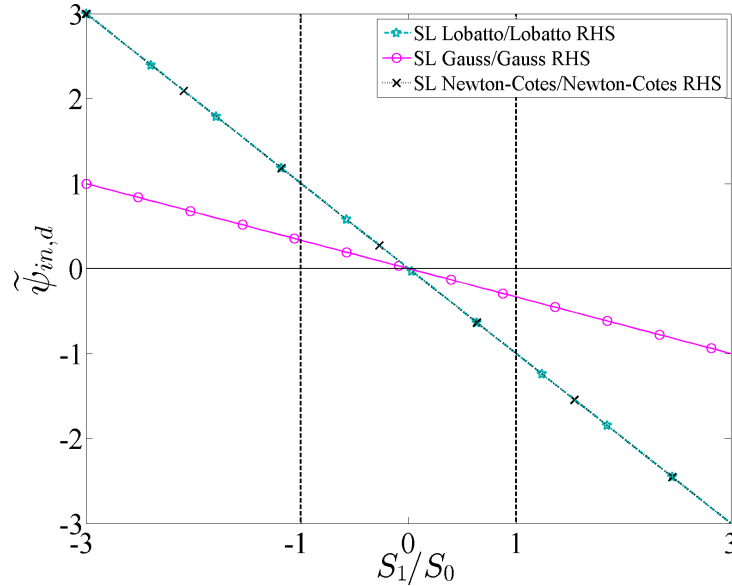


Figure 2.5: Numerical inflow values as a function of $\frac{S_1}{S_0}$ for a single cell (vacuum case) with a δ -shaped source, using linear DFEM.

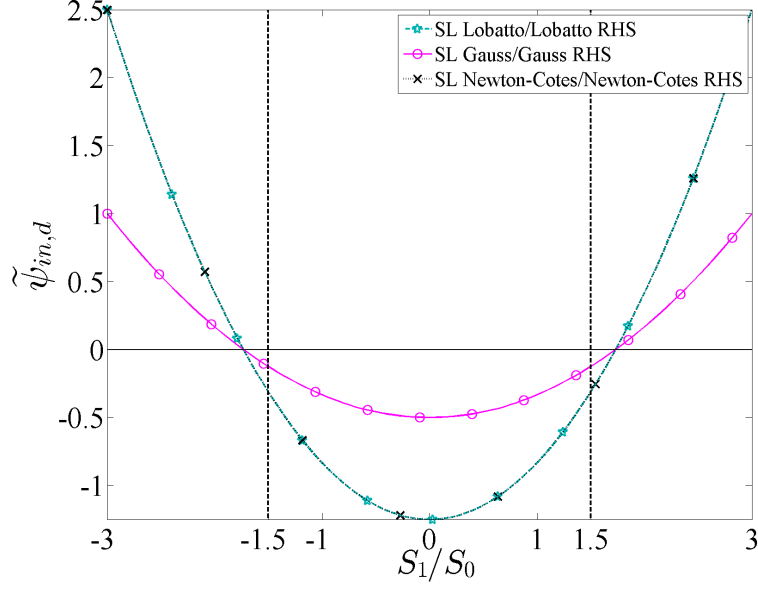


Figure 2.6: Numerical inflow values as a function of $\frac{S_1}{S_0}$ for a single cell (vacuum case) with a δ -shaped source, using quadratic DFEM.

The dotted vertical lines in Figs. 2.5-2.8 correspond to the extrema values of $\frac{S_1}{S_0}$ that yield a strictly positive polynomial source representation of degree P (indeed, the degree- P Legendre expansion of the δ -source is not everywhere positive for a wide range of possible $\frac{S_1}{S_0}$ that are physically realizable). For all trial space degrees, the Gauss scheme exhibits less negativity than either of the other two schemes. The dramatic difference between the Gauss scheme and the Lobatto scheme is solely due to the quadrature formula used to evaluate \vec{Q}_d since both schemes exactly integrate \mathbf{G} . The Newton-Cotes scheme exhibits less severe negativities than the Lobatto scheme but is less robust than the Gauss scheme. Given the results shown in Figs. 2.5-2.8, we conclude that the most robust schemes exactly integrate the source moments, Eq. (2.24).

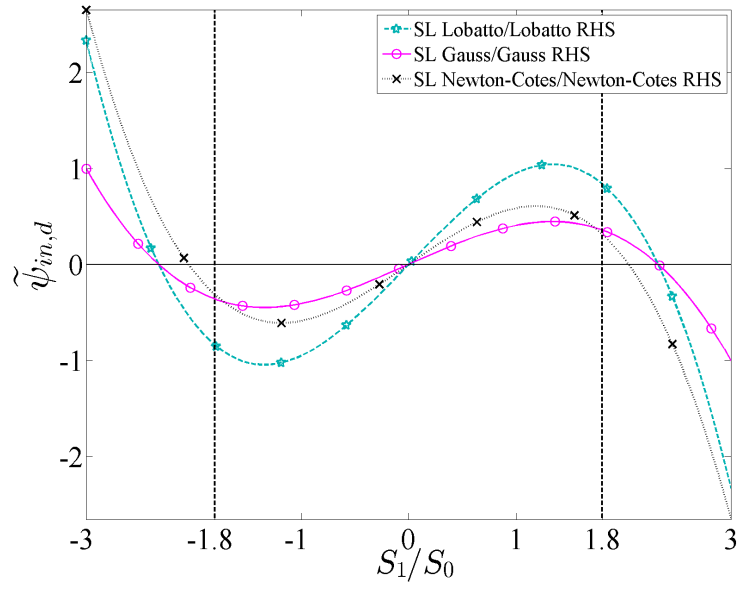


Figure 2.7: Numerical inflow values as a function of $\frac{S_1}{S_0}$ for a single cell (vacuum case) with a δ -shaped source, using cubic DFEM.

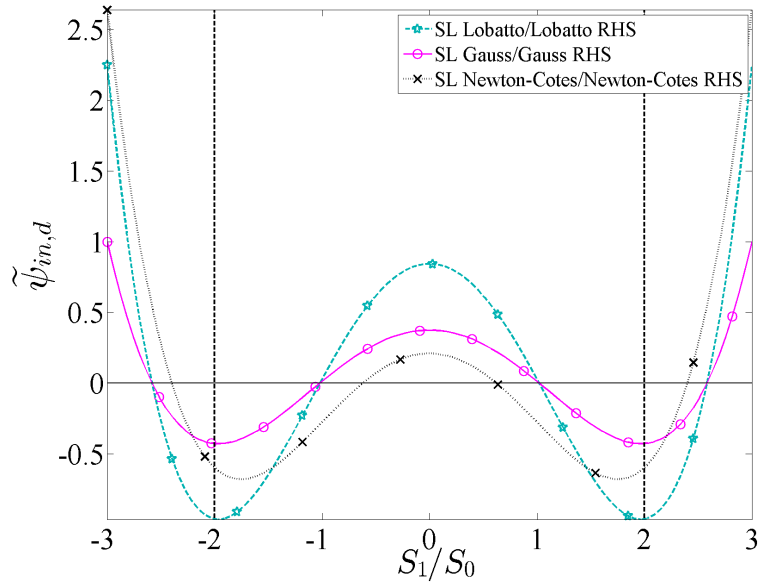


Figure 2.8: Numerical inflow values as a function of $\frac{S_1}{S_0}$ for a single cell (vacuum case) with a δ -shaped source, using quartic DFEM.

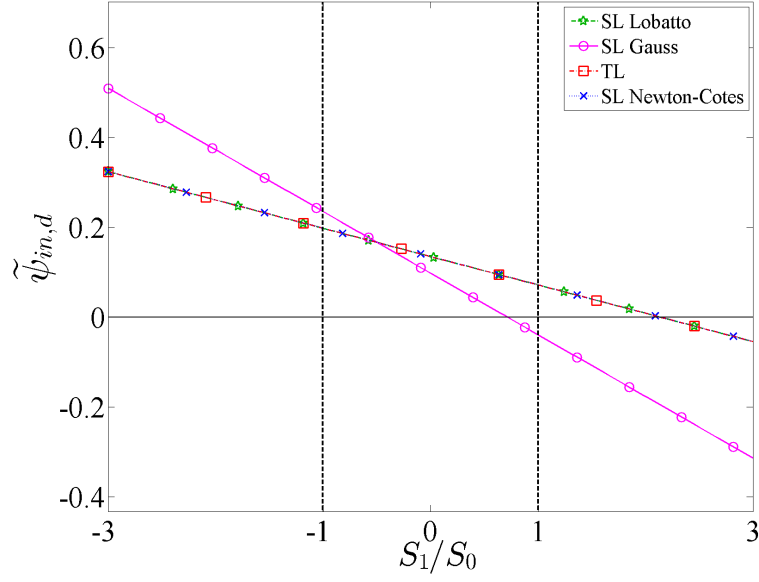


Figure 2.9: Numerical inflow values as a function of $\frac{S_1}{S_0}$, for a single cell (absorber case) with a δ -shaped source, using linear DFEM.

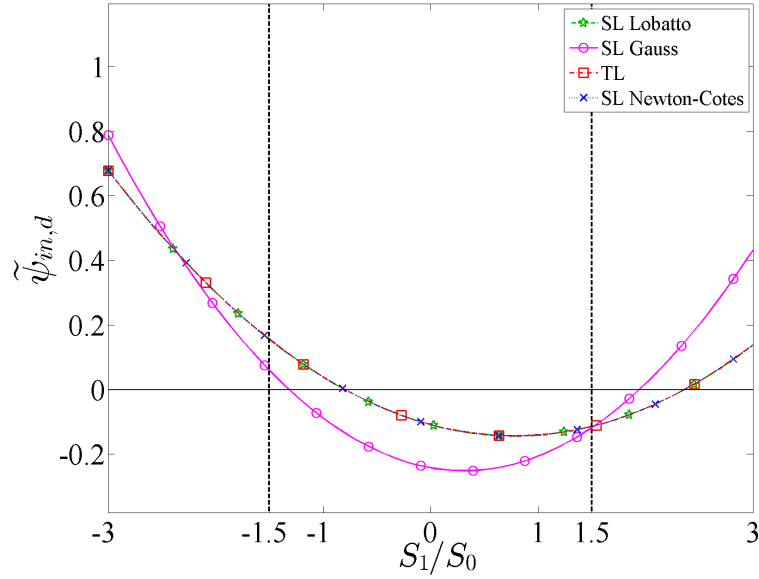


Figure 2.10: Numerical inflow values as a function of $\frac{S_1}{S_0}$, for a single cell (absorber case) with a δ -shaped source, using quadratic DFEM.

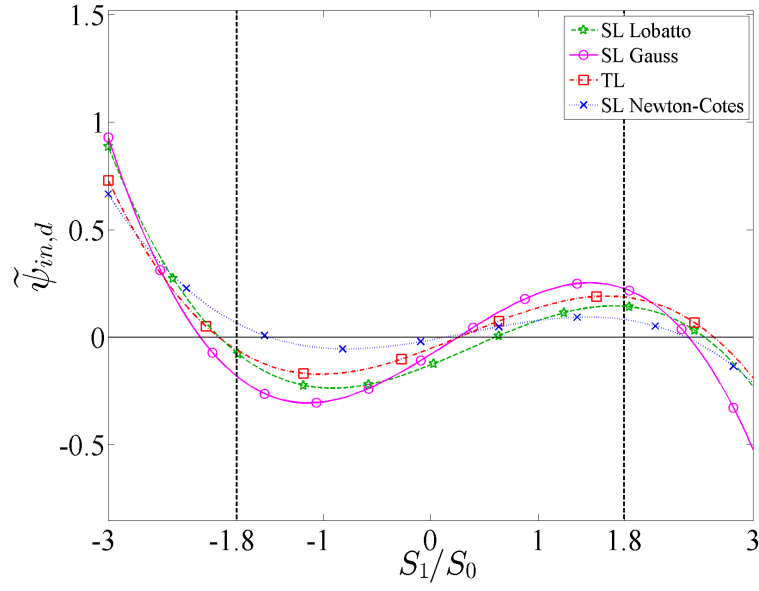


Figure 2.11: Numerical inflow values as a function of $\frac{S_1}{S_0}$, for a single cell (absorber case) with a δ -shaped source, using cubic DFEM.

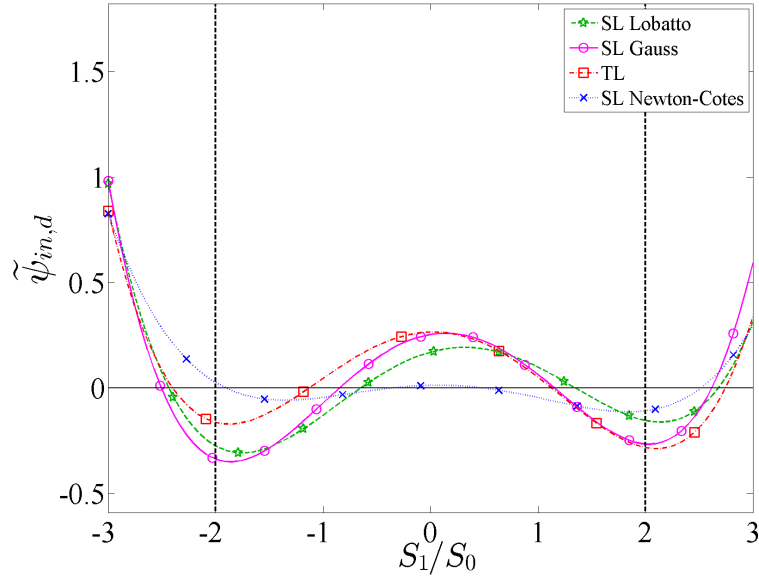


Figure 2.12: Numerical inflow values as a function of $\frac{S_1}{S_0}$, for a single cell (absorber case) with a δ -shaped source, using quartic DFEM.

In Figs. 2.9-2.12, we again examine the positivity of $\tilde{\psi}_{in,d}$, but for a non-vacuum case. Total cell optical thickness was chosen to be 5 mean free paths in Figs. 2.9-2.12 because this value led to the clearest plots. The relative behaviors observed do not change with cell optical thickness, but using a thicker domain reduces the magnitude for the values of $\tilde{\psi}_{in,d}$. All methods in Figs. 2.9-2.12 exactly integrate Eq. (2.24). Regardless of trial space chosen, all schemes exhibit some negativities, but the SL Gauss scheme exhibits the greatest negativities and oscillations. The SL Newton-Cotes scheme presents the least severe negativities.

2.5.3 Single-Cell Taylor Series Analysis

Next, we perform a local truncation error analysis by comparing the Taylor series expansions for the exact and numerical angular fluxes as a function of powers of h for the source-free, incident flux pure absorber problem. Matlab [27] has been employed to perform the symbolic Taylor series expansions about $h = 0$. We denote the Taylor-expanded quantities using the subscript T . The expansions for the analytical inflow, cell average, and outflow are given below:

$$\psi_{in,d,T} = \psi_{in,d} \tag{2.40a}$$

$$\psi_{A,d,T} = \psi_{in,d} \left(1 - \frac{h}{2} + \frac{h^2}{6} - \frac{h^3}{24} + \frac{h^4}{120} - \frac{h^5}{720} \dots \right) \tag{2.40b}$$

$$\psi_{out,d,T} = \psi_{in,d} \left(1 - h + \frac{h^2}{2} - \frac{h^3}{6} + \frac{h^4}{24} - \frac{h^5}{120} \dots \right). \tag{2.40c}$$

The Taylor expansions of the numerical analogues to the quantities in Eqs. (2.40) depend on the trial space polynomial degree, the choice of interpolatory points, and the numerical integration strategy. For brevity, we omit giving these numerical analogues. Table 2.6 gives the lowest order term for the difference between $\psi_{in,d,T}$ and the numerical analogs for the Exact DFEM and TL schemes. The same information

for the SL Newton-Cotes, SL Gauss, and SL Lobatto schemes is given in Table 2.7. The differences between $\psi_{A,d,T}$ and the respective numerical analogs are given in Table 2.8 for Exact DFEM and TL, and Table 2.9 for the SL Newton-Cotes, SL Gauss, and SL Lobatto schemes. Differences between $\psi_{out,d,T}$ and the corresponding numerical analogs are given in Table 2.10 for Exact DFEM and TL and Table 2.11 gives the lowest order difference between the SL Newton-Cotes, SL Gauss, and SL Lobatto approximations of $\psi_{out,d,T}$. In Tables 2.6-2.11 all entries are listed as $q(C)$, to be read as “the difference between the analytic Taylor expansion and the numeric analog is Ch^q with $h = \Sigma_t \Delta x / \mu$ ”. Entries of “Machine Precision” in Tables 2.6-2.11 are meant to indicate that the difference between the analytic Taylor expansion and Taylor expansion of the numerical approximation was inconclusive due to all coefficients being within machine precision.

Table 2.6: Local truncation error analysis in $\tilde{\psi}_{in,d}$ for a single cell problem with constant cross section, for Exact DFEM and TL.

Polynomial Degree of $\tilde{\psi}$	Exact DFEM	TL
1	2 (2×10^{-1})	2 (5×10^{-1})
2	3 (2×10^{-2})	3 (4×10^{-2})
3	4 (1×10^{-3})	2 (7×10^{-2})
4	5 (7×10^{-5})	3 (1×10^{-2})
5	6 (3×10^{-6})	2 (5×10^{-2})
6	7 (1×10^{-7})	3 (1×10^{-2})
7	8 (4×10^{-9})	2 (5×10^{-2})

Table 2.7: Local truncation error analysis in $\tilde{\psi}_{in,d}$ for a single cell problem with constant cross section, for SL Newton-Cotes, SL Gauss, and SL Lobatto.

Polynomial Degree of $\tilde{\psi}$	SL Newton-Cotes	SL Gauss	SL Lobatto
1	2 (5×10^{-1})	2 (2×10^{-1})	2 (5×10^{-1})
2	3 (4×10^{-2})	3 (2×10^{-2})	3 (4×10^{-2})
3	2 (1×10^{-1})	4 (1×10^{-3})	4 (3×10^{-3})
4	3 (1×10^{-2})	5 (7×10^{-5})	5 (1×10^{-4})
5	2 (6×10^{-2})	6 (3×10^{-6})	6 (7×10^{-6})
6	3 (9×10^{-3})	7 (1×10^{-7})	7 (3×10^{-7})
7	2 (4×10^{-2})	8 (4×10^{-9})	8 (8×10^{-9})

Table 2.8: Local truncation error analysis in $\tilde{\psi}_{A,d}$ for a single cell problem with constant cross section, for Exact DFEM and TL.

Polynomial Degree of $\tilde{\psi}$	Exact DFEM	TL
1	3 (1×10^{-2})	2 (2×10^{-1})
2	5 (1×10^{-4})	4 (2×10^{-3})
3	7 (7×10^{-7})	3 (3×10^{-3})
4	9 (2×10^{-9})	5 (8×10^{-5})
5	11 (5×10^{-12})	3 (1×10^{-3})
6	13 (7×10^{-15})	5 (7×10^{-5})
7	Machine Precision	3 (1×10^{-3})

Table 2.9: Local truncation error analysis in $\tilde{\psi}_{A,d}$ for a single cell problem with constant cross section, for SL Newton-Cotes, SL Gauss, and SL Lobatto

Polynomial Degree of $\tilde{\psi}$	SL Newton-Cotes	SL Gauss	SL Lobatto
1	2 (2×10^{-1})	3 (1×10^{-2})	2 (2×10^{-1})
2	4 (2×10^{-3})	5 (1×10^{-4})	4 (2×10^{-3})
3	4 (6×10^{-4})	7 (7×10^{-7})	6 (1×10^{-5})
4	6 (8×10^{-6})	9 (2×10^{-9})	8 (5×10^{-8})
5	6 (2×10^{-6})	11 (5×10^{-12})	10 (1×10^{-10})
6	8 (2×10^{-8})	13 (7×10^{-15})	12 (2×10^{-13})
7	8 (3×10^{-9})	Machine Precision	Machine Precision

Table 2.10: Local truncation error analysis in $\tilde{\psi}_{out,d}$ for a single cell with constant cross section, for Exact DFEM and TL.

Polynomial Degree of $\tilde{\psi}$	Exact DFEM	TL
1	4 (1×10^{-2})	3 (2×10^{-1})
2	6 (1×10^{-4})	5 (2×10^{-3})
3	8 (7×10^{-7})	4 (3×10^{-3})
4	10 (2×10^{-9})	6 (1×10^{-2})
5	12 (5×10^{-12})	4 (1×10^{-3})
6	14 (7×10^{-15})	6 (7×10^{-5})
7	Machine Precision	4 (1×10^{-3})

Table 2.11: Local truncation error analysis in $\tilde{\psi}_{out,d}$ for a single cell with constant cross section, for SL Newton-Cotes, SL Gauss, and SL Lobatto.

Polynomial Degree of $\tilde{\psi}$	SL Newton-Cotes	SL Gauss	SL Lobatto
1	3 (2×10^{-1})	4 (1×10^{-2})	3 (2×10^{-1})
2	5 (2×10^{-3})	6 (1×10^{-4})	5 (2×10^{-3})
3	5 (6×10^{-4})	8 (7×10^{-7})	7 (1×10^{-5})
4	7 (8×10^{-6})	10 (2×10^{-9})	9 (5×10^{-8})
5	7 (2×10^{-6})	12 (5×10^{-12})	11 (1×10^{-10})
6	9 (2×10^{-8})	14 (7×10^{-15})	13 (2×10^{-13})
7	9 (3×10^{-9})	Machine Precision	Machine Precision

This local truncation error analysis illustrates the following.

1. Exact DFEM and SL Gauss, which are equivalent, exactly integrate the mass matrix, and are the most accurate,
2. TL does not guarantee increasing order of accuracy by using higher degree polynomial trial spaces,

3. TL converges at most third or fifth order for $\tilde{\psi}_{A,d}$ and fourth or sixth order for $\tilde{\psi}_{out,d}$ for odd or even polynomial trial spaces, respectively,
4. SL Newton-Cotes increases in accuracy with higher degree polynomial trial spaces, but only for $\tilde{\psi}_{out,d}$ and $\tilde{\psi}_{A,d}$,
5. TL and SL Newton-Cotes are at most second order or third order accurate for $\tilde{\psi}_{in,d}$ for odd or even polynomial trial spaces, respectively,
6. SL Gauss is order $2P+1$ accurate in calculating $\tilde{\psi}_{A,d}$ and order $2P+2$ accurate in calculating $\tilde{\psi}_{out,d}$,
7. SL Lobatto is order $2P$ accurate in calculating $\tilde{\psi}_{A,d}$ and order $2P+1$ in calculating $\tilde{\psi}_{out,d}$,
8. SL Gauss, SL Lobatto, and Exact DFEM are accurate to order $P+1$ in calculating $\tilde{\psi}_{in,d}$, and
9. SL Gauss is more accurate than SL Lobatto (smaller error constant) in computing $\tilde{\psi}_{in,d}$, but not an order of h .

2.5.4 Convergence Rates for Spatially Discretized 1-D Domains

Here, we consider a homogeneous pure absorber material placed in a 1-D slab configuration and uniformly mesh the domain using N_{cells} cells. We use: $x \in [0, 10 \text{ cm}]$, $\Sigma_t = 1 \text{ [cm}^{-1}\text{]}$, no external sources, vacuum conditions on the right face of the slab, and a normally incident unit beam on the left face. The analytical solution to this problem is trivial to obtain:

$$\psi(x, \mu_d) = \begin{cases} \exp[-\Sigma_t x] & \mu_d = 1 \\ 0 & \text{otherwise} \end{cases}. \quad (2.41)$$

The L_2 norm of the error is:

$$E_\psi = \sqrt{\sum_{i=1}^{N_{cells}} \int_{x_{i-1/2}}^{x_{i+1/2}} \left(\psi(x, \mu_d) - \tilde{\psi}_{d,i}(x) \right)^2 dx}, \quad (2.42)$$

where we recall that $\tilde{\psi}_{d,i}(x)$ is the DFEM approximation of the angular flux in cell i . To evaluate the above integral, we use a high-order Gauss quadrature set $(x_{f,q}, w_{f,q})$ that employs a large number of quadrature points:

$$E_\psi \approx \sqrt{\sum_{i=1}^{N_{cells}} \frac{\Delta x_i}{2} \sum_{q=1}^{N_{gf}} w_{f,q} \left(\psi(x_{f,q}, \mu_d) - \tilde{\psi}_d(x_{f,q}) \right)^2}. \quad (2.43)$$

Values of E_ψ shown here are calculated using $N_{gf} = 10$. In addition to the L_2 error, we also present the cell average angular flux error, E_{ψ_A} , defined as

$$E_{\psi_A} = \sqrt{\sum_{i=1}^{N_{cells}} \Delta x_i \left(\psi_{A,d,i} - \tilde{\psi}_{A,d,i} \right)^2}, \quad (2.44)$$

and the cell outflow error, $E_{\psi_{out}}$, given by:

$$E_{\psi_{out}} = \sqrt{\sum_{i=1}^{N_{cells}} \Delta x_i \left(\psi(x_{i+1/2}, \mu_d) - \tilde{\psi}_{out,d,i} \right)^2}. \quad (2.45)$$

In Eq. (2.43), Eq. (2.44), and Eq. (2.45), Δx_i is the cell width of cell i and $\psi_{A,d,i}$ is the exact cell-averaged angular flux in cell i , which, for $\mu_d = 1$, is simply:

$$\psi_{A,d,i} = \exp[-\Sigma_t x_{i-1/2}] \frac{1}{\Delta x_i} (1 - \exp[-\Sigma_t \Delta x_i]). \quad (2.46)$$

In the plots that follow, we omit plotting the errors of Exact DFEM since the Exact DFEM solution is identical to that of SL Gauss. For linear and quadratic polynomi-

als, we plot only SL Lobatto and omit plotting TL and SL Newton-Cotes since these methods yield identical solutions for linear and quadratic trial spaces. Figures

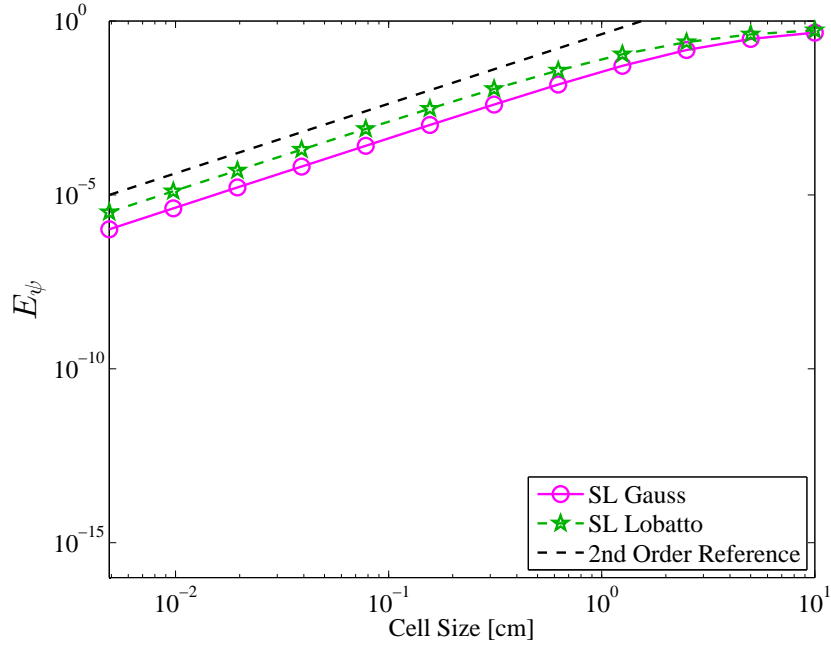


Figure 2.13: Convergence rate of the L_2 norm of the error, E_ψ , as a function of the mesh cell size for a pure absorber discretized with linear DFEM.

2.13-2.16 mirror the results of Table 2.6 and Table 2.7, which is expected since the convergence rate of E_ψ will be limited by the slowest converging local approximation which is $\tilde{\psi}_{in,d}$. Similarly, Figs. 2.17-2.20 are the multiple-cell analogue of the local truncation error analysis of $\tilde{\psi}_{A,d}$ given in Table 2.8 and Table 2.9. $E_{\psi_{out}}$, as shown in Figs. 2.21-2.24, does not converge at the local truncation error rates of Table 2.10 and Table 2.11.

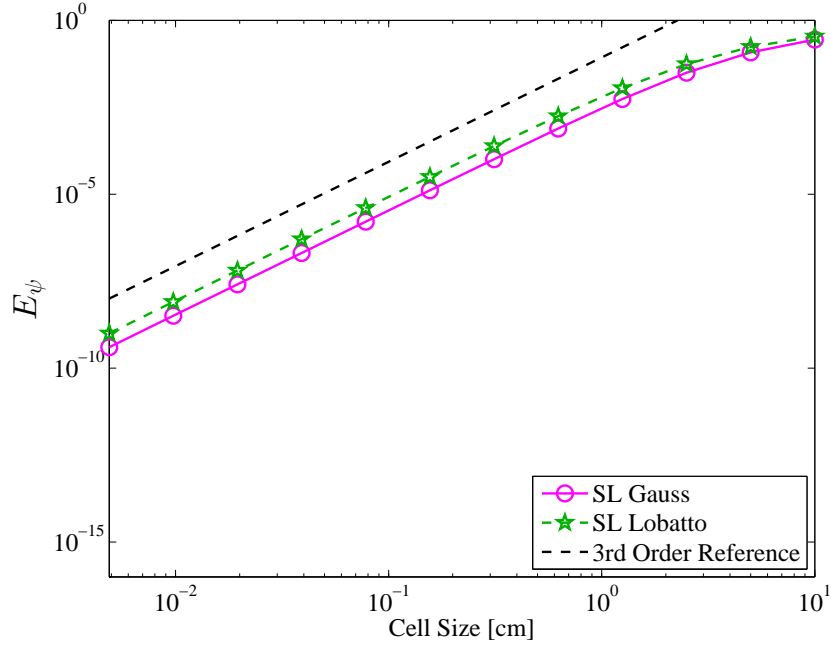


Figure 2.14: Convergence rate of the L_2 norm of the error, E_ψ , as a function of the mesh cell size for a pure absorber discretized with quadratic DFEM.

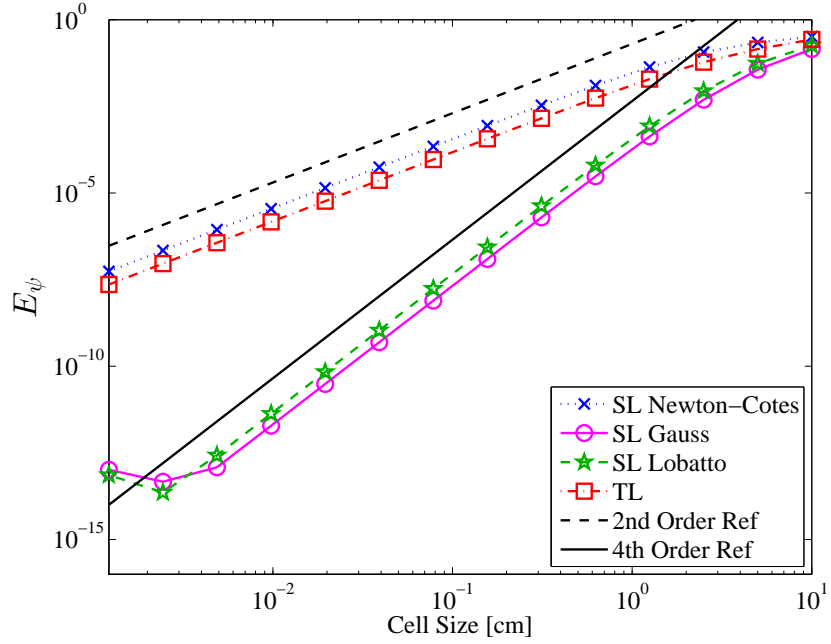


Figure 2.15: Convergence rate of the L_2 norm of the error, E_ψ , as a function of the mesh cell size for a pure absorber discretized with cubic DFEM.

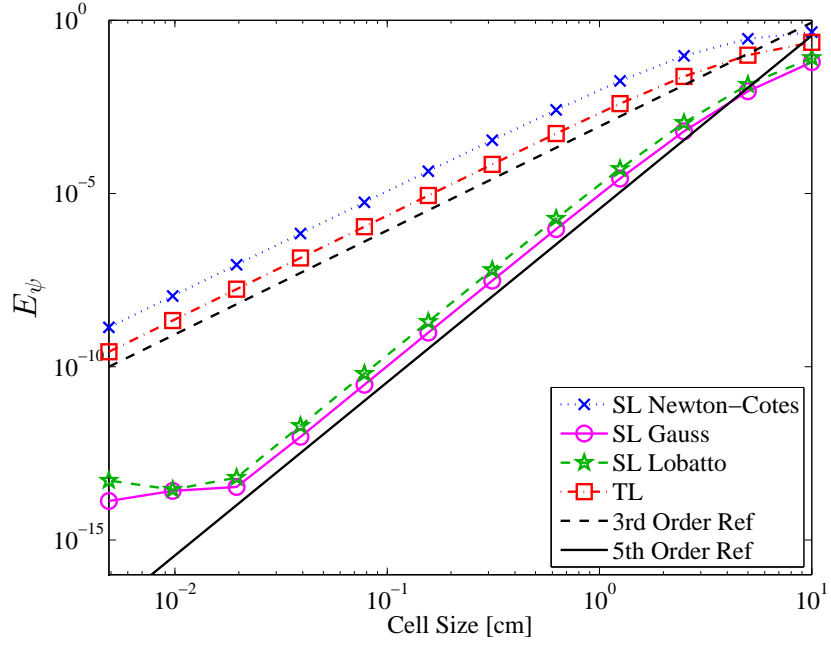


Figure 2.16: Convergence rate of the L_2 norm of the error, E_ψ , as a function of the mesh cell size for a pure absorber discretized with quartic DFEM.

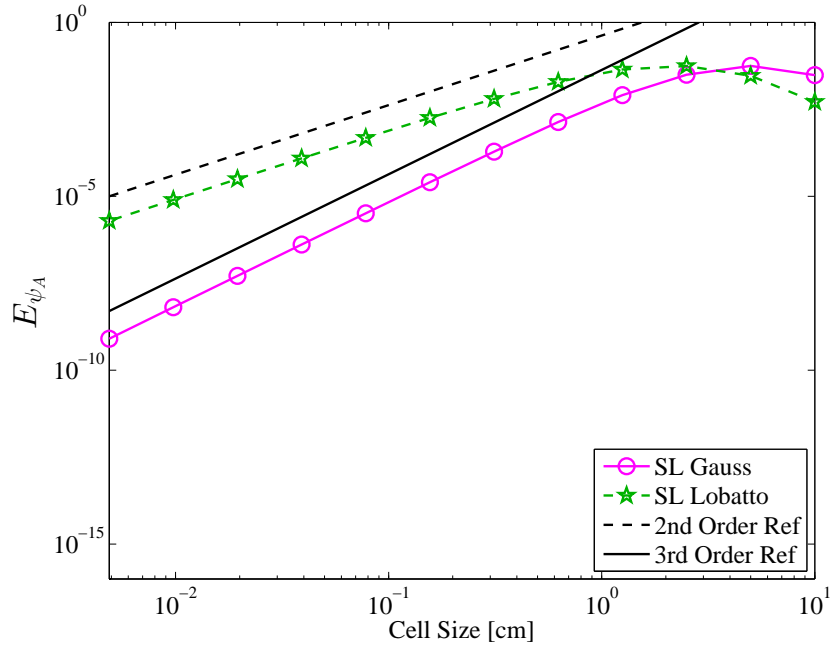


Figure 2.17: Convergence rate for $E_{\psi,A}$ as a function of the mesh cell size for a homogeneous pure absorber and linear DFEM.

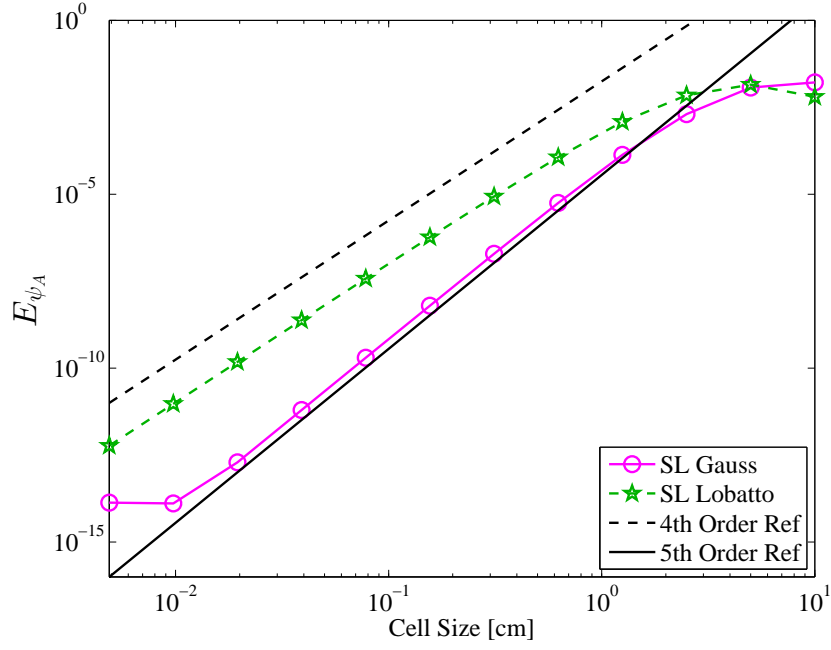


Figure 2.18: Convergence rate for $E_{\psi,A}$ as a function of the mesh cell size for a homogeneous pure absorber and quadratic DFEM.

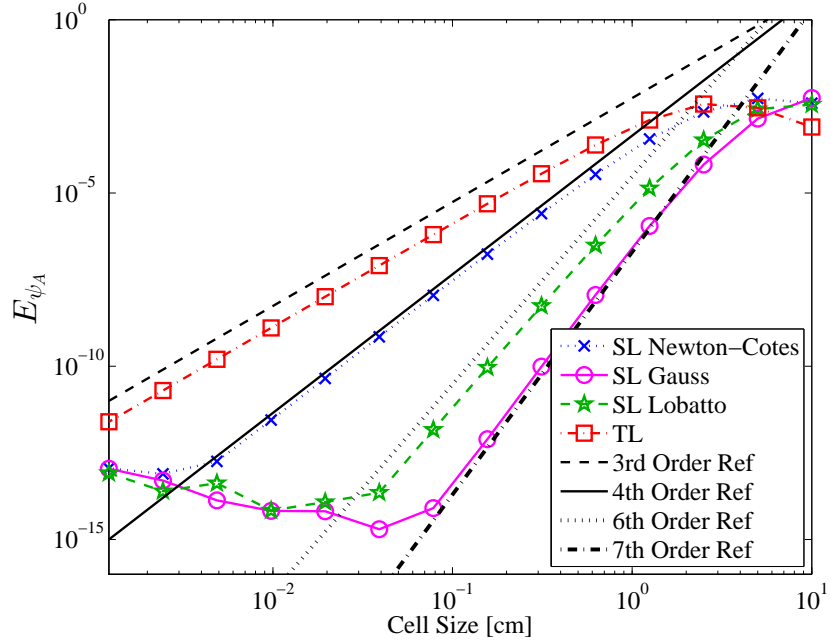


Figure 2.19: Convergence rate for $E_{\psi,A}$ as a function of the mesh cell size for a homogeneous pure absorber and cubic DFEM.

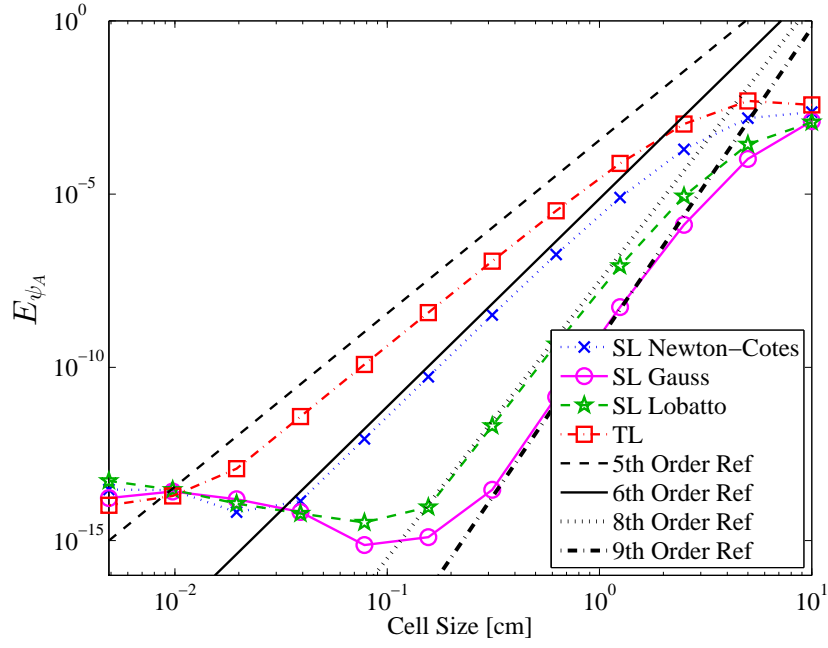


Figure 2.20: Convergence rate for $E_{\psi,A}$ as a function of the mesh cell size for a homogeneous pure absorber and quartic DFEM.

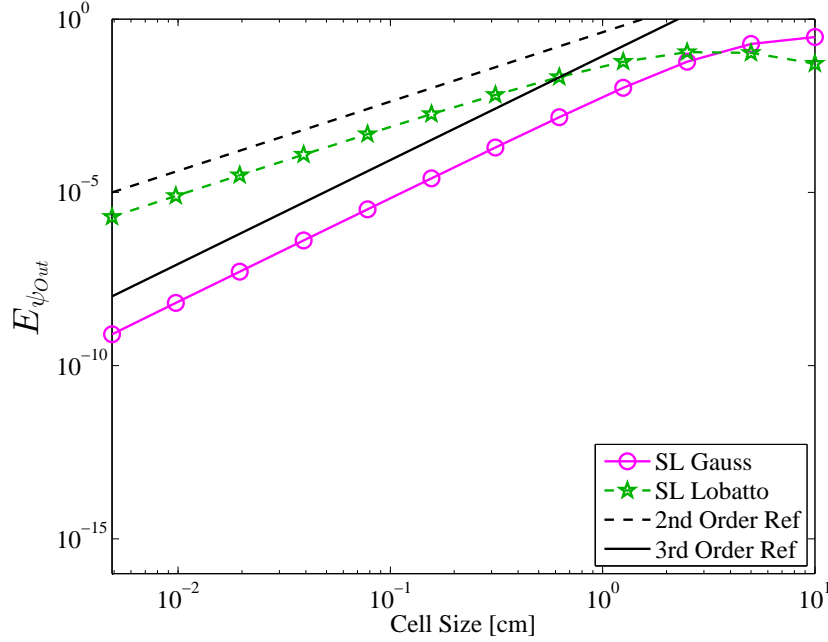


Figure 2.21: Convergence rate of $E_{\psi,out}$ as a function of the mesh cell size for a homogeneous pure absorber for linear DFEM.

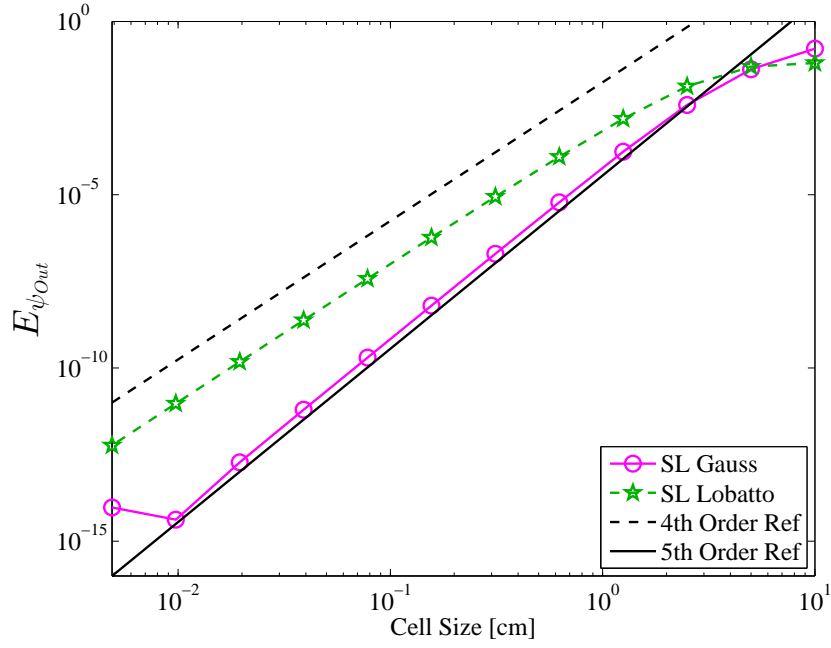


Figure 2.22: Convergence rate of $E_{\psi,out}$ as a function of the mesh cell size for a homogeneous pure absorber for quadratic DFEM.

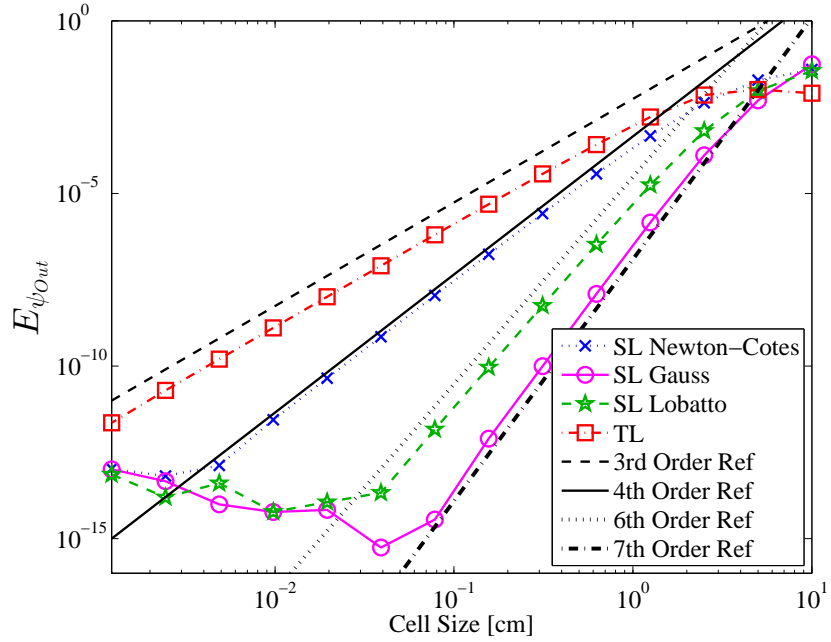


Figure 2.23: Convergence rate of $E_{\psi,out}$ as a function of the mesh cell size for a homogeneous pure absorber for cubic DFEM.

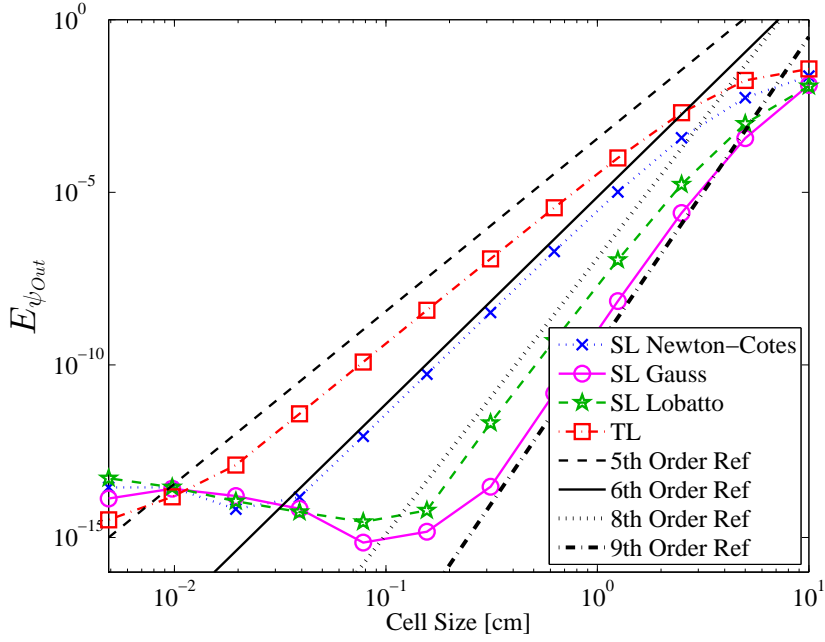


Figure 2.24: Convergence rate of $E_{\psi_{out}}$ as a function of the mesh cell size for a pure absorber for quartic DFEM.

The accumulation of errors in multiple-cell problems causes $E_{\psi_{out}}$ to globally converge one order of accuracy lower than the local truncation orders given in Table 2.10 and Table 2.11. It should be noted that the plateauing of errors E_{ψ} , E_{ψ_A} , and $E_{\psi_{out}}$ to values $\approx 10^{-14}$ in Figs. 2.13-2.16, Figs. 2.17-2.20, and Figs. 2.21-2.24, respectively, is simply a result of our numerical solutions being limited by machine precision (double precision).

2.6 Conclusions About Self-Lumping

We have shown that, for arbitrary degree polynomial trial space DFEM, a diagonal mass matrix does not necessarily ensure strictly positive angular flux outflow in a purely absorbing slab with spatially constant cross section. Indeed, the TL scheme was neither robust or accurate for polynomial trial space degree greater than linear. Also, we have shown that by using quadrature-based lumping schemes and choosing

DFEM interpolation points that are not equally spaced, robust, accurate polynomial DFEM schemes can be obtained. Based on the observed robustness, accuracy, and spatial convergence order results, we conclude that, for applications requiring robust solution techniques, the SL Lobatto scheme with odd degree polynomial trial space DFEM should be used to discretize the angular flux . If p -adaptivity is desired, software should be developed such that the ability to use either Lobatto (for odd trial space degrees) or Gauss (for even trial space degree) quadrature as the DFEM interpolation points is possible. However, given the non-monotonic behavior of the outflow angular flux as a function of the cell optical thickness when employing the SL Gauss scheme with even degree trial spaces for under-resolved problems, using SL Lobatto with an odd degree trial space would seem to be more accurate than using SL Gauss, despite SL Gauss being more accurate in the asymptotic (fine mesh) limit. Finally, though not as accurate SL Lobatto or SL Gauss, we will continue to consider SL Newton-Cotes due to its observed robustness.

3. DFEM METHODS FOR NEUTRON TRANSPORT FOR PROBLEMS WITH SPATIALLY VARYING CROSS SECTION

For many problems of interest to the nuclear science and engineering community, macroscopic cross sections in neutronics and opacities in radiative transfer calculations cannot accurately be described as piecewise constants in space. Cross sections and opacities are functions of continuously varying quantities such as temperature, density, burn-up history, etc. [28]. An examples simulation that is not be adequately described with cell-wise constant cross sections includes nuclear reactor isotopic depletion calculations. In thermal radiative transfer, interaction opacities can be rapidly varying functions of temperature. For example, consider Marshak wave problems and the canonical T^{-3} dependence [10] of absorption opacity, Across cells near the heated/cold material interface, opacity variations of several orders of magnitude are easily possible.

Historically, the neutron transport and thermal radiative transfer communities assumed interaction cross section and opacities, respectively, that were cell-wise constant [16, 29, 5]. Adams first described [6] and then presented computational results [7] for a “simple” corner balance (SCB) spatial discretization method that explicitly accounted for the spatial variation of opacity within individual spatial cells. The SCB scheme (which can be shown to be related to a LDFEM for certain geometries) accounts for opacity spatial variation within each cell via vertex-based quadrature evaluation. Similar strategies have been adapted to LDFEM radiative diffusion [10] and LDFEM TRT [30] calculations. For accurate TRT solutions, use of higher order DFEM will requires the development of corresponding higher order strategies for treating the within cell spatial variation of opacities. However, the majority of

neutron transport literature has only considered the case of cell-wise constant cross sections, see [16, 29, 31, 22]. The work of Kavenoky and Lautard [32] and more recently Santandrea and Bellier [33] are notable exceptions in neutron transport. In [32], continuous cubic finite element diffusion calculations that assume a linearly varying spatial cross section within each mesh cell were compared to results obtained using the same spatial discretization but with the assumption that cross sections are constant in each cell. Similarly, [33] compared the results of a linear characteristic scheme that assumes a linearly varying cross section in each spatial cell to those of a linear characteristic scheme that assumes a constant cross section in each cell.

In this chapter, we analyze the effects of cross-section spatial dependence on solution accuracy, as we did in our previously published work in [34]. Our work differs from [32, 33, 6, 7] by considering a discontinuous finite element (DFEM) spatial discretization of the slab geometry S_N transport equation using arbitrary degree polynomial finite element trial spaces. In addition, like [6] and [7] we do not make any approximation to the particular spatial shape of the cross-section spatial variation in each cell. We build on the quadrature integration ideas presented in Chapter 2 and employ a numerical quadrature to evaluate the mass matrix integrals that involve cross sections as a function of space. In general, the quadrature integration of the DFEM interaction term with arbitrary spatial cross section form will not be exact. However, we showed in Chapter 2 that exact computation of integrals appearing in the DFEM weak form, when cross sections are spatially constant, is not required to achieve high-order accuracy with high-order DFEM approximations. Building on this idea, we investigate the effects of using numerical quadratures to compute DFEM mass matrices, accounting for the spatial variation of cross section in space. As in Section 2 we use self-lumping numerical quadratures [24, 26], restricting quadrature integration points to the DFEM polynomial interpolation points. Results

are compared as a function of DFEM polynomial trial space degree and interpolation point type.

We demonstrate that assuming a piecewise constant cross section in each cell, when the cross section is not cell-wise constant in space, has several undesirable effects. Considering a source-free, purely absorbing medium, we show that DFEM schemes that assume a cell-wise constant cross section are at most second-order accurate for the angular flux solution and limited to at most first-order accuracy for the interaction rate solution, regardless of the DFEM polynomial trial space degree. We also show that assuming a piecewise constant cross section results in a highly discontinuous, non-monotonic spatial interaction rate. This phenomena has likely been present in published numerical results for problems with non-piecewise constant cross section but was not observed previously due to the choice of data presentation.

We then consider schemes that explicitly account for cross-section spatial variation within individual mesh cells. First, the positivity and robustness of different schemes are discussed using a source-free pure absorber problem. Next, we demonstrate that self-lumping schemes that evaluate the DFEM weak form integrals involving cross section with quadrature result in fully accurate schemes for arbitrary degree polynomial DFEM. By fully accurate we mean schemes that achieve the same order of convergence for problems with spatially varying and cell-wise constant cross section, for a given DFEM approximation order

3.1 Weak Form Derivation

We begin by repeating the DFEM neutron transport equation derived in Chapter 2:

$$\left(\mu_d \mathbf{G} + \frac{\Sigma_t \Delta x}{2} \mathbf{M} \right) \vec{\psi}_d = \frac{\Delta x}{2} \vec{Q}_d + \mu_d \psi_{in,d} \vec{f}. \quad (3.1)$$

To account for the within cell variation of cross section, we need only make one change to Eq. (3.1). We introduce the concept of a reaction matrix, \mathbf{R}_Σ where Σ is any interaction cross section or other material property:

$$\left(\mu_d \mathbf{G} + \frac{\Delta x}{2} \mathbf{R}_{\Sigma_t} \right) \vec{\psi}_d = \frac{\Delta x}{2} \vec{Q}_d + \mu_d \psi_{in,d} \vec{f}. \quad (3.2)$$

The $N_P \times N_P$ reaction matrix, \mathbf{R}_{Σ_t} is defined as:

$$\mathbf{R}_{\Sigma_t,ij} = \int_{-1}^1 \Sigma_t(s) b_i(s) b_j(s) ds. \quad (3.3)$$

Note that if Σ_t is indeed spatially constant within the mesh cell, there is no approximation in removing $\Sigma_t(s)$ from the integral of Eq. (3.3), giving

$$\mathbf{R}_{\Sigma_t,ij} = \Sigma_t \int_{-1}^1 b_i(s) b_j(s) ds, \quad (3.4)$$

which is equivalent to

$$\mathbf{R}_{\Sigma_t} = \Sigma_t \mathbf{M}. \quad (3.5)$$

3.2 Numerical Schemes

We consider two classes of numerical methods in this paper. The first class uses exact spatial integration to evaluate the integrals that define \mathbf{R}_{Σ_t} . A second class of methods uses numerical quadrature to evaluate \mathbf{R}_{Σ_t} , \mathbf{M} , and \mathbf{G} . Specifically, we limit out discussion of quadrature-based integration to so called self-lumping methods [12]. Self-lumping methods, first discussed in [24, 26] for parabolic problems, use numerical quadrature restricted to the finite element interpolation points, and thus naturally yield diagonal mass matrices. A shorthand notation is given in Table 3.1 for all of the numerical methods considered in this chapter and described in detail in the

remainder of this section.

Table 3.1: Nomenclature of numerical schemes considered for the pure absorber problem with a spatially exponential cross section.

Interpolation Point Type	\mathbf{R}_{Σ_t} Matrix Integration Strategy	Method Short Hand Name
Equally- Spaced	Exact Integration using true $\Sigma_t(x)$	EXS DFEM
Equally- Spaced	$\Sigma_t(x) \approx \hat{\Sigma}_t$, $\mathbf{R}_{\Sigma_t} \approx \hat{\Sigma}_t \mathbf{M}$ Exact Integration of \mathbf{M}	CXS DFEM
Equally- Spaced	Self-Lumping via Newton-Cotes Quadrature	SLXS Newton-Cotes
Lobatto Quadrature	Self-Lumping via Lobatto Quadrature	SLXS Lobatto
Gauss Quadrature	Self-Lumping via Gauss Quadrature	SLXS Gauss

3.2.1 Exact Spatial Integration

By exact spatial integration, we mean schemes that compute the entries of \mathbf{M} and \mathbf{G} exactly. Here, we achieve this by using equally-spaced interpolation points and employing a Gauss-Legendre quadrature rule [25] that exactly integrates the respective integrands of \mathbf{M} and \mathbf{G} . Two schemes use exact spatial integration. One approximates the spatially varying cross section as a cell-wise constant cross section. The other uses the exact cross section when integrating the weak form DFEM quantities involving cross section. The scheme that assumes a cell-wise constant cross section represents the state of the practice in the neutron transport community,

while the second scheme represents the ideal scenario for DFEM transport schemes in problems with spatially varying cross sections.

3.2.1.1 *Exact Cross Section*

The exact cross section, exact spatial integration scheme (EXS DFEM) attempts to analytically integrate the full definition of \mathbf{R}_{Σ_t} . Note that since $\Sigma_t(x)$ can be an arbitrary function, analytic integration of \mathbf{R}_{Σ_t} is in general impossible. Likewise, quadrature integration is unlikely to be exact. In our testing of the EXS DFEM scheme, we use a 20-point Gauss-Legendre quadrature to approximately integrate Eq. (3.3). Alternatively, adaptive quadrature, with a controllable tolerance, may be used such that the quadrature error in evaluating Eq. (3.3) could be reduced below some small tolerance.

3.2.1.2 *Constant Cross Section*

Historically, neutronics and some radiative transfer calculations have approximated spatially varying cross sections by assuming cell-wise constant cross sections [16, 29, 31, 5]. That is, some evaluation of the true $\Sigma_t(s)$ within a given cell is used to determine a constant value, $\hat{\Sigma}_t$, within each cell. Under this simplification, \mathbf{R}_{Σ_t} is approximated as:

$$\mathbf{R}_{\Sigma_t} = \hat{\Sigma}_t \mathbf{M}. \quad (3.6a)$$

In our test problems, the constant cross section scheme (CXS DFEM) uses the volumetric average of $\Sigma_t(s)$ to generate $\hat{\Sigma}_t$:

$$\hat{\Sigma}_t = \frac{1}{2} \int_{-1}^1 \Sigma_t(s) ds. \quad (3.7)$$

3.2.2 Self-Lumping Quadrature Integration

Schemes that are self-lumping evaluate the integrals of Eq. (3.3) using numerical quadrature. In Chapter 2 we showed that by definition, self-lumping schemes create diagonal mass matrices. Self-lumping schemes also create diagonal reaction matrices:

$$\mathbf{R}_{\Sigma_t,ij} = \begin{cases} w_i \Sigma_t(s_i) & i = j \\ 0 & \text{otherwise} \end{cases} \quad (3.8)$$

Though the choice of interpolation points does not affect exact integration schemes, as shown in Chapter 2, the choice of interpolation points was shown to influence both the robustness and accuracy of self-lumping schemes. We consider equally-spaced closed Newton-Cotes, Lobatto-Gauss-Legendre, and Gauss-Legendre quadratures as interpolation points for self-lumping schemes. We do not expect any self-lumping scheme to exactly integrate \mathbf{R}_{Σ_t} , as the integrand defining \mathbf{R}_{Σ_t} will generally not be a polynomial.

3.3 Pure Absorber Numerical Results

A beam of radiation, $\psi_{in}(\mu_d)$, is incident on the left face of the slab, the right face is a vacuum boundary, $x \in [0, x_R]$, and there are no fixed volumetric sources in the medium. We consider $\Sigma_t(x)$ to be of the form,

$$\Sigma_t(x) = c_1 e^{c_2 x}, \quad (3.9)$$

with c_1 and c_2 are constants [cm^{-1}], with $c_1 > 0$ and $c_2 \neq 0$. The analytic angular flux solution for a source-free pure absorber with an exponentially varying cross section

is:

$$\psi(x, \mu) = \begin{cases} \psi_{in}(\mu) \exp \left[\frac{c_1}{\mu c_2} (1 - e^{c_2 x}) \right] & \mu = \mu_d \\ 0 & \text{otherwise} \end{cases}. \quad (3.10)$$

By definition, the outflow angular flux from cell i , $\psi_{out,i}$ is $\psi(x_{i+1/2}, \mu_d)$ and the average angular flux within cell i , $\psi_{A,i}$ as

$$\psi_{A,i} = \frac{1}{\Delta x_i} \int_{x_{i-1/2}}^{x_{i+1/2}} \psi(x, \mu_d) dx, \quad (3.11)$$

with $\Sigma_t(x)$ defined as in Eq. (3.9). The analytical average flux value is:

$$\psi_{A,i} = \frac{\psi_{in}(\mu_d)}{\Delta x_i} \exp \left[\frac{c_1}{\mu_d c_2} \right] \left[E_1 \left(\frac{c_1 e^{c_2 x_{i+1/2}}}{\mu_d c_2} \right) - E_1 \left(\frac{c_1 e^{c_2 x_{i-1/2}}}{\mu_d c_2} \right) \right], \quad (3.12)$$

with E_1 the exponential integral [25].

3.3.1 Single Cell Outflow Comparisons

The only variable cross-section schemes that yields strictly positive angular outflows in a source-free pure absorber are the SLXS Lobatto and SLXS Newton-Cotes schemes using a linear trial space. For $\mu_d > 0$, consider a source-free, purely absorbing cell with known inflow, $\psi_{in}(\mu_d)$, of width Δx , and the total cross section at each interpolation point is $\Sigma_{t,j}$. Regardless of the actual functional form of the cross section within the cell, the linear DFEM SLXS Lobatto and SLXS Newton-Cotes schemes' numerical angular flux outflow, $\tilde{\psi}_d(1)$, is:

$$\tilde{\psi}_d(1) = \frac{2\mu_d^2 \psi_{in}(\mu_d)}{2\mu_d^2 + \Delta x^2 \Sigma_{t,1} \Sigma_{t,2} + \Delta x \mu_d \Sigma_{t,1} + \Delta x \mu_d \Sigma_{t,2}}. \quad (3.13)$$

Equation 3.13 is strictly positive when $\Sigma_t(x) \geq 0$, suggesting that the strictly positive outflow results observed in [12] might hold for an arbitrarily varying spatial cross sec-

tion. However, the results of [12] do not hold for higher-order DFEM approximations for spatially dependent cross sections.

To demonstrate that negative cell outflows are possible, we carry out the following test. In Figs. 3.1-3.4, we plot the angular flux outflow of each method as a function of trial space degree, and the parameter c_2 . We hold the total cell optical thickness to 20 mean-free-path (MFP), vary $c_2 \in [1, 10]$, fix $x_R = 1$, and $\mu_d = 1$. With an exponential cross section, the cell optical thickness in *MFP* of a cell with $x \in [0, x_R]$ is:

$$MFP = \int_0^{x_R} \Sigma_t(x) dx = \frac{c_1}{c_2} (e^{c_2 x_R} - 1) . \quad (3.14)$$

To maintain a constant optical thickness in Figs. 3.1-3.4, c_1 is required to be:

$$c_1 = \frac{c_2 MFP}{e^{c_2 x_R} - 1} . \quad (3.15)$$

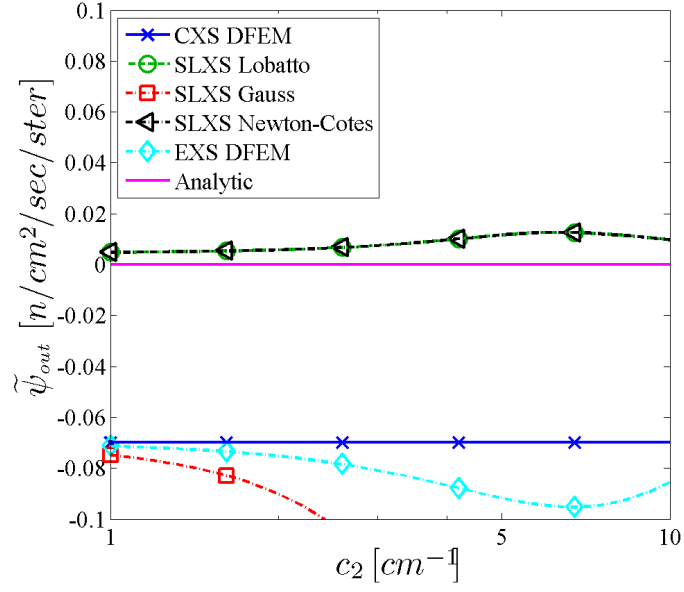


Figure 3.1: Numerical outflow from single cell pure absorber with $\Sigma_t(x) = c_1 e^{c_2 x}$, as a function of c_2 with constant optical thickness of twenty MFP, for a linear trial space.

Figures 3.1-3.4 confirms that SLXS Lobatto (and the equivalent SLXS Newton-Cotes scheme) with a linear trial space is the only scheme that explicitly accounts for the spatial variation of cross and maintains a strictly positive angular flux outflow regardless of the shape of $\Sigma_t(x)$. From Figs. 3.1-3.4 we also observe that $\tilde{\psi}_{out}$ varies for every method as a function of the shape of $\Sigma_t(x)$, with the obvious exception of CXS DFEM. Considering that the analytic angular flux outflow is only a function of total cell MFP:

$$\psi_{out,i} = \psi_{in}(\mu_d) \exp \left[- \int_0^{x_{i+1/2}} \Sigma_t(x) dx / \mu_d \right] = \psi_{in}(\mu_d) \exp [-MFP / \mu_d] , \quad (3.16)$$

it is unphysical and undesirable that $\tilde{\psi}_{out}$, for the SLXS Gauss, SLXS Lobatto, SLXS Newton-Cotes, and EXS DFEM schemes, depends on the spatial shape of $\Sigma_t(x)$.

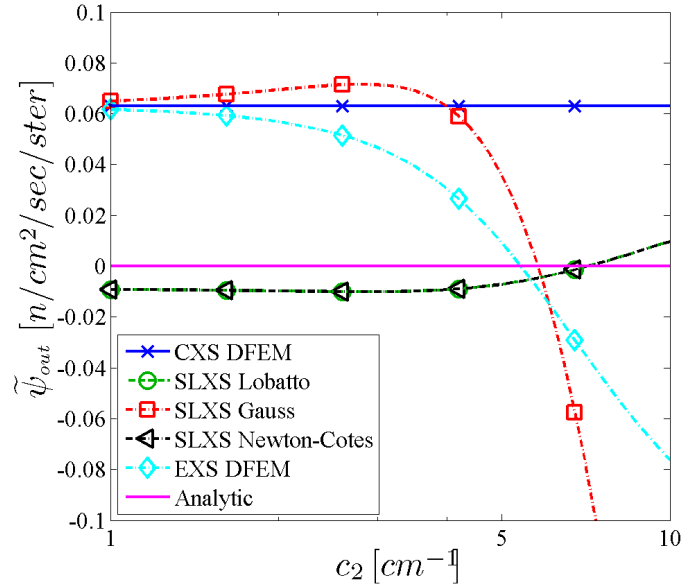


Figure 3.2: Numerical outflow from single cell pure absorber with $\Sigma_t(x) = c_1 e^{c_2 x}$, as a function of c_2 with constant optical thickness of twenty MFP, for a quadratic trial space.

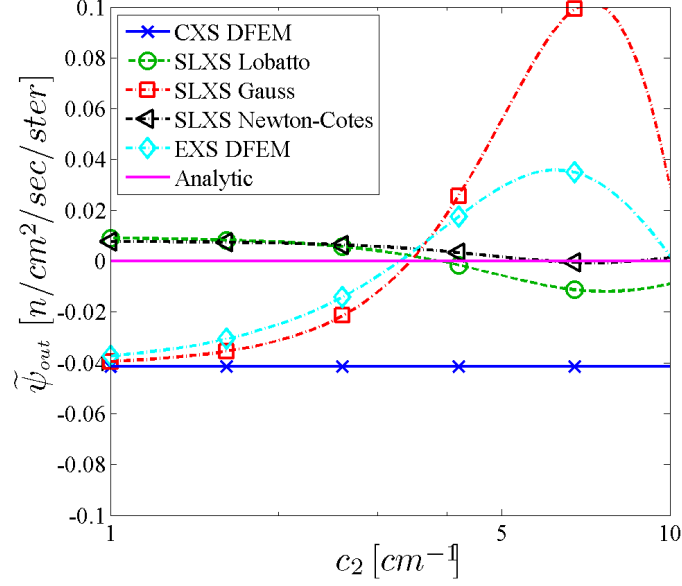


Figure 3.3: Numerical outflow from single cell pure absorber with $\Sigma_t(x) = c_1 e^{c_2 x}$, as a function of c_2 with constant optical thickness of twenty MFP, for different degree polynomial trial spaces.

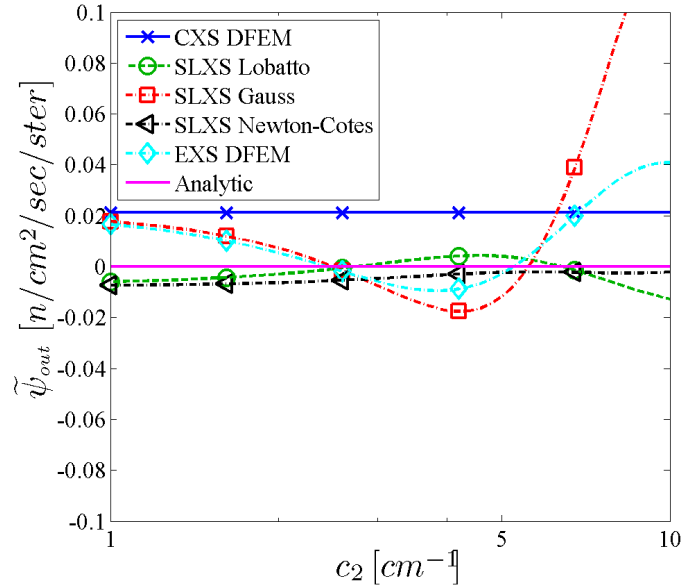


Figure 3.4: Numerical outflow from single cell pure absorber with $\Sigma_t(x) = c_1 e^{c_2 x}$, as a function of c_2 with constant optical thickness of twenty MFP, for different degree polynomial trial spaces.

3.3.2 Multiple Cell Spatial Convergence Rates

We now consider the order of spatial convergence for the following schemes: CXS DFEM, SLXS Gauss, SLXS Lobatto, and SLXS Newton-Cotes. Since exact integration of \mathbf{R}_{Σ_t} is generally not feasible, we no longer consider the EXS DFEM scheme. Convergence results of the following angular flux errors as a function of the polynomial approximation order are presented:

$$E_\psi = \sqrt{\sum_{i=1}^{N_{cells}} \int_{x_{i-1/2}}^{x_{i+1/2}} \left(\tilde{\psi}_d(x) - \psi(x, \mu_d) \right)^2 dx} \quad (3.17a)$$

$$E_{\psi_A} = \sqrt{\sum_{i=1}^{N_{cells}} \Delta x_i \left(\tilde{\psi}_{A,i} - \psi_{A,i} \right)^2} \quad (3.17b)$$

$$E_{\psi_{out}} = \sqrt{\sum_{i=1}^{N_{cells}} \Delta x_i \left(\tilde{\psi}_{out,i} - \psi(x_{i+1/2}, \mu_d) \right)^2}. \quad (3.17c)$$

In Eqs. (3.17), Δx_i is the width of cell i , cell i spans $[x_{i-1/2}, x_{i+1/2}]$, $\tilde{\psi}_d(x)$ is the DFEM numerical approximation, $\psi(x, \mu_d)$ is the analytic solution (see Eq. (3.10)). The problem is spatially discretized using N_{cells} spatial cells of equal width. We approximate the integrals defining the L^2 norm of the angular flux error, E_ψ , using a high-order Gauss quadrature set, $(w_{f,q}, s_{f,q})$, with N_{qf} points, such that:

$$\int_{x_{i-1/2}}^{x_{i+1/2}} \left(\tilde{\psi}(x) - \psi(x, \mu_d) \right)^2 dx \approx \frac{\Delta x_i}{2} \sum_{q=1}^{N_{qf}} w_{f,q} \left(\tilde{\psi}(s_{f,q}) - \psi(s_{f,q}, \mu_d) \right)^2. \quad (3.18)$$

For the results that follow, $N_{qf} = 10$. We recall the definitions for the numerical approximations of the cell average angular flux, $\tilde{\psi}_{A,i}$, and the outflow angular flux,

$\tilde{\psi}_{out,i}$:

$$\tilde{\psi}_{A,i} = \frac{1}{2} \sum_{j=1}^{N_P} w_j \psi_{i,j} \quad (3.19a)$$

$$\tilde{\psi}_{out,i} = \sum_{j=1}^{N_P} \psi_{i,j} b_j(1). \quad (3.19b)$$

We also consider the convergence of the numerical interaction rate, $\widetilde{IR}(x)$ to the true interaction rate $IR(x)$. First, we define the analytic reaction rate for our beam problem:

$$IR(x) = \Sigma_t(x) \psi(x, \mu_d). \quad (3.20)$$

Similarly, we define a cell average interaction rate as:

$$IR_{A,i} = \frac{1}{\Delta x_i} \int_{x_{i-1/2}}^{x_{i+1/2}} \Sigma_t(x) \psi(x, \mu_d) dx. \quad (3.21)$$

Defining a point-wise numerical approximation, $\widetilde{IR}(x)$, to the analytic interaction rate for the self-lumping schemes presents a unique problem, since only a numerical quadrature is used to approximate the integrand of \mathbf{R} . Quadrature integration only requires point evaluations of $\Sigma_t(x)$, not knowledge of $\Sigma_t(x)$ in between quadrature points. However, for the purpose of plotting the SLXS schemes, we define:

$$\widetilde{IR}(s) = \sum_{j=1}^{N_P} b_j(s) \psi_{j,d} \Sigma_t(s_j). \quad (3.22)$$

We approximate the cell average interaction rate in cell i as:

$$\widetilde{IR}_{A,i} = \frac{1}{2} \sum_{j=1}^{N_P} w_j \Sigma_t(s_j) \psi_{j,d}. \quad (3.23)$$

In Eq. (3.23), $\Sigma_t(s_j) = \hat{\Sigma}_t$ for the CXS DFEM scheme, and $\Sigma_t(s_j)$ is the point evaluation of the true cross section for all other schemes.

We consider two measures to assess the error of the DFEM schemes' approximation of the true interaction rate, $IR(x)$. The first, E_{IR} is an approximation of the L^2 norm of interaction rate error:

$$E_{IR} = \sqrt{\sum_{i=1}^{N_{cells}} \frac{\Delta x_i}{2} \sum_{q=1}^{N_P} w_q \left(IR(s_q) - \Sigma_t(s_q) \tilde{\psi}(s_q, \mu_d) \right)^2}. \quad (3.24)$$

We reiterate that, for the self-lumping schemes, $\tilde{IR}(s)$ is only truly defined at the DFEM interpolation points. E_{IR_A} measures the convergence of the average interaction rate:

$$E_{IR_A} = \sqrt{\sum_{i=1}^{N_{cells}} \Delta x_i (IR_{A,i} - \tilde{IR}_{A,i})^2}. \quad (3.25)$$

For our convergence study, we consider a source-free purely absorbing slab with a cross section that varies exponentially in space as in Eq. (3.9) with $c_1 = 0.1$ and $c_2 = 2 \ln(10)$. A beam of radiation is incident on the left face in the direction of $\mu_d = 1$, vacuum boundary conditions exist on the right face of the slab, and $x \in [0, 1]$. The convergence of the E_ψ , E_{ψ_A} , and $E_{\psi_{out}}$ as a function of the choice of numerical scheme for linear through quartic trial space polynomial degree are given in Figs. 3.5-3.8, Figs. 3.9-3.12, and Figs. 3.13-3.16, respectively.

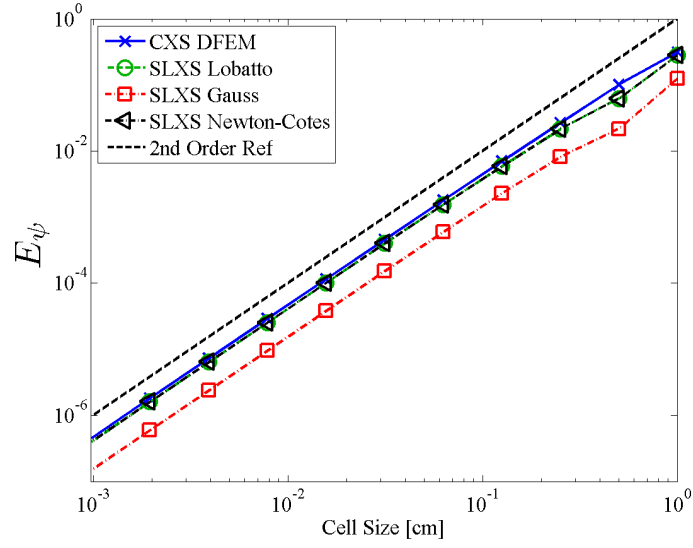


Figure 3.5: Convergence of E_ψ for a pure absorber with exponentially varying cross section discretized with linear DFEM.

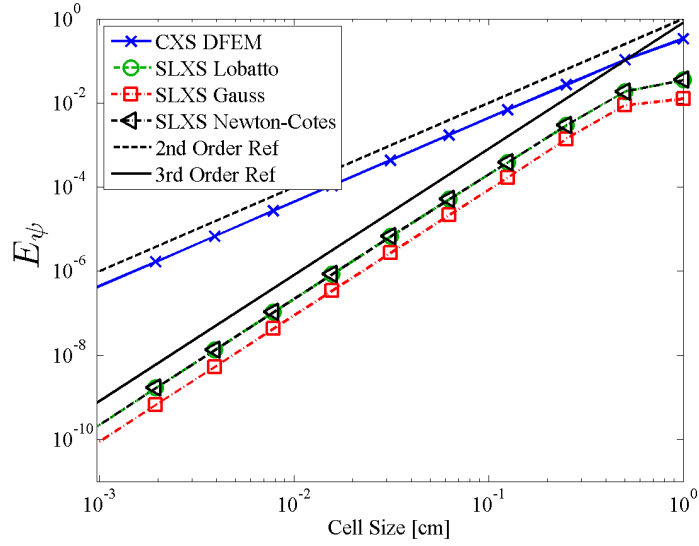


Figure 3.6: Convergence of E_ψ for a pure absorber with exponentially varying cross section discretized with quadratic DFEM.

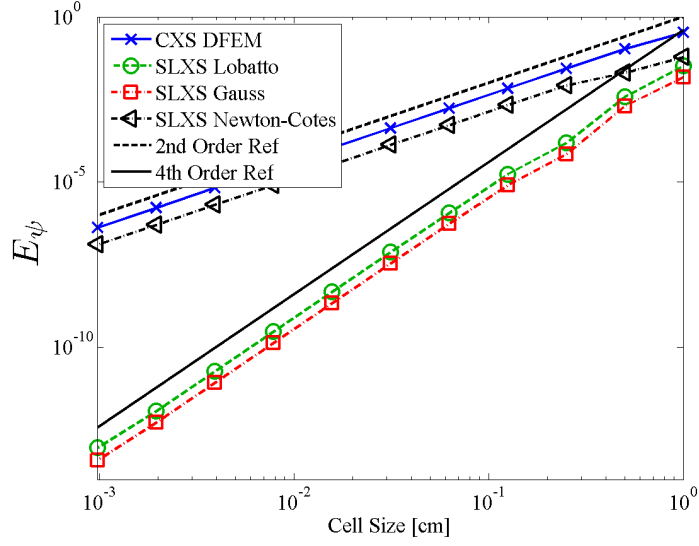


Figure 3.7: Convergence of E_ψ for a pure absorber with exponentially varying cross section discretized with cubic DFEM.

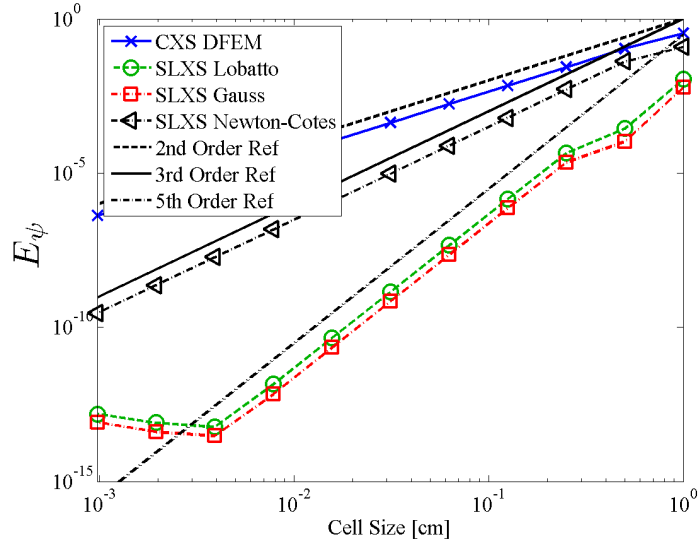


Figure 3.8: Convergence of E_ψ for a pure absorber with exponentially varying cross section discretized with quartic DFEM.

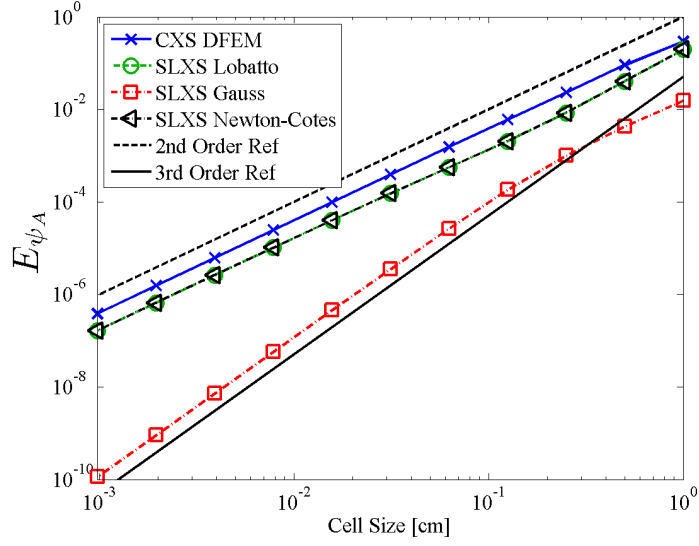


Figure 3.9: Convergence of E_{ψ_A} for a pure absorber with exponentially varying cross section discretized with linear DFEM.

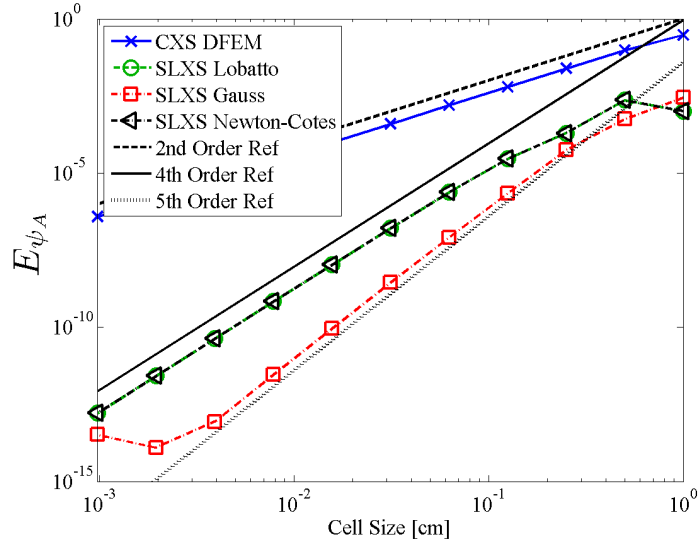


Figure 3.10: Convergence of E_{ψ_A} for a pure absorber with exponentially varying cross section discretized with quadratic DFEM.

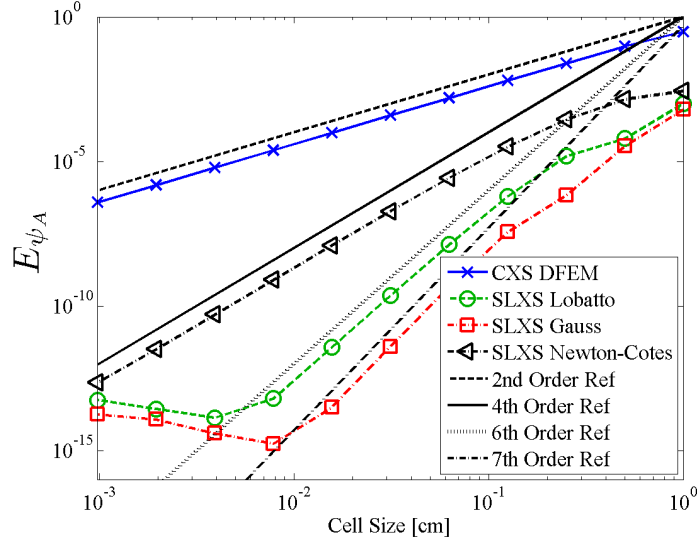


Figure 3.11: Convergence of E_{ψ_A} for a pure absorber with exponentially varying cross section discretized with cubic DFEM.

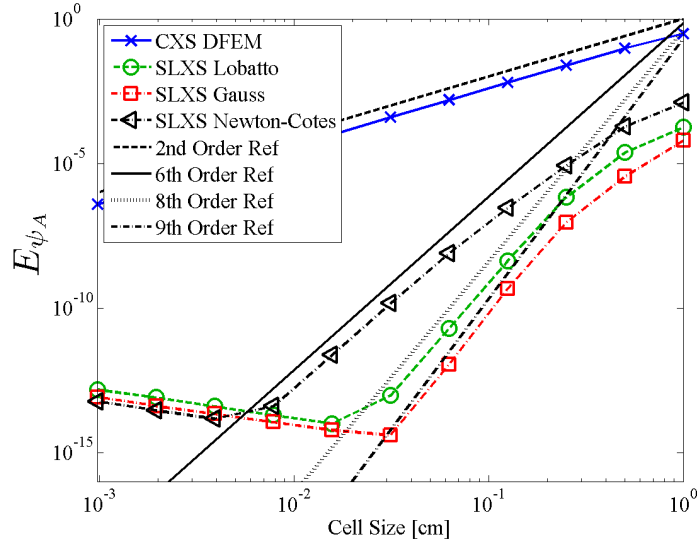


Figure 3.12: Convergence of E_{ψ_A} for a pure absorber with exponentially varying cross section discretized with quartic DFEM.

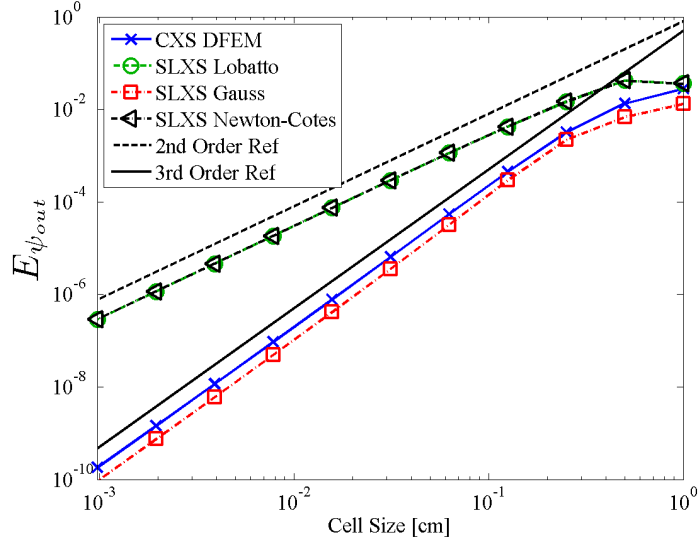


Figure 3.13: Convergence of $E_{\psi_{out}}$ for a pure absorber with exponentially varying cross section discretized with linear DFEM.

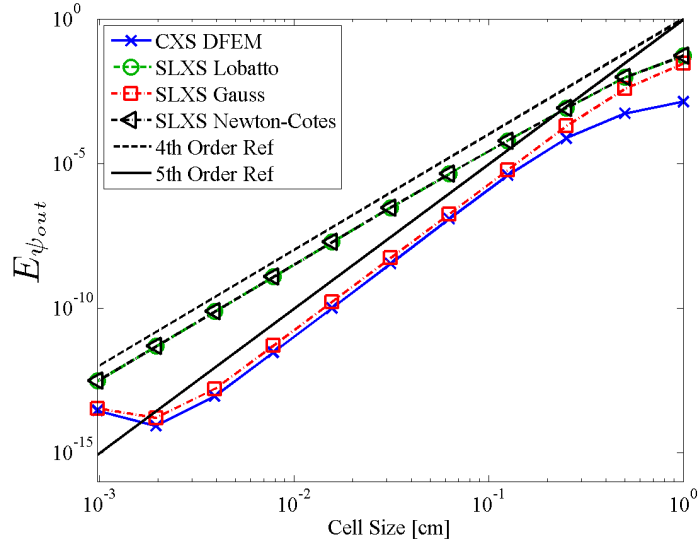


Figure 3.14: Convergence of $E_{\psi_{out}}$ for a pure absorber with exponentially varying cross section discretized with quadratic DFEM.

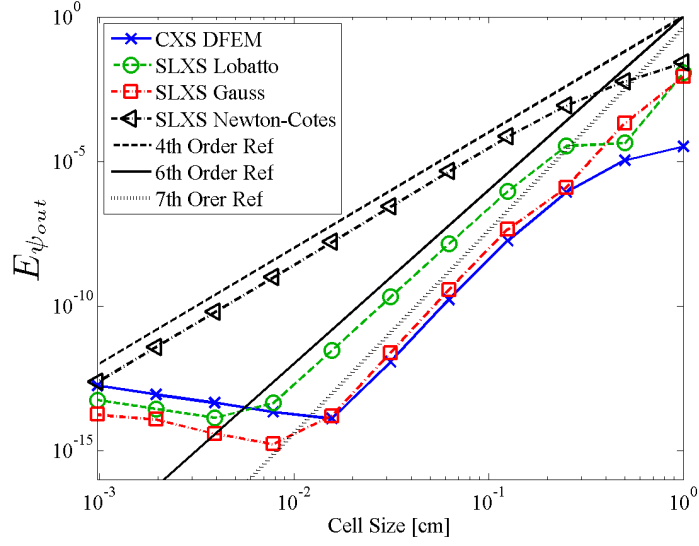


Figure 3.15: Convergence of $E_{\psi_{out}}$ for a pure absorber with exponentially varying cross section discretized with cubic DFEM.

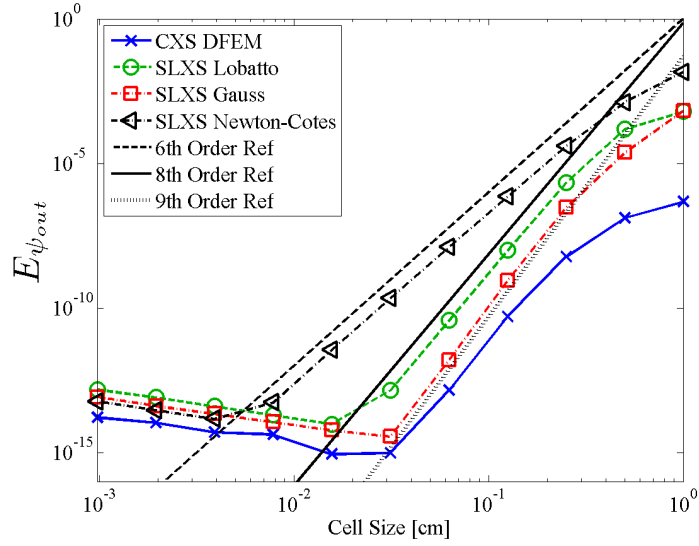


Figure 3.16: Convergence of $E_{\psi_{out}}$ for a pure absorber with exponentially varying cross section discretized with quartic DFEM.

The plateauing of numerical errors for various high-order methods using very small cell sizes in Figs. 3.5-3.16 is a consequence of having reached machine precision. The lines in Figs. 3.5-3.16 that extend to values smaller than machine precision are reference lines.

Figures 3.5-3.16 show that, for a linear angular flux trial space, CXS DFEM achieves the same orders of spatial convergence as observed with Exact DFEM in [12]. However, as the degree of the DFEM trial space is increased, the CXS DFEM scheme does not show an increase in the order of the spatial convergence rate of E_ψ and E_{ψ_A} ; the convergence rate of CXS DFEM is limited to at most second order for both E_ψ and E_{ψ_A} , regardless of the trial space polynomial degree. The increase in order of convergence of CXS DFEM for $E_{\psi_{out}}$ as trial space is increased is a result of angular flux outflow in the CXS DFEM discretization being only a function of the cell optical thickness, which is preserved exactly by our definition of $\hat{\Sigma}_t$; see Eq. (3.7).

Of the self-lumping schemes, SLXS Newton-Cotes is the least accurate. SLXS Newton-Cotes convergence of E_ψ is limited to at most second order for odd degree polynomial trial spaces and third order for even degree trial spaces. Convergence of E_{ψ_A} and $E_{\psi_{out}}$ for the SLXS Newton-Cotes scheme generally increases with an increase in the DFEM polynomial trial space degree, but is only proportional to P . Both SLXS Lobatto and SLXS Gauss converge E_ψ , E_{ψ_A} , and $E_{\psi_{out}}$ similarly to the study carried out in [12] with a spatially constant cross section.

The spatial convergence of E_ψ for SLXS Lobatto and SLXS Gauss is order $P + 1$. Though SLXS Lobatto and SLXS Gauss converge with the same order of spatial convergence for E_ψ , SLXS Gauss is more accurate than SLXS Lobatto by a constant. SLXS Gauss converges E_{ψ_A} and $E_{\psi_{out}} \propto 2P + 1$, whereas SLXS Lobatto converges both $\propto 2P$. SLXS Gauss and SLXS Lobatto converge angular flux error quantities

for the case of a spatially varying cross section with the same rates of convergence as their constant cross-section analogs did in [12]. This suggests that exactly integrating the interaction term in the DFEM moment equations is not essential for developing arbitrarily high-order accuracy DFEM schemes for radiation transport.

Convergence of E_{IR} as function of numerical scheme for linear - quartic trial spaces is given in Figs. 3.17-3.20. We observe the detrimental effect of approxi-

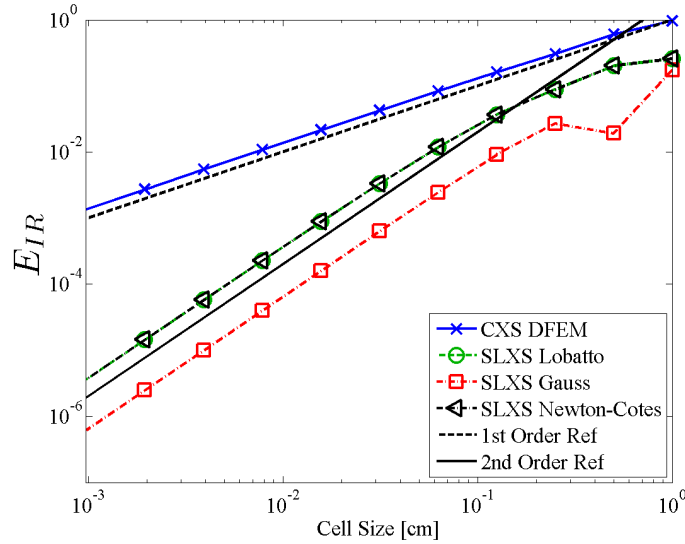


Figure 3.17: Convergence of E_{IR} for a pure absorber with exponentially varying cross section discretized with linear DFEM.

imating a spatially varying cross section with a constant in each spatial cell when we examine the L_2 convergence results for the interaction rate, E_{IR} , for the CXS DFEM scheme. Regardless of angular flux trial space polynomial degree, CXS DFEM converges E_{IR} to only first order in space. However, the self-lumping schemes exhibit the same trends in converging E_{IR} (in the L^2 -norm sense) as exhibited in converging E_ψ :

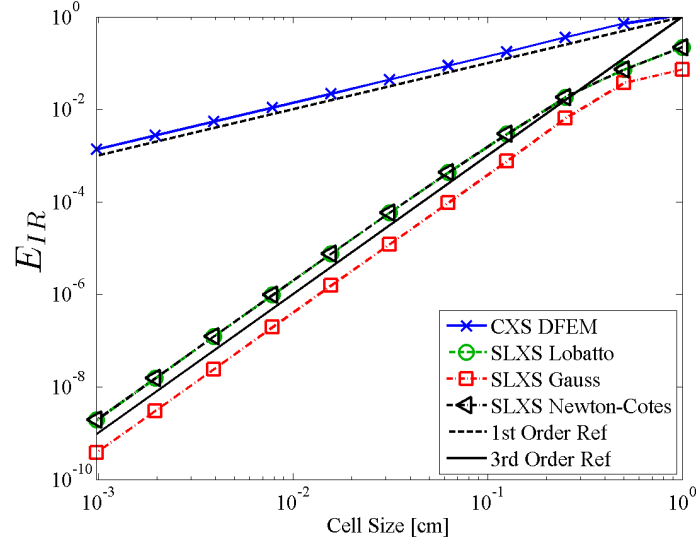


Figure 3.18: Convergence of E_{IR} for a pure absorber with exponentially varying cross section discretized with quadratic DFEM.

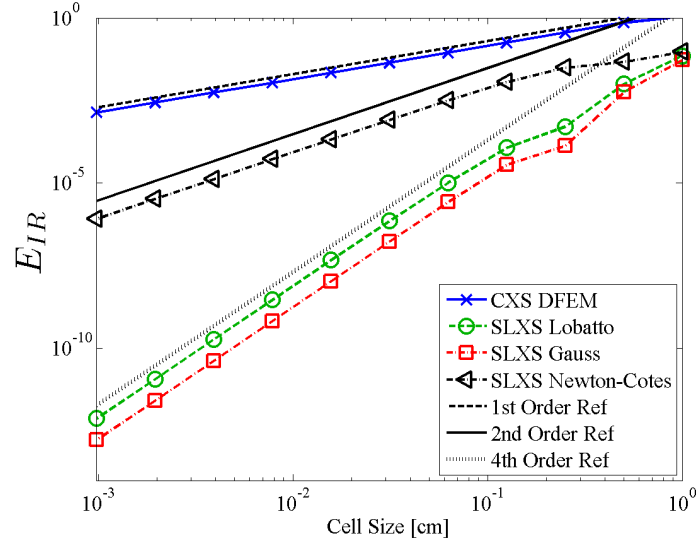


Figure 3.19: Convergence of E_{IR} for a pure absorber with exponentially varying cross section discretized with cubic DFEM.

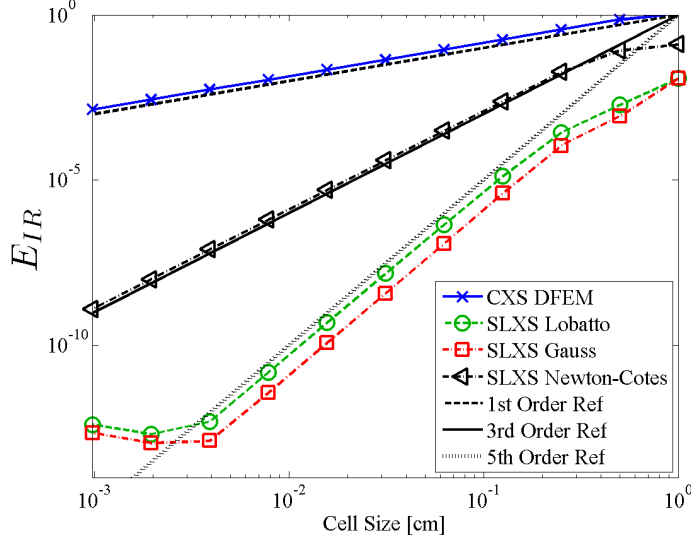


Figure 3.20: Convergence of E_{IR} for a pure absorber with exponentially varying cross section discretized with quartic DFEM.

- SLXS Lobatto and SLXS Gauss converge E_{IR} with order $P + 1$,
- SLXS Gauss is more accurate than SLXS Lobatto by a constant, and
- SLXS Newton-Cotes converges E_{IR} second order in space for odd degree trial spaces and third order in space for even degree trial spaces.

Convergence data for E_{IR_A} as function of cell size for linear - quartic trial spaces is given in Figs. 3.21-3.24. The convergence results for the error in cell average interaction rate, E_{IR_A} , shown in Figs. 3.21-3.24, do not behave as intuitively as the convergence rates for E_{IR} observed in Figs. 3.17-3.20. Given the poor performance of CXS DFEM in converging E_{IR} , one would expect that CXS DFEM would converge E_{IR_A} poorly as well. However, this is not the case and CXS DFEM converges E_{IR_A} with the same order of convergence as the best performing self-lumping scheme considered.

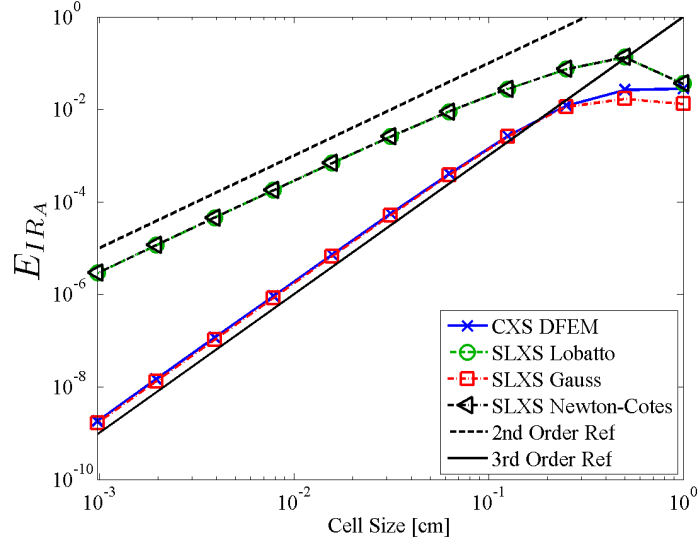


Figure 3.21: Convergence of E_{IRA} for a pure absorber with exponentially varying cross section discretized with linear DFEM.

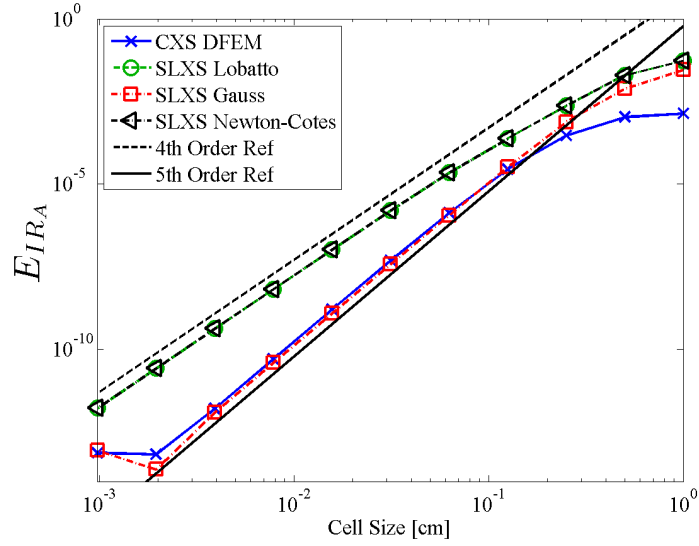


Figure 3.22: Convergence of E_{IRA} for a pure absorber with exponentially varying cross section discretized with quadratic DFEM.

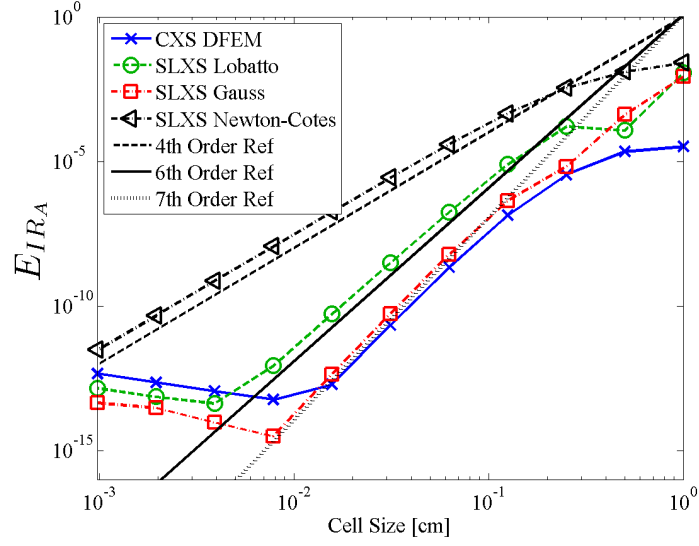


Figure 3.23: Convergence of E_{IRA} for a pure absorber with exponentially varying cross section discretized with cubic DFEM.

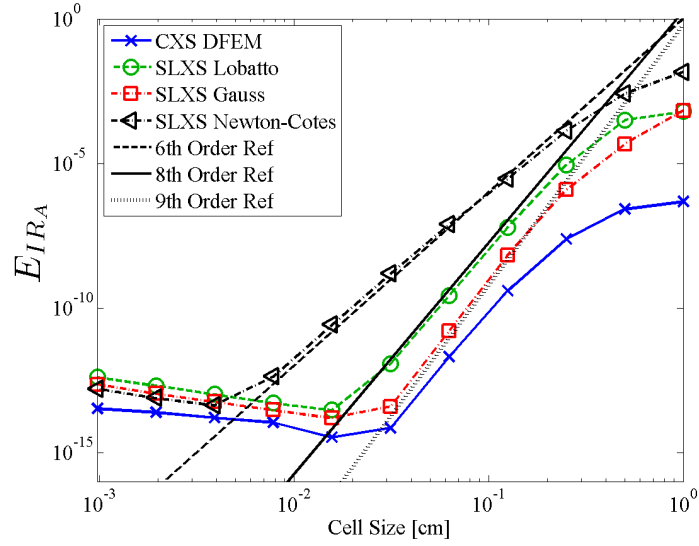


Figure 3.24: Convergence of E_{IRA} for a pure absorber with exponentially varying cross section discretized with quartic DFEM.

CXS DFEM converges E_{IR_A} with a high-order of accuracy because of the locally conservative properties of DFEM approximations, that is:

$$\text{Particles Into Cell} - \text{Particles Out of Cell} = \text{Total Interactions in Cell}. \quad (3.26)$$

As shown in Figs. 3.13-3.16, CXS DFEM converges the quantities on the left hand side of Eq. (3.26) with the same order of accuracy as any self-lumping scheme considered; CXS DFEM is at least as accurate in calculating the particles into a cell (outflow from the previous cell) and out of the cell (outflow from the current cell) as any other scheme considered. Since Eq. (3.26) holds regardless of the numerical scheme considered, it follows that CXS DFEM converges E_{IR_A} , the term in the right hand side of Eq. (3.26) summed over all cells, with the maximum order of convergence displayed by any of the DFEM schemes we consider here. Figures 3.21-3.24 validate this conclusion. CXS DFEM and SLXS Gauss exhibit the highest order of spatial convergence, converging E_{IR_A} with order $\propto 2P + 1$. SLXS Newton-Cotes and SLXS Lobatto converge E_{IR_A} with the same orders of convergence each method exhibits in converging E_{ψ_A} . For this problem SLXS Lobatto converges $E_{IR_A} \propto 2P$ and SLXS Newton-Cotes converged $E_{IR_A} \propto P$.

3.3.3 Consequences of Assuming a Cell-Wise Constant Cross Section

To more fully understand the poor convergence of point-wise error in angular flux and interaction rate, E_{ψ} and E_{IR} , associated with CXS DFEM we now examine more closely the CXS DFEM spatial approximations to $\psi(x, \mu_d)$ and $IR(x)$. We again consider a pure absorber with total absorption cross section that varies exponentially in space with $c_1 = 0.1$, and $c_2 = 2 \ln(10)$. A beam of radiation is incident on the left face in the direction of $\mu_d = 1$, vacuum boundary conditions are applied on the right face of the slab, and $x \in [0, 1]$.

In Fig. 3.25, we plot the exact $\psi(x)$ and in Fig. 3.26, we plot $IR(x)$. Additionally we plot the respective CXS DFEM numerical approximations, $\tilde{\psi}(x)$ and $\tilde{IR}(x)$, using $N_{cells} = 5$, and $\hat{\Sigma}_{t,i}$ as defined in Eq. (3.7). Also plotted in Fig. 3.25 and Fig. 3.26, we plot the analytic angular flux and reaction rate, respectively, one would obtain if the cell average cross section, $\hat{\Sigma}_{t,i}$, had been used instead of the true $\Sigma_t(x)$. We refer to these analytic solutions as $\psi_C(x)$ and $IR_C(x)$.

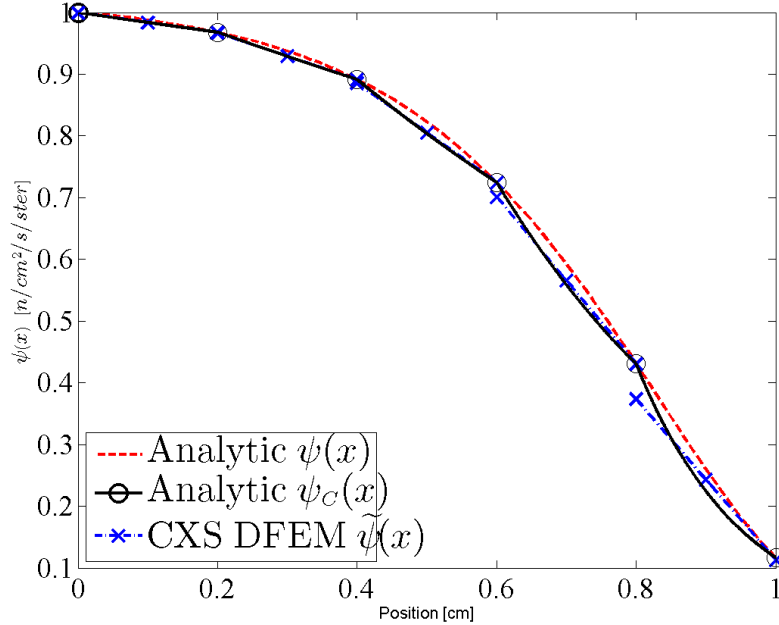


Figure 3.25: Plots of the analytic $\psi(x)$, CXS DFEM $\tilde{\psi}(x)$, and $\psi_C(x)$ for the pure absorber with exponential cross section.

Since CXS DFEM is a discontinuous scheme, some discontinuity is expected in the plot of $\tilde{\psi}$ in Fig. 3.25 and of \tilde{IR} in Fig. 3.26. However, the discontinuities present in Fig. 3.26 are highly disconcerting. The analytic $IR(x)$ is smooth and does not vary rapidly within individual mesh cells, yet there are significant, non-monotonic

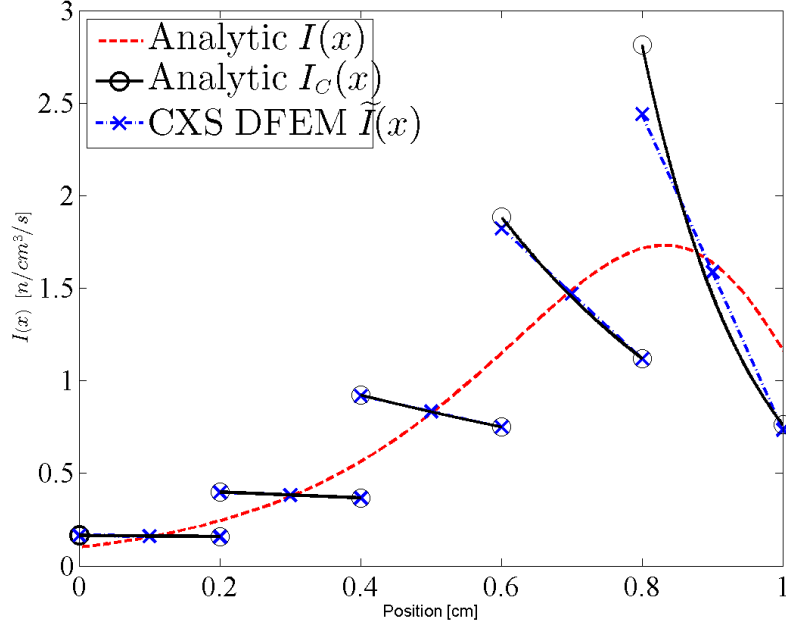


Figure 3.26: Plots of the analytic $IR(x)$, CXS DFEM $\widetilde{IR}(x)$, and $IR_C(x)$ for the pure absorber with exponential cross section.

discontinuities in the CXS DFEM interaction rate solution to the pure absorber problem with exponentially varying cross section. The noticeably poor behavior of $\widetilde{IR}(x)$ in Fig. 3.26 is inherent to the assumption of a cell-wise constant cross section. This inherent error of approximating a continuously varying cross section with cell-wise constants is clearly visible when comparing the plots of $IR_C(x)$ to $IR(x)$ in Fig. 3.26. The only difference and source of error in $IR_C(x)$ is the assumption of a cell-wise constant cross section. Figure 3.25 and Fig. 3.26 does not suggest that linear DFEM is unsuitable for use in problems with spatially varying cross sections. Rather, comparing the CXS DFEM $\widetilde{\psi}(x)$ to $\psi_C(x)$ in Fig. 3.25 and $\widetilde{IR}(x)$ to $IR_C(x)$ in Fig. 3.26, we see that CXS DFEM is very accurate when compared to the analytic solution of the problem CXS DFEM is solving, a pure absorber with cell-wise constant cross section. Given the poor accuracy of CXS DFEM in approximating the true $\psi(x)$

and $IR(x)$, we wish to see how a scheme that does not assume a cell-wise constant cross section behaves. Consider $\tilde{\psi}(x)$ and $\tilde{IR}(x)$ obtained with SLXS Lobatto using a linear DFEM trial space and five spatial cells, shown in Fig. 3.27 and Fig. 3.28, for the same problem. In Fig. 3.27, the differences between the angular flux solutions obtained using (1) a cell-wise constant cross section (CXS DFEM) and (2) evaluating cross section values at quadrature points (SLXS Lobatto) are small.

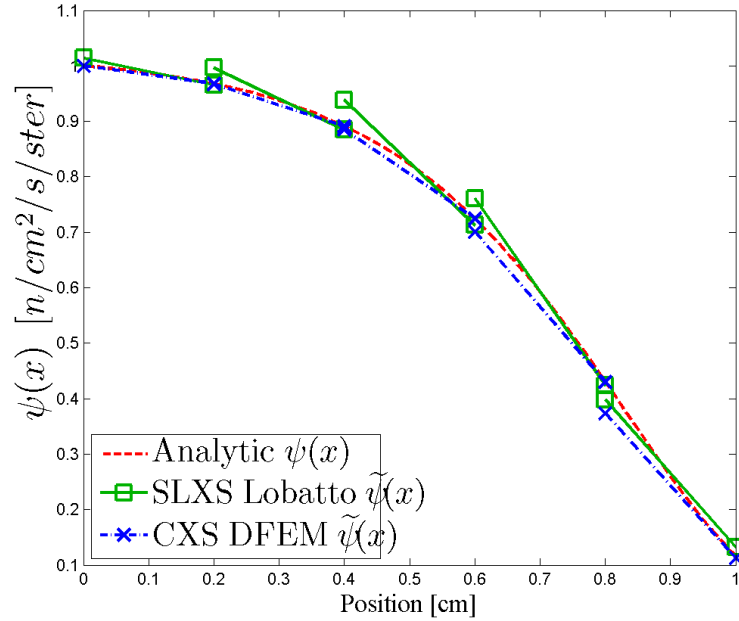


Figure 3.27: Plot of the linear trial space SLXS Lobatto and CXS DFEM approximations of $\tilde{\psi}(x)$ for the pure absorber problem with exponentially varying cross section.

This is not the case when comparing the different approximations to the interaction rate, $\tilde{IR}(x)$, in Fig. 3.28. Though there are discontinuities in the SLXS Lobatto $\tilde{IR}(x)$, the discontinuities are smaller and the SLXS Lobatto $\tilde{IR}(x)$ is monotonic unlike the CXS DFEM $\tilde{IR}(x)$. The SLXS Lobatto $\tilde{IR}(x)$ is clearly more accurate

than the CXS DFEM $\widetilde{IR}(x)$.

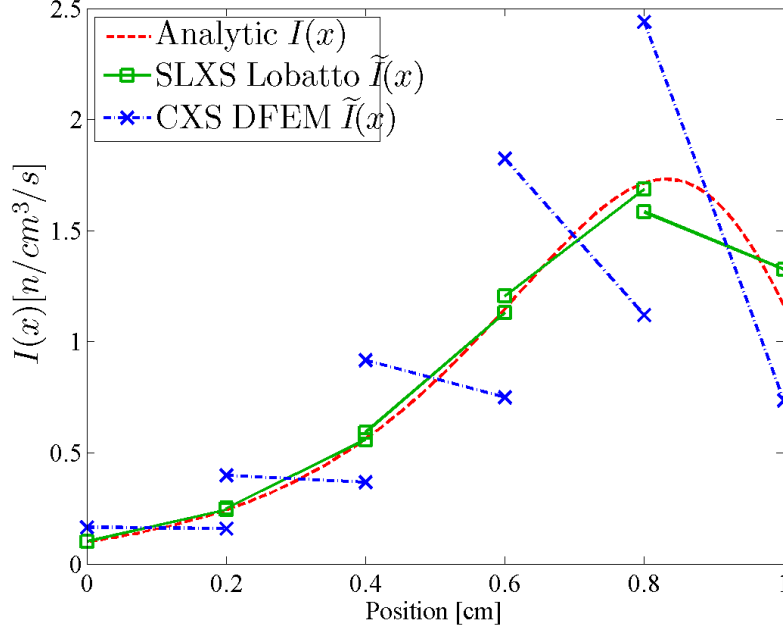


Figure 3.28: Plot of the linear trial space SLXS Lobatto and CXS DFEM approximations of $\widetilde{IR}(x)$ for the pure absorber problem with exponentially varying cross section.

In this problem, there are two possible sources of error that could cause a DFEM to be inaccurate: inexact matrix evaluation and not incorporating cross-section spatial variation into the scheme. By definition, CXS DFEM exactly integrates the mass matrix, and we showed in Section 2 that schemes that exactly integrate the mass matrix are more accurate than schemes that only approximately integrate the mass matrix, like SLXS Lobatto. Thus, the poor accuracy of CXS DFEM relative to SLXS Lobatto is entirely caused by the approximation of a spatially varying cross section with a cell-wise constant value.

The “blading” in $\widetilde{IR}(x)$ has not previously been reported in neutron transport

or thermal radiative transport community literature. We are likely not the first to have generated these large, non-monotonic discontinuities. In fact, we believe that blading has frequently been present in DFEM both neutron transport and thermal radiative transfer simulations but has likely gone unnoticed due to the prevalence of linear DFEM and simplified data visualization using cell midpoint values.

To demonstrate that a minor choice in data presentation can obscure the existence of blading, consider Fig. 3.29 and Fig. 3.30 that linearly interpolate between $\tilde{\psi}_{A,i}$ and $\tilde{I}\tilde{R}_{A,i}$ plotted at cell centers. Though Fig. 3.29 is visually indistinguishable from Fig. 3.25, the blading of $\tilde{I}\tilde{R}(x)$ present in Fig. 3.26 is not at all visually present in Fig. 3.30. Interaction rate terms are present in other radiation transport physics. In particular, we think of the radiative transfer analog to the neutronics interaction rate, absorption rate density. We hypothesize here, and will show in Section 6 that assuming cell-wise constant opacities for problems with temperature dependent opacities introduces blading in thermal radiative transfer temperature solutions, via the material energy equation's absorption rate density term.

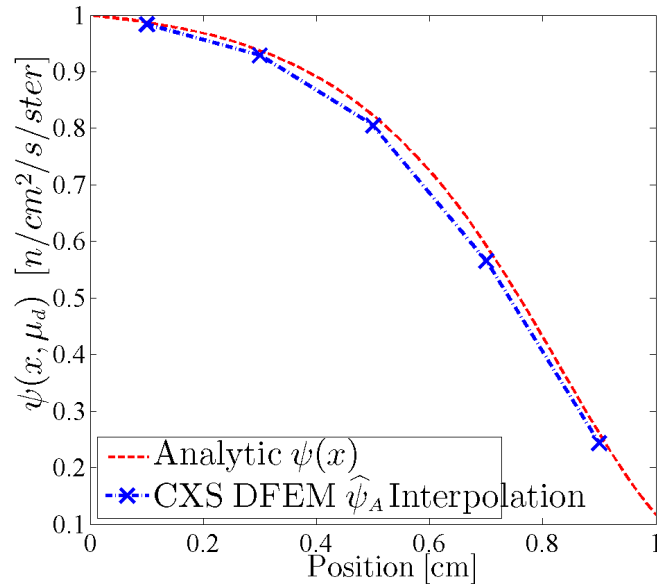


Figure 3.29: Plot of the CXS DFEM cell average angular flux at cell centers with linear interpolation for a pure absorber with exponentially varying spatial cross section.

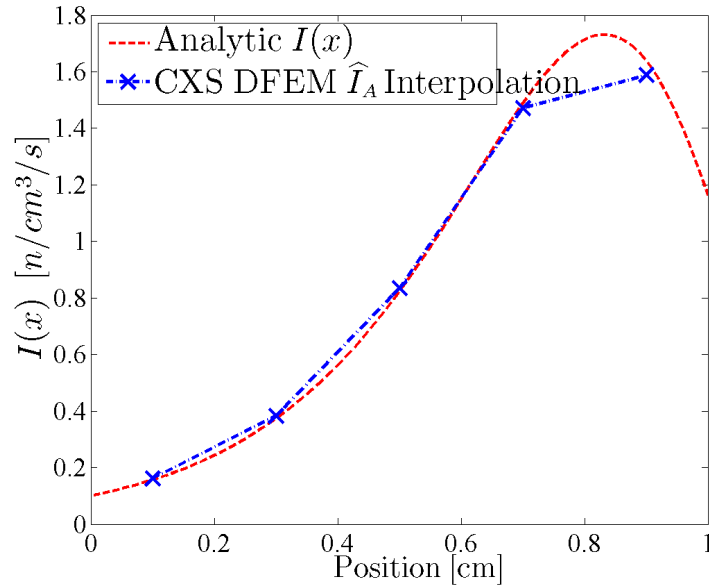


Figure 3.30: Plot of the CXS DFEM cell average interaction rate at cell centers with linear interpolation for a pure absorber with exponentially varying spatial cross section.

3.3.4 Flux-Weighting versus Volume-Averaged Cross Sections

In our results thus far, we have only considered volume-averaged cell-wise cross sections (the CXS DFEM scheme). However, in reactor physics problems, a flux-weighted cross section is often used to generate spatially averaged cross sections [35]. We now introduce the flux-weighted cell-wise constant cross section scheme (FW CXS), which differs from the CXS DFEM scheme only by how $\hat{\Sigma}$ is defined in each cell. For the FW CXS scheme, we define $\hat{\Sigma}_i$ as

$$\hat{\Sigma}_i = \frac{\int_{x_{i-1/2}}^{x_{i+1/2}} \Sigma(x) \psi(x, \mu_d) dx}{\int_{x_{i-1/2}}^{x_{i+1/2}} \psi(x, \mu_d) dx}. \quad (3.27)$$

In practice, flux-weighting is often done using the scalar flux in order not to have angle-dependent total cross section. However for the beam problem we consider here, $\psi(x, \mu_d)$ is proportional to the scalar flux.

We first compare the accuracy of FW CXS versus volume-averaged CXS DFEM for a cubic DFEM trial space, as shown in Figs. 3.31-3.34. In Figs. 3.31-3.34, we omit a plot of $E_{\psi_{out}}$ as we have already demonstrated that the accuracy of any method in calculating $E_{\psi_{out}}$ determines the method's accuracy in calculating E_{IRA} . That is, if E_{IRA} converges at a given rate, $E_{\psi_{out}}$ converges at the same rate.

Figures 3.31-3.33 show that FW CXS scheme is more accurate than CXS DFEM when comparing E_{ψ} , E_{ψ_A} and, at low resolutions, E_{IR} . However, though designed to preserve cell average interaction rates, FW CXS scheme is not only less accurate than CXS DFEM in calculating cell average interaction rates, it converges E_{IRA} at most second order in space, whereas a volume-averaged cross section converges $E_{IRA} \propto 2P + 1$ for the pure absorber problem.

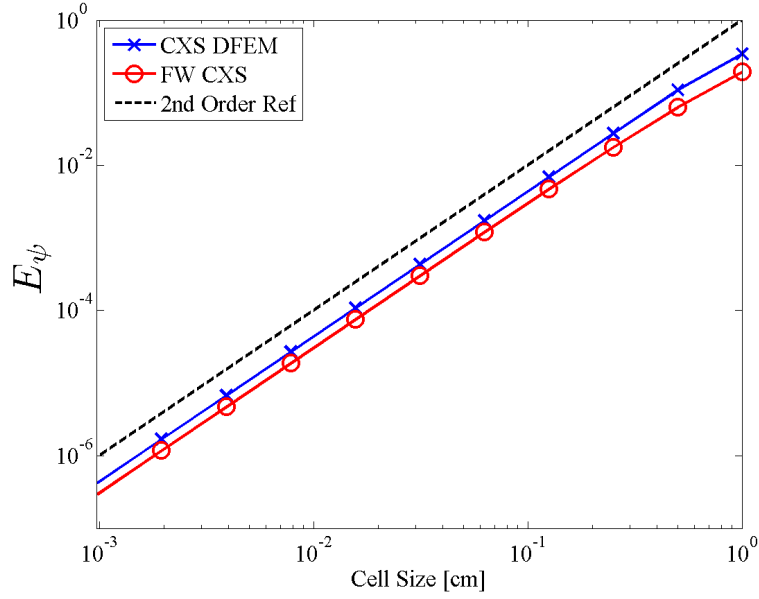


Figure 3.31: Accuracy comparison of flux weighted constant cross section scheme (FW CXS) to volume averaged cross section scheme (CXS DFEM) with a cubic angular flux trial space.

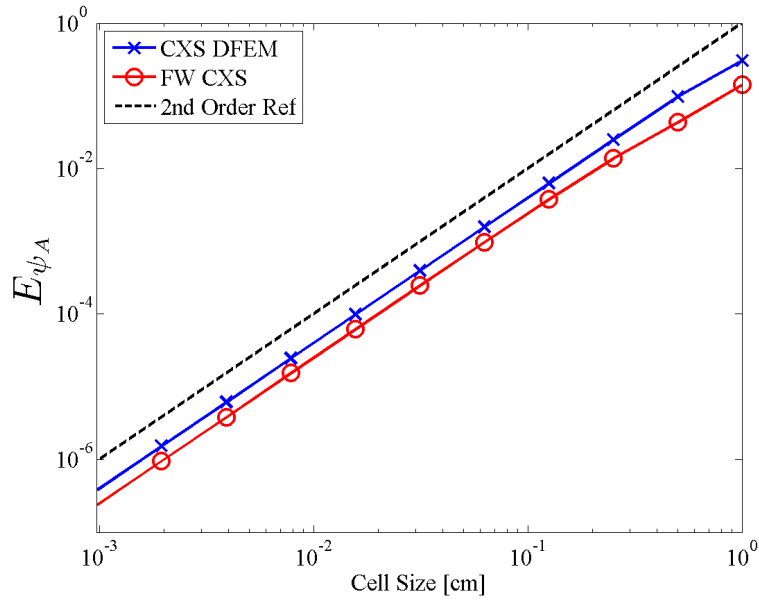


Figure 3.32: Accuracy comparison of flux weighted constant cross section scheme (FW CXS) to volume averaged cross section scheme (CXS DFEM) with a cubic angular flux trial space.

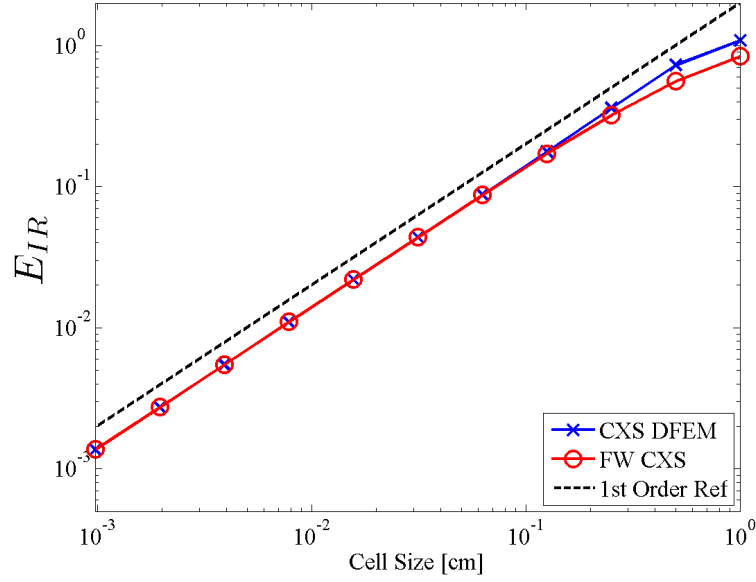


Figure 3.33: Accuracy comparison of flux weighted constant cross section scheme (FW CXS) to volume averaged cross section scheme (CXS DFEM) with a cubic angular flux trial space.

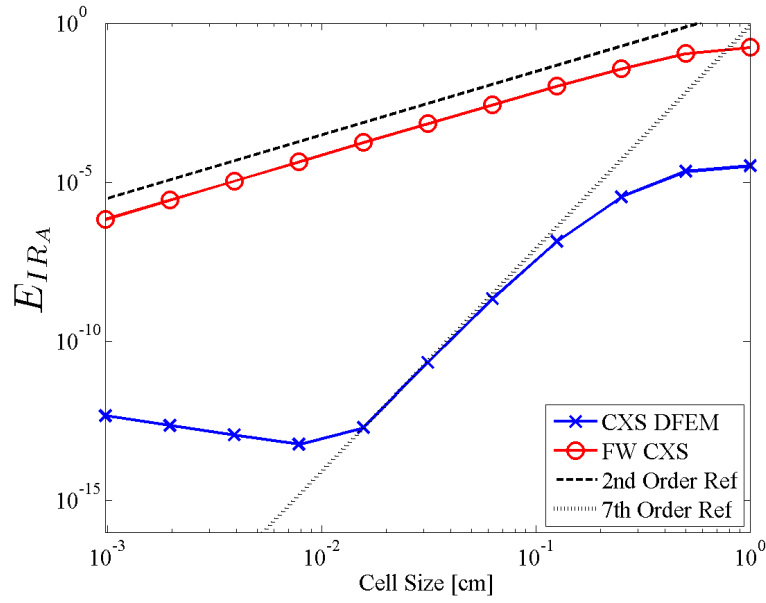


Figure 3.34: Accuracy comparison of flux weighted constant cross section scheme (FW CXS) to volume averaged cross section scheme (CXS DFEM) with a cubic angular flux trial space.

Finally, we consider the FW CXS scheme's $\tilde{\psi}(x)$ in Fig. 3.35 and $\tilde{IR}(x)$ in Fig. 3.36. In Fig. 3.35 and Fig. 3.36 the FW CXS and CXS DFEM schemes calculate slightly different solution representations. However, the FW CXS scheme exhibits the same interaction rate blading phenomena as the CXS DFEM scheme, reiterating that blading is a result of approximating a spatially varying cross section as a cell-wise constant. The choice of cell-wise cross section does not eliminate blading.

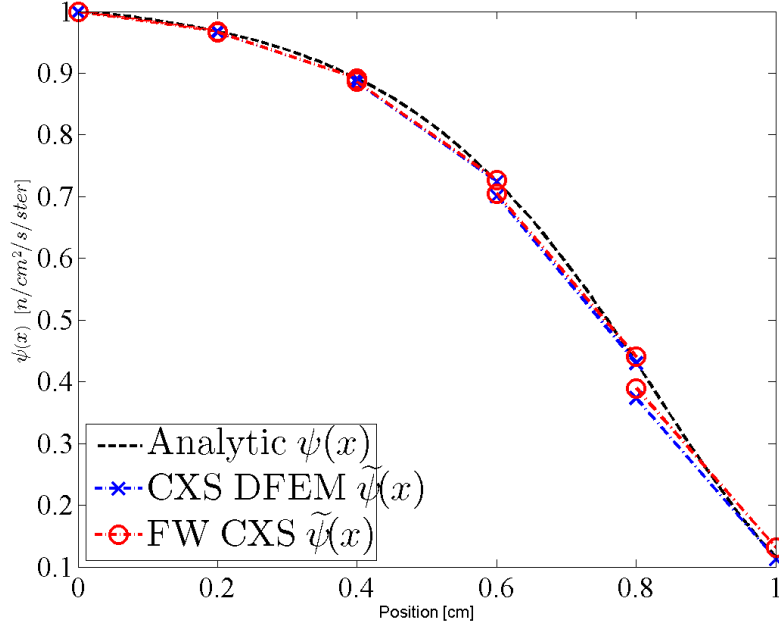


Figure 3.35: Plot of the linear trial space FW CXS and CXS DFEM approximations to $\tilde{\psi}(x)$ for the pure absorber problem with exponentially varying spatial cross section.

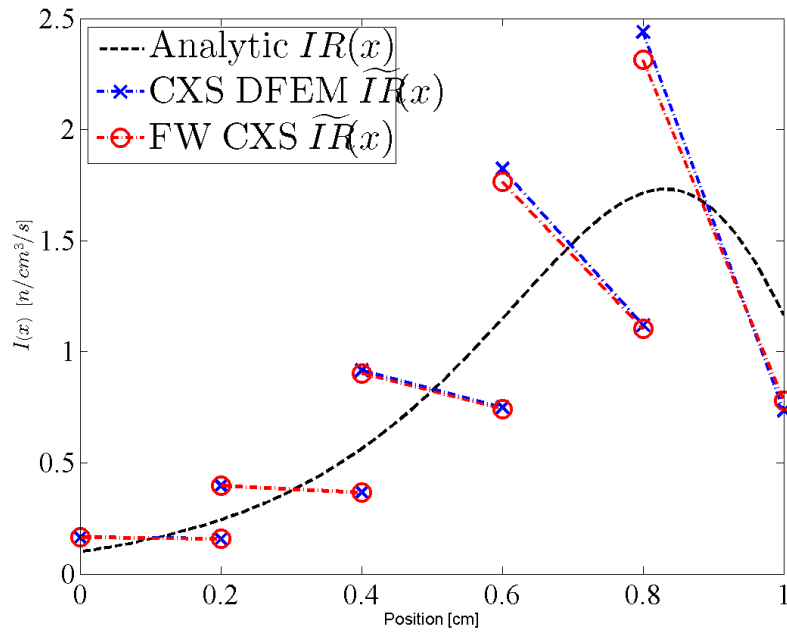


Figure 3.36: Plot of the linear trial space FW CXS and CXS DFEM approximations to $\widetilde{IR}(x)$ for the pure absorber problem with exponentially varying spatial cross section.

Table 3.2: Shorthand notation of different cell spacing schemes.

Spacing Label	Spacing Type
EQUAL	Equally-spaced cells
MFP	Constant optical thickness cells
LOG	Logarithmically spaced cells

3.3.5 Effects of Mesh Spacing

In practice, computational domains are not necessarily discretized with uniform grids; rather cells are concentrated in regions where the solution is known or assumed to vary rapidly. For the pure absorber problem, we compare two alternative methods of mesh spacing (logarithmic grids and equal optical thickness grids, see Table 3.2), to the results obtained with equally-spaced mesh cells.

We derive how to generate these meshes in Section 3.3.5.1 and give results in Section 3.3.5.2

3.3.5.1 Generating Improved Spatial Meshes

Two alternative meshing strategies are compared to equally-spaced meshed. In the following, we will use a shorthand notation, given in Table 3.2. With the MFP meshing strategy, we find each cell width by determining the width of each cell from $i = 1$ (leftmost cell) to $i = N_{cell}$ as outlined by Eq. (3.28): First, we determine the average cell optical thickness:

$$\bar{h} = \frac{\int_{x_{1/2}}^{x_{N_{cell}+1/2}} \Sigma_t(x) dx}{N_{cell}}. \quad (3.28a)$$

Then, we solve the following equation for $x_{i+1/2}$:

$$\int_{x_{i-1/2}}^{x_{i+1/2}} \Sigma_t(x) dx - \bar{h} = 0, \quad (3.28b)$$

yielding

$$x_{i+1/2} = \frac{1}{c_2} \log \left[\frac{c_2(\bar{h} + \Sigma_t(x_{i-1/2}))}{c_1} \right]. \quad (3.28c)$$

There are several ways to specify LOG spacing, but we elected to set a ratio, 0.6, between adjacent cell sizes with the caveat that we would set a minimum cell size, Δx_{min} . In our convergence testing, at the first refinement when $\Delta x_{N_{cell}} < \Delta x_{min}$, the grid is “fixed” and all further refinements uniformly refine the ”fixed” grid. $\Delta x_{N_{cell}}$ is the cell width for the right most cell where, for $R < 1$,

$$\Delta x_i = \Delta x_1 R^{i-1}, \quad i \in [1, N_{cell}]. \quad (3.29)$$

Δx_1 is determined by requiring that the geometric series of cell widths completely fill the space:

$$\Delta x_1 = (x_{N_{cell}+1/2} - x_{1/2}) \frac{1 - R}{1 - R^{N_{cell}}} \quad (3.30)$$

The grid is “fixed” by resetting the width of every cell whose width, if set to the value required for a purely logarithmically spaced grid with R would be below Δx_{min} , to Δx_{min} . After imposing this, cell widths are determined by requiring the cells that were not reset to fill the problem space logarithmically using R . If there is no minimum cell width, at high mesh refinements, most cells will be infinitesimally small and the large cells will never be refined, causing error to stagnate. Logarithmic spacing represents the “smart” meshing strategy most likely to be employed in engineering practice as it requires the least amount of solution information prior to problem execution. For all of our calculations, we set $R = 0.6$ and $\Delta x_{min} = 10^{-3} [cm]$.

3.3.5.2 Mesh Spacing Results

We begin our discussion of alternate mesh spacing by first noting that the choice of mesh spacing method does not alter asymptotic convergence rates, as shown in

Figs. 3.37-3.40. The convergence of the SLXS Lobatto scheme using a quadratic trial space for E_ψ is shown in Fig. 3.37, for E_{ψ_A} in Fig. 3.38, for E_{IR} in Fig. 3.39, and for E_{IR_A} in Fig. 3.40. Regardless of the choice of mesh spacing, all error quantities of interest converge at the same asymptotic rate. Plots showing other trial space degrees and DFEM schemes are omitted for brevity. We also omit showing the convergence of $E_{\psi_{out}}$ as we have already demonstrated that the convergence rate of $E_{\psi_{out}}$ and E_{IR_A} are related and identical.

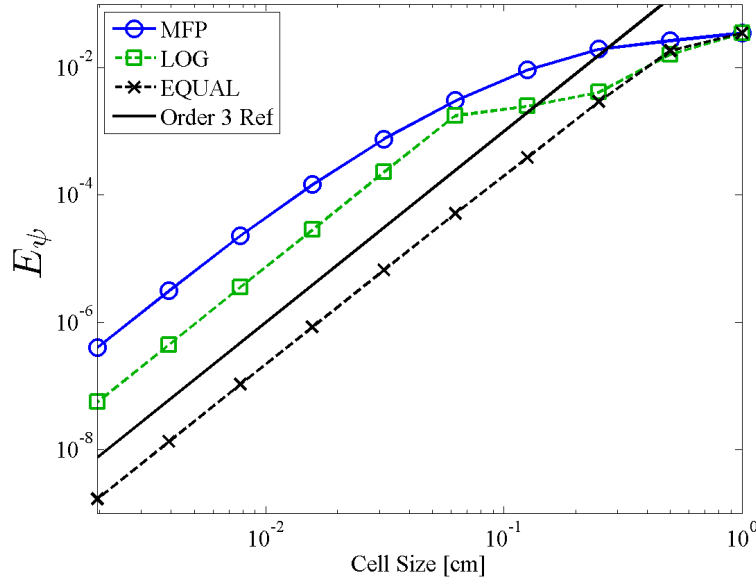


Figure 3.37: Asymptotic convergence of the SLXS Lobatto scheme using a quadratic trial space, for different mesh spacing methodologies.

At mesh refinements that are not in the asymptotic convergence regime, the intelligent meshing can result in a significant reduction in error. Consider the difference smart meshing has in reducing E_ψ , E_{ψ_A} , E_{IR} , and E_{IR_A} for the SLXS Lobatto scheme in Figs. 3.41-3.44, for the SLXS Gauss scheme in Figs. 3.45-3.48, and for CXS DFEM

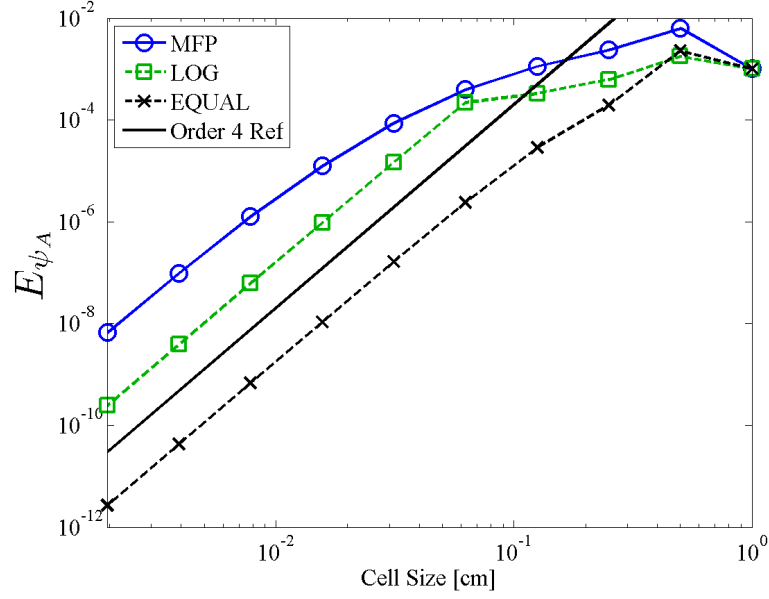


Figure 3.38: Asymptotic convergence of the SLXS Lobatto scheme using a quadratic trial space, for different mesh spacing methodologies.

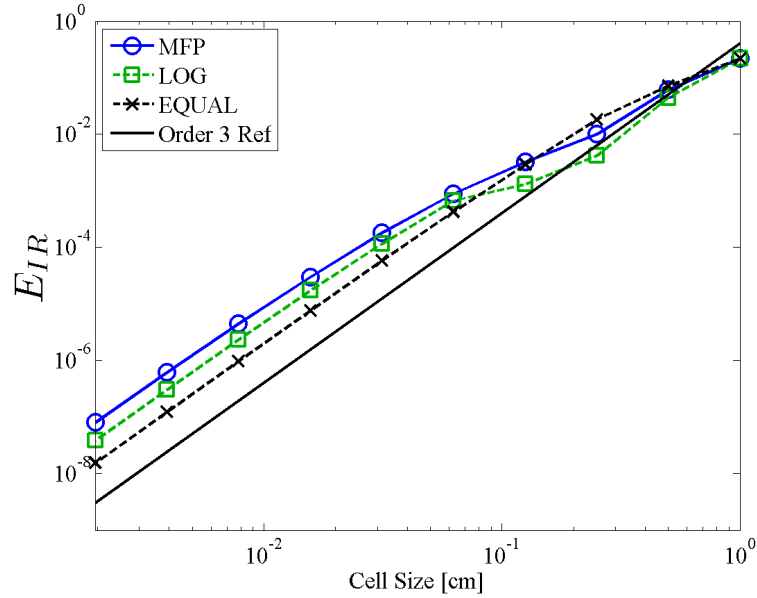


Figure 3.39: Asymptotic convergence of the SLXS Lobatto scheme using a quadratic trial space, for different mesh spacing methodologies.

in Figs. 3.49-3.52 when using a quadratic DFEM trial space at low cell resolutions.

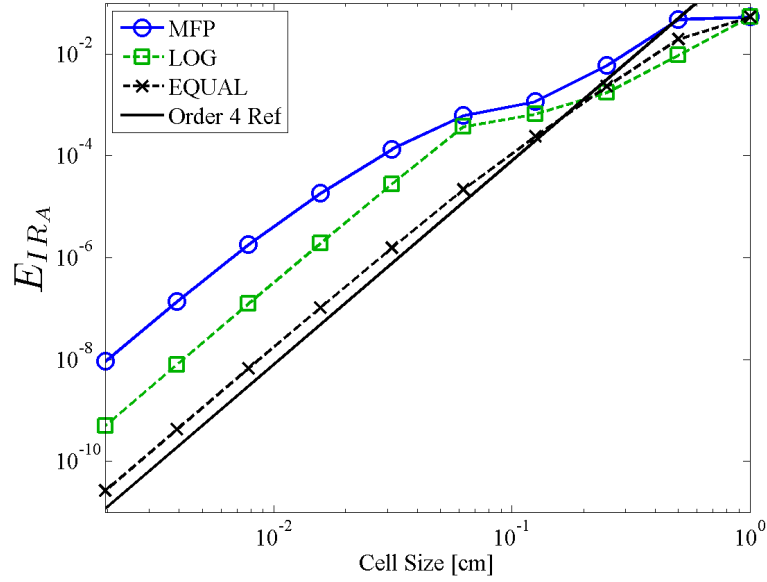


Figure 3.40: Asymptotic convergence of the SLXS Lobatto scheme using a quadratic trial space, for different mesh spacing methodologies.

First, we note that any particular mesh spacing methodology results in solutions that are more accurate for certain quantities, but not for all quantities. For example, with the self-lumping schemes, E_ψ is generally smaller when using an equally spaced mesh, but using the LOG mesh results in orders of magnitude improvement in E_{IR} and E_{IRA} on coarse meshes. Figures 3.49-3.52, CXS DFEM again illustrates that LOG spacing is more accurate in calculating interaction rate quantities than an equally-spaced mesh and equally-spaced meshes are generally more accurate than other meshing strategies for calculating angular flux quantities. However, CXS DFEM shows a two order of magnitude reduction in calculating E_{IRA} when using a mesh that has a uniform optical thickness in each cell. This is a direct result of CXS DFEM converging as $\hat{\sigma}\Delta x \rightarrow 0$.

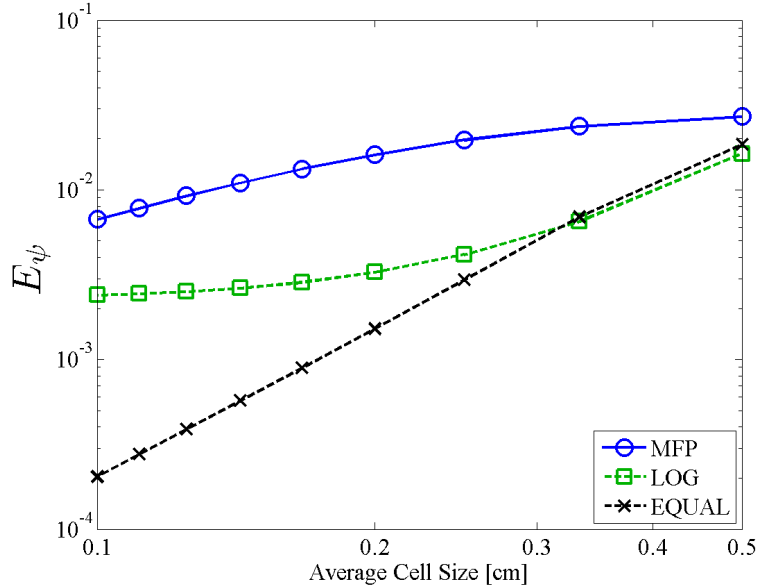


Figure 3.41: Errors associated with the SLXS Lobatto using a quadratic trial space scheme for different mesh spacing methodologies, at low resolutions.

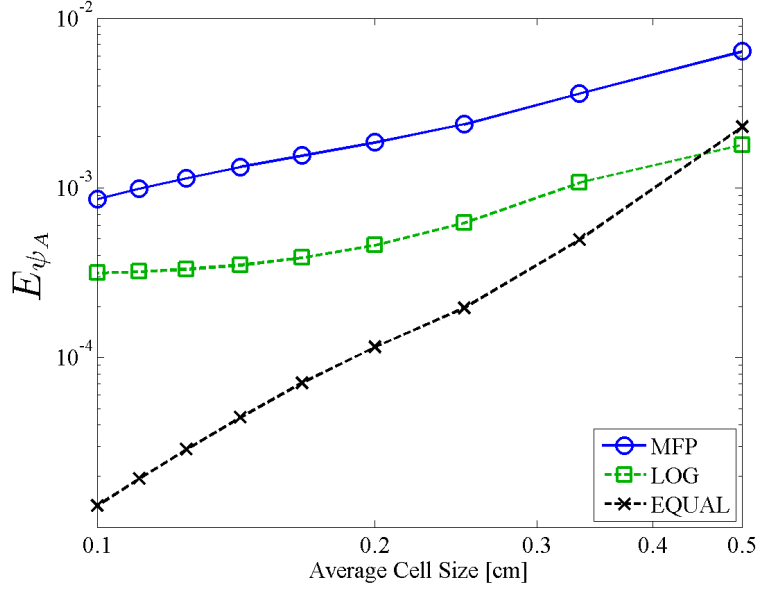


Figure 3.42: Errors associated with the SLXS Lobatto using a quadratic trial space scheme for different mesh spacing methodologies, at low resolutions.

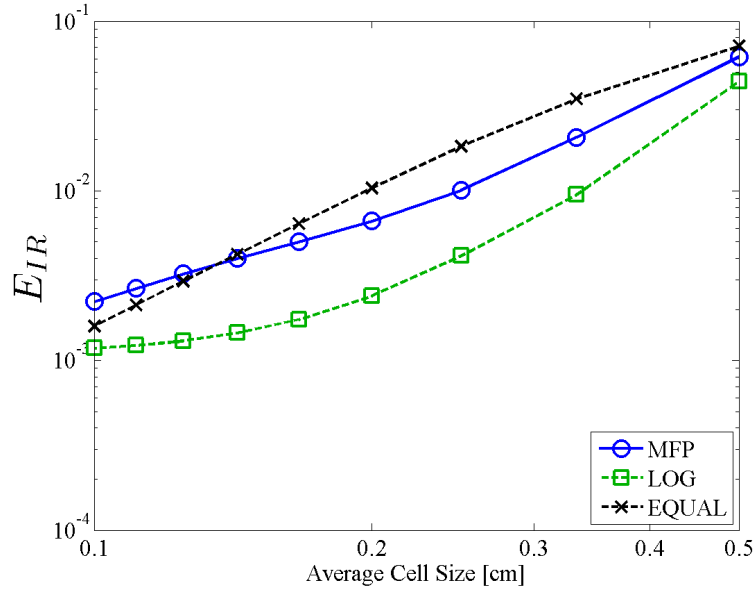


Figure 3.43: Errors associated with the SLXS Lobatto using a quadratic trial space scheme for different mesh spacing methodologies, at low resolutions.

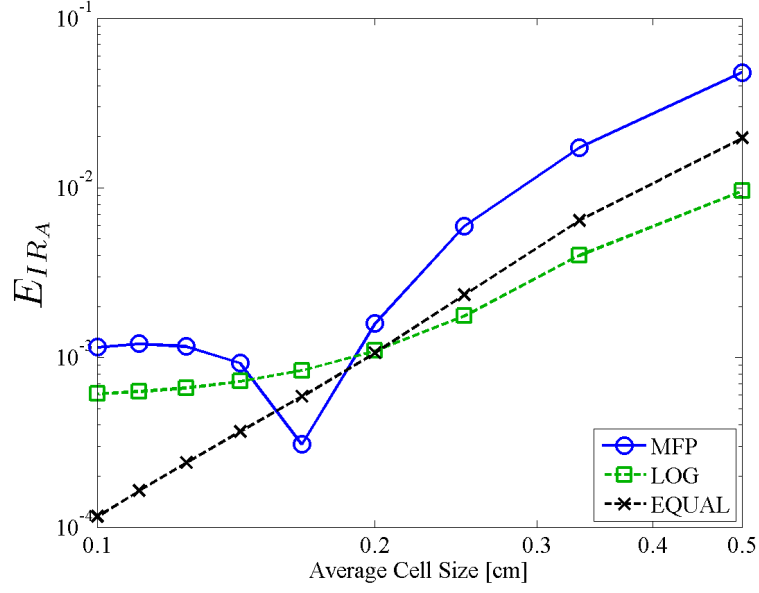


Figure 3.44: Errors associated with the SLXS Lobatto using a quadratic trial space scheme for different mesh spacing methodologies, at low resolutions.

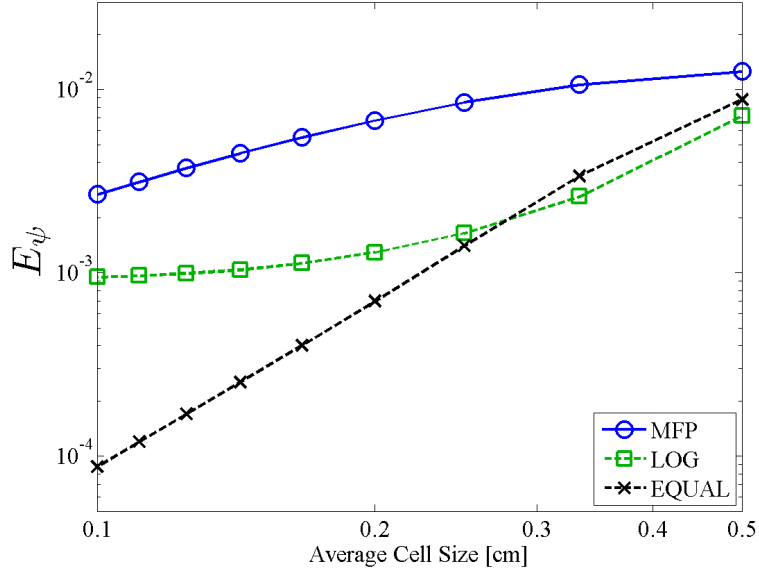


Figure 3.45: Errors associated with the SLXS Gauss using a quadratic trial space for different mesh spacing methodologies, at low resolutions.

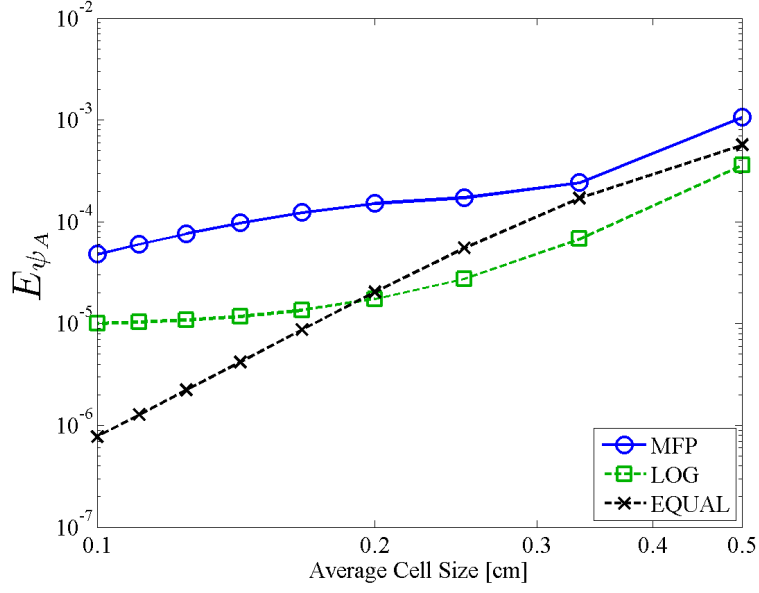


Figure 3.46: Errors associated with the SLXS Gauss using a quadratic trial space for different mesh spacing methodologies, at low resolutions.

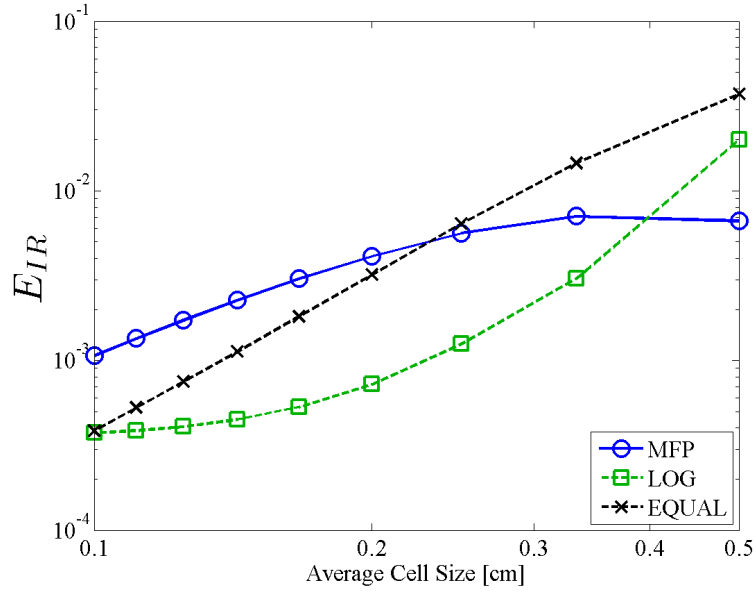


Figure 3.47: Errors associated with the SLXS Gauss using a quadratic trial space for different mesh spacing methodologies, at low resolutions.

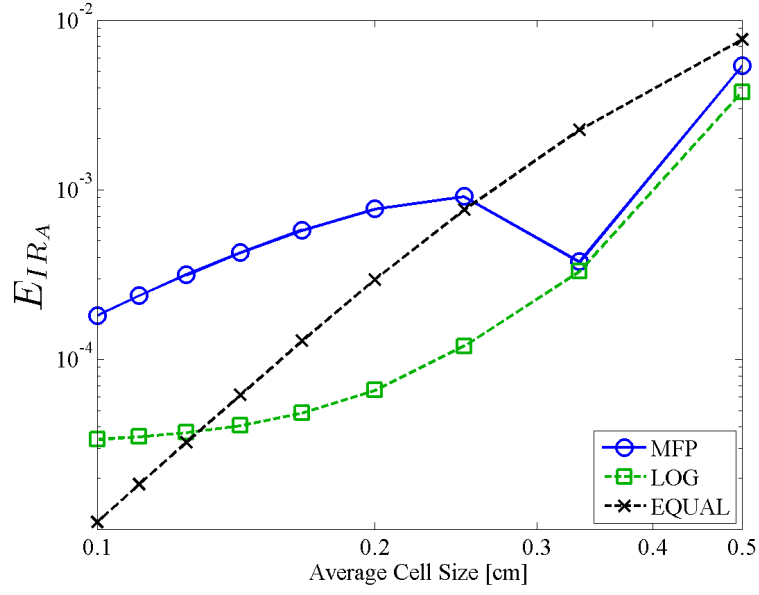


Figure 3.48: Errors associated with the SLXS Gauss using a quadratic trial space for different mesh spacing methodologies, at low resolutions.

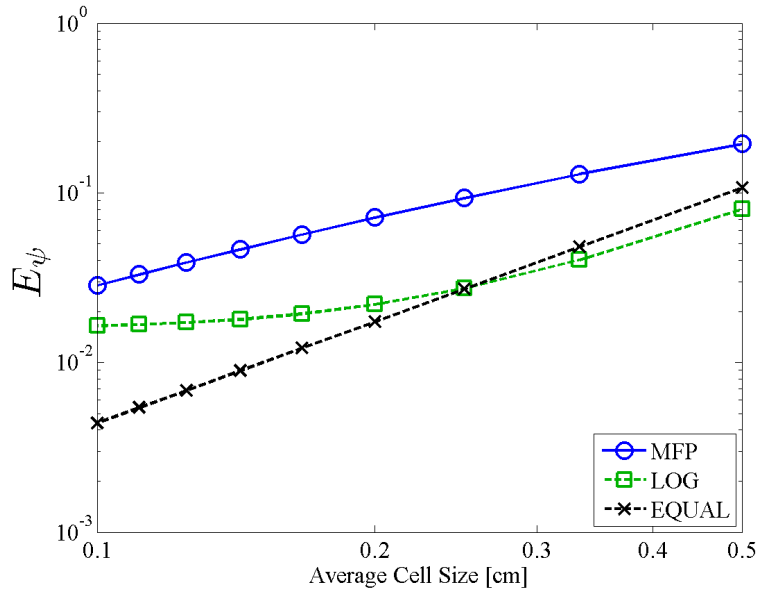


Figure 3.49: Errors associated with CXS DFEM using a quadratic trial space for different mesh spacing methodologies, at low resolutions.

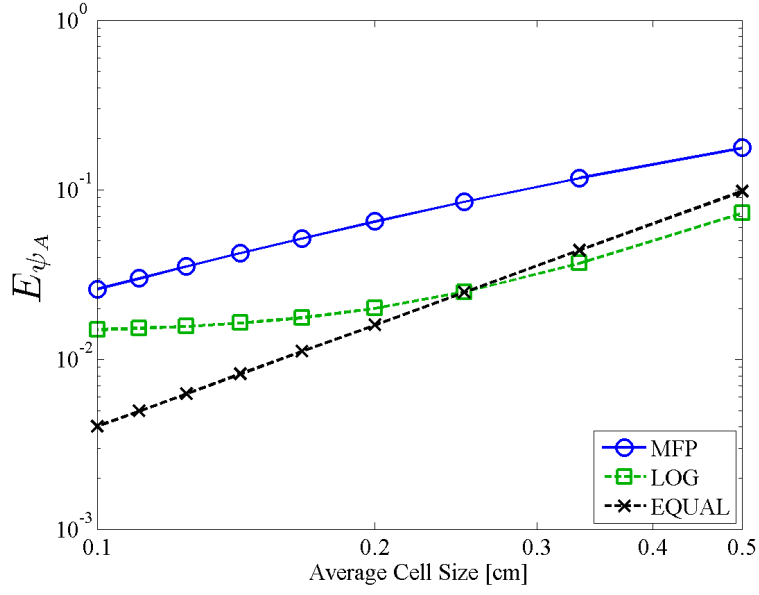


Figure 3.50: Errors associated with CXS DFEM using a quadratic trial space for different mesh spacing methodologies, at low resolutions.

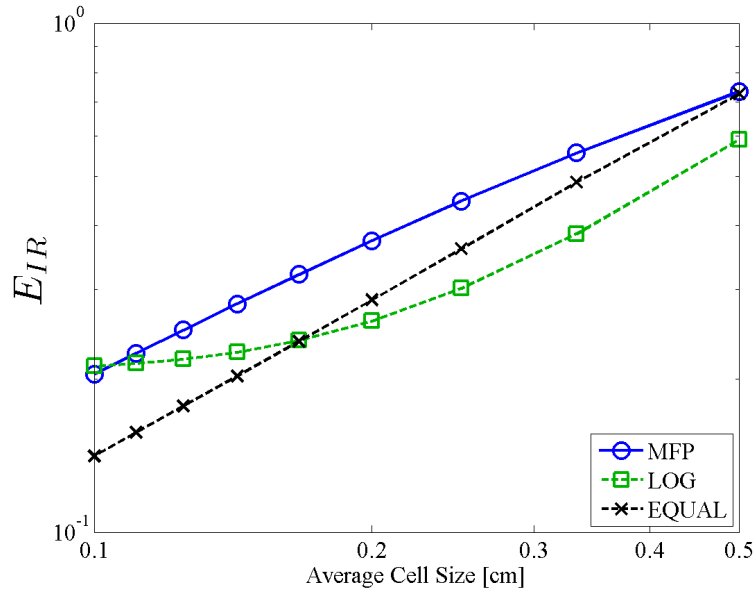


Figure 3.51: Errors associated with CXS DFEM using a quadratic trial space for different mesh spacing methodologies, at low resolutions.

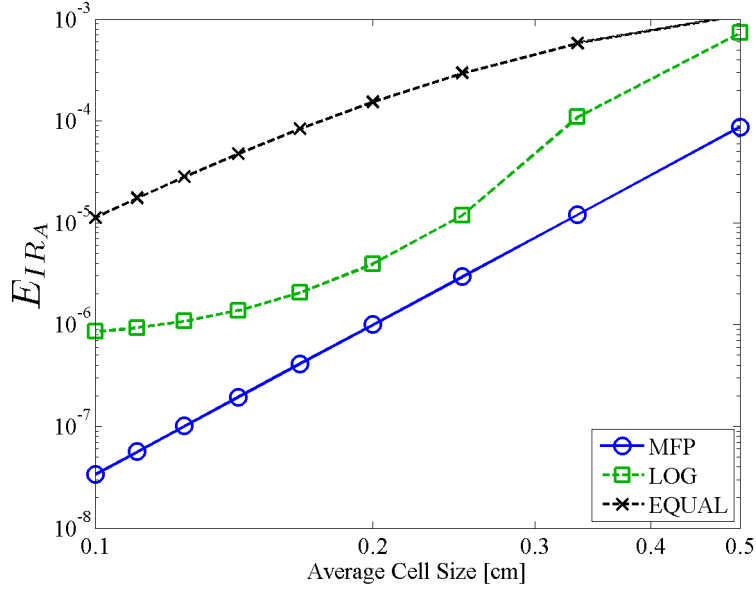


Figure 3.52: Errors associated with CXS DFEM using a quadratic trial space for different mesh spacing methodologies, at low resolutions.

3.4 Conclusions to Be Carried Forward

The SLXS Lobatto and SLXS Gauss schemes are accurate for problems S_N neutron transport problems that have within cell spatial interaction cross-section variation. Though no less accurate for the case of spatially varying cross sections that for cell-wise constant cross section, the SLXS Newton-Cotes scheme will no longer be considered. SLXS Newton-Cotes is not as asymptotically accurate as SLXS Gauss or SLXS Lobatto schemes, though the SLXS Lobatto scheme is equivalent to SLXS Newton-Cotes for linear and quadratic trial spaces. We continued to consider a self-lumping scheme using closed Newton-Cotes quadrature due to the robustness SL Newton-Cotes exhibited for the cell-wise constant cross section case. However, since no scheme other than linear SLXS Lobatto will yield strictly positive angular flux outflow for arbitrary within cell cross section variation, we conclude that SLXS

Lobatto has all of the strengths exhibited by SLXS Newton-Cotes, with the added benefit of increased accuracy.

We have also documented that the assumption of a cell-wise constant cross section will in general be less accurate than using a self-lumping scheme that explicitly accounts for within cell variation of cross section when constructing \mathbf{R} . Though the CXS DFEM was more accurate than self-lumping schemes in calculating E_{IR_A} , and $E_{\psi_{out}}$, when using the exact volume average cross section, this is more a function of our test problem being a pure absorber than an inherent benefit of CXS DFEM that will be true for all problems of interest.

4. ITERATIVE ACCELERATION FOR THE S_N NEUTRON TRANSPORT EQUATIONS

Thus far, we have failed to discuss how the global system of angular flux unknowns is solved, focusing instead on the solution of a single system of equations that describes the unknowns of a single spatial cell, for a single discrete direction. We have implicitly assumed the existence and knowledge of a transport sweep [29], or single, un-accelerated Richardson iteration. In Section 4.1, we explain the fundamental iterative techniques used to solve the neutron transport equation for scatter media. In Section 4.2 we discuss the choices available for synthetic accelerators. Finally we two different synthetic acceleration techniques compatible with our chose spatial discretizations of the transport equation. In Section 4.3 we derive the S_2 synthetic acceleration (S2SA) technique[36] and in Section 4.4 we derive a modified interior penalty (MIP) diffusion synthetic acceleration[37] (DSA) operator. Finally, in Section 4.5, we compare verify the implementation of each for a set of test problems with spatially constant and and spatially varying interaction cross sections.

4.1 Iterative Solution of the Neutron Transport S_N Equations

To describe the the iterative process by which the discrete ordinates neutron transport equations are solved, we re-visit the spatially analytic, steady-state, mono-energetic discrete ordinates neutron transport equation, but do not have a monolithic right hand side source. Rather, we treat the right hand side as having both an isotropic scattering component, and a fixed source component, as given in Eq. (4.1).

$$\mu \frac{d\psi_d}{dx} + \Sigma_t \psi_d = \frac{\Sigma_s}{4\pi} \phi + S_d. \quad (4.1)$$

The traditional practice is to solve Eq. (4.1) iteratively with Richardson iteration. Each Richardson iteration is referred to as a transport sweep, where for a fixed right hand side, ψ_d is update direction by direction, cell by cell, sweeping from each direction's incident boundary through the entire mesh with upwinding at cell interfaces. Introducing iteration index ℓ , this process can be written as:

$$\mu \frac{d\psi_d^{(\ell+1)}}{dx} + \Sigma_t \psi_d^{(\ell+1)} = \frac{\Sigma_s}{4\pi} \phi^{(\ell)} + S_d. \quad (4.2)$$

After we find, $\psi_d^{(\ell+1)}$, we update $\phi^{(\ell+1)}$ using the discrete ordinates definition:

$$\phi^{(\ell+1)} = 2\pi \sum_{d=1}^{N_{dir}} w_d \psi_d^{(\ell+1)}. \quad (4.3)$$

Though convergent, the source iteration process can converge arbitrarily slow, as shown by Larsen[38], when $\frac{\Sigma_s}{\Sigma_t} \rightarrow 1$. To accelerate the convergence of source iteration, the diffusion synthetic acceleration (DSA) technique was developed [39]. DSA is best explained through example.

To start, we consider to a single source iteration:

$$\mu \frac{d\psi_d^{(\ell+1/2)}}{dx} + \Sigma_t \psi_d^{(\ell+1/2)} = \frac{\Sigma_s}{4\pi} \phi^{(\ell)} + S_d. \quad (4.4)$$

Subtracting Eq. (4.1) from Eq. (4.4) yields:

$$\mu \frac{d\delta\psi_d^{(\ell+1/2)}}{dx} + \Sigma_t \delta\psi_d^{(\ell+1/2)} = \frac{\Sigma_s}{4\pi} \delta\phi^{(\ell)}, \quad (4.5)$$

where we have defined the iterative error of the angular flux, $\delta\psi_d^{(\ell+1)}$,

$$\psi_d = \psi_d^{(\ell+1)} + \delta\psi_d^{(\ell+1)}, \quad (4.6)$$

and scalar flux iterative error, $\delta\phi^{(\ell)}$

$$\phi = \phi^{(\ell)} + \delta\phi_d^{(\ell)} . \quad (4.7)$$

Subtracting $\frac{\Sigma_s}{4\pi}\delta\phi^{(\ell)}$ from both sides of Eq. (4.5), we see arrive at:

$$\mu \frac{d\delta\psi_d^{(\ell+1/2)}}{dx} + \Sigma_t \delta\psi_d^{(\ell+1/2)} - \frac{\Sigma_s}{4\pi} \delta\phi^{(\ell+1/2)} = \frac{\Sigma_s}{4\pi} \delta\phi^{(\ell)} - \frac{\Sigma_s}{4\pi} \delta\phi^{(\ell+1/2)} . \quad (4.8)$$

Recognizing:

$$\phi = \phi^{(\ell+1/2)} + \delta\phi^{(\ell+1/2)} , \quad (4.9a)$$

$$\phi = \phi^{(\ell)} + \delta\phi^{(\ell)} , \quad (4.9b)$$

$$\phi^{(\ell+1/2)} - \phi^{(\ell)} = (\phi - \delta\phi^{(\ell+1/2)}) - (\phi - \delta\phi^{(\ell)}) , \text{ and} \quad (4.9c)$$

$$\delta\phi^{(\ell)} - \delta\phi^{(\ell+1)} = \phi^{(\ell+1/2)} - \phi^{(\ell)} , \quad (4.9d)$$

Eq. (4.8) becomes:

$$\mu \frac{d\delta\psi_d^{(\ell+1/2)}}{dx} + \Sigma_t \delta\psi_d^{(\ell+1/2)} = \frac{\Sigma_s}{4\pi} \delta\phi^{(\ell+1/2)} + \frac{\Sigma_s}{4\pi} (\phi^{(\ell+1/2)} - \phi^{(\ell)}) . \quad (4.10)$$

Equation 4.10 indicates that if we could solve a transport problem with a driving source equal to the difference between two scattering iterates,

$$\frac{\Sigma_s}{4\pi} (\phi^{(\ell+1/2)} - \phi^{(\ell)}) , \quad (4.11)$$

then we could get the iterative error of iteration $\delta\phi^{(\ell+1/2)}$, add to the $\phi^{(\ell+1/2)}$ we already have, and then have the exact solution, ϕ . However, solving Eq. (4.10) is as difficult as solving the original problem in Eq. (4.1). Alternatively, if we could solve

an approximation to Eq. (4.10), perhaps the result would satisfy:

$$\phi \approx \Delta\phi^{(\ell+1/2)} + \phi^{(\ell+1/2)}, \quad (4.12)$$

where $\Delta\phi^{(\ell+1/2)}$ comes from the lower order approximation to Eq. (4.10). The idea of using a lower order approximation to approximately solve Eq. (4.10) is central to synthetic acceleration.

4.2 Qualitative Comparison of Different Synthetic Acceleration Techniques

Three types of synthetic acceleration have received significant attention in the neutron transport and thermal radiative transfer literature, S_2 synthetic acceleration (S2SA)[36], diffusion synthetic acceleration (DSA)[39], and transport synthetic acceleration (TSA)[40]. Each method of synthetic acceleration has both advantages and disadvantages relating to the computational efficiency and iterative effectiveness of each method.

The S2SA method was shown to be iteratively effective in both slab and 1-D spherical geometries. Additionally, it is easily compatible with any DFEM spatial discretization of the S_N neutron transport equations. S2SA solves for ψ_+ and ψ_- using a single global matrix solve, rather than a direction by direction solve for the full transport ψ_d unknowns. The full transport scalar flux iterative update is then defined as,

$$\Delta\phi^{(\ell+1/2)} \approx 2\pi [w_+\psi_+ + w_-\psi_-], \quad (4.13)$$

where w_+ , w_- correspond to the weights of a direction quadrature (typically Gauss) set with corresponding discrete direction μ_+ and μ_- , and ψ_+ , ψ_- are the fundamental unknowns of the S_2 discretization. Thus, S2SA uses the same local matrices of Eq. (3.2) as the full transport operator. However, the S2SA global matrix that must be

inverted to solve for ψ_+ and ψ_- is extremely difficult to invert for multiple spatial dimensions. This makes S2SA iteratively effective, but computationally expensive, limiting the extensibility of any techniques that require S2SA. But since our initial focus is on solving slab geometry neutron transport problems, we will continue to consider S2SA as it is readily adaptable to our new spatial discretization schemes. [36]

Gelbard and Hageman first showed in [39] that diffusion synthetic acceleration (DSA) could be used to accelerate the convergence of source iteration in neutron transport since the diffusion operator effectively attenuates the slowly varying error modes that hinder the convergence of source iteration. To be unconditionally effective, Larsen showed that DSA needed to be derived in a method consistent with the spatial and angular discretization of the transport equation [38]. Adams and Martin first showed that partially consistent diffusion discretizations could be used to effectively accelerate DFEM spatial discretizations of the neutron transport equation [41]. Though shown to be unconditionally stable for certain geometries the M4S DSA proposed in [41] has been shown to be unstable for unstructured multi-dimensional geometries [42]. To allow more general applicability, we wish to consider a more advanced DSA discretization. Alternative DSA discretizations that have been applied successfully to unstructured multi-dimensional geometries include: the partially consistent WLA DSA proposed in [43], the fully consistent DSA (FCDSA) proposed in [42], and the partially consistent MIP DSA proposed in [37]. WLA DSA produces a symmetric positive definite (SPD) diffusion matrix and is unconditionally stable, but the spectral radius of the WLA DSA scheme increases on distorted mesh cells and for optically thick cells with scattering ratios very close to unity [43, 42]. While the FCDSA scheme remains effective in optically thick cells, it creates a diffusion operator that is very difficult and costly to invert[42]. The MIP DSA discretization

[37] of Wang and Ragusa generates a SPD diffusion operator, remains effective for all cell optical thicknesses, has been successfully applied to high order DFEM S_N transport, and can be used with adaptive mesh refinement. Further, it was shown in [44] that the MIP DSA diffusion operator can be inverted very quickly using advanced preconditioners such as algebraic multi-grid. Thus, if we can ideally define a MIP discretization that is iteratively effective for neutron transport problems discretized with our higher order DFEM methods that account for the spatial variation of cross section within each cell, we will have found a scheme that is most likely to prove useful in more meaningful (multiple spatial dimension) thermal radiative transfer simulations.

TSA differs from S2SA and DSA in that it does not attempt to invert a single matrix that describes a lower order operator approximation to the full discrete ordinates neutron transport equation. The main advantage of TSA over DSA is that using TSA allows for re-use of much of the software already developed for transport sweeps, greatly lowering software development overhead as compared to using DSA. However, TSA is in general not as effective as DSA, and finding the most efficient set of parameters that complete the description of the TSA scheme is problem dependent. [40]

We will derive an S2SA operator in Section 4.3 and MIP DSA operator in Section 4.4. We elect not to derive a TSA operator as it occupies a middle ground between, a S2SA operator that re-uses a lot of the transport sweep capability we have already developed with our high order DFEM of Chapter 3, but is computationally challenging to invert (in multiple spatial dimensions), and the MIP DSA operator we define in Section 4.4 that requires significant derivation independent of the DFEM neutron transport methodology we have already derived, but that is computationally efficient to invert.

4.3 S_2 Synthetic Acceleration

We begin our derivation by repeating the S_2 , spatially analytic angular flux update equations from Morel[36] (Eqs. (12a) and (12b)), noting that we have elected to use ψ^+ and ψ^- instead of c^+ and c^- , Σ represents macroscopic interaction cross sections rather than σ , and we define $\phi = 2\pi \sum_d w_d \psi_d$:

$$\mu_+ \frac{d\psi^+}{dx} + \Sigma_t \psi^+ = \frac{\Sigma_s}{4\pi} \Delta \phi + \frac{\Sigma_s}{4\pi} (\phi^{(\ell+1/2)} - \phi^{(\ell)}) \quad (4.14a)$$

$$\mu_- \frac{d\psi^-}{dx} + \Sigma_t \psi^- = \frac{\Sigma_s}{4\pi} \Delta \phi + \frac{\Sigma_s}{4\pi} (\phi^{(\ell+1/2)} - \phi^{(\ell)}) . \quad (4.14b)$$

In Eqs. (4.14) we are assuming scattering is isotropic only. Spatially discretizing with a P degree DFEM as in Chapter 3, for an interior cell, c , we have:

$$\left(\mu_+ \mathbf{G}_+ + \frac{\Delta x_c}{2} \mathbf{R}_{\Sigma_t} \right) \vec{\psi}_c^+ = \frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} \Delta \vec{\phi}_c + \frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_c^{(\ell+1/2)} - \vec{\phi}_c^{(\ell)} \right) + \mu_+ \psi_{in,+} \vec{f}_+ \quad (4.15a)$$

$$\left(\mu_- \mathbf{G}_- + \frac{\Delta x_c}{2} \mathbf{R}_{\Sigma_t} \right) \vec{\psi}_c^- = \frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} \Delta \vec{\phi}_c + \frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_c^{(\ell+1/2)} - \vec{\phi}_c^{(\ell)} \right) + \mu_- \psi_{in,-} \vec{f}_- . \quad (4.15b)$$

In Eqs. (4.15), we note that $\mu_+ > 0$ and $\mu_- < 0$, and as such use the \pm subscripts to define the appropriate \mathbf{G} and \vec{f} , defined in Eqs. (4.16) and Eqs. (4.17), respectively:

$$\mathbf{G}_{+,i,j} = b_i(1)b_j(1) - \int_{-1}^1 \frac{db_i}{ds} b_j(s) ds \quad (4.16a)$$

$$\mathbf{G}_{-,i,j} = -b_i(-1)b_j(-1) - \int_{-1}^1 \frac{db_i}{ds} b_j(s) ds , \quad (4.16b)$$

$$\vec{f}_{+,i} = b_i(-1) \quad (4.17a)$$

$$\vec{f}_{-,i} = -b_i(1) . \quad (4.17b)$$

Noting that the inflow to cells on the interior is the outflow from the appropriate cell, for and using the definitions of Eqs. (2.7b) and Eqs. (2.8b), we define $\psi_{in,+}\vec{f}_+$ and $\psi_{in,-}\vec{f}_-$ entirely in terms of $\vec{\psi}_{c-1}^+$ and $\vec{\psi}_{c+1}^-$,

$$\psi_{in,+}\vec{f}_+ = \mathbf{U}_+\vec{\psi}_{c-1}^+ \quad (4.18a)$$

$$\psi_{in,-}\vec{f}_- = \mathbf{U}_-\vec{\psi}_{c+1}^-, \quad (4.18b)$$

where

$$\mathbf{U}_+ = \begin{bmatrix} b_1(-1) \\ \vdots \\ b_{N_P}(-1) \end{bmatrix} [b_1(1) \dots b_{N_P}(1)] \quad (4.19a)$$

$$\mathbf{U}_- = \begin{bmatrix} b_1(1) \\ \vdots \\ b_{N_P}(1) \end{bmatrix} [b_1(-1) \dots b_{N_P}(-1)] . \quad (4.19b)$$

Finally, assuming a symmetric quadrature,

$$\frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} \Delta \vec{\phi}_c = \frac{\Delta x_c}{4} \mathbf{R}_{\Sigma_s} (\vec{\psi}_c^+ + \vec{\psi}_c^-) , \quad (4.20)$$

we may write Eqs. (4.15) as:

$$\left(\mu_+ \mathbf{G}_+ + \frac{\Delta x_c}{2} \mathbf{R}_{\Sigma_t} \right) \vec{\psi}_c^+ - \frac{\Delta x_c}{4} \mathbf{R}_{\Sigma_s} (\vec{\psi}_c^+ + \vec{\psi}_c^-) - \mu_+ \mathbf{U}_+ \vec{\psi}_{c-1}^+ = \frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} (\vec{\phi}_c^{(\ell+1/2)} - \vec{\phi}_c^{(\ell)}) \quad (4.21a)$$

$$\left(\mu_- \mathbf{G}_- + \frac{\Delta x_c}{2} \mathbf{R}_{\Sigma_t} \right) \vec{\psi}_c^- - \frac{\Delta x_c}{4} \mathbf{R}_{\Sigma_s} (\vec{\psi}_c^+ + \vec{\psi}_c^-) - \mu_- \mathbf{U}_- \vec{\psi}_{c+1}^- = \frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} (\vec{\phi}_c^{(\ell+1/2)} - \vec{\phi}_c^{(\ell)}) . \quad (4.21b)$$

Thus, the S2SA scheme uses all of the same matrices, in particular we think of \mathbf{R}_{Σ_t} and \mathbf{R}_{Σ_s} , that we have already defined in our higher fidelity transport model. To find ψ^+ and ψ^- , we must then solve a system of $2 \times N_P \times N_{cell}$ linear equations with $2 \times N_P \times N_{cell}$ unknowns. It is important to note that S2SA can accelerate not only the scalar flux, but also the first angular moment, J of the S_N neutron transport equations

To complete our derivation of the S2SA scheme, we must now define appropriate boundary conditions. We will focus only on the leftmost boundary, though similar equations for the right boundary can be defined analogously. It is sufficient for our purposes to consider problems only with specified incident flux boundary conditions and reflective boundaries. With incident flux conditions, we wish for the accelerated iterate to maintain the same inflow current as the specified boundary condition. Allowing for non-isotropic incident fluxes, the incident current, J^+ specified by our problem is:

$$\sum_{d=N_{dir}/2+1}^{N_{dir}} w_d \mu_d \psi_{in,d}. \quad (4.22)$$

Given the S2SA equations were derived via the assumption of a P_1 angular flux, the additive angular flux correction for direction d is:

$$\Delta\phi = 2\pi (\psi^+ + \psi^-) \quad (4.23a)$$

$$\Delta J = 2\pi (\mu_+ \psi^+ + \mu_- \psi^-) \quad (4.23b)$$

$$\Delta\psi_d = \frac{\Delta\phi}{4\pi} + \mu_d \frac{3\Delta J}{4\pi}. \quad (4.23c)$$

Wishing to maintain J^+ , we have:

$$J^+ = 2\pi \sum_{d=N_{dir}/2+1}^{N_{dir}} w_d \mu_d \left[\psi_{in,d} + \frac{\Delta\phi}{4\pi} + \mu_d \frac{3\Delta J}{4\pi} \right], \quad (4.24)$$

which implies

$$0 = \sum_{d=N_{dir}/2+1}^{N_{dir}} w_d \mu_d \left[\frac{\Delta \phi}{4\pi} + \mu_d \frac{3\Delta J}{4\pi} \mu_d \right]. \quad (4.25)$$

Inserting the definitions of Eqs. (4.23), and allowing for DFEM interpolation points that do not exist at the left boundary:

$$0 = \sum_{d=N_{dir}/2+1}^{N_{dir}} w_d \mu_d \left[\frac{1}{2} \left(\psi_{in}^+ + \vec{L} \vec{\psi}_1^- \right) + \frac{3\mu_d}{2} \left(\mu_+ \psi_{in}^+ + \mu_- \vec{L} \vec{\psi}_1^- \right) \right], \quad (4.26)$$

where

$$\vec{L} = [b_1(-1) \dots b_{N_P}(-1)] . \quad (4.27)$$

Defining constants dependent on the S_N quadrature used,

$$\langle \mu^+ \rangle = \sum_{\mu_d > 0} w_d \mu_d \quad (4.28a)$$

$$\langle \mu^+ \rangle_2 = \sum_{\mu_d > 0} w_d \mu_d^2, \quad (4.28b)$$

Eq. (4.26) becomes

$$0 = \frac{\langle \mu^+ \rangle}{2} \psi_{in}^+ + \frac{\langle \mu^+ \rangle}{2} \vec{L} \vec{\psi}_1^- + \frac{3}{2} \langle \mu^+ \rangle_2 \left(\mu_+ \psi_{in}^+ + \mu_- \vec{L} \vec{\psi}_1^- \right). \quad (4.29)$$

Solving Eq. (4.29) for ψ_{in}^+ ,

$$\psi_{in}^+ = - \left(\frac{\langle \mu^+ \rangle}{2} + \frac{3}{2} \langle \mu^+ \rangle_2 \mu_+ \right)^{-1} \left(\frac{\langle \mu^+ \rangle}{2} + \frac{3}{2} \langle \mu^+ \rangle_2 \mu_- \right) \vec{L} \vec{\psi}_1^-. \quad (4.30)$$

Defining a constant, C_{inc} ,

$$C_{inc} = - \left(\frac{\langle \mu^+ \rangle}{2} + \frac{3}{2} \langle \mu^+ \rangle_2 \mu_+ \right)^{-1} \left(\frac{\langle \mu^+ \rangle}{2} + \frac{3}{2} \langle \mu^+ \rangle_2 \mu_- \right), \quad (4.31)$$

and substituting into Eqs. (4.15), we have

$$\left(\mu_+ \mathbf{G}_+ + \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_t}\right) \vec{\psi}_1^+ = \frac{\Delta x_1}{8\pi} \mathbf{R}_{\Sigma_s} \Delta \vec{\phi}_1 + \frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_c^{(\ell+1/2)} - \vec{\phi}_c^{(\ell)}\right) + \mu_+ C_{inc} \vec{f}_+ \vec{L} \vec{\psi}_1^- \quad (4.32a)$$

$$\left(\mu_- \mathbf{G}_- + \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_t}\right) \vec{\psi}_1^- = \frac{\Delta x_1}{8\pi} \mathbf{R}_{\Sigma_s} \Delta \vec{\phi}_1 + \frac{\Delta x_c}{8\pi} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_1^{(\ell+1/2)} - \vec{\phi}_c^{(\ell)}\right) + \mu_- \psi_{in,-} \vec{f}_- . \quad (4.32b)$$

Noting that $\vec{f}_+ \vec{L}$ creates an $N_P \times N_P$ matrix, and inserting all of our other definitions, we have the equations for cell 1 for incident angular flux boundary conditions

$$\left(\mu_+ \mathbf{G}_+ + \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_t}\right) \vec{\psi}_1^+ - \frac{\Delta x_1}{4} \mathbf{R}_{\Sigma_s} \left(\vec{\psi}_1^+ + \vec{\psi}_1^-\right) - \mu_+ C_{inc} \vec{f}_+ \vec{L} \vec{\psi}_1^- = \frac{\Delta x_1}{8\pi} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_1^{(\ell+1/2)} - \vec{\phi}_1^{(\ell)}\right) \quad (4.33a)$$

$$\left(\mu_- \mathbf{G}_- + \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_t}\right) \vec{\psi}_1^- - \frac{\Delta x_1}{4} \mathbf{R}_{\Sigma_s} \left(\vec{\psi}_1^+ + \vec{\psi}_1^-\right) - \mu_- \mathbf{U}_- \vec{\psi}_2^- = \frac{\Delta x_1}{8\pi} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_1^{(\ell+1/2)} - \vec{\phi}_1^{(\ell)}\right) . \quad (4.33b)$$

For reflective conditions, we have a zero current on the left edge:

$$0 = 2\pi \sum_d w_d \mu_d \psi_d^{(\ell+1/2)} \quad (4.34)$$

$$0 = 2\pi \sum_d w_d \mu_d \left[\psi_d^{(\ell+1/2)} + \frac{\Delta \phi}{4\pi} + \mu_d \frac{3\Delta J}{2} \right] . \quad (4.35)$$

Equation 4.34 implies

$$0 = \sum_d w_d \mu_d \left[\frac{1}{2} \left(\psi_{in,+} + \vec{L} \vec{\psi}_1^- \right) + \frac{3\mu_d}{2} \left(\mu_+ \psi_{in,+} + \mu_- \vec{L} \vec{\psi}_1^- \right) \right] . \quad (4.36)$$

Expanding our earlier quadrature definitions,

$$\langle \mu \rangle = \sum_d w_d \mu_d \quad (4.37a)$$

$$\langle \mu^2 \rangle = \sum_d w_d \mu_d^2 \quad (4.37b)$$

we now solve for $\psi_{in,+}$ as a function of $\vec{\psi}_1^-$:

$$\psi_{in,+} = - \left(\frac{\langle \mu \rangle}{2} + \frac{3\mu_+}{2} \langle \mu^2 \rangle \right)^{-1} \left(\frac{\langle \mu \rangle}{2} + \frac{3\mu_-}{2} \langle \mu^2 \rangle \right) \vec{L} \vec{\psi}_1^-. \quad (4.38)$$

Defining another constant, C_{ref} ,

$$C_{ref} = - \left(\frac{\langle \mu \rangle}{2} + \frac{3\mu_+}{2} \langle \mu^2 \rangle \right)^{-1} \left(\frac{\langle \mu \rangle}{2} + \frac{3\mu_-}{2} \langle \mu^2 \rangle \right) \quad (4.39)$$

, the leftmost cell equations with a reflective boundary condition are:

$$\left(\mu_+ \mathbf{G}_+ + \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_t} \right) \vec{\psi}_1^+ - \frac{\Delta x_1}{4} \mathbf{R}_{\Sigma_s} \left(\vec{\psi}_1^+ + \vec{\psi}_1^- \right) - \mu_+ C_{ref} \vec{f}_+ \vec{L} \vec{\psi}_1^- = \frac{\Delta x_1}{8\pi} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_1^{(\ell+1/2)} - \vec{\phi}_1^{(\ell)} \right) \quad (4.40a)$$

$$\left(\mu_- \mathbf{G}_- + \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_t} \right) \vec{\psi}_1^- - \frac{\Delta x_1}{4} \mathbf{R}_{\Sigma_s} \left(\vec{\psi}_1^+ + \vec{\psi}_1^- \right) - \mu_- \mathbf{U}_- \vec{\psi}_2^- = \frac{\Delta x_1}{8\pi} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_1^{(\ell+1/2)} - \vec{\phi}_1^{(\ell)} \right). \quad (4.40b)$$

4.4 Modified Interior Penalty Diffusion Synthetic Acceleration

We now derive the modified interior penalty diffusion synthetic acceleration (MIP DSA) operator introduced by Ragusa and Wang [37], but also considered by Turcksin and Ragusa [44]. To accelerate the convergence of $N_P \times N_{cell}$ spatial unknowns of the scalar flux, we will need to solve a system of $N_P \times N_{cell}$ linear equations with $N_P \times N_{cell}$ unknowns. Adapted from [37], the MIP DSA update will attempt to solve

the diffusion approximation of Eq. (4.10),

$$-\nabla \cdot D \nabla \Delta \phi + \Sigma_a \Delta \phi = \Sigma_s (\phi^{(\ell+1/2)} - \phi^{(\ell)}) , \quad (4.41)$$

where we use the standard definitions [35],

$$D = \frac{1}{3\Sigma_t} \quad (4.42)$$

$$\Sigma_A = \Sigma_t - \Sigma_s . \quad (4.43)$$

MIP DSA is presented in the standard finite element bilinear form

$$b_{MIP}(\Delta \phi, b_*) = l_{MIP}(b_*) , \quad (4.44)$$

with

$$\begin{aligned} b_{MIP}(\Delta \phi, b_*) &= (\Sigma_a \Delta \phi, b_*)_{\mathcal{D}} + \left(D \vec{\nabla} \Delta \phi, \vec{\nabla} b_* \right)_{\mathcal{D}} \\ &\quad + (\kappa_e \llbracket \Delta \phi \rrbracket, \llbracket b_* \rrbracket)_{E_h^i} \\ &\quad + (\llbracket \Delta \phi \rrbracket, \{\!\{ D \partial_n b_* \}\!\})_{E_h^i} \\ &\quad + (\{\!\{ D \partial_n \Delta \phi \}\!\}, \llbracket b_* \rrbracket)_{E_h^i} \\ &\quad + (\kappa_e \Delta \phi, b_*)_{\partial \mathcal{D}^d} - \frac{1}{2} (\Delta \phi, D \partial_n b_*)_{\partial \mathcal{D}^d} \\ &\quad - \frac{1}{2} (D \partial_n \Delta \phi, b_*)_{\partial \mathcal{D}^d} , \end{aligned} \quad (4.45)$$

and

$$l_{MIP}(b_*) = (\Sigma_s (\phi^{(\ell+1/2)} - \phi^{(\ell)}), b_*) . \quad (4.46)$$

In Eqs. (4.44)-(4.46), b_* is any/every basis function (also referred to as test functions), the $(f, g)_{E_h^i}$ operator acting on quantities f and g is a sum over all cell interior

edges:

$$(f, g)_{E_h^i} = \sum_{c=1}^{N_{cell}-1} (f, g)_{c+1/2}, \quad (4.47)$$

the jump operator, $\llbracket f \rrbracket_{c+1/2}$, is defined as

$$\llbracket f \rrbracket_{c+1/2} = f(x_{c+1/2}^+) - f(x_{c+1/2}^-), \quad (4.48)$$

where $x_{c+1/2}$ is the position of cell edge $c + 1/2$, the average operator, $\{f\}_{c+1/2}$, is

$$\{f\}_{c+1/2} = \frac{1}{2} \left[f(x_{c+1/2}^+) + f(x_{c+1/2}^-) \right], \quad (4.49)$$

and ∂_n is the edge directed normal dotted into the gradient operator. We assume the edge normal always pointing from left to right, thus $\partial_n = \frac{d}{dx}$. Also in Eqs (4.45)-(4.46), the $(f, g)_{\mathcal{D}}$ operator is an integration of quantities f and g over the entire domain \mathcal{D} :

$$(f, g)_{\mathcal{D}} = \sum_{c=1}^{N_{cell}} (f, g)_c \quad (4.50)$$

$$(f, g)_c = \int_{x_{c-1/2}}^{x_{c+1/2}} f g \, dx. \quad (4.51)$$

Finally, κ_e is defined on edge $c + 1/2$ as

$$\kappa_e = \kappa_{c-1/2} = \max \left(\frac{1}{4}, \kappa_{c-1/2}^{IP} \right), \quad (4.52)$$

and $\kappa_{c-1/2}^{IP}$ is defined as:

$$\kappa_{c-1/2}^{IP} = \frac{Z_{MIP}}{2} p_c (p_c + 1) \frac{D_c}{\Delta x_c} \Big|_{x_{c-1/2}^+} + \frac{Z_{MIP}}{2} p_{c-1} (p_{c-1} + 1) \frac{D_{c-1}}{\Delta x_{c-1}} \Big|_{x_{c-1/2}^-}, \quad (4.53)$$

where Z_{MIP} is an arbitrarily chosen constant. Unless otherwise stated, we use $Z_{MIP} = 2$, as recommended by Wang and Ragusa[37].

Focusing now on an interior mesh cell, c , and the N_P b_i that are non-zero in that cell, we now go about defining all of the terms of Eq. (4.45). First, we consider the volumetric integration terms, defining

$$(\Sigma_a \Delta\phi, b_*) = \frac{\Delta x_c}{2} \mathbf{R}_{\Sigma_a} \vec{\Delta\phi}_c, \quad (4.54)$$

and

$$(D\vec{\nabla} \Delta\phi, \vec{\nabla} b_*) = \frac{2}{\Delta x_c} \mathbf{S}. \quad (4.55)$$

In Eq. (4.55), we have defined:

$$\mathbf{S}_{ij} = \int_{-1}^1 \frac{1}{3\Sigma_t(s)} \frac{db_i}{ds} \frac{db_j}{ds} ds. \quad (4.56)$$

Now we treat the edge terms. We begin by defining $[[\Delta\phi]]$ on each edge,

$$[[\Delta\phi]]_{c-1/2} = \vec{L} \vec{\Delta\phi}_c - \vec{R} \vec{\Delta\phi}_{c-1}, \quad (4.57a)$$

$$[[\Delta\phi]]_{c+1/2} = \vec{L} \vec{\Delta\phi}_{c+1} - \vec{R} \vec{\Delta\phi}_c, \quad (4.57b)$$

where we've defined

$$\vec{L} = [b_1(-1) \dots b_{N_P}(-1)] \quad (4.58a)$$

$$\vec{R} = [b_1(1) \dots b_{N_P}(1)]. \quad (4.58b)$$

Treating all test functions for cell c at once, we now define $\llbracket b_* \rrbracket$:

$$\llbracket b_* \rrbracket_{c-1/2} = \vec{L}^T - \vec{0} \quad (4.59a)$$

$$\llbracket b_* \rrbracket_{c+1/2} = \vec{0} - \vec{R}^T, \quad (4.59b)$$

where $\vec{0}$ is a length N_P column vector whose entries are all identically zero. Now we define the average operator on the edges $\{D\partial_n\Delta\phi\}$ and $\{D\partial_nb_*\}$. On edges $c-1/2$ and $c+1/2$,

$$\{D\partial_n\Delta\phi\}_{c-1/2} = \frac{D(x_{c-1/2}^+)}{\Delta x_c} \vec{L}_s \Delta \vec{\phi}_c + \frac{D(x_{c-1/2}^-)}{\Delta x_{c-1}} \vec{R}_s \Delta \vec{\phi}_{c-1} \quad (4.60a)$$

$$\{D\partial_n\Delta\phi\}_{c+1/2} = \frac{D(x_{c+1/2}^+)}{\Delta x_{c+1}} \vec{L}_s \Delta \vec{\phi}_{c+1} + \frac{D(x_{c+1/2}^-)}{\Delta x_c} \vec{R}_s \Delta \vec{\phi}_c, \quad (4.60b)$$

where

$$\vec{L}_s = \left[\left. \frac{db_1}{ds} \right|_{s=-1} \cdots \left. \frac{db_{N_P}}{ds} \right|_{s=-1} \right] \quad (4.61a)$$

$$\vec{R}_s = \left[\left. \frac{db_1}{ds} \right|_{s=1} \cdots \left. \frac{db_{N_P}}{ds} \right|_{s=1} \right]. \quad (4.61b)$$

On edges $c-1/2$ and $c+1/2$,

$$\{D\partial_nb_*\}_{c-1/2} = \frac{D(x_{c-1/2}^+)}{\Delta x_c} \vec{L}_s^T \quad (4.62a)$$

$$\{D\partial_nb_*\}_{c+1/2} = \frac{D(x_{c-1/2}^+)}{\Delta x_c} \vec{R}_s^T. \quad (4.62b)$$

Combining Eq. (4.54), Eq. (4.55), Eqs. (4.57), Eqs. (4.59), Eqs. (4.60), and Eqs.

(4.62), we have the left hand side of the cell c MIP DSA equations:

$$\begin{aligned}
& \frac{\Delta x_c}{2} \mathbf{R}_{\Sigma_a} \vec{\Delta} \phi_c + \frac{2}{\Delta x_c} \mathbf{S} \vec{\Delta} \phi_c \\
& + \left\{ \left[\kappa_{c-1/2} \vec{L}^T \left(\vec{L} \vec{\Delta} \phi_c - \vec{R} \vec{\Delta} \phi_{c-1} \right) \right] - \left[\kappa_{c+1/2} \vec{R}^T \left(\vec{L} \vec{\Delta} \phi_{c+1} - \vec{R} \vec{\Delta} \phi_c \right) \right] \right\} \\
& + \left\{ \frac{D(x_{c-1/2}^+)}{\Delta x_c} \vec{L}_s^T \left(\vec{L} \vec{\Delta} \phi_c - \vec{R} \vec{\Delta} \phi_{c-1} \right) + \frac{D(x_{c-1/2}^-)}{\Delta x_c} \vec{R}_s^T \left(\vec{L} \vec{\Delta} \phi_{c+1} - \vec{R} \vec{\Delta} \phi_c \right) \right\} \\
& + \left\{ \vec{L}^T \left(\frac{D(x_{c-1/2}^+)}{\Delta x_c} \vec{L}_s \vec{\Delta} \phi_c + \frac{D(x_{c-1/2}^-)}{\Delta x_{c-1}} \vec{R}_s \vec{\Delta} \phi_{c-1} \right) \dots \right. \\
& \quad \left. - \vec{R}^T \left(\frac{D(x_{c+1/2}^+)}{\Delta x_{c+1}} \vec{L}_s \vec{\Delta} \phi_{c+1} + \frac{D(x_{c+1/2}^-)}{\Delta x_c} \vec{R}_s \vec{\Delta} \phi_c \right) \right\} . \quad (4.63)
\end{aligned}$$

The right hand side on the interior is obviously

$$\frac{\Delta x_c}{2} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_c^{(\ell+1/2)} - \vec{\phi}_c^{(\ell)} \right) . \quad (4.64)$$

We now consider the leftmost cell. We will derive equations for reflective and incident current boundary conditions. Analogous equations can be derived for the rightmost cell. Starting with incident flux boundary conditions, we must define the $(\kappa_{1/2} \Delta \phi, b_*)$, $(\Delta \phi, D \partial_n b_*)$, and $(D \partial_n \Delta \phi, b_*)$ operators. They are:

$$(\kappa_{1/2} \Delta \phi, b_*) = \kappa_{1/2} \vec{L}^T \vec{L} \vec{\Delta} \phi_1 \quad (4.65a)$$

$$(\Delta \phi, D \partial_n b_*) = \frac{2D(x_{1/2}^+)}{\Delta x_1} \vec{L}_s^T \vec{L} \vec{\Delta} \phi_1 \quad (4.65b)$$

$$(D \partial_n \Delta \phi, b_*) = \frac{2D(x_{1/2}^-)}{\Delta x_1} \vec{L}^T \vec{L}_s \vec{\Delta} \phi_1 . \quad (4.65c)$$

Combing with the cell integral quantities, for problems with an incident flux bound-

ary condition, the leftmost cell equation is

$$\begin{aligned} \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_a} \vec{\Delta \phi}_1 + \frac{2}{\Delta x_1} \mathbf{S} \vec{\Delta \phi}_1 \\ \kappa_{1/2} \vec{L}^T \vec{L} \vec{\Delta \phi}_1 - \frac{D(x_{1/2}^+)}{\Delta x_1} \vec{L}_s^T \vec{L} \vec{\Delta \phi}_1 - \frac{D(x_{1/2})}{\Delta x_1} \vec{L}^T \vec{L}_s \vec{\Delta \phi}_1 \\ = \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_1^{(\ell+1/2)} - \vec{\phi}_1^{(\ell)} \right) . \end{aligned} \quad (4.66)$$

If the left edge satisfies a reflective condition, the leftmost cell equations are:

$$\frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_a} \vec{\Delta \phi}_1 + \frac{2}{\Delta x_1} \mathbf{S} \vec{\Delta \phi}_1 = \frac{\Delta x_1}{2} \mathbf{R}_{\Sigma_s} \left(\vec{\phi}_1^{(\ell+1/2)} - \vec{\phi}_1^{(\ell)} \right) + C_{MIP} \vec{L}^T , \quad (4.67)$$

where C_{MIP} is the incident partial current,

$$C_{MIP} = 2\pi \sum_{\mu_d > 0} w_d \mu_d \psi_d^{(\ell+1/2)} . \quad (4.68)$$

4.5 Numerical Results Verifying Implementation and Performance of S2SA and MIP DSA

We will consider two test problems to compare the effectiveness of S2SA and MIP DSA compared against source iteration, abbreviated as SI in the following results. To measure effectiveness, we will numerically compare the spectral radius, ρ of each scheme. The spectral radius of an iterative technique is the largest magnitude eigenvalue of the iterative operator. For an iteration scheme to be convergent, it must be true $d < 1$. After sufficiently many iterations, the following is true:

$$\|\delta \phi^{(\ell+1)}\| \leq \rho \|\delta \phi^{(ell)}\| , \quad (4.69)$$

where $\|\cdot\|$ is a valid norm and $\delta\phi^{(\ell)}$ is the error as defined in Eq. (4.7). We will numerically test our methods by considering a trivial solution problem. That is to say we will solve a problem with vacuum boundary conditions on all edges and no volumetric solutions. However, rather than initialize with a zero solution, we set the values of $\phi^{(0)}$ to be random numbers $\in [0, 1]$. We then use 150 iterations to estimate ρ , taking the last value to be the converged estimate of ρ . In Section 4.5.1 we present results for a scattering medium with a spatially constant cross section and in Section 4.5.2, we consider a scattering medium with spatially varying interaction cross section.

In the plots of Section 4.5.1 and Section 4.5.2 we will examine ρ as a function of scattering ratio, c , S_N order, DFEM trial space degree, iterative technique, DFEM interpolation point type, and cell optical thickness, $\Sigma_t\Delta x$.

4.5.1 Spatially Constant Cross Section Scattering Problem

Our first test problem with is a 20 [cm] slab with spatially homogeneous cross section. We use at least 10 cells in all simulations in Figs. 4.1-4.9. For $\Sigma_t\Delta x \geq 2$, we increase Σ_t to the necessary value to obtain the desired $\Sigma_t\Delta x$, while holding $\Delta x = 2[\text{cm}]$ constant. For $\Sigma_t\Delta x < 2$, we hold $\Sigma_t = 1[\text{cm}^{-1}]$ constant, while increasing the number of cells to achieve the desired $\Sigma_t\Delta x$.

The purpose of the following section is to

1. verify that results from our S2SA implementation for linear SL Gauss are qualitatively similar to those presented in [36],
2. determine whether S2SA is an effective iterative acceleration technique, for SL Gauss and SL Lobatto DFEM with $P \in [1, 4]$,
3. verify that results from our MIP DSA implementation used in conjunction with

SL Gauss, gives results similar to those presented in [37] for $P \in [1, 4]$, and

4. determine whether MIP DSA is an effective iterative acceleration technique for SL Lobatto DFEM.

In Fig. 4.1, we compare SI, S2SA, and MIP DSA for linear SL Gauss, with S_8 angular quadrature, and $c = 0.999$. We see that as expected[38], ρ for source iteration $\approx c$. Additionally, we see that MIP DSA achieves a ρ similar to that given in Figs. 6 and 7 of [37]. Likewise, our S2SA implementation estimates $\rho \leq \approx 0.2$, which agrees well with the Fourier analysis of [36] a true spectral radius of $0.2127c$. Since the analysis of [36] is for an infinite medium, we expect our estimate of ρ to be less.

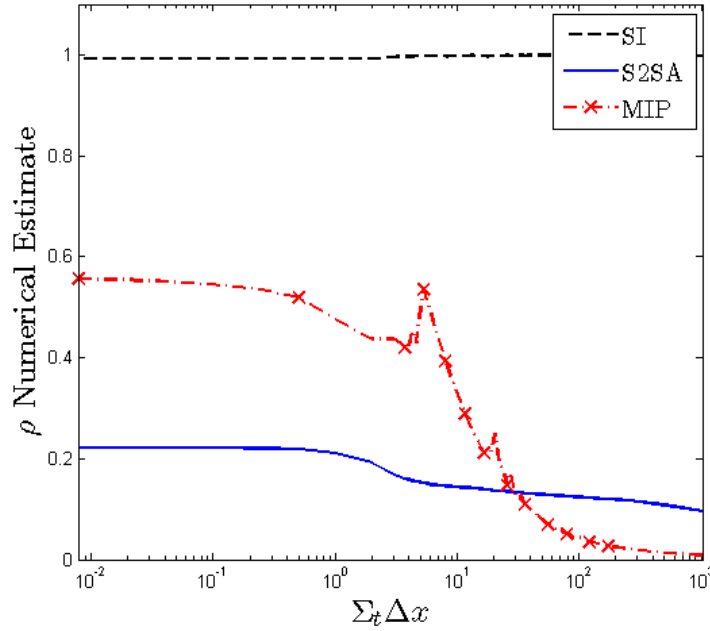


Figure 4.1: Estimates of ρ for different iterative techniques for S_8 , $c = 0.999$, linear SL Gauss.

In Fig. 4.2, we compare the ρ estimates of SI, S2SA, and MIP DSA for linear SL Lobatto differencing. Results indicate that both S2SA and MIP DSA are compatible with SL Lobatto neutron transport. MIP DSA for linear SL Lobatto exhibits the same peaking as observed with linear SL Gauss, though the peak is slightly smoother. The abrupt dips in the ρ estimate of MIP DSA for $\Sigma_t \Delta x \in [0, 10]$ are caused by our numerical estimation oscillating between two different values of ρ . Consider Fig. 4.3, which again plots ρ for linear SL Lobatto iterative schemes, but plots the average value of ρ obtained while iterating. The dips of Fig. 4.2 are gone in Fig. 4.3. If we could have iterated further, Fig. 4.2 would mirror Fig. 4.3. However, numerical precision prevented further iteration. After many iterations, point-wise values of $|\phi| \rightarrow \ll 10^{-154}$, since we are estimating ρ after iteration ℓ as:

$$\rho^{(\ell)} \approx \frac{\|\phi^{(\ell+1)}\|_{L^2}}{\|\phi^{(\ell)}\|_{L^2}}, \quad (4.70)$$

we have

$$\phi_{cell,i}^2 < 10^{-308}, \quad (4.71)$$

where 10^{-308} is approximately the smallest magnitude double precision floating point number [45]. Thus, to floating point precision, $\|\phi^{(\ell+1)}\| = 0$, and we can no longer estimate ρ . We now examine the sensitivity of S2SA and MIP DSA to different values of c , S_N , and DFEM trial space degree. From Fig. 4.1 and Fig. 4.2, we expect that the choice of DFEM interpolation point will have nearly negligible effect on ρ , but continue to compare SL Lobatto and SL Gauss solutions to one another for completeness.

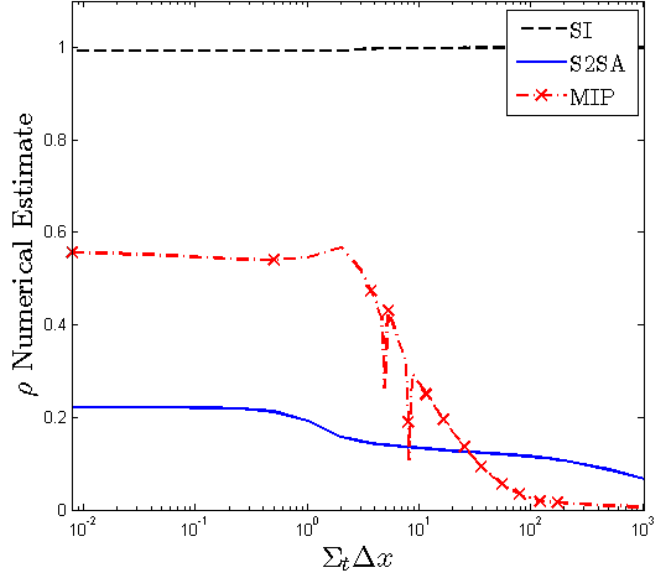


Figure 4.2: Estimates of ρ for different iterative techniques for S_8 , $c = 0.999$, linear SL Lobatto.

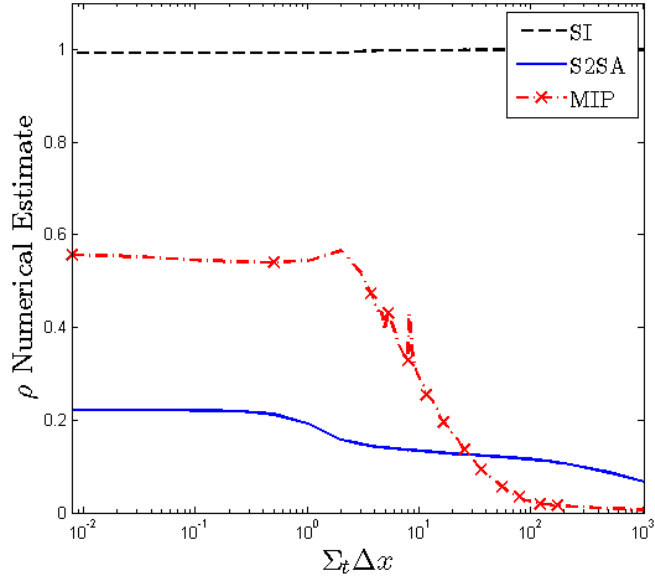


Figure 4.3: Average value of ρ during estimation, for different iterative techniques for S_8 , $c = 0.999$, with linear SL Lobatto spatial discretization.

4.5.1.1 S2SA Spectral Radius Sensitivity

In Fig. 4.4 and Fig. 4.5, we compute ρ for S2SA as a function of S_N order for linear SL Gauss and linear SL Lobatto, respectively. In both Figs. 4.4-4.5, the higher the S_N order, the larger ρ , across all cell optical thicknesses, though there is little difference between S_8 and S_{16} . Comparing our estimates of ρ of S2SA, as a function

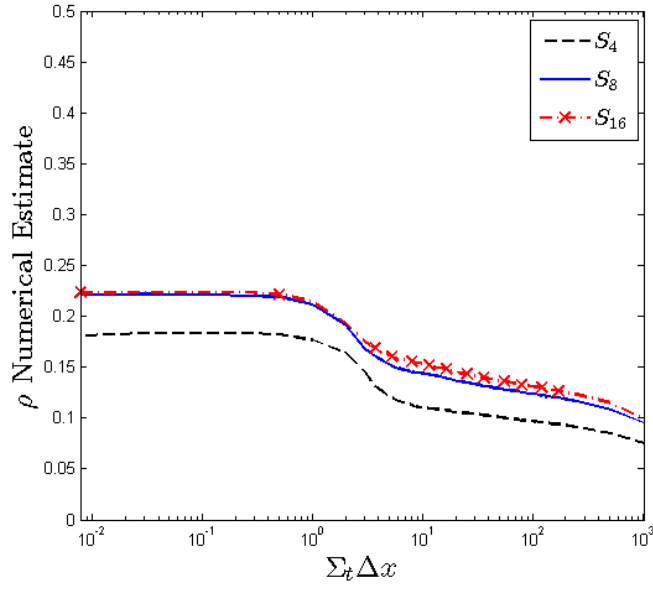


Figure 4.4: Estimate of ρ for S2SA as a function of S_N order for $c = 0.999$ for linear SL Gauss.

of c , for linear SL Gauss and SL Lobatto in Fig. 4.6 and Fig. 4.7, respectively, we conclude that the closer c is to unity, the larger, ρ , but there is negligible increase from $c = 0.999$ to $c = 0.9999$, except in very optically thick cells, but ρ is not largest with the S2SA scheme for large values of $\Sigma_t \Delta x$. Finally, in Fig. 4.8 and Fig. 4.9, we compare ρ for S_8 , $c = 0.999$ S2SA with SL Gauss and SL Lobatto differencing, respectively, as a function of DFEM trial space degree. ρ remains nearer the optically

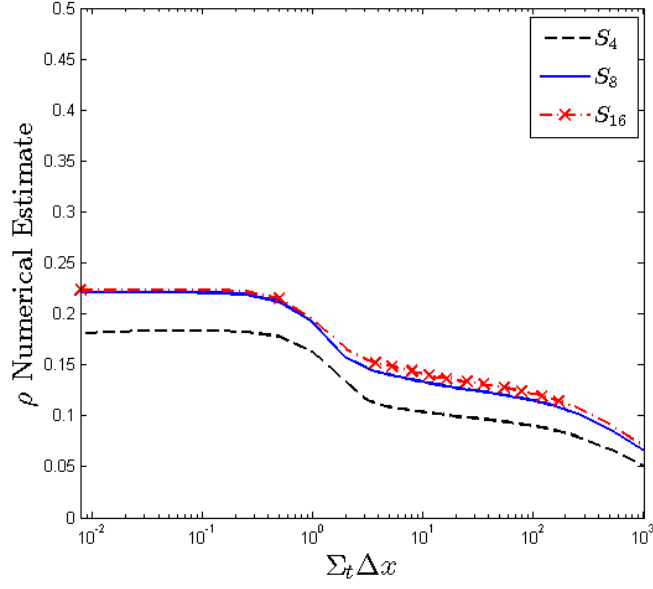


Figure 4.5: Estimate of ρ for S2SA as a function of S_N order for $c = 0.999$ for linear SL Lobatto.

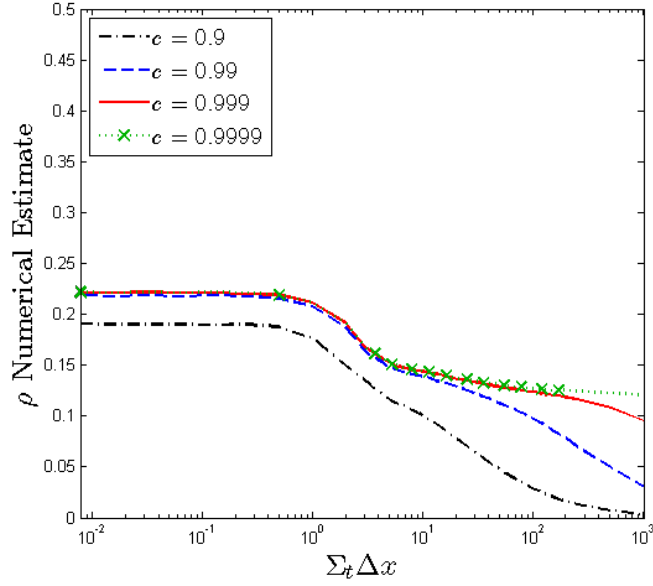


Figure 4.6: Estimates of ρ for S2SA as a function of c for S_8 linear SL Gauss.

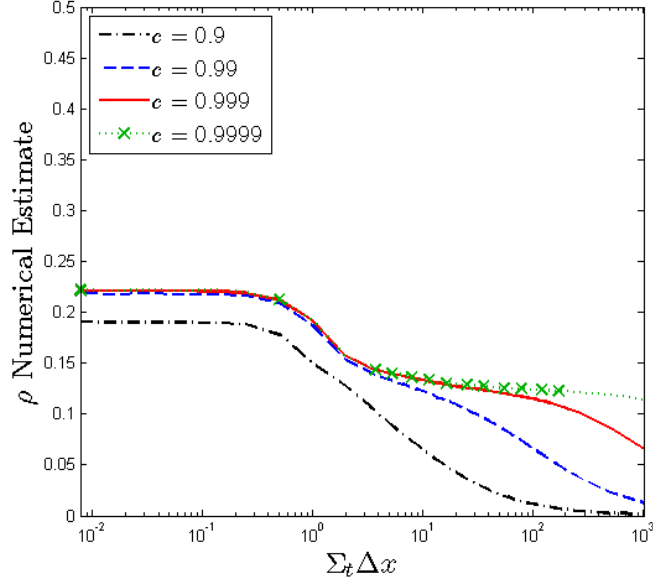


Figure 4.7: Estimates of ρ for S2SA as a function of c for S_8 linear SL Lobatto.

thin value for larger values of $\Sigma_t \Delta x$ the higher the DFEM trial space degree, but S2SA remains stable for all P . Regardless of c , P , $\Sigma_t \Delta x$, and S_N order, we numerically verified that the S2SA matrix that must be inverted was neither symmetric or positive definite, requiring direct (Gaussian) elimination to invert the matrix.

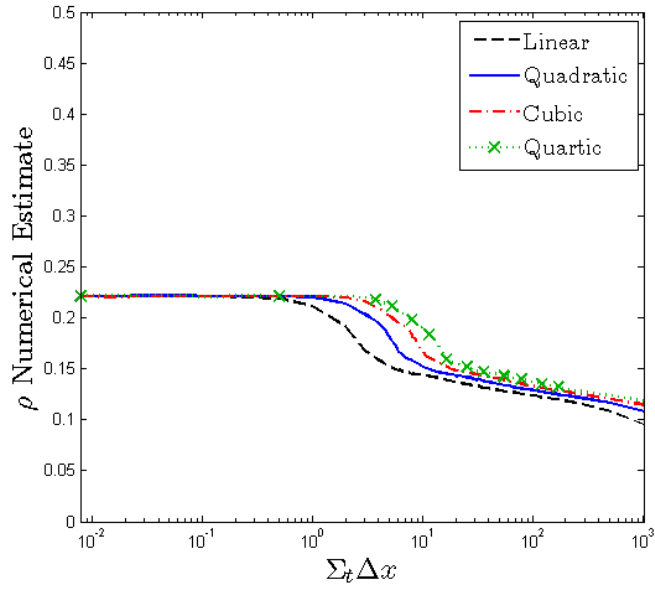


Figure 4.8: Estimates of ρ for S2SA as a function of S_8 , $c = 0.999$, SL Gauss as a function of trial space degree.

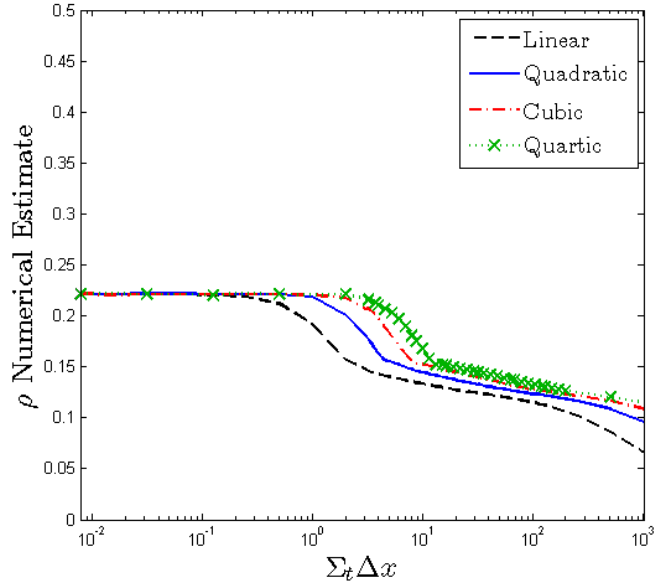


Figure 4.9: Estimates of ρ for S2SA as a function of S_8 , $c = 0.999$, SL Lobatto as a function of trial space degree.

4.5.1.2 MIP DSA Spectral Radius Sensitivity

In Fig. 4.10 and Fig. 4.11 we compare ρ for MIP DSA as a function of S_N order, for linear SL Gauss and linear SL Lobatto, respectively, with $c = 0.999$. As with

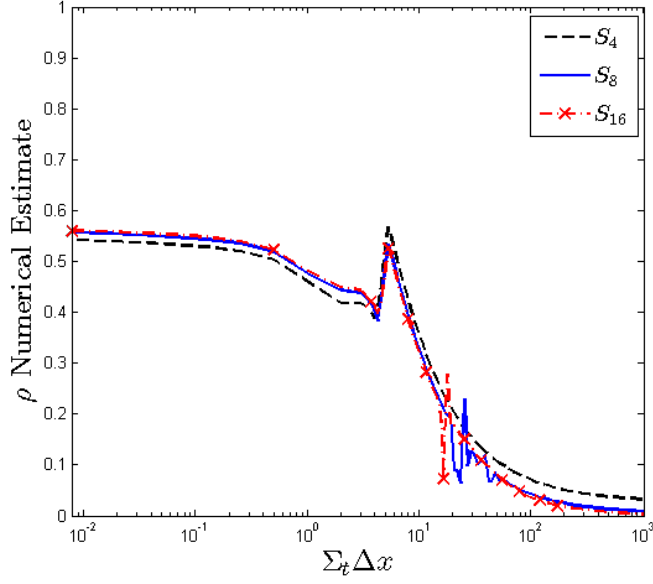


Figure 4.10: Estimate of ρ for MIP DSA as a function of S_N order for $c = 0.999$ linear SL Gauss.

S2SA, the higher S_N order, the larger ρ is for the MIP DSA scheme. However, the value of ρ obtained with S_8 is only slightly smaller than the ρ of S_{16} , and as such we prefer the reduced computational work of S_8 simulations to S_{16} in examining iterative effectiveness. Figures 4.10-4.11 also exhibit ρ dips associated with insufficient iteration to converge to the correct ρ .

Figure 4.12 and Fig. 4.13 examine the effect of c on ρ for MIP DSA with the linear SL Gauss and SL Lobatto schemes. The general trend is that as $c \rightarrow 1$, the

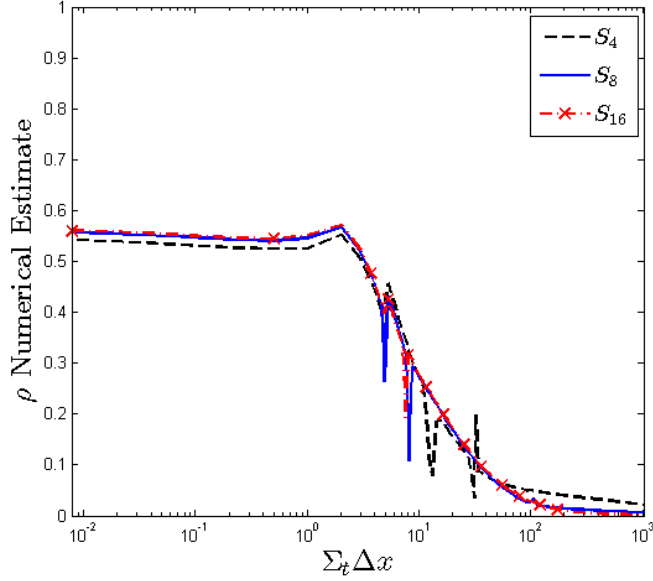


Figure 4.11: Estimate of ρ for MIP DSA as a function of S_N order for $c = 0.999$ linear SL Lobatto.

larger the estimate of ρ . However, we are most concerned about the large spikes in ρ for $c = 0.9999$ for $\Sigma_t \Delta x \in [10, 100]$. Allowing for the possibility that the ending ρ estimate was not converged, we plot the average ρ found while iterating for SL Gauss in Fig. 4.14 and for SL Lobatto in Fig. 4.15. While the averaging of ρ does eliminate the most worrisome spike in ρ , the spike SL Lobatto exhibits in Fig. 4.13 that approaches unity, not all spikes are eliminated. It is at this point that we remember that the MIP DSA scheme uses a constant, κ_{IP} , which depends on an arbitrarily chosen constant. In all of the results that have presented to this point, we have assumed that made an assumption regarding Z_{MIP} of Eq. (4.53), $Z_{MIP} = 2$. Just as Wang did in his dissertation[46], we now consider $Z_{MIP} = 4$.

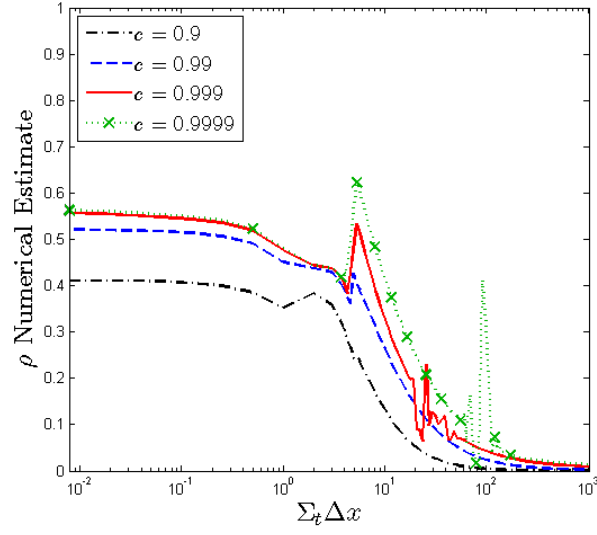


Figure 4.12: Estimate of ρ for MIP DSA as a function of c for S_8 linear SL Gauss.

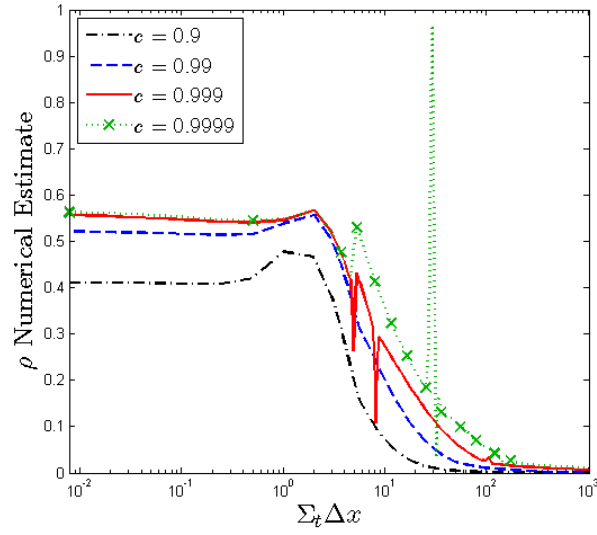


Figure 4.13: Estimate of ρ for MIP DSA as a function of c for S_8 linear SL Lobatto.

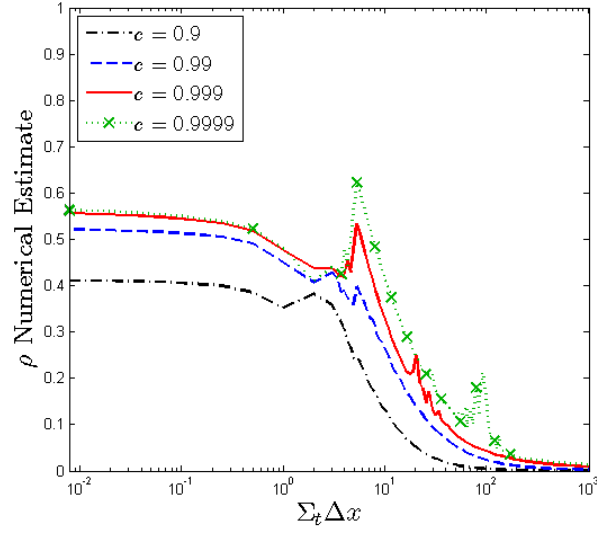


Figure 4.14: Average estimate of ρ for MIP DSA as a function of c for S_8 linear SL Gauss during iteration process.

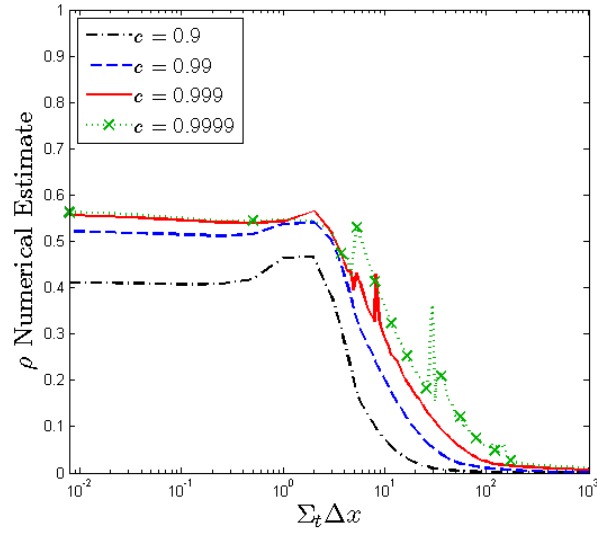


Figure 4.15: Average estimate of ρ for MIP DSA as a function of c for S_8 linear SL Lobatto during iteration process.

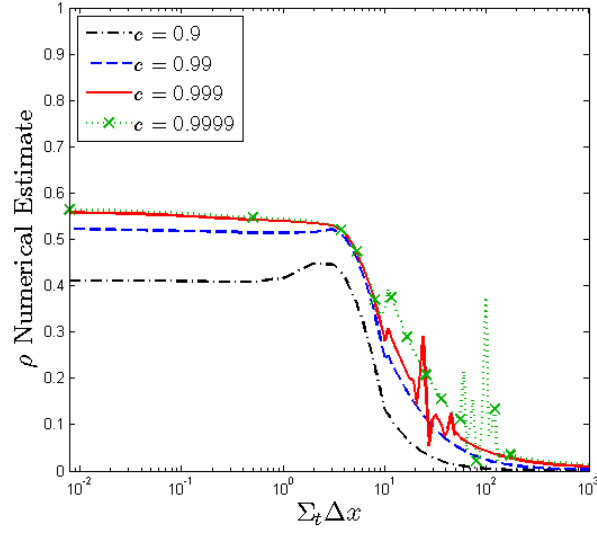


Figure 4.16: ρ for MIP DSA as a function of c for S_8 linear SL Gauss, $Z_{MIP} = 4$.

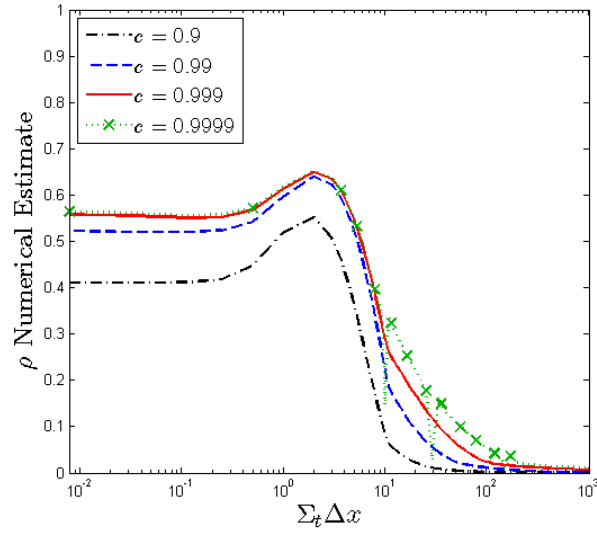


Figure 4.17: ρ for MIP DSA as a function of c for S_8 linear SL Lobatto, $Z_{MIP} = 4$.

Estimates of ρ for MIP DSA as a function of c for SL Gauss and SL Lobatto, with $Z_{MIP} = 4$ are given in Fig. 4.16 and Fig. 4.17, respectively. The differences between Figs. 4.12-4.13 and Figs. 4.16-4.17 is relatively small. With $Z_{MIP} = 4$, SL Gauss maintains the optically thin limit of ρ for larger values of $\Sigma_t \Delta x$, but does not display the large spike before $\Sigma_t \Delta x = 1$. With $Z_{MIP} = 4$, SL Lobatto displays a modest, smooth increase in ρ over a large range of $\Sigma_t \Delta x$, but eliminates the spikes in ρ present in the optically thick regions of $\Sigma_t \Delta x$ for $Z_{MIP} = 2$ in Fig. 4.13 and Fig. 4.15. To determine whether the small spikes in ρ for $\Sigma_t \Delta x \geq 100$ in Fig. 4.16 are a spurious result of our iterative process, we plot the average ρ achieved during the iteration process for SL Gauss with $Z_{MIP} = 4$ in Fig. 4.18. The averaging of

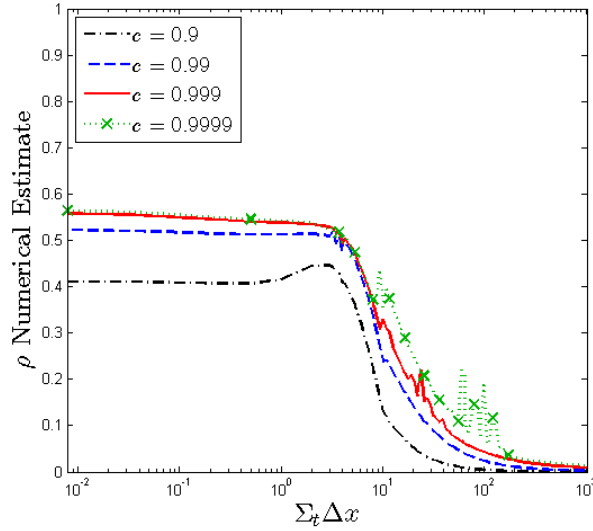


Figure 4.18: Average estimate of ρ during iteration for MIP DSA as a function of c for S_8 linear SL Gauss, with $Z_{MIP} = 4$.

Fig. 4.18 mitigates some of the spikes seen in Fig. 4.16, but does not eliminate them.

Finally, having determined that $c = 0.999$ and S_8 will yield nearly maximal ρ , we

calculate ρ as a function of P for MIP DSA acceleration of SL Gauss and SL Lobatto transport spatial discretizations in Fig. 4.19 and Fig. 4.20, respectively, using $Z_{MIP} = 2$. Unfortunately, it appears that MIP DSA is not guaranteed to be convergent for

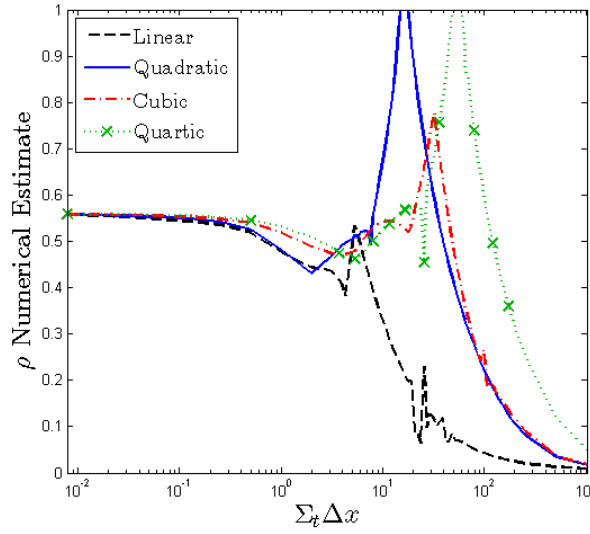


Figure 4.19: Estimate of ρ for MIP DSA as a function of P for S_8 , $c = 0.999$, SL Gauss with $Z_{MIP} = 2$.

all P for SL Lobatto and SL Gauss. Hoping that this is a result of the end iterate for ρ is not yet converged, we plot the average ρ observed during the iteration process in Fig. 4.21 and Fig. 4.22 for SL Gauss and SL Lobatto, respectively. Figures 4.21-4.22 unfortunately confirm that convergence is not guaranteed for $Z_{MIP} = 2$. Seeking a solution that guarantees $\rho < 1$ for all values of $\Sigma_t \Delta x$ and P , we now examine the effect of choosing $Z_{MIP} = 4$ in Figs. 4.23-4.24. In Figs. 4.23-4.24, both SL Gauss and SL Lobatto remain convergent for all values of $\Sigma_t \Delta x$ for $P \in [1, 4]$, when $Z_{MIP} = 4$. As such, moving forward we will assume $Z_{MIP} = 4$.

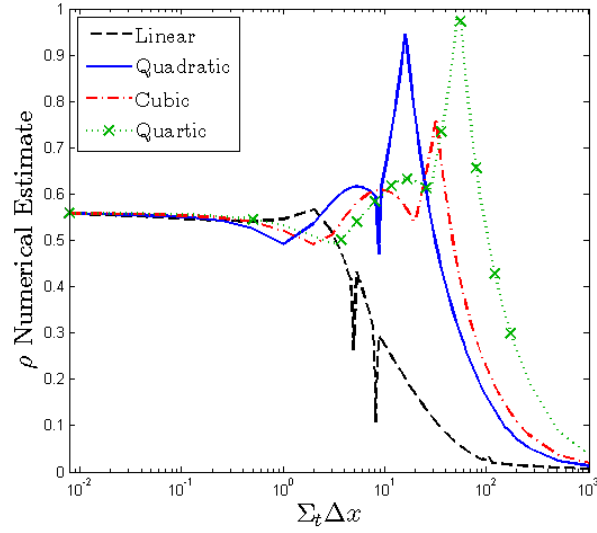


Figure 4.20: Estimate of ρ for MIP DSA as a function of P for S_8 , $c = 0.999$, SL Lobatto with $Z_{MIP} = 2$.

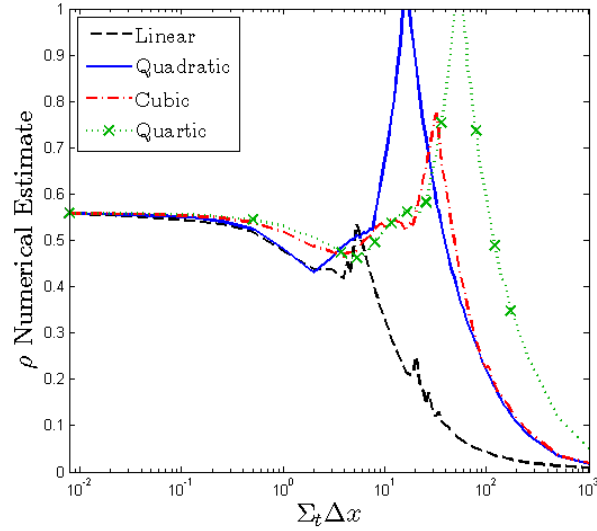


Figure 4.21: Average estimate of ρ for MIP DSA during iteration, as a function of P for S_8 , $c = 0.999$, SL Gauss with $Z_{MIP} = 2$.

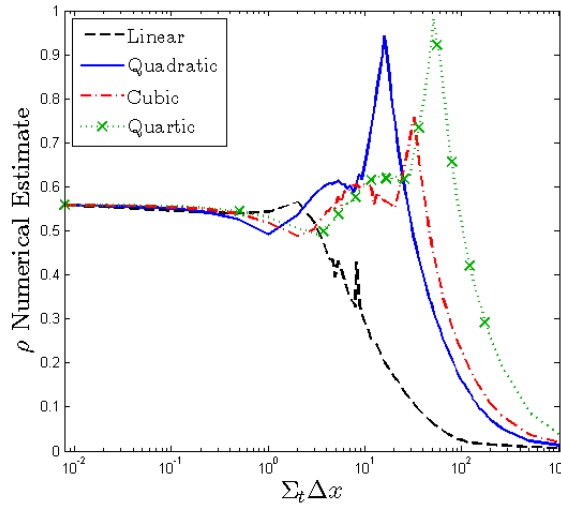


Figure 4.22: Average estimate of ρ for MIP DSA during iteration, as a function of P with S_8 , $c = 0.999$, SL Lobatto with $Z_{MIP} = 2$.

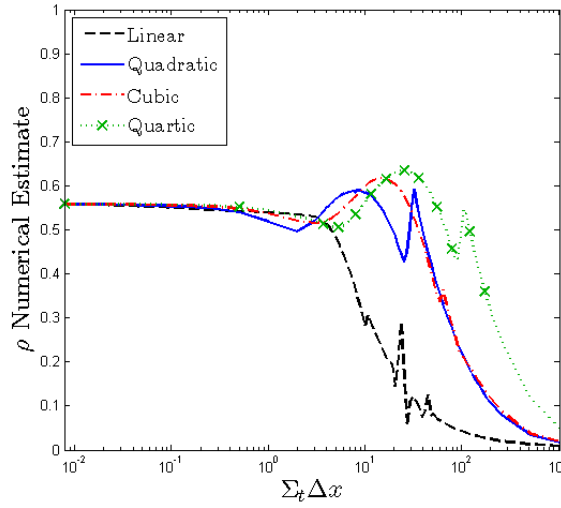


Figure 4.23: ρ for MIP DSA as a function of P for S_8 , $c = 0.999$, SL Gauss with $Z_{MIP} = 4$.

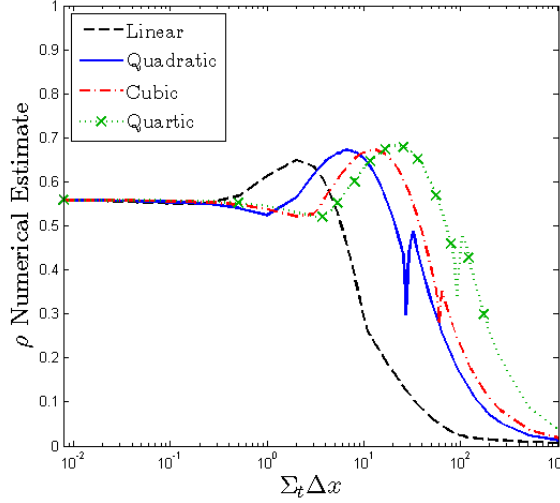


Figure 4.24: Estimates of ρ for MIP DSA as a function of P for S_8 , $c = 0.999$, SL Lobatto with $Z_{MIP} = 4$.

4.5.2 Spatially Varying Cross Section Scattering Problem

To test the effectiveness of MIP DSA and S2SA for a problem with a spatially varying cross section, we again consider a slab $x \in [0, 20[cm]]$. We impose

$$\Sigma_t(x) = \Sigma_{t,0} \exp \left[\frac{|(10 - x)|}{2} \right]. \quad (4.72)$$

We hold c constant in space. We will estimate ρ as a function of $\overline{\Sigma_t \Delta x}$, the average optical thickness of each mesh cell. For values of $\overline{\Sigma_t \Delta x} > 2$, we will use 10 mesh cells, and adjust $\Sigma_{t,0}$ to achieve the desired optical thickness. For values of $\overline{\Sigma_t \Delta x} < 2$, we will hold $\Sigma_{t,0}$ constant, and increase the number of mesh cells. We wish to maintain a total slab optical thickness of at least 20 mean free paths. Since,

$$\text{Total Mean Free Path} = 2 \int_0^{10 \text{ cm}} \Sigma_{t,0} \exp \left[\frac{10 - x}{2} \right] dx \quad (4.73a)$$

$$\text{Total Mean Free Path} = 4 \Sigma_{t,0} (\exp[5] - 1), \quad (4.73b)$$

the minimum value of $\Sigma_{t,0}$ is:

$$\Sigma_{t,0} = \frac{5}{\exp[5] - 1} . \quad (4.73c)$$

For values of $\overline{\Sigma_t \Delta x} > 2$,

$$10\overline{\Sigma_t \Delta x} = 4\Sigma_{t,0} (\exp[5] - 1) \quad (4.73d)$$

$$\Sigma_{t,0} = \frac{5\overline{\Sigma_t \Delta x}}{2 (\exp[5] - 1)} . \quad (4.73e)$$

4.5.2.1 S2SA

In Fig. 4.25 and Fig. 4.26, we plot the estimate of ρ for the variable cross section problem, with $c = 0.999$ and S_8 angular quadrature for S2SA using SL Gauss and SL Lobatto schemes, respectively, as a function of trial space degree. S2SA remains

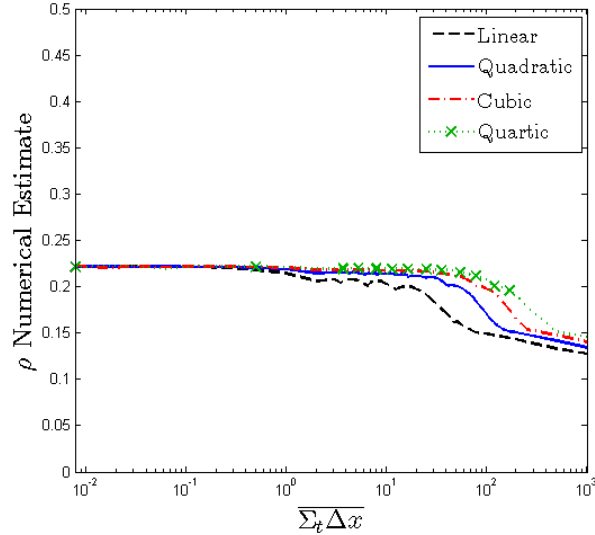


Figure 4.25: Estimate of ρ for S2SA as a function of $\overline{\Sigma_t \Delta x}$ and P , with SLXS Gauss.

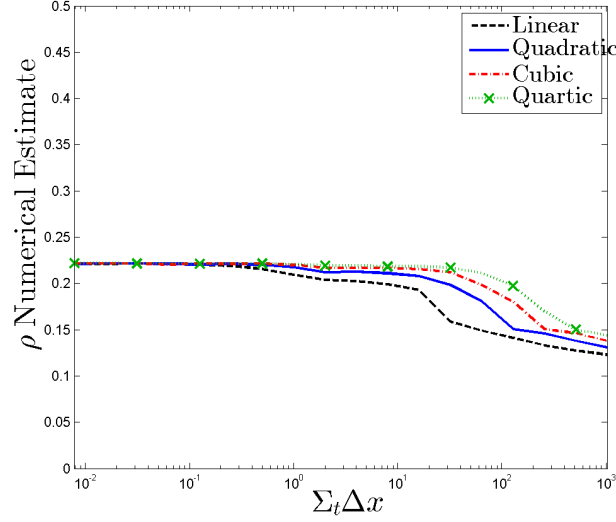


Figure 4.26: Estimate of ρ for S2SA as a function of $\overline{\Sigma_t \Delta x}$ and P , with SLXS Lobatto.

unconditionally stable for the case of a spatially varying cross section, for all P considered, for both SLXS Gauss and SLXS Lobatto schemes.

4.5.2.2 MIP DSA

We now examine ρ for the MIP DSA scheme as a function of P for the variable cross section problem, with $c = 0.999$, S_8 angular quadrature, and $Z_{MIP} = 4$. Results for SLXS Gauss are given in Fig. 4.27 and ρ for MIP DSA with SLXS Lobatto scheme is given in Fig. 4.28. To verify that the oscillations present in Figs. 4.27-4.28 are not a result of the numerical estimates to ρ not converging, we plot the iteration average ρ in Fig. 4.29 and Fig. 4.30 for SLXS Gauss and SLXS Lobatto, respectively.

Figures 4.29-4.30 verify that the observed variations in Fig. 4.27 and Fig. 4.28 are not numerical noise. Further, since $\rho < 1$ in Figs. 4.27-4.28, we conclude that MIP DSA is a stable iterative scheme for SLXS Gauss and SLXS Lobatto, for all P considered.

The MIP DSA matrix was verified to remain SPD under the assumption of a

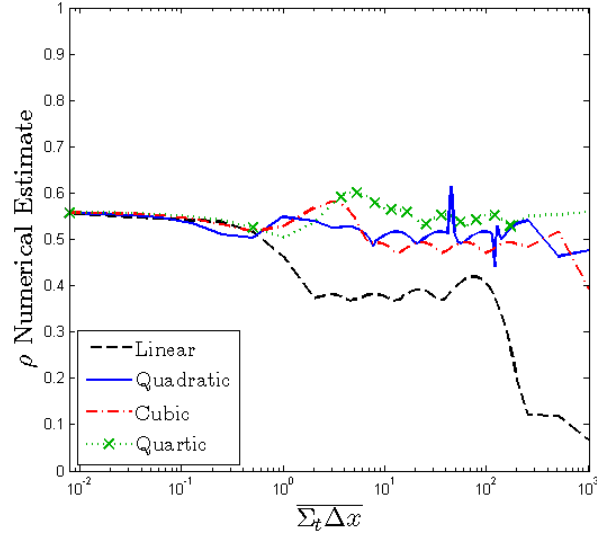


Figure 4.27: Estimate of ρ for MIP DSA with SLXS Gauss as a function of $\overline{\Sigma_t \Delta x}$ and P .

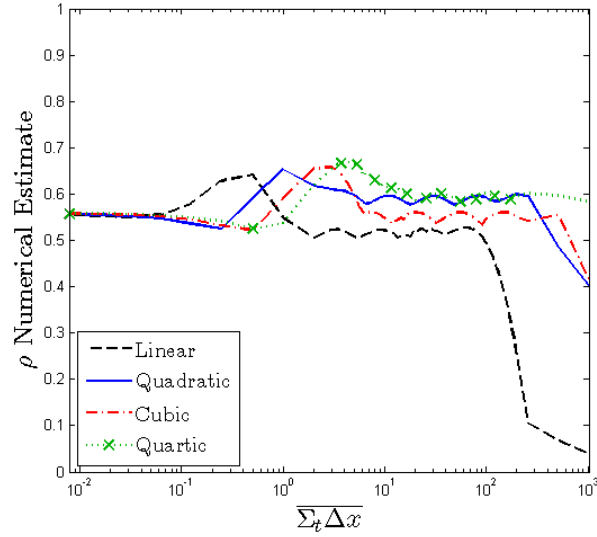


Figure 4.28: Estimate of ρ for MIP DSA with SLXS Lobatto as a function of $\overline{\Sigma_t \Delta x}$ and P .

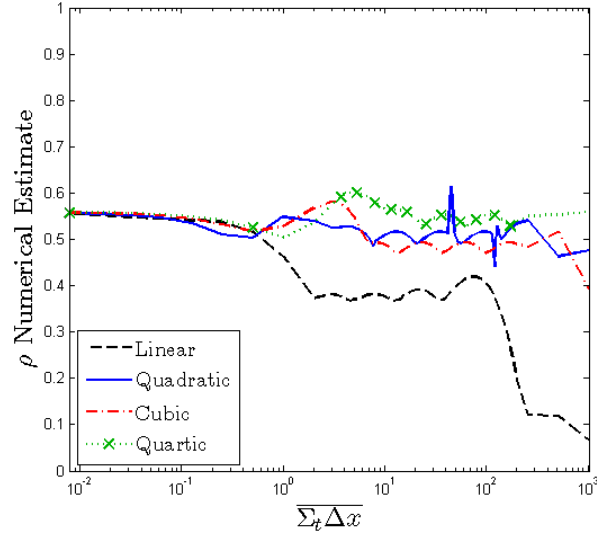


Figure 4.29: Estimate of ρ for MIP DSA with SLXS Gauss as a function of $\overline{\Sigma_t \Delta x}$ and P .

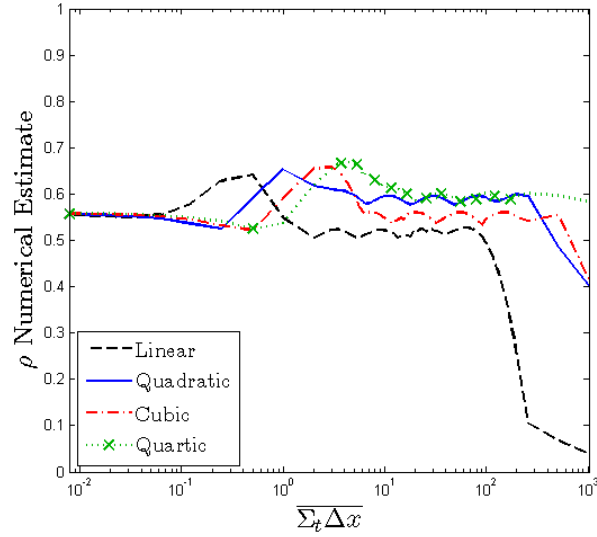


Figure 4.30: Estimate of ρ for MIP DSA with SLXS Lobatto as a function of $\overline{\Sigma_t \Delta x}$ and P .

spatially varying cross section.

4.6 Conclusions

Both S2SA and MIP DSA are stable iterative acceleration techniques for problems with spatially constant and spatially varying cross section. We have verified our implementations of S2SA and MIP DSA by comparing to published, spatially constant cross section results. While the S2SA technique is more effective than MIP DSA (smaller ρ), the S2SA matrix that must be inverted is not symmetric or positive definite. Thus, S2SA requires direct matrix inversion which in general is not feasible for large systems of equations. Conversely, though MIP DSA is not as effective as S2SA, the MIP DSA matrix is SPD, thus it can be inverted much more efficiently [faster] [44] than the S2SA matrix can be inverted. As such, to demonstrate that self-lumping schemes that account for within cell spatial variation of quadrature can be used for more realistic problems, we first employ S2SA to accelerate neutron transport within a model fuel depletion problem in Chapter 5. However, since S2SA is really only practical for small, one dimensional problems, we will apply a variant of MIP DSA to accelerate the iterative process required for thermal radiative transfer problems in Chapter 6.

5. FUEL DEPLETION PROBLEM

We now consider a fuel depletion problem to illustrate that self-lumping DFEM schemes remain accurate for more complex physics than simply a purely absorbing medium. The depletion method we use [time stepping scheme, time step size, etc.] is chosen for its simplicity, not for its fidelity relative to state-of-the art depletion methodologies. Our goal is to assess the accuracy of spatial discretization methods for problems with spatially varying cross sections, not to propose a new depletion method. However, we stress that self-lumping methods can be implemented with any time depletion method or time stepping scheme since implementation of self-lumping only requires changes pertaining to the spatial discretization.

5.1 Problem Physical Description

We will consider a 1-D slab fuel depletion problem. The dimensions and layout of the fuel and moderator layers lattice is shown in Fig. 5.1.

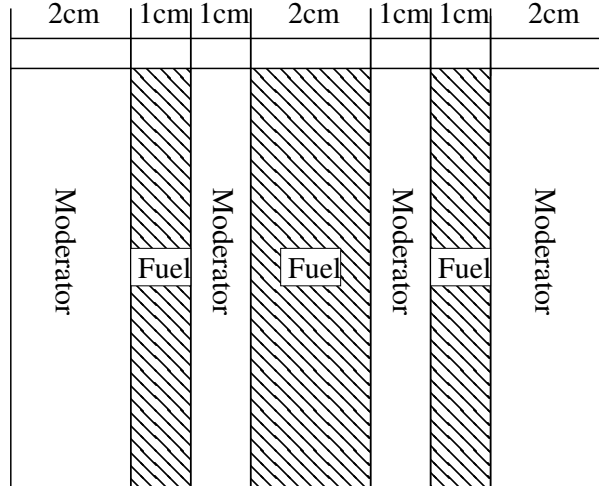


Figure 5.1: Depletion problem fuel/moderator lattice.

Initially, each fuel region is a homogeneous mixture containing only fissile ^{235}U and fertile ^{238}U nuclei, with compositions given in Table 5.1. As fuel depletion

Table 5.1: Fuel atom density data.

Fuel Density [g/cm^3]	10.97
Atom Fraction ^{235}U	0.05
Atom Fraction ^{238}U	0.95
Fuel Molecular Weight [amu]	270.03

progresses, the isotopic composition of fuel changes. The moderator is light water and its composition, given in Table 5.2, does not change with irradiation. We track

Table 5.2: Water atom density data.

Water Density [g/cm^3]	1
Atom Fraction ^1H	$\frac{2}{3}$
Atom Fraction ^{16}O	$\frac{1}{3}$
Water Molecular Weight [amu]	18.02

five nuclide types in the fuel during the fuel depletion problem: fissile, fertile, parasitic absorber fission product, scattering fission product, and inert, whose spatial nuclide densities, [$atom/cm^3$], are respectively denoted as N_{FS} , N_{FT} , N_{FP-A} , N_{FP-S} , and N_I .

We use a two-energy-group approximation with the standard numbering convention- the lower the group number, the faster the neutron. All neutrons are born fast, there is no thermal upscattering, and we assume all scattering and fission is isotropic in angle. We also assume that if neutron absorption leads to isotopic transmutation,

the transmutation occurs at the time of absorption, there are no radioactive decay chains. All possible transmutation paths are shown in Fig. 5.2.

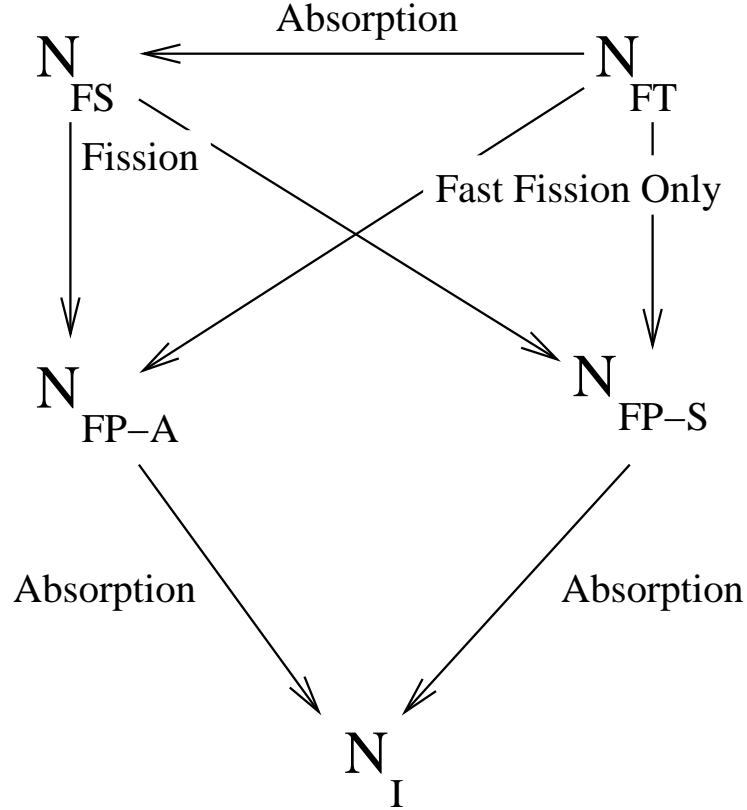


Figure 5.2: Depletion problem possible transmutation paths and mechanisms.

Under these assumptions, and given the transmutation paths given in Fig. 5.2, the fully analytic, nonlinear depletion equations are:

$$\mu \frac{\partial \psi_1}{\partial x} + \Sigma_{t,1} \psi_1 = \frac{\Sigma_{s,1 \rightarrow 1}}{2} \phi_1 + \frac{1}{k} \left(\frac{\nu \Sigma_{f,1}}{2} \phi_1 + \frac{\nu \Sigma_{f,2}}{2} \phi_2 \right) \quad (5.1a)$$

$$\mu \frac{\partial \psi_2}{\partial x} + \Sigma_{t,2} \psi_2 = \frac{\Sigma_{s,2 \rightarrow 2}}{2} \phi_2 + \frac{\Sigma_{s,1 \rightarrow 2}}{2} \phi_1 \quad (5.1b)$$

$$\begin{aligned} \frac{\partial N_{FS}}{\partial t} = & -N_{FS} [(1 - \gamma_{FS,1})\sigma_{a,FS,1}\phi_1 + (1 - \gamma_{FS,2})\sigma_{a,FS,2}\phi_2] \\ & + N_{FT} [\gamma_{FT,1}\sigma_{a,FT,1}\phi_1 + \gamma_{FT,2}\sigma_{a,FT,2}\phi_2] \end{aligned} \quad (5.1c)$$

$$\frac{\partial N_{FT}}{\partial t} = -N_{FT} [\sigma_{a,FT,1}\phi_1 + \sigma_{a,FT,2}\phi_2] \quad (5.1d)$$

$$\begin{aligned} \frac{\partial N_{FP-S}}{\partial t} = & N_{FS} [(1 - p_{FS,1})y_{FS,1}\sigma_{a,FS,1}\phi_1 + (1 - p_{FS,2})y_{FS,2}\sigma_{a,FS,2}\phi_2] \\ & + N_{FT} [(1 - p_{FT,1})y_{FT}\sigma_{a,FT,1}\phi_1] - N_{FS} [\sigma_{a,FP-S,1}\phi_1 + \sigma_{a,FP-S,2}\phi_2] \end{aligned} \quad (5.1e)$$

$$\begin{aligned} \frac{\partial N_{FP-A}}{\partial t} = & N_{FS} [p_{FS,1}y_{FS,1}\sigma_{a,FS,1}\phi_1 + p_{FS,2}y_{FS,2}\sigma_{a,FS,2}\phi_2] \\ & + N_{FT} p_{FT,1}y_{FT,1}\sigma_{a,FT,1}\phi_1 \\ & - N_{FP-A} [(1 - \xi_{FP-A,1})\sigma_{a,FP-A,1}\phi_1 + (1 - \xi_{FP-A,2})\sigma_{a,FP-A,2}\phi_2] \end{aligned} \quad (5.1f)$$

$$\begin{aligned} \frac{\partial N_I}{\partial t} = & N_{FP-A} [\xi_{FP-A,1}\sigma_{a,FP-A,1}\phi_1 + \xi_{FP-A,2}\sigma_{a,FP-A,2}\phi_2] \\ & + N_{FP-S} [\sigma_{a,FP-S,1}\phi_1 + \sigma_{a,FP-S,2}\phi_2] . \end{aligned} \quad (5.1g)$$

In Eq. (5.1a) and Eq. (5.1b) ϕ_g is the group g scalar flux [$n/cm^2/sec$], $\Sigma_{t,g}$ is the total macroscopic cross section [cm^{-1}] of group g , $\Sigma_{s,g \rightarrow g'}$ [cm^{-1}] is the macroscopic cross section for neutrons scattering from group g to group g' , k is the system multiplication factor, and $\nu\Sigma_{f,g}$ is the average number of neutrons released per fission (ν) for a fission induced by a neutron in group g , multiplied by the group g macroscopic fission cross section. The nuclide production destruction equations, Eqs. (5.1c) - (5.1g), use the following notation: $\gamma_{m,g}$ is the probability that a neutron absorption

in nuclide m results in the production of a fissile isotope, $y_{m,g}$ is the ratio of nuclide m 's fission cross section to total cross section for group g , $p_{m,g}$ is the probability that the fission of nuclide m yields a parasitic absorber fission product, and ξ_g is the probability that when a parasitic absorber fission product absorbs a neutron, another parasitic absorber fission product is produced. Though not explicitly noted in Eqs. (5.1), all scalar fluxes ϕ_g , macroscopic cross sections Σ_g , and nuclide densities N , are functions of position. We use Gauss-Legendre S_2 angular quadrature, with weights that sum to 2, to approximate the scalar fluxes.

Macroscopic cross sections are generated from nuclide density and microscopic cross section data. As an example, $\Sigma_{t,g}$ is calculated as shown in Eq. (5.2):

$$\begin{aligned} \Sigma_{t,g,i} = N_{FS}\sigma_{t,FS,g} + N_{FT}\sigma_{t,FT,g} + N_{FP-A}\sigma_{t,FP-A,g} \\ + N_{FP-S}\sigma_{t,FP-S,g} + N_I\sigma_{t,I,g} . \end{aligned} \quad (5.2)$$

Macroscopic fission cross section and average neutrons per fission products are found in a similar fashion, but we can limit our consideration to the fissile and fertile nuclide densities as shown in Eq. (5.3),

$$\nu\Sigma_{f,g,i} = N_{FS}\nu_{FS,g}\sigma_{f,FS,g} + N_{FT}\nu_{FT,g}\sigma_{f,FT,g} . \quad (5.3)$$

5.1.1 Microscopic Cross Section and Yield Data

We complete the specification of the depletion problem by giving the physical data used to solve the problem. Microscopic cross section data for the water is given in Table 5.3. Absorption and scattering cross sections for the fertile and fissile nuclides are given in Table 5.4, and fission cross sections and average neutrons per fission are given in Table 5.5. Radiative capture fractions and probability of an absorbed

neutron inducing fission are given in Table 5.6. Cross-section data for the fission products and inert nuclides are given in Table 5.7. Fission product yields and the parasitic absorber fission product regeneration fraction, ξ , are given in Table 5.8.

Table 5.3: Water microscopic cross section, in barns [10^{-24} cm^2]. Moderator is composed only of H_2O .

Nuclide	$\sigma_{a,1}$	$\sigma_{a,2}$	$\sigma_{s,1 \rightarrow 1}$	$\sigma_{s,2 \rightarrow 2}$	$\sigma_{s,1 \rightarrow 2}$
^1H	0	0.332	0	20.47	3.926
^{16}O	0	0	2.739	3.780	0

Table 5.4: Fuel microscopic cross sections, in barns [10^{-24} cm^2].

Nuclide	$\sigma_{a,1}$	$\sigma_{a,2}$	$\sigma_{s,1 \rightarrow 1}$	$\sigma_{s,2 \rightarrow 2}$	$\sigma_{s,1 \rightarrow 2}$
$^{235}_{92}\text{U}$	1.325	683.21	4.566	15.04	0
$^{238}_{92}\text{U}$	0.374	2.717	4.804	9.36	0

Table 5.5: Average neutron yield per fission, and fission microscopic cross section [10^{-24} cm^2].

Nuclide	ν_1	ν_2	$\sigma_{f,1}$	$\sigma_{f,2}$
$^{235}_{92}\text{U}$	2.6	2.4	1.235	584.4
$^{238}_{92}\text{U}$	2.8	N/A	0.308	0

Table 5.6: Radiative capture fraction, and fission probability for fissile and fertile nuclides.

Nuclide	γ_1	γ_2	y_1	y_2
$^{235}_{92}\text{U}$	$\frac{0.09}{1.325} = 0.068$	$\frac{98.81}{683.21} = 0.145$	$\frac{1.235}{1.325} = 0.932$	$\frac{584.4}{683.21} = 0.855$
$^{238}_{92}\text{U}$	$\frac{0.066}{0.374} = 0.177$	1	$\frac{0.308}{0.374} = 0.823$	0

Table 5.7: Parasitic absorber fission product, scattering fission product, and inert nuclide microscopic cross section data in barns [10^{-24} cm^2].

Nuclide	$\sigma_{a,1}$	$\sigma_{a,2}$	$\sigma_{s,1}$	$\sigma_{s,2}$	$\sigma_{s,1 \rightarrow 2}$
FP-A	15	1000	0.5	5	0
FP-S	0.5	5	15	100	0
Inert	1	5	1	5	0

Table 5.8: Fission product branch ratios and parasitic absorber fission product re-generation fraction.

$p_{FS,1}$	0.3
$p_{FS,2}$	0.3
$p_{FT,1}$	0.3
$p_{FT,2}$	0.3
$\xi_{FP-A,1}$	0.3
$\xi_{FP-A,2}$	0.5

5.1.2 Reactor Power Levels and Normalization

Vacuum boundary conditions are imposed on both sides of the slab. We normalize reactor scalar flux values so that the reactor produces a constant fission power level of 2000 [W] for the duration of the burn-up cycle. The burn-up cycle length consists of 600 full-power days and we use a time step of 10 days to update the scalar fluxes. A typical beginning-of-cycle flux profile is shown in Fig. 5.3 and an end-of-cycle scalar flux profile is shown in Fig. 5.4.

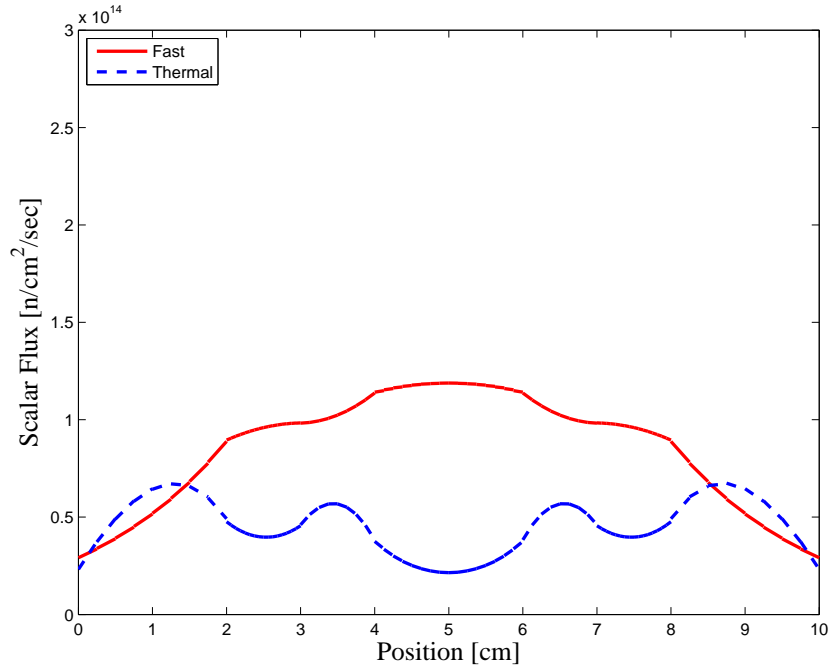


Figure 5.3: Example normalized scalar flux profiles at beginning and end of fuel burn-up cycle.

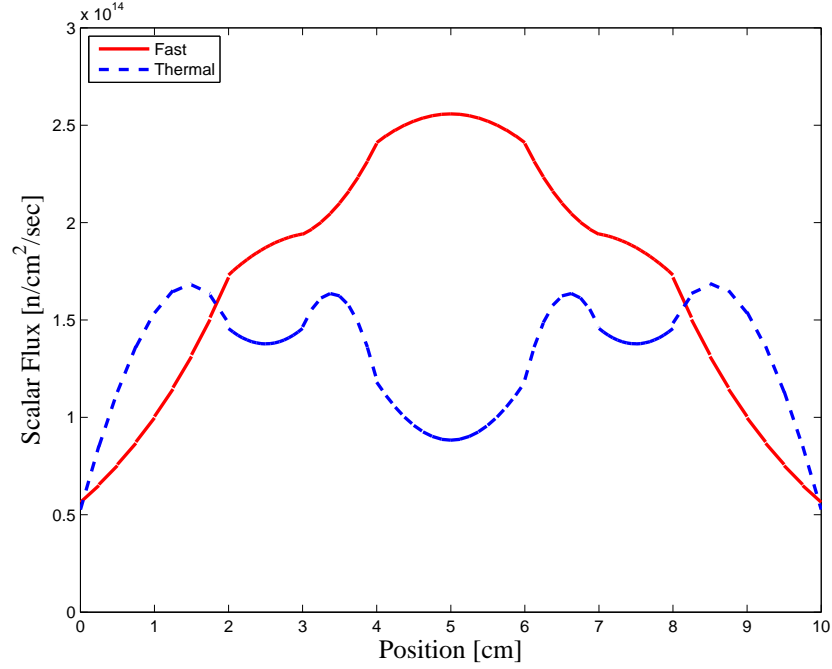


Figure 5.4: Example normalized scalar flux profiles at beginning and end of fuel burn-up cycle.

5.2 Spatial Discretization

We solve Eqs. (5.1) using a semi-static approach [35] assuming the flux distribution at the start of the time step remains constant throughout the time step. Nuclide densities are advanced in time using explicit Euler time differencing, then a corresponding radiation solution is found. We will consider three DFEM schemes to spatially discretize Eqs. (5.1).

1. AD DFEM: expands the angular flux in a P degree polynomial trial space using equally-spaced interpolation points, uses exact spatial integration, assumes cell-wise constant cross sections for solving the radiation equations, and tracks only cell average nuclide densities.
2. SL Collapse: expands both the angular flux and nuclide densities in a P degree

polynomial trial space, uses self-lumping quadrature to approximate integrals, and assumes cell-wise constant cross sections for solving the radiation equations. Lobatto quadrature is used as the DFEM interpolation points for odd degree trial spaces and Gauss quadrature as the DFEM interpolation points for even degree trial spaces.

3. SL Full: expands both the angular flux and nuclide densities in a P degree polynomial trial space, uses self-lumping quadrature to approximate integrals, and explicitly accounts for the variation of macroscopic cross section within each spatial cell. Lobatto quadrature is used as the DFEM interpolation points for odd degree trial spaces and Gauss quadrature as the DFEM interpolation points for even degree trial spaces.

5.2.1 Radiation Solution

Using the nomenclature developed in Chapter 2 and Chapter 3, the spatially discretized radiation equations are:

$$\mu_d \mathbf{G} \vec{\psi}_{d,1} - \mu_d \psi_{in,d,1} \vec{f} + \frac{\Delta x}{2} \mathbf{R}_{\Sigma_{t,1}} \vec{\psi}_{d,1} = \frac{\Delta x}{4} \mathbf{R}_{\Sigma_{s,1 \rightarrow 1}} \vec{\phi}_1 + \frac{1}{k_\tau} \frac{\Delta x}{4} \left(\mathbf{R}_{\nu \Sigma_{f,1}} \vec{\phi}_1 + \mathbf{R}_{\nu \Sigma_{f,2}} \vec{\phi}_2 \right), \quad (5.4)$$

and

$$\mu_d \mathbf{G} \vec{\psi}_{d,2} - \mu_d \psi_{in,d,2} \vec{f} + \frac{\Delta x}{2} \mathbf{R}_{\Sigma_{t,2}} \vec{\psi}_{d,2} = \frac{\Delta x}{4} \left(\mathbf{R}_{\Sigma_{s,2 \rightarrow 2}} \vec{\phi}_2 + \mathbf{R}_{\Sigma_{s,1 \rightarrow 2}} \vec{\phi}_1 \right). \quad (5.5)$$

In Eq. (5.4) and Eq. (5.5) k_τ is the multiplication factor at time index τ and $\mathbf{R}_{\Sigma_{s,1 \rightarrow 1}}$, $\mathbf{R}_{\Sigma_{s,1 \rightarrow 2}}$, $\mathbf{R}_{\Sigma_{s,2 \rightarrow 2}}$, $\mathbf{R}_{\nu \Sigma_{f,1}}$, and $\mathbf{R}_{\nu \Sigma_{f,2}}$ are defined analogously to \mathbf{R}_{Σ_t} , as in Eq. (3.3), replacing $\Sigma_t(s)$ with $\Sigma_{s,1 \rightarrow 1}(s)$, $\Sigma_{s,1 \rightarrow 2}(s)$, $\Sigma_{s,2 \rightarrow 2}(s)$, $\Sigma_{f,1}(s)$, and $\Sigma_{f,2}(s)$,

respectively.

We solve Eq. (5.4) and Eq. (5.5) following the standard power iteration procedure, described for DFEM neutron transport in [47]. Convergence is checked after each power iteration. Using ℓ as the iteration index, convergence after the $\ell + 1$ iterate is said to occur when:

$$\delta_k = \left| \frac{k^{(\ell+1)} - k^{(\ell)}}{k^{(\ell)}} \right| < \epsilon_k, \quad (5.6)$$

and

$$\delta_\phi = \max_{g=1,2} \max_{i=1,\dots,N_{cell}} \max_{j=1,\dots,N_P} \left| \frac{\phi_{g,i,j}^{(\ell+1)} - \phi_{g,i,j}^{(\ell)}}{\phi_{g,i,j}^{(\ell)}} \right| < \epsilon_\phi. \quad (5.7)$$

In our computational results we use $\epsilon_k = 10^{-12}$ and $\epsilon_\phi = 10^{-10}$. For each power iteration, the within group components of Eq. (5.4) and Eq. (5.5) are solved separately with a single transport sweep and S2SA iteration. Since we are using an S_2 angular quadrature, a single source iteration with a single S2SA step exactly solves a given within group neutron transport problem.

The converged scalar flux is normalized such that the desired power level of $P_{Total} = 2000 [W/cm^2]$, is achieved. All fission energy is assumed to be deposited only in the fuel. For the SL Full scheme, we calculate the normalization factor, F_P , as:

$$F_P = E_f \sum_{g=1}^2 \left[\sum_{i=1}^{N_{Fuel}} \frac{\Delta x_i}{2} \sum_{j=1}^{N_P} w_j \Sigma_{f,g,i,j} \phi_{g,i,j} \right], \quad (5.8)$$

where in E_f is the energy released per fission, assumed to be 200 [MeV], w_j is the quadrature weight associated with the j -th DFEM interpolation point, and N_{Fuel} is the total number of spatial cells in the fuel region. The SL Collapse and AD DFEM

schemes calculate F_P as

$$F_P = E_f \sum_{g=1}^2 \left[\sum_{i=1}^{N_{Fuel}} \frac{\Delta x_i \hat{\Sigma}_{f,g,i}}{2} \sum_{j=1}^{N_P} w_j \Sigma_{f,g,i,j} \phi_{g,i,j} \right]. \quad (5.9)$$

$\tilde{\phi}^{(\ell+1)}$ is then scaled as:

$$\tilde{\phi}_g^{(\ell+1)} \leftarrow \frac{P_{Total}}{F_P} \tilde{\phi}_g^{(\ell+1)}. \quad (5.10)$$

5.2.2 Nuclide Density

We consider three spatial discretization schemes to solve the nuclide production/destruction components of Eqs. (5.1). The first, AD DFEM, tracks only cell average nuclide densities, approximating the true spatial distribution of nuclide m , $N_m(x, t)$, as being a constant in each cell, equal to the cell average density of nuclide m . Denoting the average nuclide density in cell i for nuclide m as $\bar{N}_{m,i}$ and the average group g scalar flux in cell i as $\bar{\phi}_{g,i}$, we give the fissile nuclide update equation for the AD DFEM scheme in Eq. (5.11):

$$\begin{aligned} \frac{\bar{N}_{FS,i}^{\tau+1} - \bar{N}_{FS,i}^{\tau}}{\Delta t} = & \bar{N}_{FT,i}^{\tau} [\gamma_{FT,1} \sigma_{a,FT,1} \bar{\phi}_{1,i}^{\tau} + \gamma_{FT,2} \sigma_{a,FT,2} \bar{\phi}_{2,i}^{\tau}] \\ & - \bar{N}_{FS,i}^{\tau} ((1 - \gamma_{FS,1}) \sigma_{a,FS,1} \bar{\phi}_{1,i}^{\tau} + (1 - \gamma_{FS,2}) \sigma_{a,FS,2} \bar{\phi}_{2,i}^{\tau}). \end{aligned} \quad (5.11)$$

In Eq. (5.11) superscript τ denotes time index τ quantities and Δt is the time step size. The update equations for N_{FT} , N_{FP-A} , N_{FP-S} , and N_I can be derived analogously to Eq. (5.11). AD DFEM calculates $\bar{\phi}_{g,i}$ using closed Newton-Cotes quadrature with the quadrature points limited to the DFEM interpolation points:

$$\bar{\phi}_{g,i} = \frac{1}{2} \sum_{j=1}^{N_P} w_j \phi_{g,j}. \quad (5.12)$$

The averaging in Eq. (5.12) is exact since an N_P point closed Newton-Cotes quadrature can exactly integrate any P degree polynomial. Equation (5.11) locally updates all average nuclide densities, $\bar{N}_{FS,i}$, $\bar{N}_{FT,i}$, $\bar{N}_{FP-A,i}$, $\bar{N}_{FP-S,i}$, and $\bar{N}_{I,i}$, simultaneously via a 5×5 matrix-vector multiply.

The SL Full and SL Collapse schemes approximate the true spatial density of nuclide m as a P degree Lagrange polynomial, $\tilde{N}_m(s)$, in each cell:

$$\tilde{N}_m(s) = \sum_{j=1}^{N_P} N_{m,j} b_j(s), \quad (5.13)$$

where b_j are the $N_P = P + 1$ Lagrange interpolatory polynomials in the interval $s \in [-1, 1]$. We require the set of nuclide density DFEM interpolation points to be the same set of N_P points as the angular flux DFEM interpolation points. Following a Galerkin procedure, we multiply each production/destruction nuclide equation of Eqs. (5.1) by basis function b_j and integrate generating $5(P + 1)$ equations. The system of update equations for $\vec{N}_{FS,i}$ is shown in Eqs. (5.14):

$$\begin{aligned} \frac{1}{\Delta t} \frac{\Delta x}{2} \mathbf{M} \left(\vec{N}_{FS,i}^{\tau+1} - \vec{N}_{FS,i}^{\tau} \right) = & -(1 - \gamma_{FS,1}) \sigma_{a,FS,1} \frac{\Delta x}{2} \widehat{\mathbf{M}}_{\phi_{1,i},\tau} \vec{N}_{FS,i}^{\tau} \\ & - (1 - \gamma_{FS,2}) \sigma_{a,FS,2} \frac{\Delta x}{2} \widehat{\mathbf{M}}_{\phi_{2,i},\tau} \vec{N}_{FS,i}^{\tau} \\ & + \gamma_{FT,1} \sigma_{a,FT,1} \frac{\Delta x}{2} \widehat{\mathbf{M}}_{\phi_{1,i}^{\tau}} \vec{N}_{FT,i}^{\tau} + \gamma_{FT,2} \sigma_{a,FT,2} \frac{\Delta x}{2} \widehat{\mathbf{M}}_{\phi_{2,i}^{\tau}} \vec{N}_{FT,i}^{\tau}. \end{aligned} \quad (5.14)$$

The equations for $\vec{N}_{FT,i}$, $\vec{N}_{FP-A,i}$, $\vec{N}_{FP-S,i}$, and $\vec{N}_{I,i}$ are derived in a similar fashion to Eq. (5.14). In Eq. (5.14) we have defined:

$$\widehat{\mathbf{M}}_{\phi_{g,i}^{\tau},jk} = \int_{-1}^1 b_j(s) b_k(s) \tilde{\phi}_{g,\tau,i}(s) ds, \text{ and} \quad (5.15)$$

$$\vec{N}_{m,i}^\tau = \begin{bmatrix} N_{m,1}^\tau \\ \vdots \\ N_{m,j}^\tau \\ \vdots \\ N_{m,P+1}^\tau \end{bmatrix}. \quad (5.16)$$

Given that we track five nuclides in the fuel region, each expanded in a P degree polynomial in each cell, there are $5(P+1)$ unknowns in each cell, thus Eq. (5.14) is a closed, $5(P+1) \times 5(P+1)$ system of linear equations for the $5(P+1)$ unknown nuclide densities, $N_{m,i,j}$ in each cell.

Using self-lumping quadrature to approximate Eq. (5.15) causes $\widehat{\mathbf{M}}_{\phi_{g,i}^\tau}$ to be a diagonal matrix. Recalling that \tilde{N}_m uses the same interpolation points as $\tilde{\phi}_g$, approximating the integration of Eq. (5.15) with numerical quadrature restricted to the DFEM interpolating points, results in

$$\widehat{\mathbf{M}}_{\phi_{g,i}^\tau}^{jk} = \begin{cases} w_j \phi_{g,\tau,i,j} & j = k \\ 0 & \text{otherwise} \end{cases}. \quad (5.17)$$

Macroscopic cross sections are generated from nuclide density and microscopic cross section data. For the AD DFEM scheme, each cell has a single macroscopic cross section (per reaction type), and a single value of nuclide density for each nuclide type. Thus, interaction cross sections are easily tabulated. As an example, the cell average total interaction cross section in cell i is calculated in Eq. (5.18):

$$\begin{aligned} \widehat{\Sigma}_{t,g,i} = & N_{FS,i} \sigma_{t,FS,g} + N_{FT,i} \sigma_{t,FT,g} + N_{FP-A,i} \sigma_{t,FP-A,g} \\ & + N_{FP-S,i} \sigma_{t,FP-S,g} + N_{I,i} \sigma_{t,I,g}. \end{aligned} \quad (5.18)$$

Macroscopic fission cross section and average neutrons per fission products are found in a similar fashion, but we can limit our consideration to the fissile and fertile nuclide densities as shown in Eq. (5.19),

$$\widehat{\nu\Sigma}_{f,g,i} = N_{FS,i}\nu_{FS,g}\sigma_{f,FS,g} + N_{FT,i}\nu_{FT,g}\sigma_{f,FT,g}. \quad (5.19)$$

The SL Full and SL Collapse schemes calculate macroscopic cross sections in a similar fashion to Eq. (5.18) and Eq. (5.19), but instead of calculating a cell average, $\widehat{\Sigma}_{g,i}$, they calculate macroscopic values at each DFEM interpolation point. SL Collapse then averages the macroscopic cross section at each DFEM interpolation point to estimate the cell average cross section, as shown in Eq. (5.20) for $\widehat{\Sigma}_{t,i}$:

$$\widehat{\Sigma}_{t,i} = \frac{1}{2} \sum_{j=1}^{N_P} w_j \Sigma_{t,g,i,j}. \quad (5.20)$$

5.3 Numerical Results

Since an analytic solution to this depletion problem is not available we employ a fine spatial mesh to obtain the reference solution. We use a fine mesh of 10,240 cells and the SL Full scheme with a quartic polynomial trial space as our reference numerical solution. We present L_2 spatial error measures for

1. the total scalar flux (E_ϕ),
2. the fissile nuclide density ($E_{N_{FS}}$),
3. the fertile nuclide density ($E_{N_{FT}}$), and
4. the parasitic absorber fission product ($E_{N_{FP-A}}$).

To allow for easier comparison, we normalize each error to the reference solution quantity. We define E_ϕ as:

$$E_\phi = \frac{\sqrt{\sum_{g=1}^2 \sum_{i=1}^{N_{ref}} \frac{\Delta x_i}{2} \sum_{q=1}^{N_{qf}} w_q \left(\tilde{\phi}_{ref,i,g}(s_q) - \tilde{\phi}_{num,i,g}(s_q) \right)^2}}{\sqrt{\sum_{g=1}^2 \sum_{i=1}^{N_{ref}} \frac{\Delta x_i}{2} \sum_{q=1}^{N_{qf}} w_q \tilde{\phi}_{ref,i,g}(s_q)^2}}, \quad (5.21)$$

where N_{ref} is the number of reference cells, $\tilde{\phi}_{ref,i,g}(s)$ is the reference solution group g scalar flux in cell i , and $\tilde{\phi}_{num,i,g}(s)$ is the coarse mesh numerical scheme's approximation of the group g scalar flux in cell i . Error measures for N_{FS} , N_{FT} , and N_{FP-A} are derived similarly.

Convergence of E_ϕ is shown in Figs. 5.5-5.8 as a function of DFEM trial space degree and DFEM scheme.

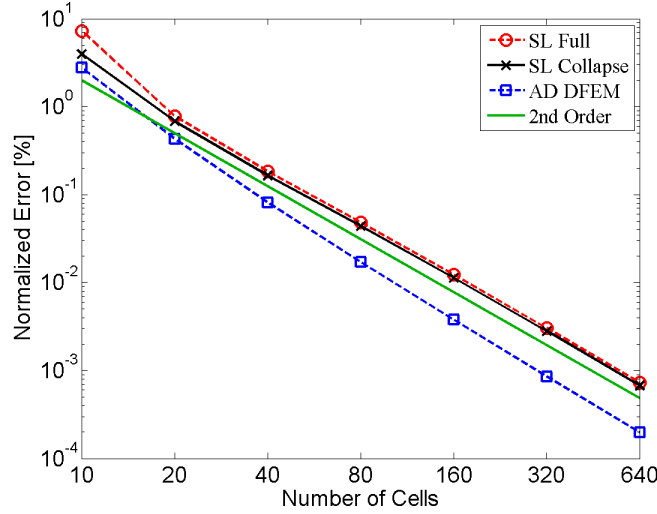


Figure 5.5: Normalized total scalar flux error for the depletion problem at end of cycle, for linear DFEM.

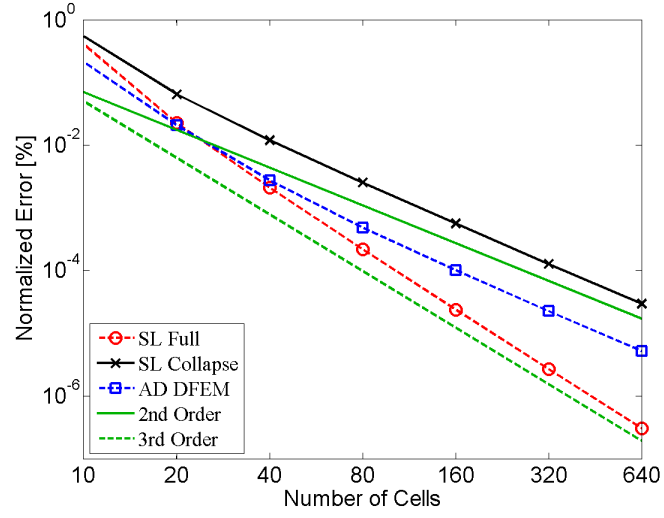


Figure 5.6: Normalized total scalar flux error for the depletion problem at end of cycle, for quadratic DFEM.

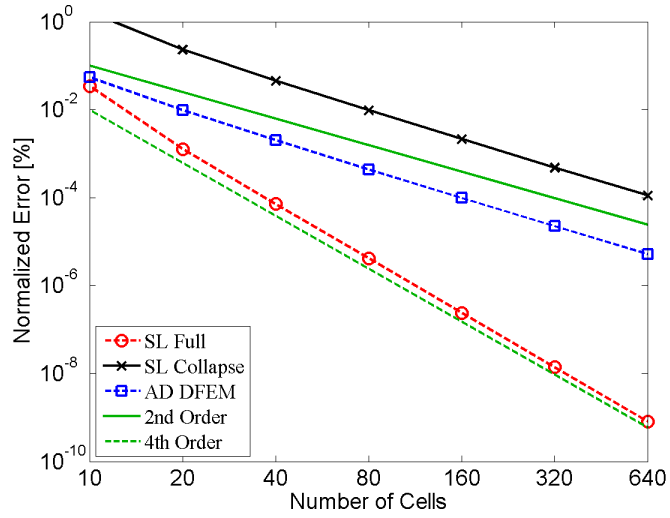


Figure 5.7: Normalized total scalar flux error for the depletion problem at end of cycle, for cubic DFEM.

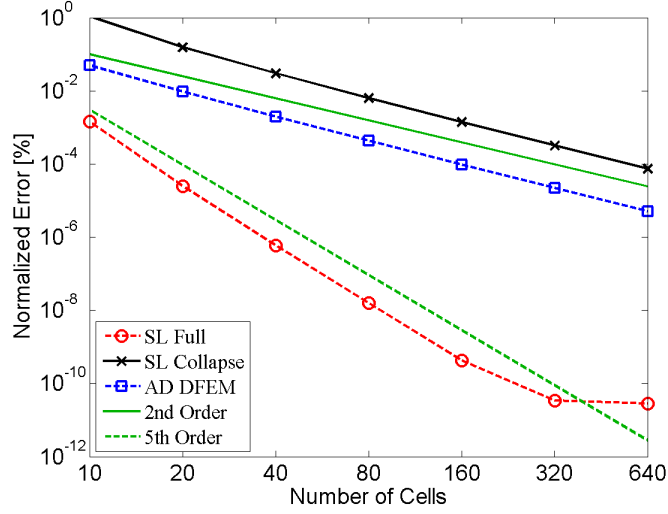


Figure 5.8: Normalized total scalar flux error for the depletion problem at end of cycle, for quartic DFEM.

Figures 5.5-5.8 re-emphasize two key results observed in the case of a pure absorber. First, when employing cell-wise constant cross sections, angular / scalar flux convergence is at most second order in space, regardless of the DFEM trial space polynomial degree. Second, exact integration of the interaction terms in the DFEM moment equations is not required to achieve high-order accuracy. In the depletion term, the DFEM interaction term is a degree $3P$ polynomial, and self-lumping schemes using Gauss or Lobatto quadrature only integrate $2P + 1$ and $2P - 1$ degree polynomials, respectively.

Convergence of $E_{N_{FS}}$, $E_{N_{FT}}$, and $E_{N_{FP-A}}$ are given in Figs. 5.9-5.12, Figs. 5.13-5.16, and Figs. 5.17-5.20, respectively for linear through quartic trial space degrees.

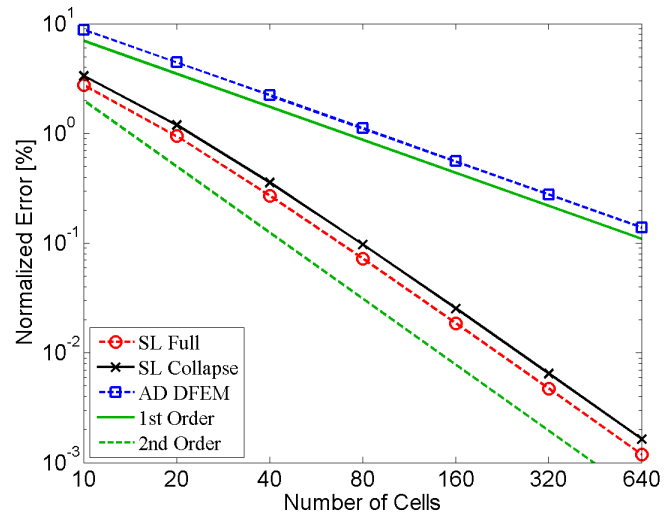


Figure 5.9: Normalized fissile nuclide density error for the depletion problem at end of cycle, for linear DFEM

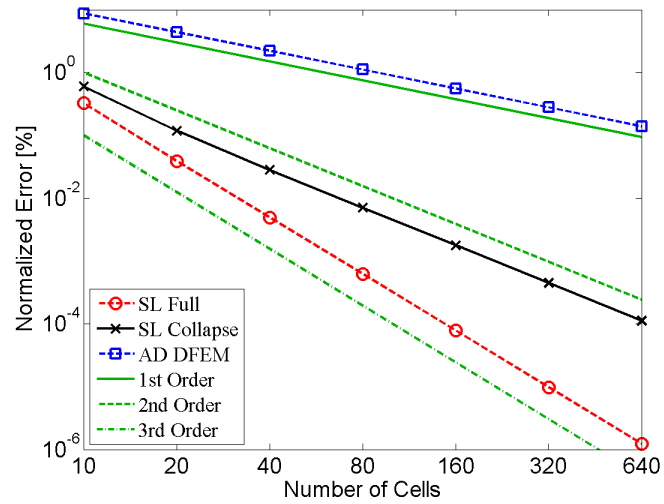


Figure 5.10: Normalized fissile nuclide density error for the depletion problem at end of cycle, for quadratic DFEM.

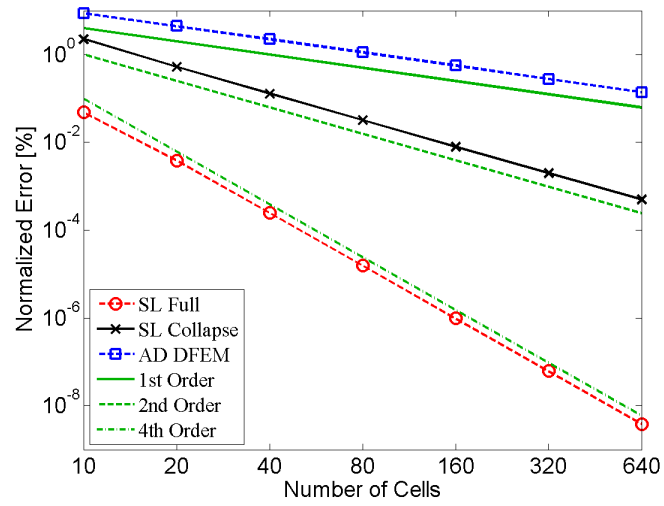


Figure 5.11: Normalized fissile nuclide density error for the depletion problem at end of cycle, for cubic DFEM.

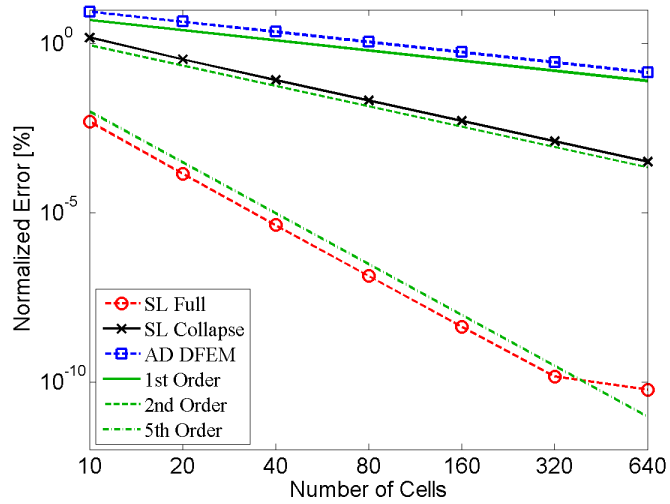


Figure 5.12: Normalized fissile nuclide density error for the depletion problem at end of cycle, for quartic DFEM.

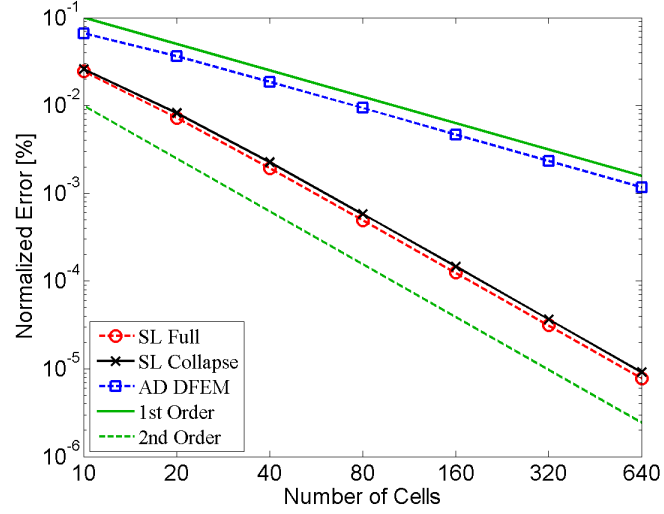


Figure 5.13: Normalized fertile nuclide density error for the depletion problem at end of cycle, for linear DFEM.

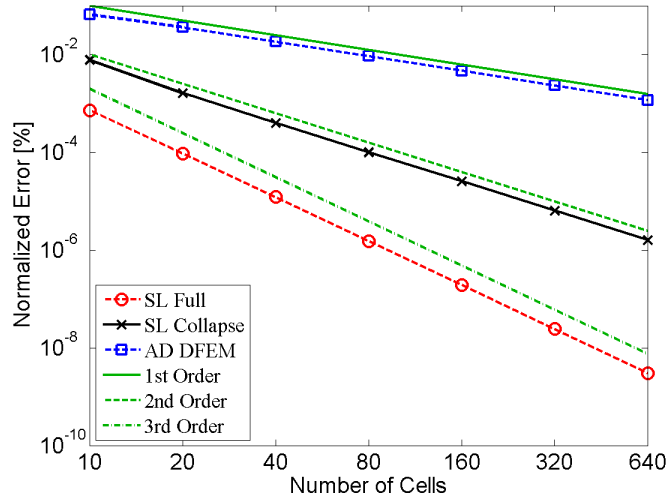


Figure 5.14: Normalized fertile nuclide density error for the depletion problem at end of cycle, for quadratic DFEM..

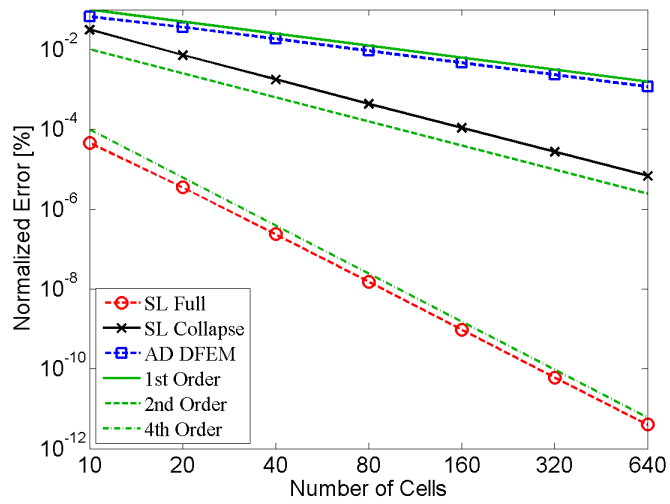


Figure 5.15: Normalized fertile nuclide density error for the depletion problem at end of cycle, for cubic DFEM.

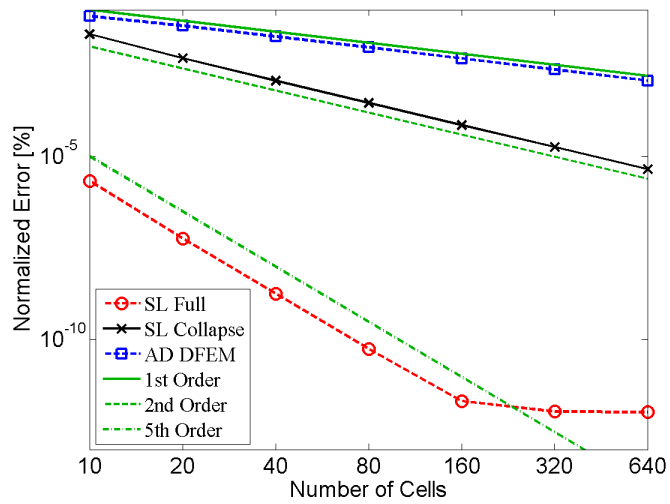


Figure 5.16: Normalized fertile nuclide density error for the depletion problem at end of cycle, for quartic DFEM.

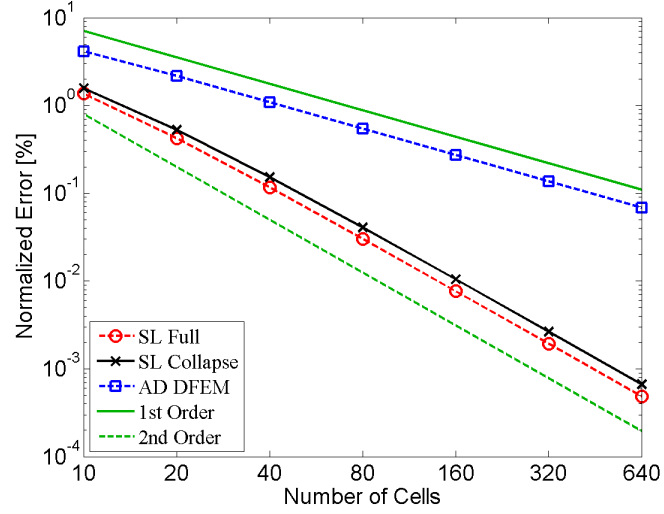


Figure 5.17: Normalized parasitic absorber fission product error for the depletion problem at end of cycle, for linear DFEM.

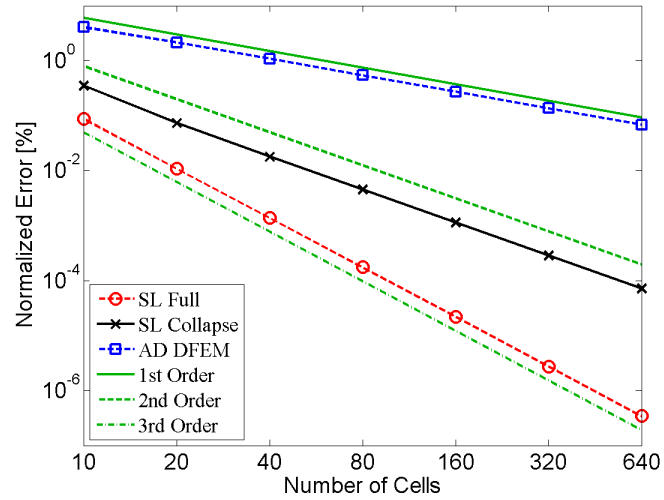


Figure 5.18: Normalized parasitic absorber fission product error for the depletion problem at end of cycle, for quadratic DFEM.

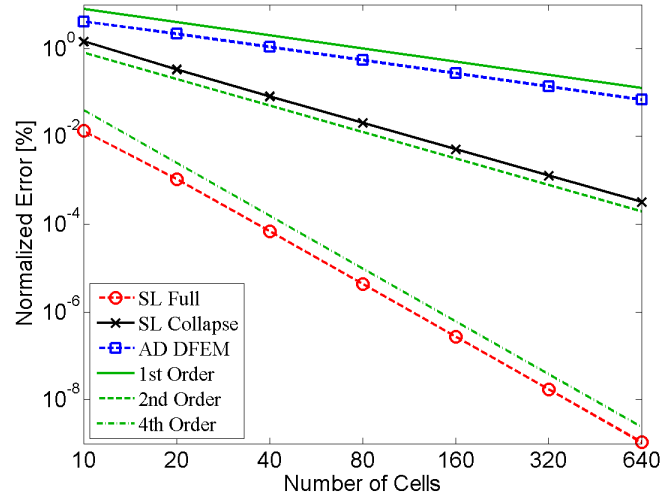


Figure 5.19: Normalized parasitic absorber fission product error for the depletion problem at end of cycle, for cubic DFEM.

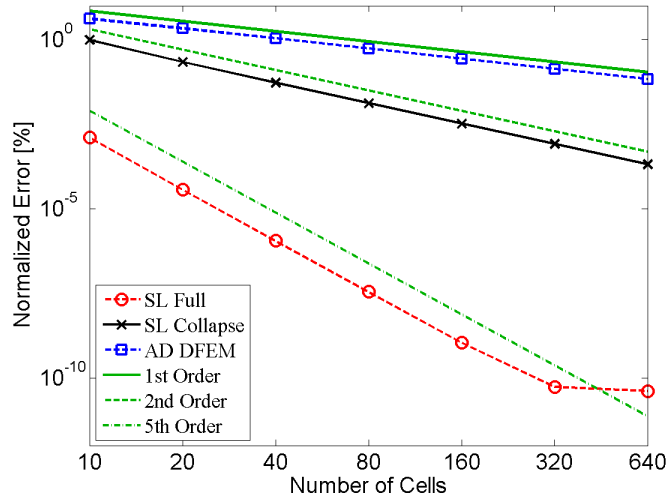


Figure 5.20: Normalized parasitic absorber fission product error for the depletion problem at end of cycle, for quartic DFEM.

Examining the spatial convergence in nuclide densities, we make several observations. First, we note that the AD DFEM scheme (cell-wise average cross section, cell average nuclide density) achieves at most first-order convergence for all spatial nuclide density errors, regardless of the angular flux trial space degree. The AD DFEM scheme is limited to at most first-order convergence of the error in the spatial distribution of nuclides because the scalar flux is updated using only a cell-wise average cross section and only the cell average nuclide density is tracked. Second, though the SL Collapse scheme expands nuclide density in a P degree polynomial DFEM trial space, it achieves at most second-order L^2 convergence of the error in nuclides spatial distribution, for all trial space polynomial degrees. SL Collapse is limited to at most second-order convergence of the spatial nuclide density solely because the scheme assumes a constant cross section in each cell when updating the scalar flux. The respective first-order and second-order convergence of the error in nuclide spatial distribution of the AD DFEM and SL Collapse scheme verifies the result observed in the pure absorber problem: assuming a cell-wise average cross section for coupled radiation transport problems limits the order of convergence of any quantity that depends on an interaction rate. SL Full achieves $P + 1$ order convergence of the error in spatial nuclide density for the fuel depletion problem, showing that coupled systems of equations involving radiation transport can be solved with arbitrary order of accuracy using high-order DFEM polynomial trial spaces and self-lumping numerical quadrature that explicitly accounts for the spatial variation of cross section within each cell.

6. LAST CHAPTER: THE IMPORTANCE OF RESEARCH

6.1 New Section

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