

The Consortium for Enabling Technologies and Innovation

Virtual Summer Meeting for Young Researchers

INTRODUCTORY BAYESIAN APPROACH TO GAMMA SPECTRA ANALYSIS FOR ISOTOPIC IDENTIFICATION

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Disclaimer: This work was done as part of an AI for Engineering workshop and was not funded by ETI.

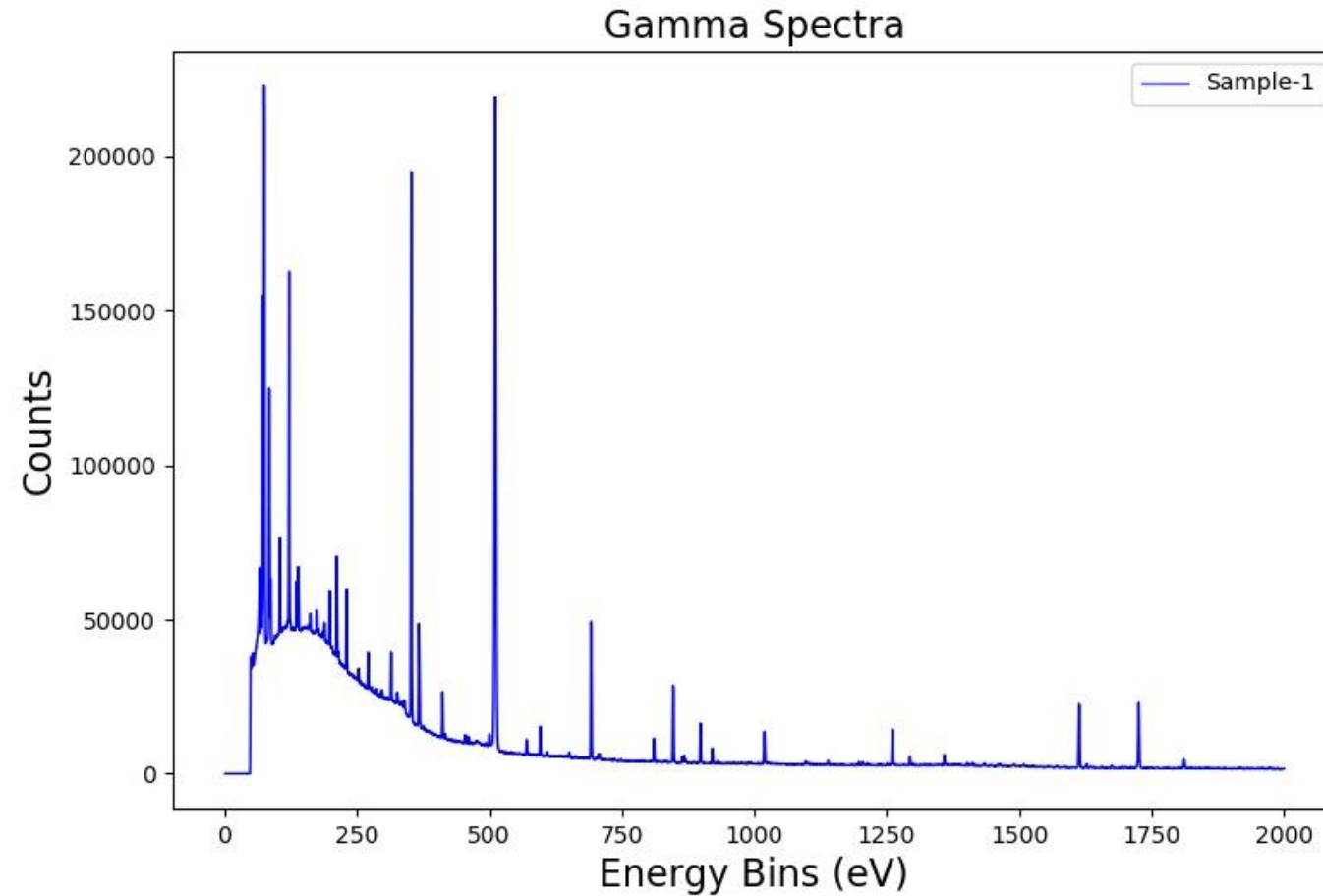
What is PGAA?

- Concept
- Challenges

Step 1:

Establish Background

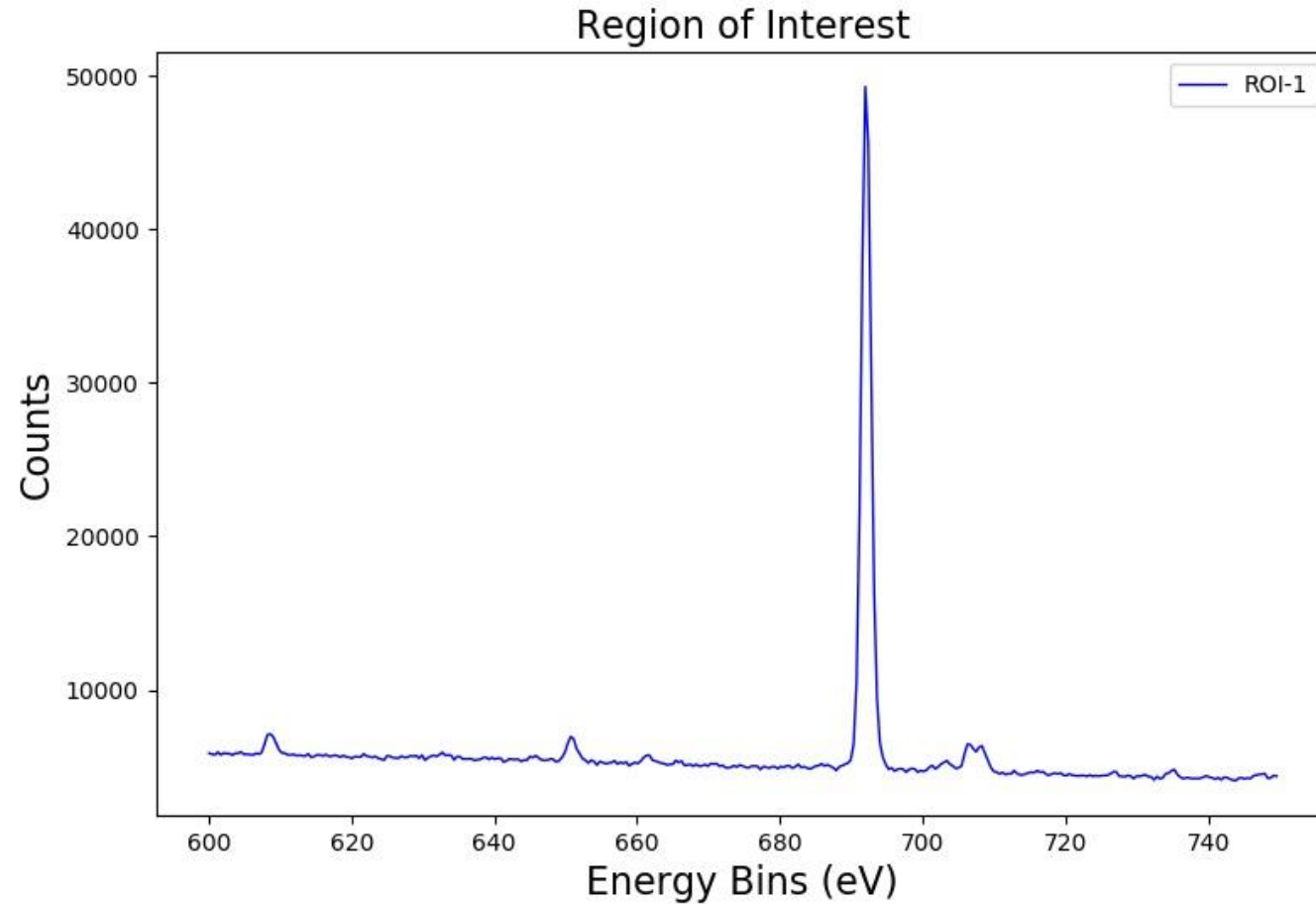
- Loading and parsing dataset



Step 2:

Establish ROI

- Defining an energy window around a Region of Interest (ROI)
- Plot showing



Step 2a:

Materials

PGAA Database

- Material selection
- Materials present
- List of most likely cross-section and material

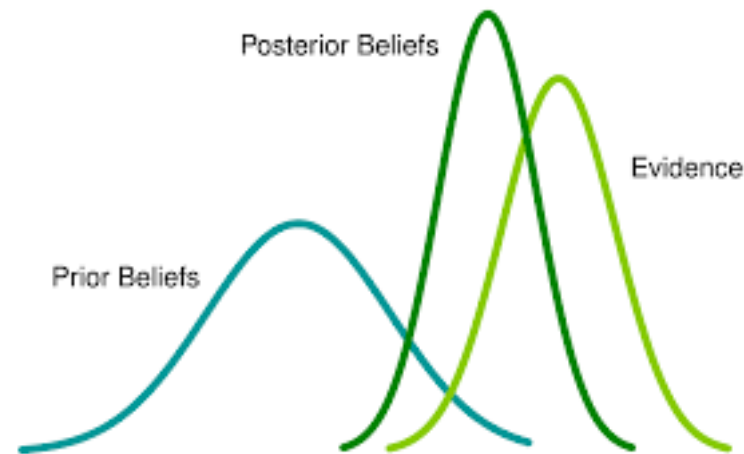
	A	isotope	Z	T	Gamma Energy	Gamma Energy Value	Sigma	Sigma Value	Uncertainty	Knaught	Knaught value	Knaught unc
0	1	2-H	1	p	2223.24835(9)	2223.248	0.3326(7)	3.326000e-01	7.000000e-04	1.0000(21)	1.000000e+00	2.100000e-03
1	2	3-H	1	p	6250.243(3)	6250.243	0.000519(7)	5.190000e-04	7.000000e-06	0.001560(21)	1.560000e-03	2.100000e-05
2	3	4-He	2	p	20520.5	20520.460	4.2E-11(12)	4.200000e-11	1.200000e-11	3.2E-11(9)	3.180000e-11	9.090000e-12
3	6	7-Li	3	p	477.595(3)	477.595	0.00153(8)	1.530000e-03	8.000000e-05	0.00067(4)	6.680000e-04	3.490000e-05
4	6	7-Li	3	p	6768.81(4)	6768.810	0.00151(9)	1.510000e-03	9.000000e-05	0.00066(4)	6.590000e-04	3.930000e-05

	A	isotope	Z	T	Gamma Energy	Gamma Energy Value	Sigma	Sigma Value	Uncertainty	Knaught	Knaught value	Knaught unc
31611	238	239-U	92	p	4118.54(5)	4118.54	0.00148(15)	0.00148	0.00015	1.88E-05(19)	1.880000e-05	1.910000e-06
31612	238	239-U	92	p	4612.40(5)	4612.40	0.0031(3)	0.00310	0.00030	3.9E-05(4)	3.950000e-05	3.820000e-06
31613	238	239-U	92	p	4660.62(5)	4660.62	0.0034(3)	0.00340	0.00030	4.3E-05(4)	4.330000e-05	3.820000e-06
31614	238	239-U	92	p	4672.59(5)	4672.59	0.00116(13)	0.00116	0.00013	1.48E-05(17)	1.480000e-05	1.660000e-06
31615	238	239-U	92	p	4806.38(5)	4806.38	7E-05(7)	0.00007	0.00007	9E-07(9)	8.910000e-07	8.910000e-07

Step 3: Bayesian Statistics

$$P(A | B) = \frac{P(B|A)P(A)}{P(B)}$$

$$\text{Posterior} = \frac{\text{Likelihood} * \text{Prior}}{\text{Marginal}}$$



Step 4: Define Priors

- A prior tells the code that something will always lie within a certain range or at a specific number
- Potential automation— have code “learn” priors

```
##### Means #####
if N == 1:

    pass

else:

    for i in range(N-1):

        if means[i] > means[i+1]:

            return -np.inf

    for i in range(N):

        if means[i] > E[-1] or means[i] < E[0]:

            return -np.inf
```


Step 5: Markov Chain Monte Carlo

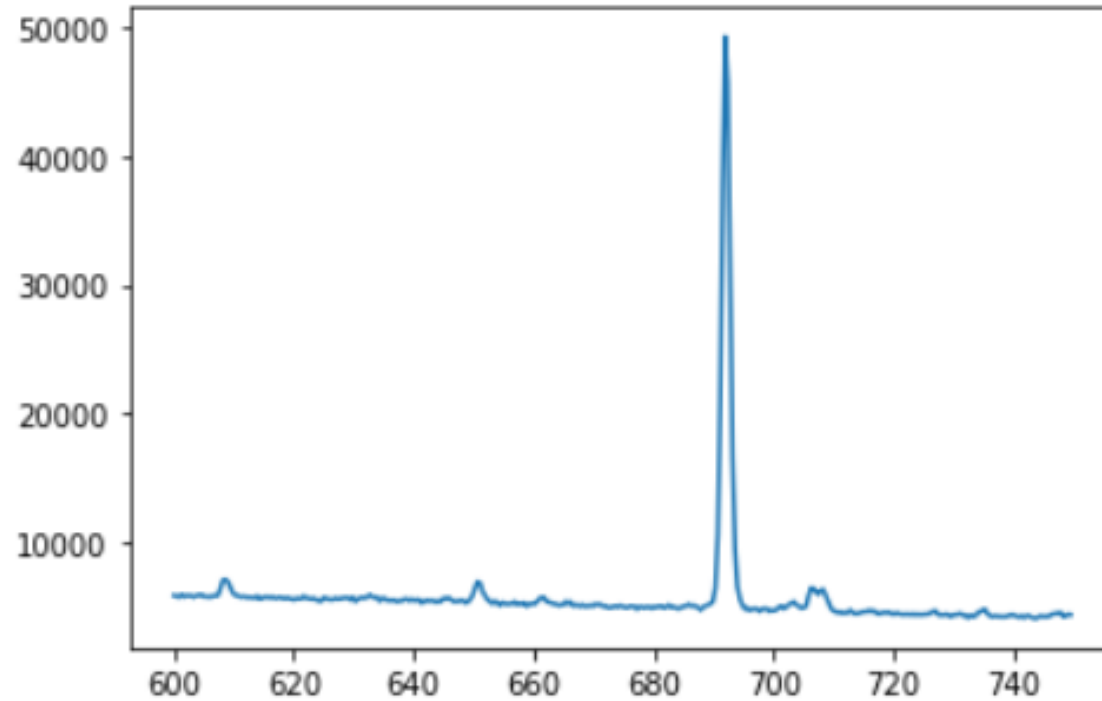
- Primarily used for numerical approximations of multi-dimensional integrals
- Create arbitrary starting points for “walkers” a sufficient distance from each other
 - “walkers” spend most time in regions of higher probability
 - Produce a cluster of points where the posterior probability is largest
- These points are used to compute expected values efficiently

Step 5:

Initialization

- Slope
- Background bias
- Peak location (mean)
- Amplitudes
- Peak widths
- Background noise

Then setting 150 Walkers to each take 1000 steps



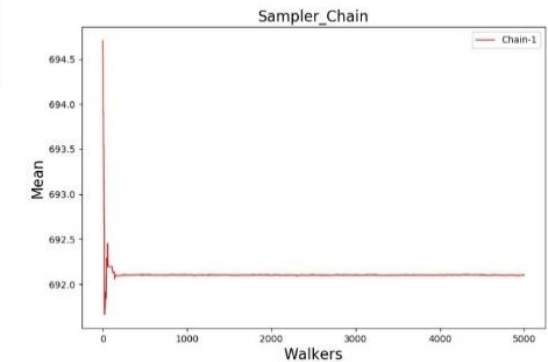
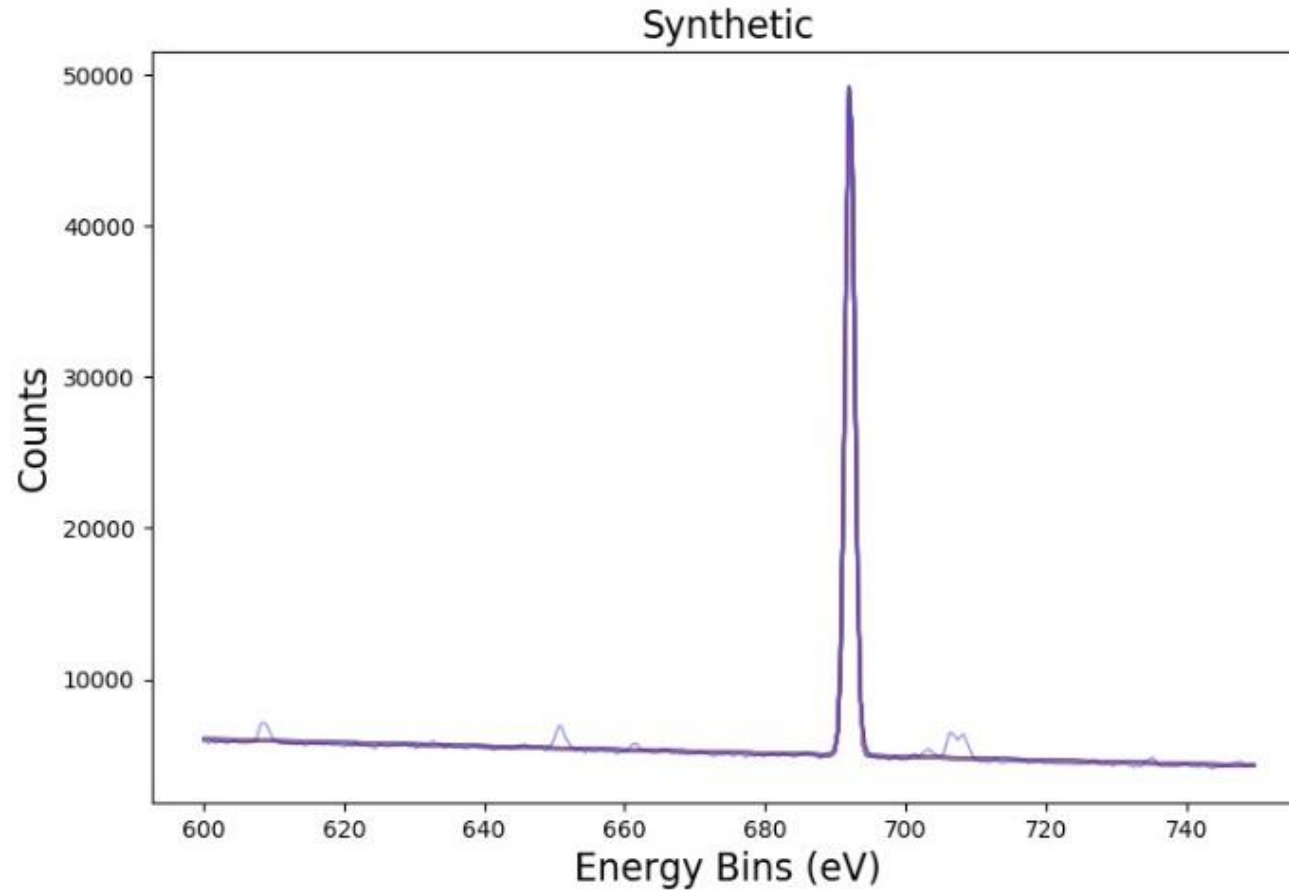
```

What value should I use for mean?1 (done to quit) 692
What value should I use for mean?2 (done to quit) done
Number of Parameters:
6

```

Step 5:

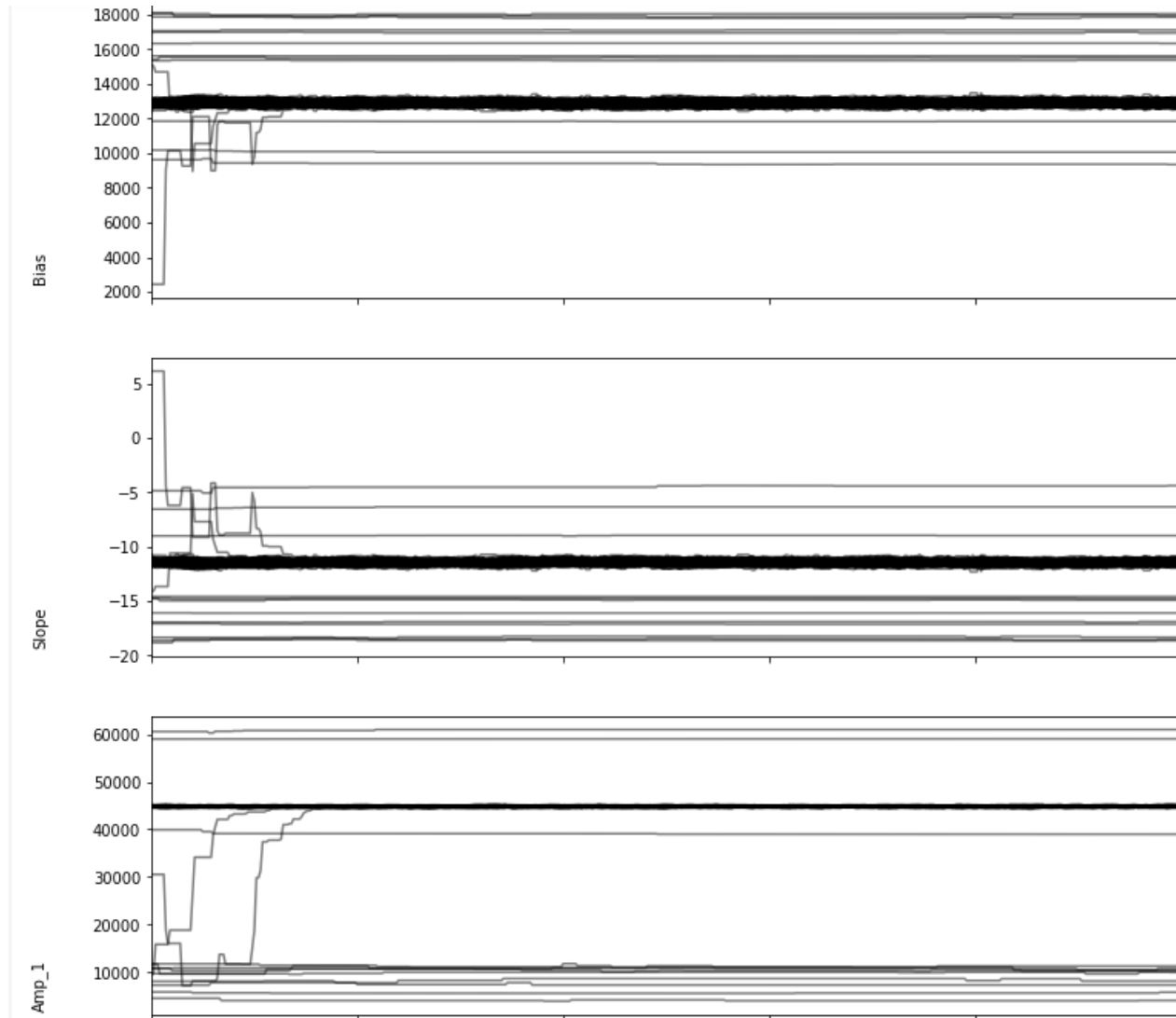
- Overlay of original plot and a “Synthetic” consisting of 10 walkers' paths
- The goal is for the walkers to fit the original path perfectly purely based on the priors



Step 6:

Sanity Check/Distribution

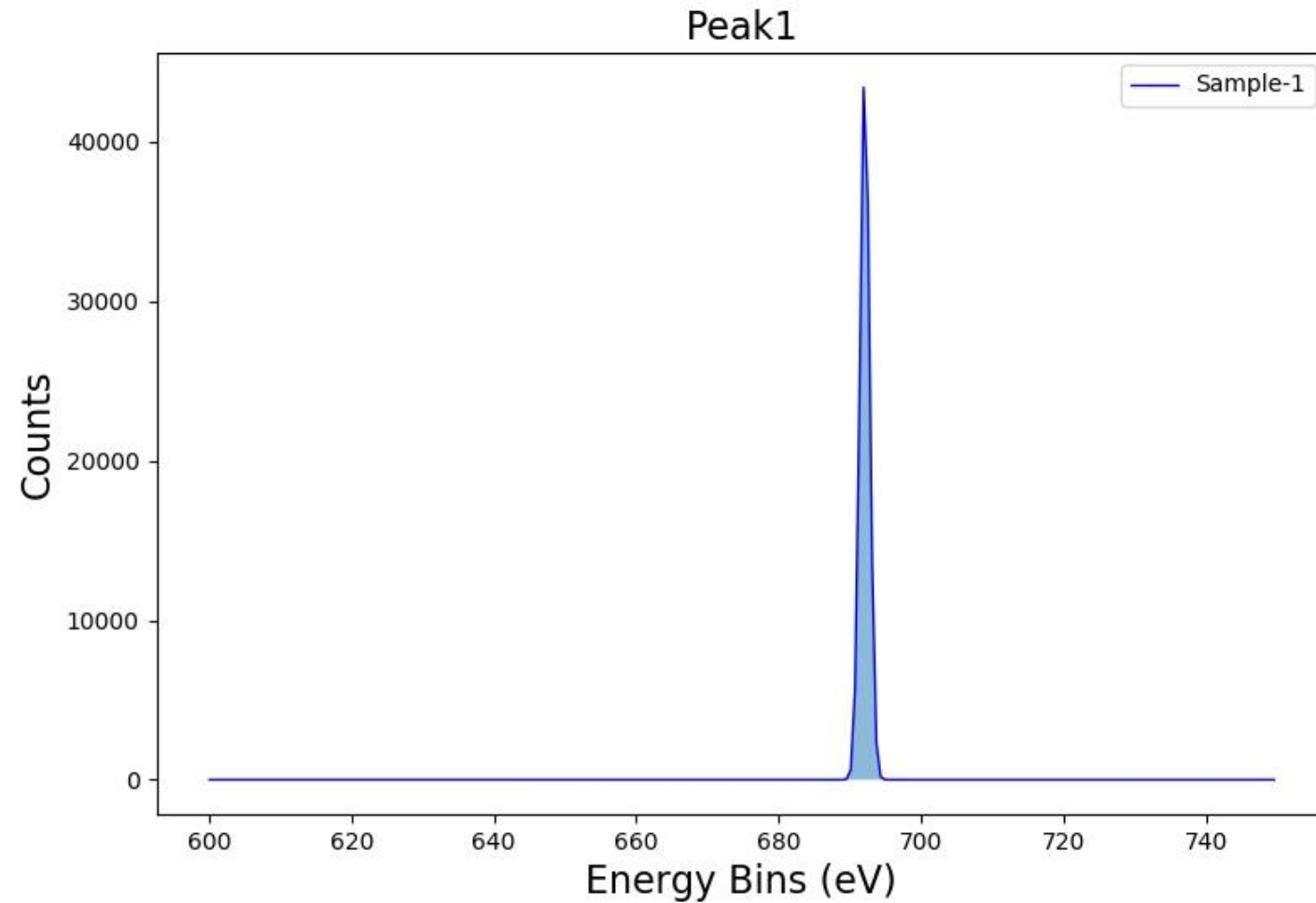
- Darker the lines mean higher cluster of walkers.
- All other lines represent paths a walker takes.
- There should be a visible trend that the walkers are trying to merge to a singular point



Step 7:

Plotting

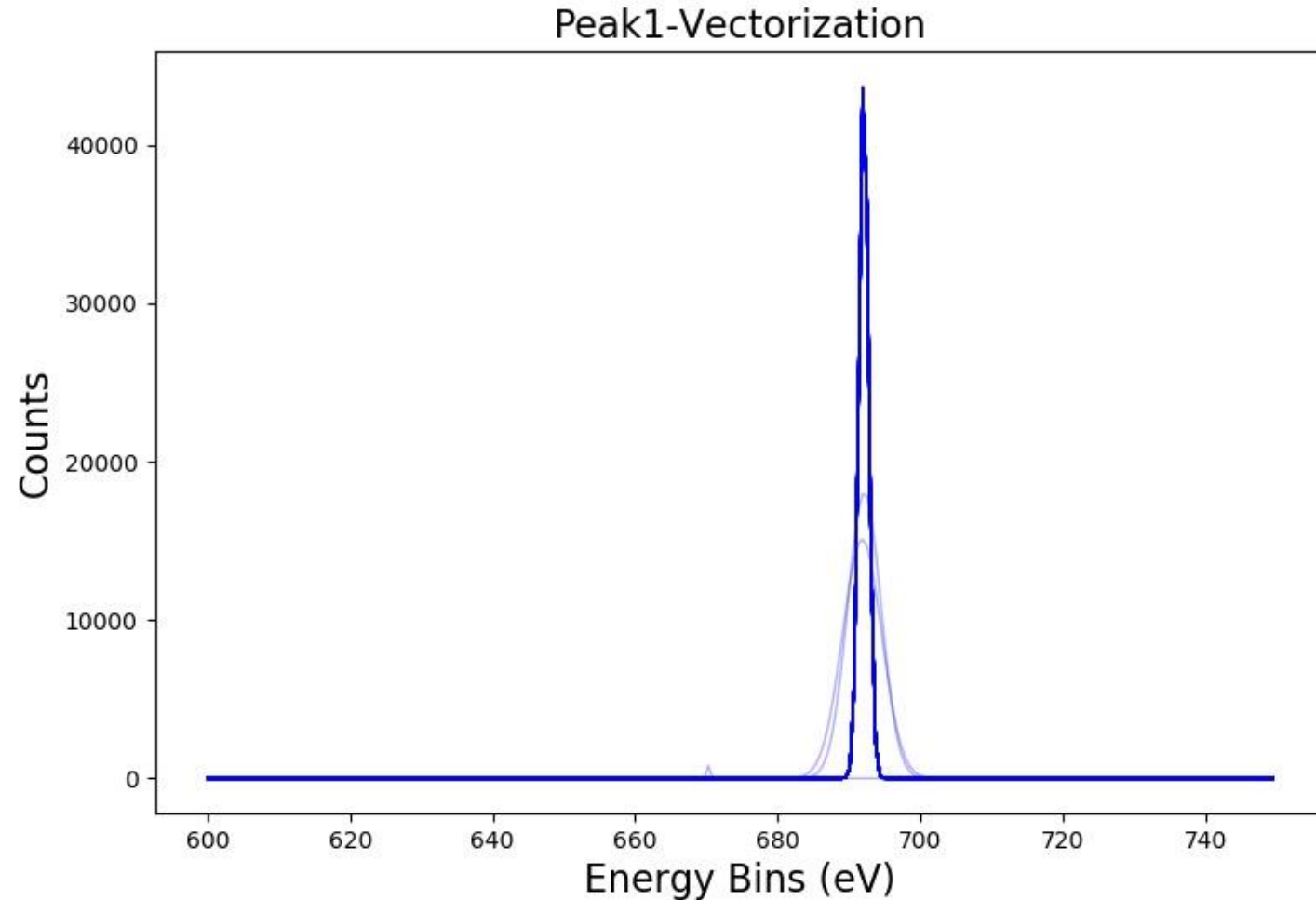
- This is now a fit of the peak of interest.



Step 7:

Vectorization

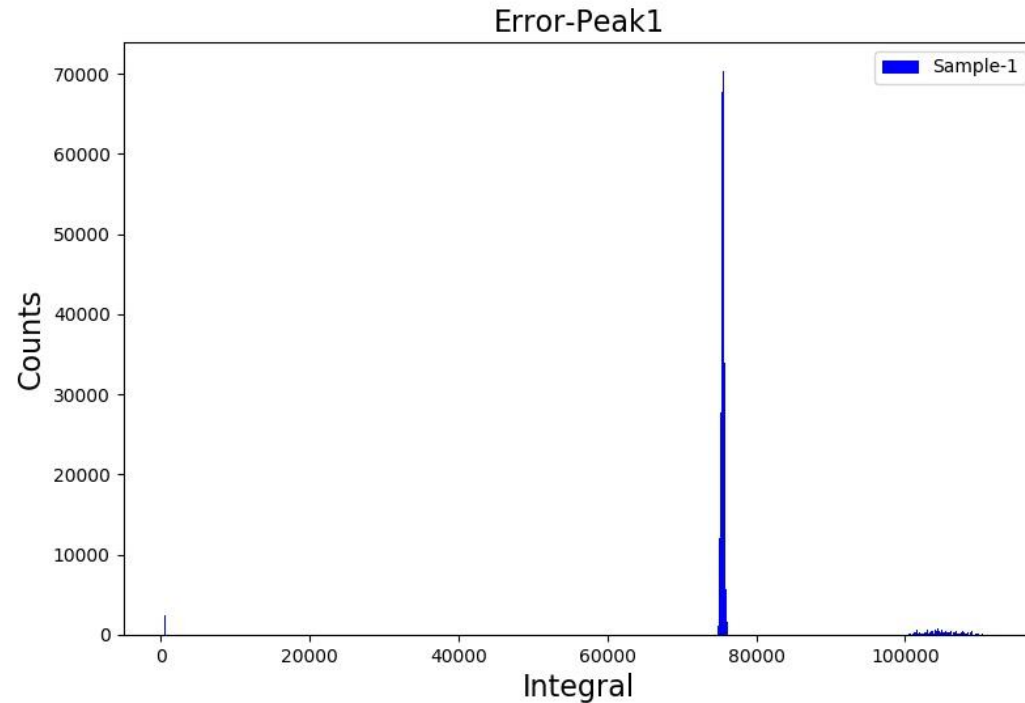
- This plot depicts a gradient for which walkers are fitting the peak, the darker lines meaning more walkers on that path, lighter meaning less. Fewer lines meaning less variation



Step 7:

Error

- Y-axis, number of samples saying this is the right integral.
- X-axis, area under the curve.
- Then computing the error from our bounds we get great results.



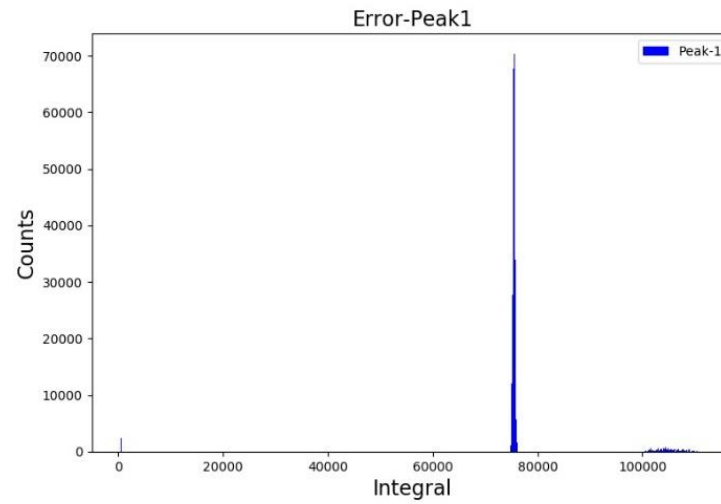
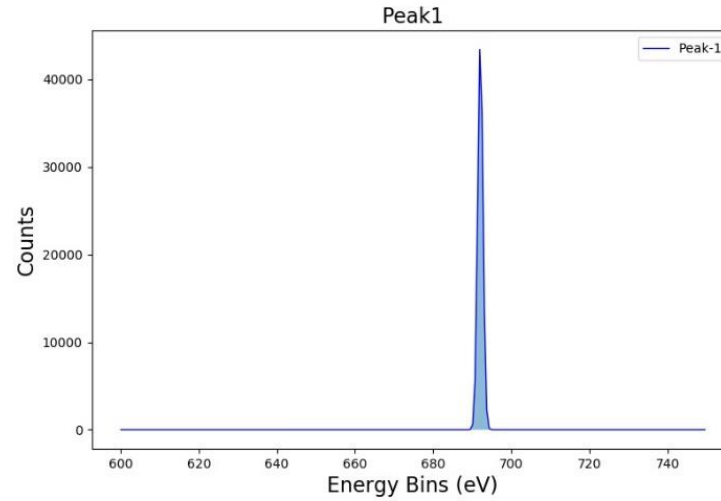
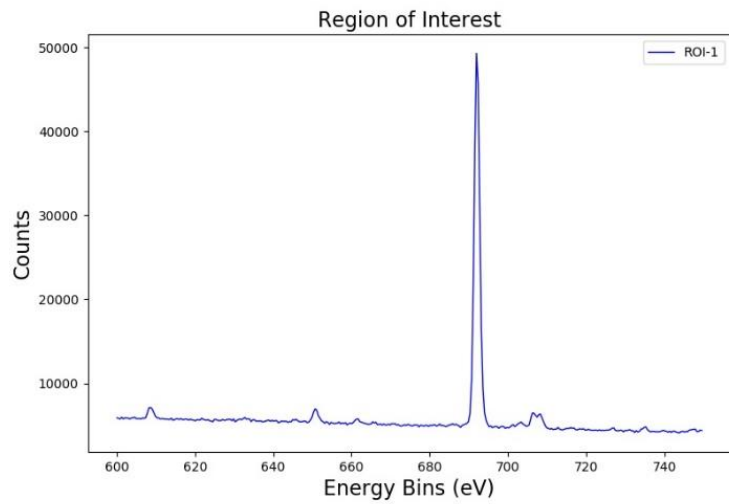
```
print('error in P1 lower:', (((fp-sp)+fp)-fp)/fp)*100)
print('error in P1 upper:', (((ep-fp)+fp)-fp)/fp)*100)
print()
# print('error in P2 lower:', (((fp1-sp1)+fp1)-fp1)/fp1)*100)
# print('error in P2 upper:', (((ep1-fp1)+fp1)-fp1)/fp1)*100)
```

```
error in P1 lower: 0.2869768750046384
error in P1 upper: 0.30498320321024125
```

Applications of Bayesian Approach

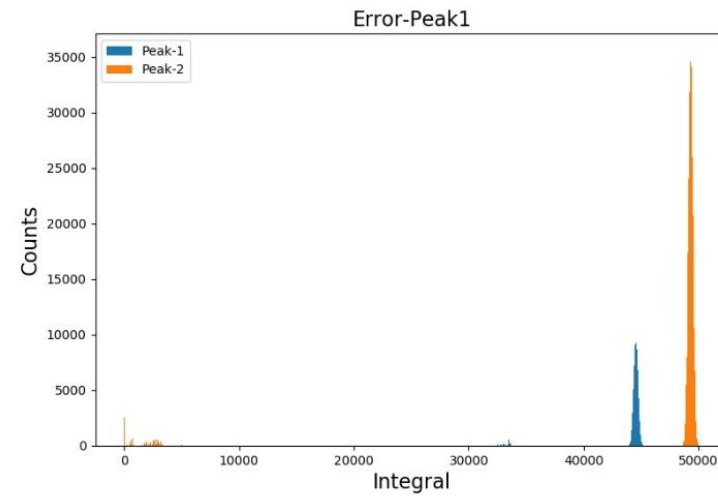
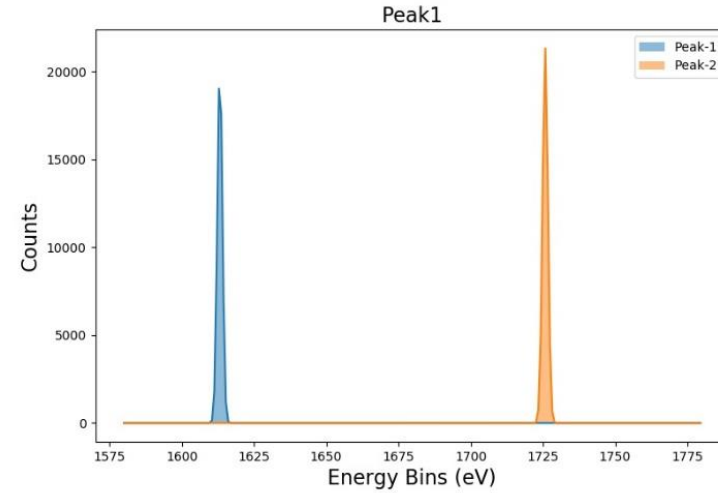
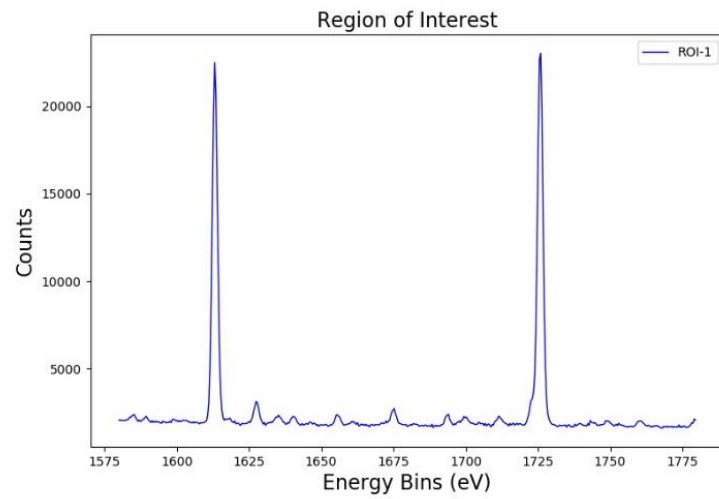
- Primarily used for numerical approximations of multi-dimensional integrals
 - Single, multi, overlapping peaks
 - Custom solutions possible
- Limitations and sources of error

Single Peak



error in P1 lower: 0.28697
error in P1 upper: 0.30498

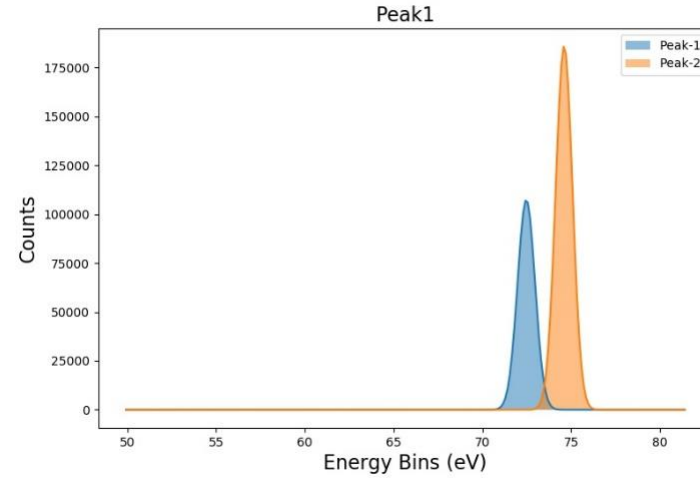
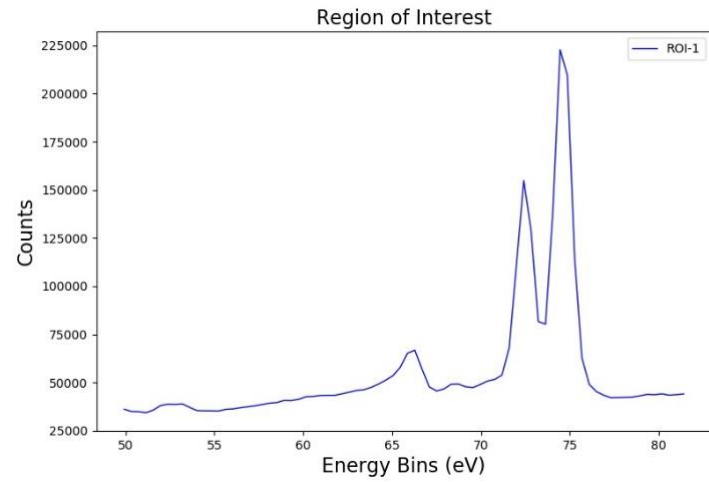
Multi Peak



error in P1 lower: 0.43563:
error in P1 upper: 0.42188:

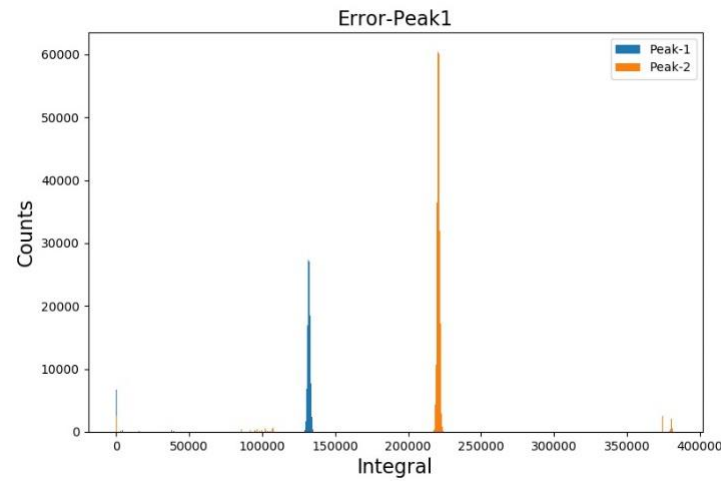
error in P2 lower: 0.40227:
error in P2 upper: 0.40272:

Overlapping Peaks



error in P1 lower: 0.71501
 error in P1 upper: 0.89306

error in P2 lower: 0.54862
 error in P2 upper: 0.42021



Notes & Next Steps

- What can we do to help predict the shape of ugly peaks?
- Data Sampling: provide hundreds of previous spectra to help code “learn”
- Define priors in a generalized way

Conclusions

- Decrease in error
- Assists reading of spectra
- Flexible approach
- Beneficial to PGAA field

Thank you! Questions?

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