Outline

Validation of Nucldar Data using MC Codes Using MC codes to valdiate codes & transport data

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- Overview
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- 2 Script

- Script
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- Script analysis
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Aims

The purpose of this work is to determine how different MC codes treat the nuclear data they use

- Determine response of each MC code to nuclear data on the same geometry and data (ENDF/B-VIIR1)
- 2 Each code using the same data, i.e. the same ace file
- Oetermine the reason for any discrepancy, return reason to evalator/data producer/code developer
- I downloaded the data only 4 days ago...

Script

Outline

The system is setup to make our lives as easy as possible, breaking the system into three main parts

- benchmark builds and runs input deck for each code
- analysis creates a neutron spectra plot for each nuclide
- build_report writes tex file containing each plot

- Script scans mcnp xsdir file and converts to other codes formats (Serpent/OpenMC)
- creates an input deck for each code for each nuclide
- submits each job to server
- wait.... for about 3 days

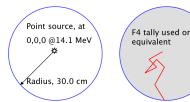
Script - analysis

- For each nuclides in the problem loops over the output data and produces spectra plots
- Produces meta-deta for each nuclide, like fraction of flux out of acceptable range
- Runs the R statistical analysis package to produce graphs

Script - build_report

- Builds LATEX document contains full and zoomed plots
- Includes tables of total flux ratios and statistical errors





Spectra binned in 616 TART format, the volume of the cell set to 1.0 cm3 and the density is set to 1.0 g/cm3

- Simulations ran for 100 Million histories
- Tracklength estimator in the sphere

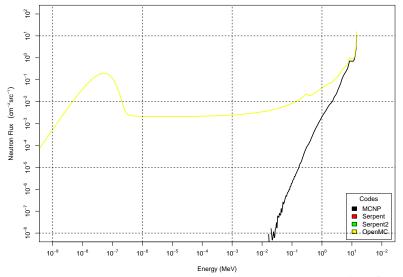
- MCNP5 v1.6
- Serpent 1 & 2
- OpenMC
- No $S(\alpha,\beta)$

Firstly

- I have no agenda, other than I want to see good agreement
- I am not claiming any one code is correct
- Unless I have good reason for doing so..



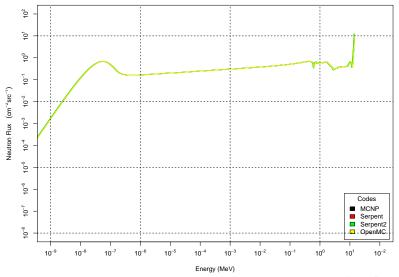
Wrong - ⁷Be



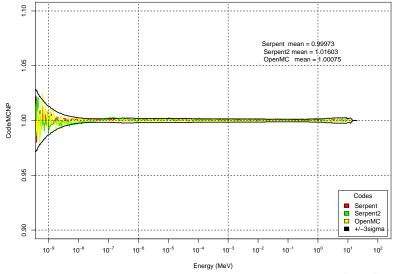
Wrong - ⁷Be

- In this case, dah dah dah..... it looks like its MCNP
- Serpent 1 & 2 both failed to read the data
- ACE File Gen @293.6K for n-004_Be_007 (R. Arcilla, NNDC)
- MCNP no warnings
- Serpent Warning: Nuclide is missing (n,gamma) channel
- Also thinks the data are truncated at 8.1 MeV

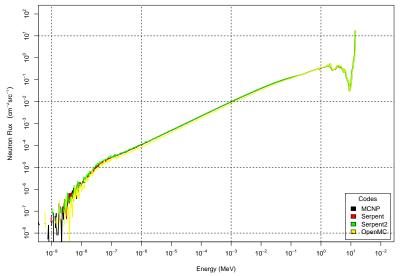
Right - ⁹Be



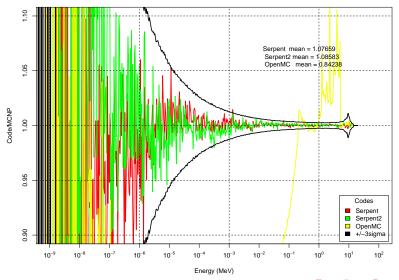
Right - ⁹Be



Right - ¹⁵N



Right - ¹⁵N



Wrong - ¹⁵N

- OpenMC ??
- Seen previously with other development codes, issue with the anisotropy of inelastic scattering

Wrong - ²²Na

Outline

Serpent Processing XS data:

Isotope 11022.71c (Na-22)...

Warning: 59 negative ures ptable values.

OpenMC ERROR STOP

WARNING: Negative value(s) found on probability

table for nuclide 11022.71c

ERROR: Too many rejections on evaporation

spectrum.

Batch: 1

Generation: 0

- MCNP ran to completion
- ENDF/B-VII.1: ACE File Gen @293.6K for n-011_Na_022 (R. Arcilla, NNDC) mat1122 04/20/12 probability tables used from 1.5000E-02 to 1.0000E-01 mev.



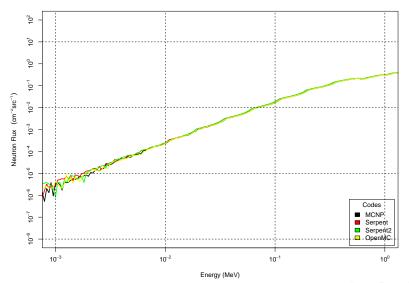
Wrong - ²³Na

• OpenMC ERROR: Too many rejections on evaporation spectrum.

Batch: 1

Generation: 0

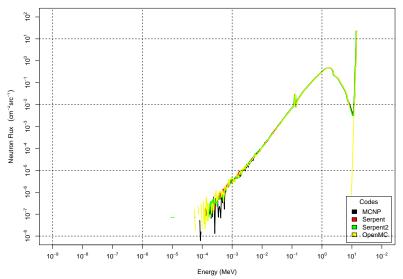
MCNP, and the Serpents ran to completion



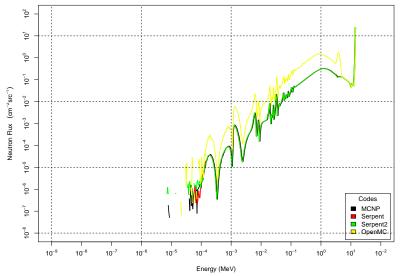
Wrong - ³⁶Ar

Negative Ures again

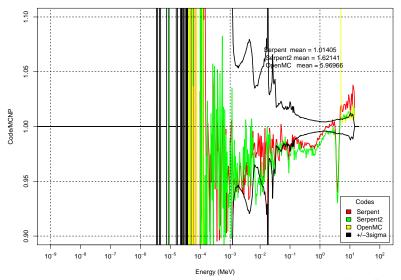
Weird - ³⁸Ar



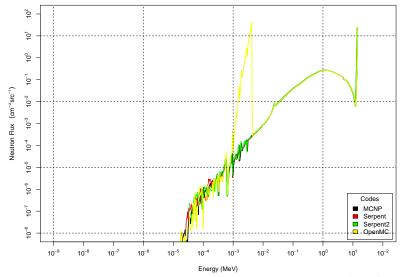
Weird - ⁵⁵Mn



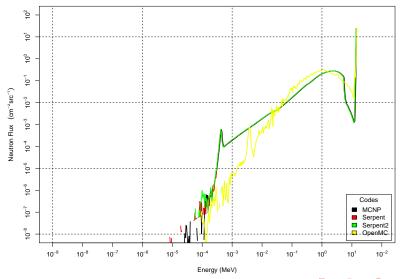
Weird - ⁵⁵Mn



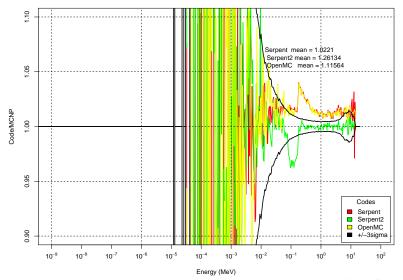
Weird - ⁵⁸Co



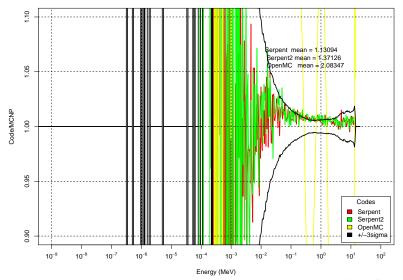
Weirder - ⁵⁹Co



Wrong - ⁷⁴As



Poor - ¹⁴⁹Pm



Poor Misc

- 44,46,48 Ca OpenMC overestimate
- Sc, Ti, V, Cr, Fe Agreement within appropriate range
- ⁵⁹Ni made OpenMC Segfault, others were ok
- If nuclide not mentioned its ok

Conclusions

- Total fluxes agree well and if had been used without spectra the differences would not have been noticed
- Most nuclides agree well, however several results were suspect
- Several bizarre misinterpretations of the data, ¹⁵N

Extensions to this work

- Would like to be able to process my own data with NJOY
- Would like to be able to convert my NJOY processed data to Partisn format
- Would like to add PARTISN to the mix
- Would like to add Geant4 and Phits



Any questions?