

Verification of the Cardinal Multiphysics Solver for 1-D Coupled Heat Transfer and Neutron Transport

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ABSTRACT

Cardinal is a multiphysics software tool that couples OpenMC Monte Carlo transport and NekRS Computational Fluid Dynamics (CFD) to the Multiphysics Object-Oriented Simulation Environment (MOOSE). This work verifies Cardinal for coupled neutron transport and heat conduction using a 1-D analytical solution from previous work by the Naval Nuclear Laboratory. This numerical benchmark includes S_2 transport, Doppler-broadened cross sections, thermal conduction and expansion, and convective boundary conditions. The goal of this work is to verify Cardinal's basic multiphysics modeling capabilities for coupled neutronics and heat conduction. The benchmark provides analytical solutions for the temperature and flux distributions, as well as the k -eigenvalue. Using these solutions, an L_2 error norm was computed for each spatial discretization: namely finite element heat conduction mesh and Monte Carlo cells. The temperature error showed linear convergence on a log-log plot of error vs. mesh element number, with a slope of -0.9986 ($R^2 \approx 1.0$). Nearly all spatial flux predictions, except a few points in the $N = 250$ case, space were within 2σ of the analytical solution, for Monte Carlo cell counts between 50 and 1000. The eigenvalue k_{eff} also agrees well with the benchmark value for each mesh size. The outcome of this work is verification of coupled Monte Carlo-thermal conduction modeling using Cardinal.

KEYWORDS: Cardinal, MOOSE, OpenMC, multiphysics, verification

1. INTRODUCTION

With recent advancements in methods, software, and computing, high-fidelity, multiphysics Modeling and Simulation (M&S) is becoming an important component of the nuclear engineer's "toolbox." These high-fidelity models substitute more conservative safety factors with physics-based simulation. This can reduce uncertainty in analyses, enabling tighter margins to realize improved economics and licensing certainty. However, analytical benchmarks and comparison to experimental data are required to assess the stability, convergence, and accuracy of these high-fidelity models for reactor design and analysis.

Cardinal* is an open-source code [1] that wraps OpenMC [2] Monte Carlo particle transport and NekRS [3] CFD within the MOOSE [4] framework. This coupling brings high-fidelity multiphysics feedback to the MOOSE "ecosystem." Cardinal couples OpenMC and NekRS to MOOSE simulations by copying data between the internal code data structures (e.g. a vector of tally results in OpenMC) and a `MooseMesh`, or the unstructured mesh class in MOOSE. MOOSE's mesh-to-mesh interpolation system then communicates between the `MooseMesh` "mirror" of the external code's solution and an arbitrary coupled MOOSE application in the form of boundary conditions (such as for conjugate heat transfer with NekRS) or source terms (such as for volumetric heating with OpenMC). Convergence is obtained with Picard iteration.

*<https://cardinal.cels.anl.gov/>

For coupled neutronics-thermal-fluid simulations with OpenMC, each Picard iteration consists of several steps: 1) a MOOSE application (e.g. BISON, Pronghorn, NekRS via Cardinal, ...) solves for temperatures and densities; 2) Cardinal transfers temperatures and densities to the OpenMC model; 3) OpenMC solves for the nuclear heating; and 4) Cardinal transfers the tally values back to the `MooseMesh` “mirror.” These steps continue until convergence criteria are achieved. In this work, we pursue verification of these multiphysics aspects of Cardinal using a 1-D analytical benchmark from the Naval Nuclear Laboratory [5]. This work does not require CFD, and thus NekRS will be left out of the discussion from this point on. However, we note that a companion paper in this conference does use NekRS’s heat conduction solver for verification on a different, fixed-source benchmark [6].

The remainder of this paper is organized as follows. In Section 2, we summarize the analytical benchmark modeled in this work. Section 3 then describes the Cardinal computational model of the benchmark. Section 4 presents comparisons between Cardinal and the analytical benchmark. Finally, Section 5 presents conclusions and outlines ongoing and future efforts in the verification and validation of Cardinal.

2. BENCHMARK PROBLEM DESCRIPTION

The analytical benchmark couples three physics: S_2 neutron transport with Doppler broadening, heat conduction, and thermal expansion. S_2 transport restricts the neutron direction to only the $\pm x$ direction. A summary of the governing Ordinary Differential Equations (ODEs) and boundary conditions in the 1-D slab is shown in Figure 1. The assumed model for thermal conductivity has already been inserted into the energy conservation equation.

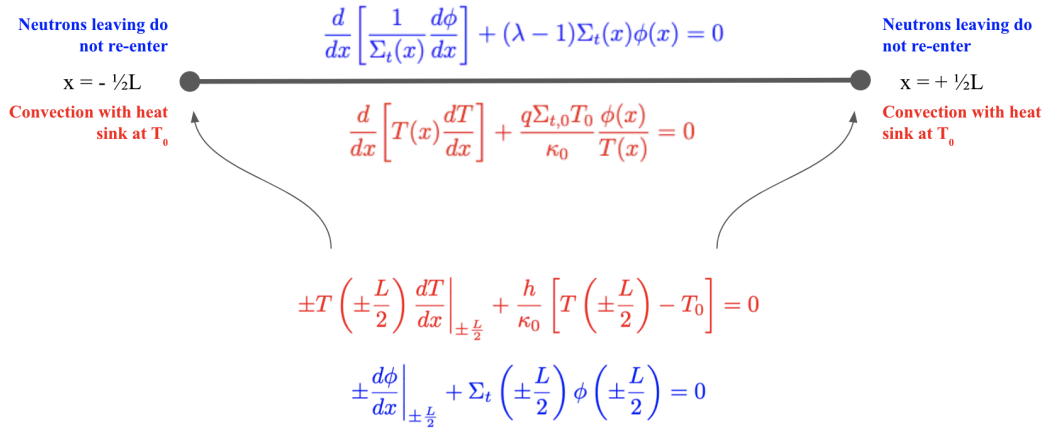


Figure 1: The domain, ODEs, and boundary conditions for the slab.

This benchmark uses a one-group assumption for the neutron cross sections. As neutrons transport, heat from fission events deposits volumetric power in the slab, causing thermal expansion and affecting the temperature distribution; thermal expansion is restricted to only the x -direction. This slab elongation feeds back into neutronics and heat conduction by influencing the domain length and material density. The slab has convective boundary conditions at the endpoints $x = \pm L/2$ with the heat sink temperature T_0 . The Doppler-broadened, total, microscopic cross section follows an inverse-root temperature relationship,

$$\sigma_t(x) = \sigma_{t,0} \sqrt{\frac{T_0}{T(x)}}. \quad (1)$$

Due to thermal expansion, the slab density varies as

$$\rho(x) = \rho_0 \sqrt{\frac{T_0}{T(x)}}. \quad (2)$$

This gives a Doppler-broadened, macroscopic, total cross section that accounts for changes in density due

to temperature as

$$\begin{aligned}\Sigma_t(x) &= \frac{\rho_0 \sigma_{t,0} N_A}{A} \frac{T_0}{T(x)} \\ &= \Sigma_{t,0} \frac{T_0}{T(x)},\end{aligned}\tag{3}$$

where $\sigma_{t,0}$ is the total microscopic cross section at T_0 , N_A is Avogadro's number, and A is the mass number of the medium. The conduction equation governs energy conservation in the slab and can be described in terms of the thermal conductivity κ , the energy released per reaction q , the total macroscopic cross section Σ_t , and the neutron flux ϕ ,

$$\frac{d}{dx} \left[\kappa(T) \frac{dT(x)}{dx} \right] + q \Sigma_t(x) \phi(x) = 0.\tag{4}$$

Note that typically q represents the energy per fission, but in this benchmark, q is per total reaction.

By defining the problem in this way, the differential equations for thermal conduction and neutron transport have similar forms, suggesting that it may be possible that the temperature and flux are related by a constant, $T(x) = f\phi(x)$. This assumption is satisfied by manipulating the ODEs, inserting the cross section temperature dependence, and matching coefficients so that the two ODEs are of the same form. The matching of coefficients imposes two constraints that give equations for the total, microscopic cross section $\sigma_{t,0}$ and the heat transfer coefficient h , in terms of the slab parameters. An important consequence of this problem is that the fission power is uniform throughout the geometry (but the flux is not). For full details on the analytical solution's derivation, see [5].

3. COMPUTATIONAL MODEL

This section describes the OpenMC and MOOSE computational models, followed by the coupling's convergence criteria and a description of the mapping between OpenMC and MOOSE geometries. When the benchmark model was completed, Cardinal did not yet support moving geometries in OpenMC. Instead of modeling thermal expansion using thermomechanics, our simulation uses the analytical solution for the equilibrium length L from [5] that accounts for all three physics effects. This simulation accounts for changes in density due to temperature by using the total, macroscopic cross section – which absorbs density to become a $\frac{1}{T}$ dependence – as was mentioned in Section 2. We note, in a companion paper in this conference [7], that Cardinal now supports mesh-based geometry (and deforming-mesh problems) through the DAGMC plugin, and hence incorporating the thermomechanics feedback will be included in our future work.

3.1. OpenMC Model

The slab is divided evenly into N identically-shaped, rectangular cells along the x -dimension; the cases here are $N = [5, 10, 25, 50, 100, 250, 500, 1000]$ cells. The geometry used in OpenMC must be three-dimensional, despite the benchmark being a one-dimensional problem. In general, reflective boundary conditions can be used to remove dimensions from a problem. Thus, this benchmark can be represented with vacuum boundary conditions at $x = \pm \frac{L}{2}$ and reflective boundary conditions at the y and z boundaries. To simplify normalization of the power integral in Equation (19) of the benchmark, the geometry has 1 cm dimensions in the y and z directions [5]. Figure 1 shows a diagram of the 1-D geometry, governing equations, and boundary conditions for the different physics.

The benchmark's one-group assumption was satisfied using OpenMC's multigroup mode. Since the benchmark uses a fictitious material with a known function for the temperature dependence, the simulation took advantage of OpenMC's capability for user-defined cross sections via the `XSdata` class. The cross section for each reaction was specified for 50 evenly spaced temperatures between 308 K and 358 K (a range slightly larger than, but including the analytical temperature range from the benchmark) and was exported to a library. Then, when running OpenMC in multigroup mode, that library is loaded to determine the appropriate cross section in each region as temperature changes from iteration to iteration. For a temperature T between two library data points, we use the nearest temperature available.

OpenMC required slight source code modifications to accommodate S_2 -like transport. While reflective boundary conditions in the y and z directions are necessary, in general, to establish a 1-D problem, in this benchmark particles only move in the $\pm x$ direction. In typical OpenMC simulations, the physics of each reaction describes the scattering dynamics. The Monte Carlo algorithm is agnostic to the direction particles move, but it is not typically constrained to a discrete directional distribution. However, when modeling this benchmark, any history with a particle moving perpendicular to the x -direction would attenuate particles in less x -distance than if particles were constrained to either $\pm x$. To address this, we first use OpenMC's `PolarAzimuthal` distribution to restrict the birth direction of particles to the $\pm x$ direction. We then also modified OpenMC in a patched branch to mimic S_2 transport in two ways. First, when determining the angular cosine of scattering events, μ , particles either continue forward ($\mu = 1$) or are back-scattered ($\mu = -1$) with equal probability, as opposed to sampling the reaction physics for μ . Second, since the simulation uses k -eigenvalue mode, neutrons born in subsequent generations of the simulation also need to have their angular birth distributions restricted to $\pm x$.

As with any Monte Carlo eigenvalue method, each Picard iteration requires sufficient inactive batches to converge the fission source, followed by sufficient active batches to accumulate the relevant tally results. A Shannon Entropy [8] study for this system was conducted on the 1000 cell case and found that the Shannon Entropy converges after about 5 inactive batches. While this may seem like a low number, the problem is 1-D and the initial guess provided to OpenMC is a uniform distribution, which is the same shape as the actual converged solution. The k -eigenvalue simulation used 50,000 particles per batch, with 50 inactive batches and 100 active batches for every case.

For simplicity, we always run 200 Picard iterations, though more advanced metrics in Cardinal can be used for programmatically evaluating coupled physics convergence. Once the temperature was converged following 200 Picard iterations, a final OpenMC run was performed using the temperature distribution produced from the last Picard iteration with 250,000 particles per batch and the same number of batches.

Cardinal kept track of a few tallies in order to compare to the analytical solution: a flux cell tally, a fission heating (κ -fission) cell and global tally, and the eigenvalue k_{eff} . Though flux and the eigenvalue are the only neutronics quantities to compare with the benchmark, these other tallies are needed in order to compute a source strength. For an eigenvalue problem, the tally normalization relies on a source strength, S , related to the total system power:

$$S = \frac{P}{H} \left[\frac{\text{sp}}{\text{s}} \right] \quad (5)$$

where $P = 10^{22} \frac{\text{eV}}{\text{s}}$ is the slab integrated power from the benchmark and $H \left[\frac{\text{eV}}{\text{sp}} \right]$ is the total tallied fission heating tally, where “sp” indicates “source particle.” Each tally must be multiplied by this source strength to convert to physically meaningful units. Since OpenMC does not report volume-normalized tallies, it is necessary to divide by the cell volume for quantities such as neutron flux and volumetric heating.

3.2. MOOSE Heat Conduction Model

MOOSE is used to solve for the temperature distribution within the slab via the Heat Conduction Module. In this MOOSE model, a mesh is used that has identical dimensions to the cells used in the OpenMC model. While this is not required by Cardinal, it allows for 1:1 feedback between the temperature computed in MOOSE and the heat source computed in OpenMC (and is adequate for this geometrically simple benchmark). MOOSE solves the conduction equation using the Finite Element Method (FEM):

$$-\nabla \cdot [\kappa_s(\mathbf{r}, T_s) \nabla T_s(\mathbf{r})] = \dot{q}_s, \quad (6)$$

where κ_s is the thermal conductivity in the solid and \dot{q}_s is the heat source, in this case from fission. The boundary conditions used by MOOSE match the temperature boundary conditions in red shown in Figure 1. MOOSE is coupled to OpenMC by receiving the heat source in each element from the fission heating tally. During each iteration, MOOSE recomputes the temperature distribution from the heat source and boundary conditions, and then sends the temperatures back to OpenMC for the next transport solve.

3.3. Convergence Criteria

The MOOSE heat conduction solve used a Jacobian Free Newton Krylov (JFNK) iterative method, with 10^{-7} for absolute tolerance and 10^{-9} for relative tolerance. The Monte Carlo method is a stochastic method, so convergence is achieved by using an appropriate number of batches and histories per batch. The configurations used here were discussed in Section 3.1.

In terms of converging global iterations across all single physics, each case used 200 Picard iterations. To assist with convergence of Monte Carlo quantities, Robbins-Monro relaxation was applied to the flux and fission heating tallies. This updates the quantities of interest for the $n + 1$ th iteration as an average over the previous n Monte Carlo solutions [9]. So the flux at iteration n would be given by

$$\Phi_n = \frac{1}{n+1} \sum_{i=0}^n \phi_i, \quad (7)$$

where Φ_n is the relaxed flux at step n and ϕ_i is the flux output from the i th Monte Carlo solve. In order to assess the convergence of each physics, we plotted error versus mesh element size and compared numerical versus analytical solutions in each mesh element. These will be shown in Section 4.

In this benchmark, a high number of Picard iterations was used in order to ensure convergence, but other modelers may not want to incur the computational cost of running 200 Picard iterations. MOOSE provides steady-state detection as an alternate way of detecting convergence. By setting a steady-state tolerance, MOOSE will iterate until the solution norm changes by less than this tolerance. In Cardinal, OpenMC and NekRS variables are “external” to other MOOSE non-linear variables. These auxiliary variables may include some variables that should not be included in the steady-state detection (such as field variables storing the OpenMC cell ID, which does not change with iteration but are sometimes used for diagnostics). This can trick the steady-state detection into “detecting” convergence faster than it should. Thus, this feature was not adopted for this work, though a new issue was opened in the MOOSE repository to allow the user to control which auxiliary variables are used to assess convergence. As long as the user removes any auxiliary variables not used for coupling, they could rely on steady-state detection instead.

3.4. Data Mapping

Cardinal does not require the geometry models used in each single-physics sub-application to exactly match the meshes/geometry used in other coupled codes. At the beginning of every Cardinal simulation, a mapping between MOOSE’s mesh and other geometry representations is established. For a MOOSE-OpenMC coupling, the centroid of every MOOSE mesh element is mapped to a corresponding OpenMC cell using OpenMC’s find-cell routines. Cardinal then creates a tally for every user-requested type in each of the regions of OpenMC that a MOOSE centroid falls in. This mapping is used for transferring relevant quantities back and forth between Picard iterations.

For this simulation, the MOOSE geometry and OpenMC geometries were intentionally created to have the same dimensions so that the mapping is 1:1. Each region in the `MooseMesh` computes a temperature and sends it to a geometrically identical OpenMC region. This same region in an OpenMC solve computes a heat source via the fission heating tally and sends it back to MOOSE for the next conduction solve. Again, we want to emphasize that this strict 1:1 mapping is *not* mandatory in Cardinal, and rather a general centroid-based, non-conformal, many-to-one mapping from elements to cells can be used. In order to explore the convergence with spatial resolution, eight different cases were analyzed with $N = [5, 10, 25, 50, 100, 250, 500, 1000]$ spatial elements/cells in both MOOSE and OpenMC.

4. RESULTS

The results to compare with the benchmark are the temperature distribution, flux distribution, and k -eigenvalue. The benchmark provides analytical solutions for all of these quantities. An example temperature and flux distribution is shown in Figure 2 for the 50 mesh element case. In order to assess the accuracy of the numerical quantities and verify that refining the mesh increases the accuracy, the analytical

solution was used to compute an L_2 error norm for temperature and flux. The error, ε_T , is given by

$$\varepsilon_T = \frac{\|T_a - T_x\|_2}{\|T_a\|_2}, \quad (8)$$

where T_a is the analytical solution evaluated at the x -centroid of each mesh element and T_x is the temperature computed for that voxel from the multiphysics simulation. The same convergence test was used on the flux solution as well. Using the same notation convention, the flux error norm, ε_ϕ , is defined as

$$\varepsilon_\phi = \frac{\|\phi_a - \phi_x\|_2}{\|\phi_a\|_2}. \quad (9)$$

Figure 3 shows the temperature and flux error norms as a function of the number of mesh elements or cells.

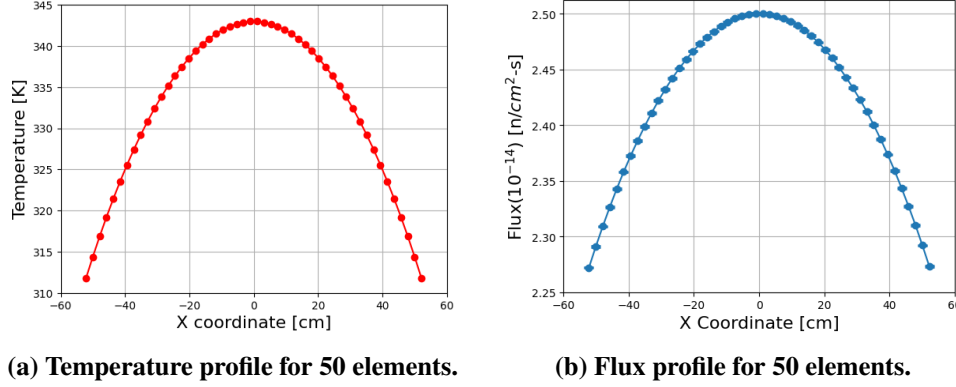


Figure 2: Numerical solutions for 50 mesh elements. The error bars show the relative error of the flux, which are nearly smaller than the circular marker sizes.

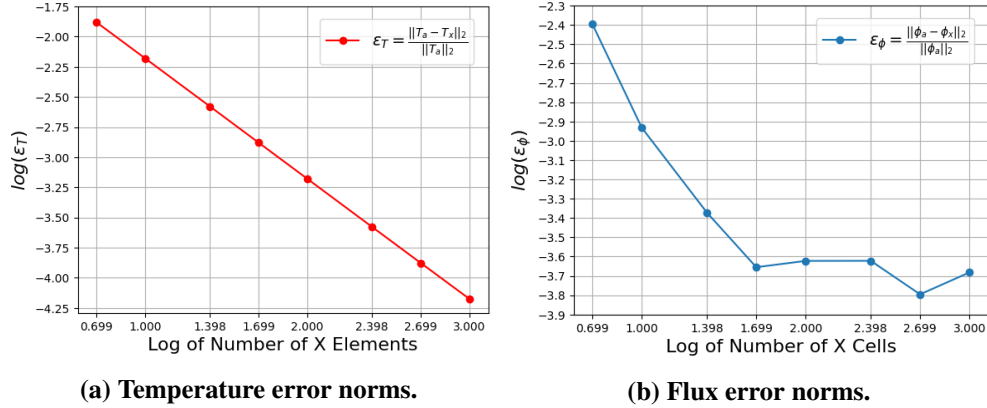


Figure 3: Error norms as a function of heat conduction mesh element count and OpenMC cell count, respectively.

For the temperature distributions, linear convergence is achieved, as the slope of the temperature line in Figure 3 is -0.9986 ($R^2 \approx 1.0$). The flux error does not appear to follow a linear pattern for the whole domain like the temperature error does. The first four points decrease as the number of cells used increases, but after that the errors do not continue to behave linearly. The slope of the first four points is -1.381 ($R^2 = 0.9814$). The flux error appears to stagnate after 50 cells, suggesting that the error will not improve considerably by increasing cell number because other error components dominate (e.g. stochastic noise, error from a cross section library at 50 temperatures, etc.). At this point, the cases will be categorically referred in two groups: coarse (5, 10, 25) and fine (50, 100, 250, 500, 1000).

While the L_2 error norm of the flux does not exhibit convergence in the same way that the temperature

does, the results do compare favorably with the analytical results throughout the domain and for all spatial resolutions in the fine group. The ratio between the computed quantity, C , and the expected answer, E , for the temperature in each element and flux in each cell are shown for all fine cases in Figure 4. A black line at $C/E = 1$ is marked in each plot indicating perfect agreement. Note the scales of the y -axes - the temperature is everywhere being predicted to within 0.006% (for the coarsest temperature mesh) and flux is everywhere being predicted to within 0.05%.

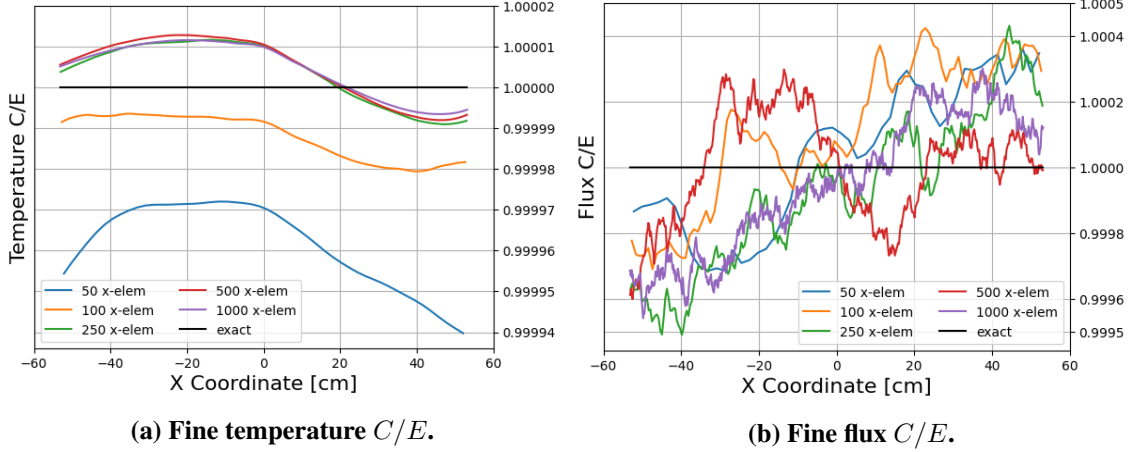


Figure 4: C/E for fine cases.

As the mesh refinement increases, the C/E trend towards to the ideal. For comparison, the coarser cases' C/E are included in Figure 5. The C/E errors are a few orders of magnitude larger than the fine cases, and a significant improvement in agreement can be seen between each coarse case.

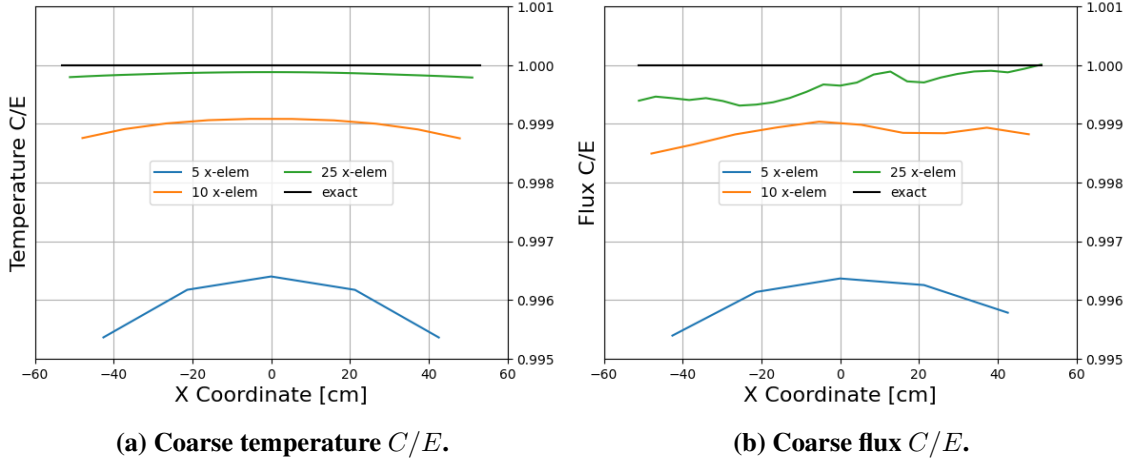


Figure 5: C/E for coarse cases.

The temperature C/E ratio is computed using FEM, but since the flux uses a stochastic approach, its C/E depends on two quantities with uncertainty. The flux from OpenMC $\hat{\phi}$ has uncertainty $\sigma_{\hat{\phi}}$ and the fission heating tally \hat{H} has uncertainty $\sigma_{\hat{H}}$. Writing the flux C/E explicitly gives

$$\begin{aligned}
 C/E &= \frac{P\hat{\phi}}{V_{cell}\hat{H}\phi_A} \\
 &= \frac{S\hat{\phi}}{V_{cell}\phi_A},
 \end{aligned} \tag{10}$$

where V_{cell} is the tallied region's volume and ϕ_A is the analytical flux. Propagating both errors:

$$\sigma_{C/E}^2 = \left(\frac{\partial C/E}{\partial \hat{\phi}} \right)^2 \sigma_{\hat{\phi}}^2 + \left(\frac{\partial C/E}{\partial \hat{H}} \right)^2 \sigma_{\hat{H}}^2. \quad (11)$$

Now denote ϕ , σ_ϕ , H , and σ_H as the quantities with physical units, i.e. the OpenMC outputs multiplied by the source strength and divided by volume. Evaluating the derivatives and simplifying gives

$$\sigma_{C/E}^2 = \left(\frac{\sigma_\phi}{\phi_A} \right)^2 + \left(\frac{\phi}{\phi_A} \frac{V_{cell}}{P} \sigma_H \right)^2. \quad (12)$$

Figure 6 shows individual C/E with 2σ error bars (95% confidence interval) for all the fine cases. Nearly all points are within 2σ , indicating that Cardinal is computing a correct flux distribution, within statistical uncertainty. Only a handful of points in the $N = 250$ case appear to be outside the error bars, but all other cases are fully contained. The coarse cases are excluded, as they are, expectedly, outside the 2σ error bars.

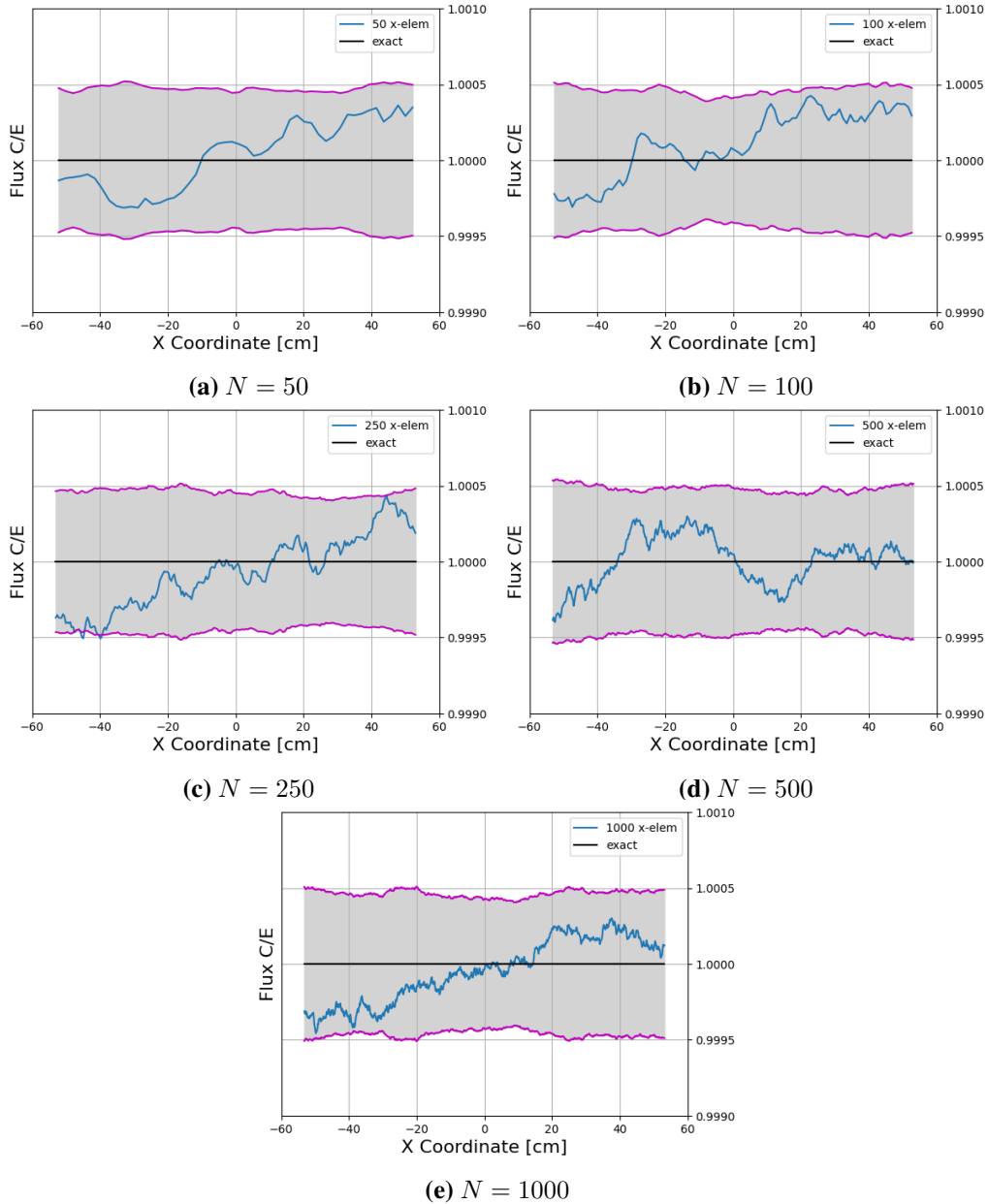


Figure 6: C/E in blue with two-sigma error bars (gray bounded by purple).

At this point, the observant reader may be wondering why the fine cases all appear to have about the same errors in C/E despite using smaller and smaller tally regions. In Monte Carlo, decreasing the volume of a tally region generally causes there to be greater relative error with the same number of histories. This seemingly counter-intuitive result is rooted in the benchmark itself. The slab domain length is $L = 106.47$ cm; however, the Mean Free Path (MFP) ranges between 239 cm for $T = 308$ K and 278 cm for $T = 358$ K. This means the average particle track will transport a particle from the source birth position to a vacuum boundary condition. Pair that with a spatially uniform birth distribution, and this essentially causes each cell to gain an equal number of contributions to its neighbors; since this happens in each case, the relative error in each case stays about constant.

The k -eigenvalue for each simulation is reported in Table 1. The results show close agreement with the analytical solution, as all fine cases agree within 4 pcm. The coarse cases range from 6 to 67 pcm difference. The eigenvalues generally agree better as more mesh elements are used. The only exception being that the $N = 100$ case agrees the best (less than 1 pcm difference from the analytical). The quality agreement is due to k_{eff} being a system-wide parameter which is less sensitive to spatial discretization of temperature feedback than the flux distribution itself.

Table 1: Eigenvalue with uncertainty for each mesh size, and the difference from the analytical.

Resolution	k_{eff}	(numerical - analytical) [pcm]
analytical	0.29557	-
5	0.29624 ± 0.00003	67 ± 3
10	0.29581 ± 0.00004	24 ± 4
25	0.29563 ± 0.00004	6 ± 4
50	0.29553 ± 0.00004	-4 ± 4
100	0.29557 ± 0.00003	0 ± 3
250	0.29561 ± 0.00004	4 ± 4
500	0.29561 ± 0.00004	4 ± 4
1000	0.29558 ± 0.00004	1 ± 4

5. CONCLUSIONS

Overall, the numerical results show agreement with the analytical solutions. The convergence is first order for the temperature and the C/E for the fluxes contain the analytical solution for nearly every point within two-sigma across all fine cases. The eigenvalue also agrees very well for all cases.

While experimental benchmarking is critical for nuclear M&S, agreement with analytical benchmarks also serves an important purpose by increasing confidence in multiphysics coupling. Though typical industry-grade simulations would not run S_2 transport, this modification allows Cardinal to be compared against a theoretical problem. A companion paper in this conference with Cardinal [6] couples OpenMC Monte Carlo transport to NekRS heat conduction. In future work, we will extend our computational model to include NekRS providing the heat conduction solution.

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