

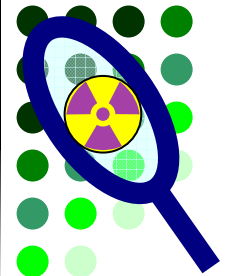
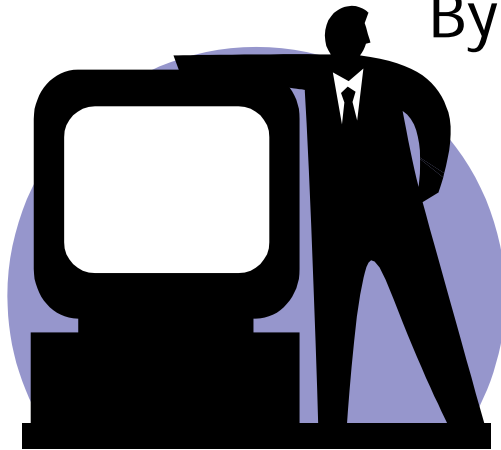


Spectral Analysis Options

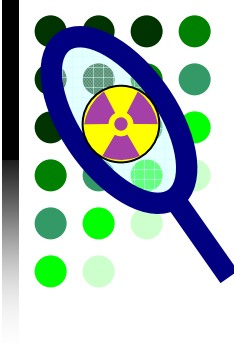
Radio Nuclide Identification for Mindless
Automatons

By Karl Einar Nelson, PhD

UCRL-PRES-226143



Goals

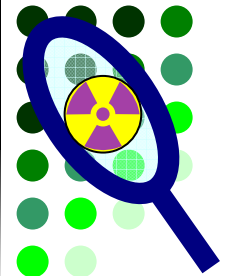


- Present the challenges in automated nuclide identification
- Define common terms required to discuss identification algorithms
- Demonstrate the need for Metrics and Test Benches

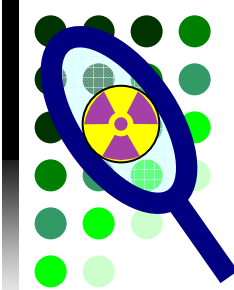


Problem Statement

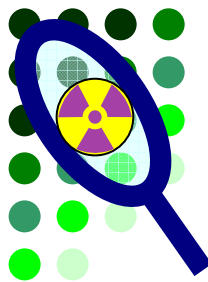
What challenges we face.



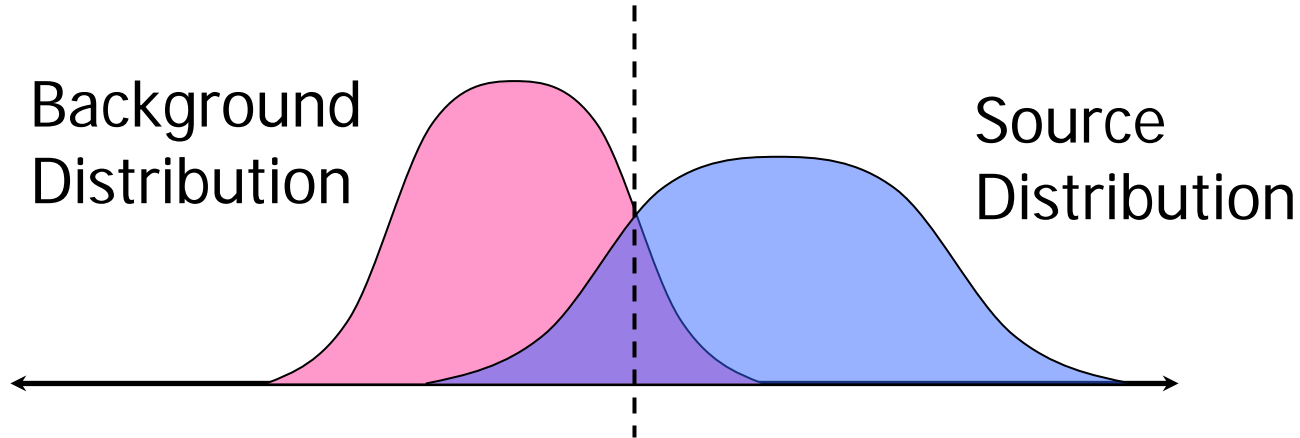
What is the value of nuclide identification?



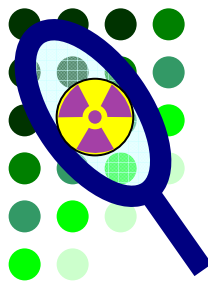
- The goal of any protection system is to recommend a course of action to maximize the probability that a threat will be caught.
 - Minimize unnecessary searches of non-threatening material
 - Raising response appropriately on potentially threatening sources
- The limiting resource is the number of secondary inspections
- The brightest sources are either RDD potentials or medical sources
- Nuclide Identification is simply a means to achieve our ends



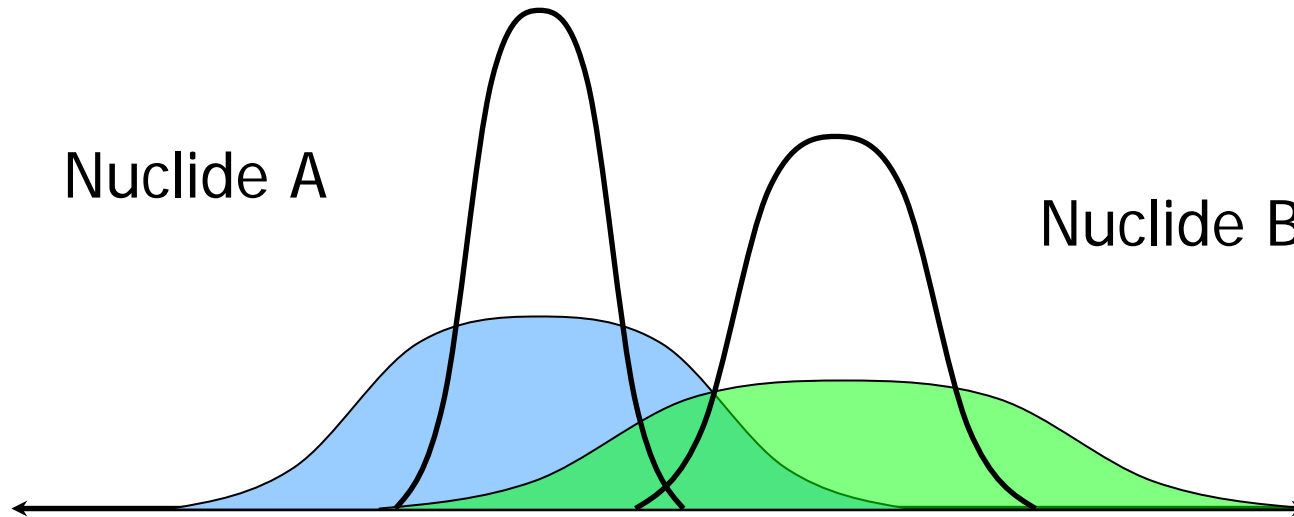
Errors



- False Negative
 - Incorrectly omitting a nuclide that is present
 - Allows a threat to pass
- False Positive
 - Incorrectly reporting a nuclide that is not present
 - Results in an unnecessary search
- Errors
 - Result from overlap in statistical distributions
 - Can be minimized but never eliminated

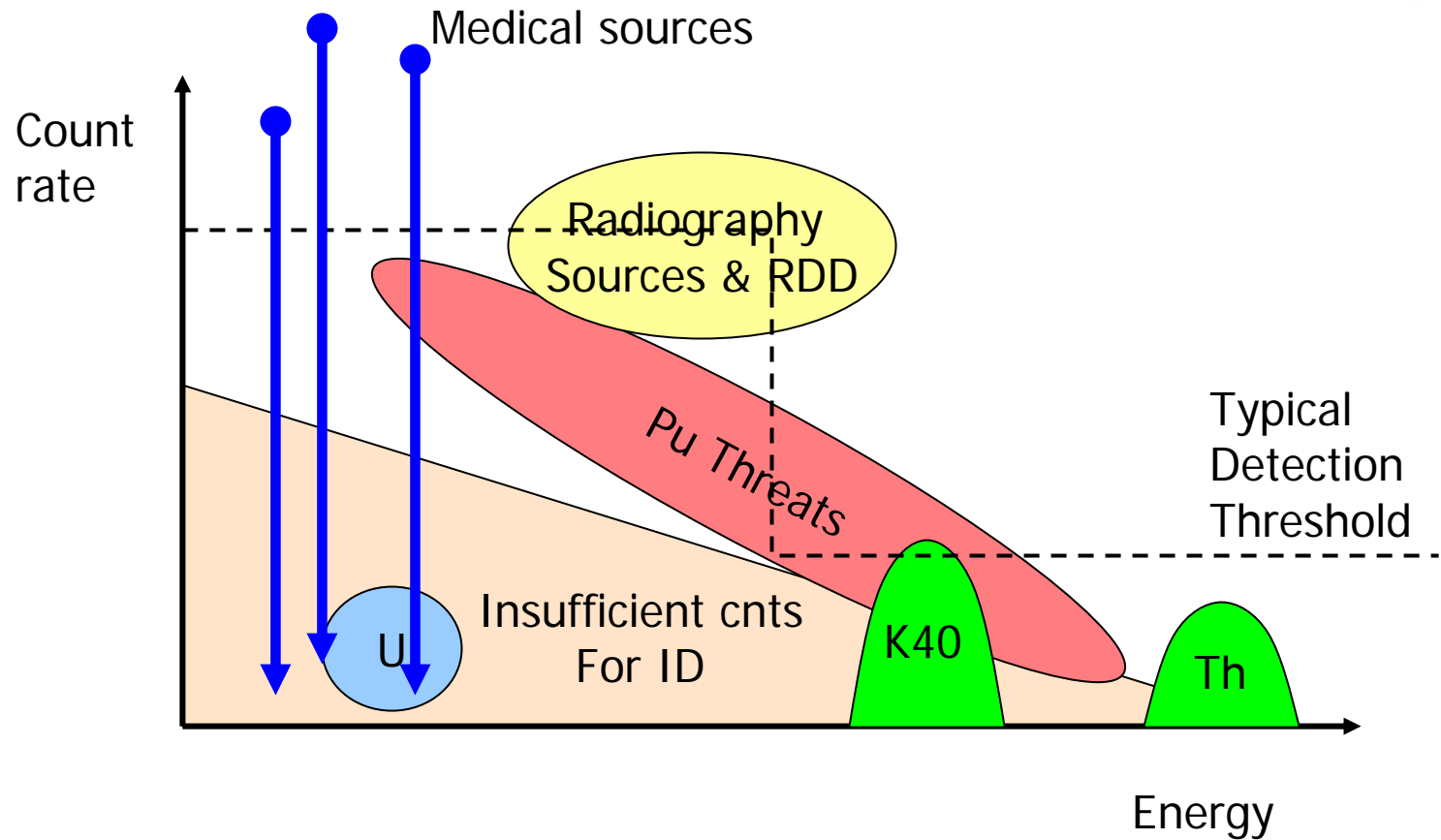
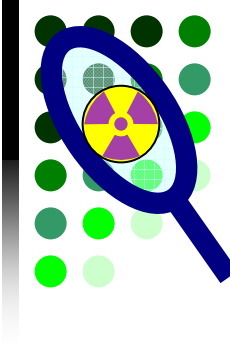


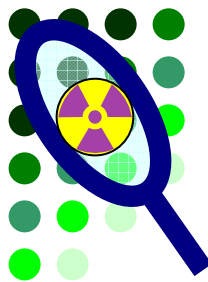
Ambiguity



- Ambiguity – an observation can be reasonably interpreted in more than one way.
 - As the source counts are reduced distributions grow wider
 - Nuclides become more difficult to separate
 - Numbers of errors increase

Domain Source Sample Space





Risk Categories

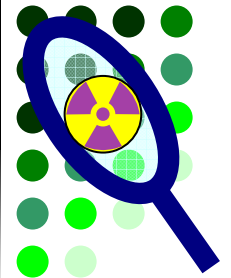
HIGH Must do secondary inspection	MEDIUM	LOW May avoid secondary inspection
<ul style="list-style-type: none">•High confidence threatening ID•Highly shielded ID (RDD)•High count rate, no ID•Suspect Mixture	<ul style="list-style-type: none">•Ambiguous ID with more counts than expected from background	<ul style="list-style-type: none">•Apparent Background•High confidence non-threatening sources

Improvements in spectral id => Fewer errors + Minimize ambiguous cases
=> Reduced number of unnecessary secondary inspections

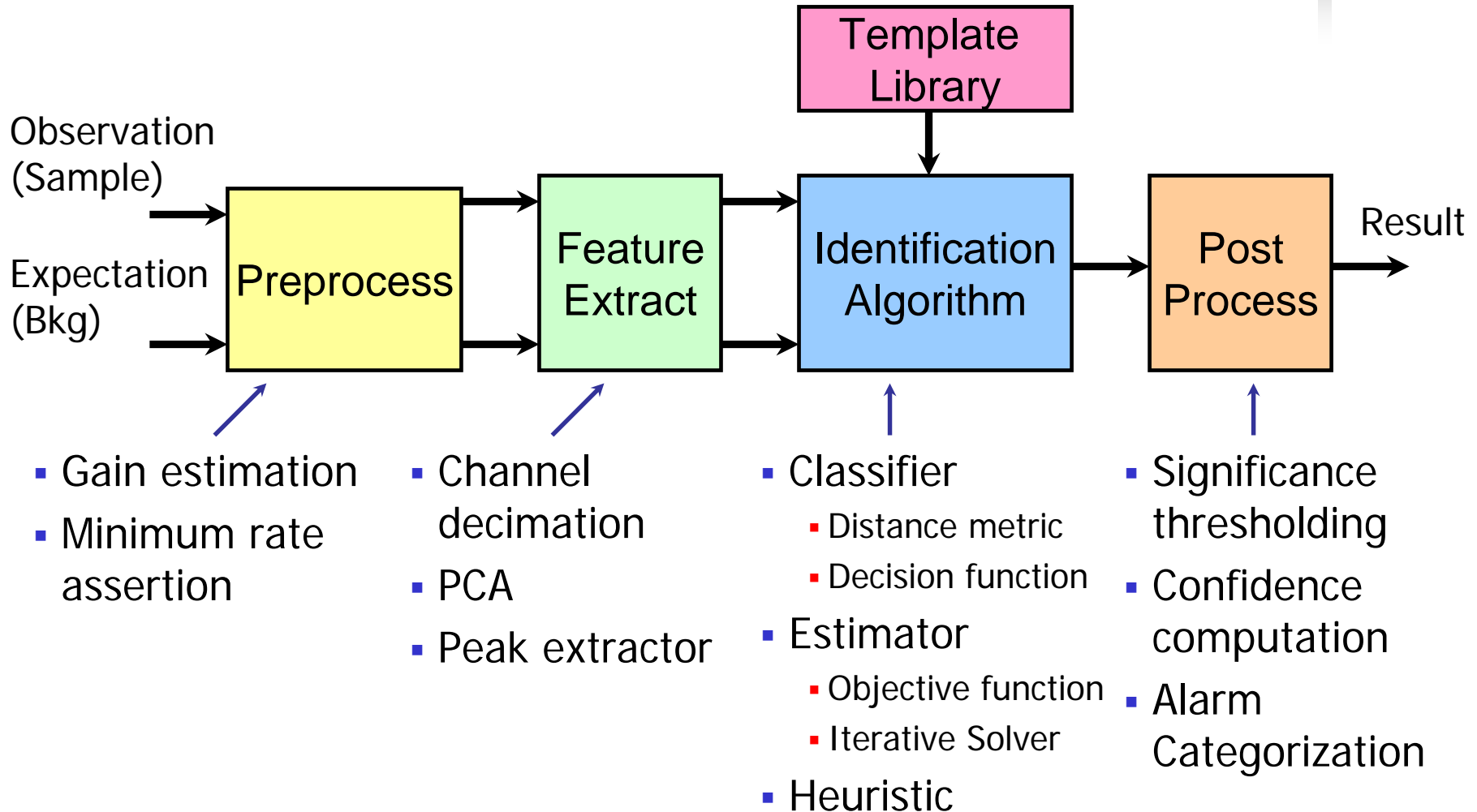
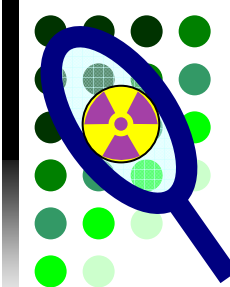


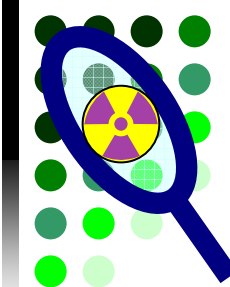
Identification Components

What makes mindless automatons tick?



Anatomy of Nuclide Identifier





Template Algorithm Types

CLASSIFIERS

- Asks
What template best fits the observation?
- Properties
 - Produces a list of templates ranked by fitness to the observation
 - Template library must fully span input domain, including naturally occurring mixtures

ESTIMATORS

- Asks
What weighted sum of templates best represents the observation?
- Properties
 - Produces a weighting vector
 - Mixture can span into novel and unanticipated situations
 - Best solution may not represent the true solution (No ranking)

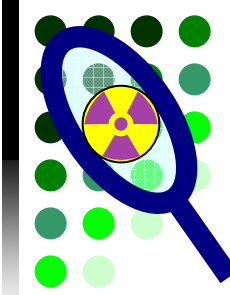
CLASSIFIER

- Has Ranking
- Limited Solutions

ESTIMATOR

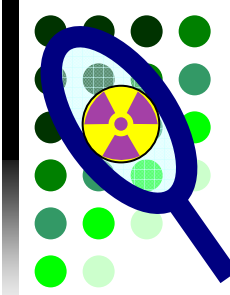
- Lacks Ranking
- Unlimited Solutions

Identification Methods

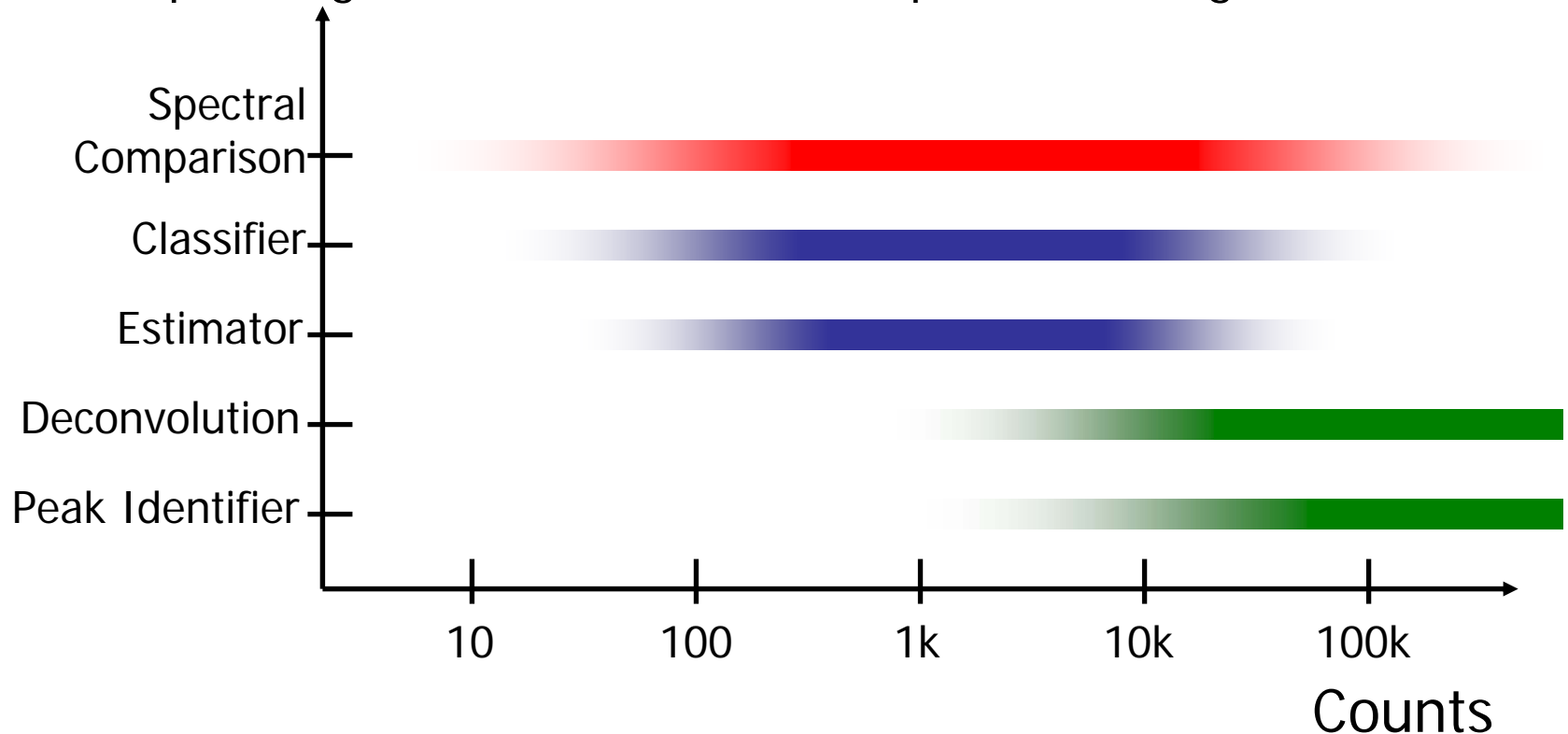


- Feature Extractors
 - **Channel Decimation**
 - **PCA**
 - Peak Extractor
- Feature enhancers
 - **Deconvolution**
- Other
 - **Multi-detector Spectral Comparison**
 - **Asymmetric Detector Nuclide Identification/Deconvolution**
- Identification Algorithms
 - Classifiers
 - Neural Networks
 - Bayes Classifier
 - **Nearest Neighbor**
 - Estimators
 - Multiple linear regression (Fittodb)
 - **Gauss-Newton**
 - **Expectation Maximization**
 - Heuristics
 - Energy Banding
 - Peak identification
 - Other
 - Shielding Estimators (MBS)

Domain of Nuclide Identifiers



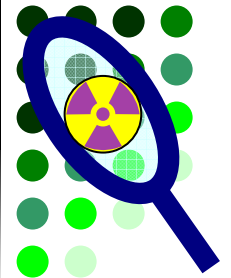
- As number of counts increase, un-modeled effects become visible
 - Ground Bounce, Scattering in Environment, Age, Trace isotopes
- Template algorithms fail due to inadequate modeling

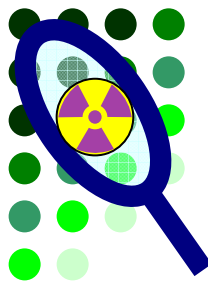




Performance Testing

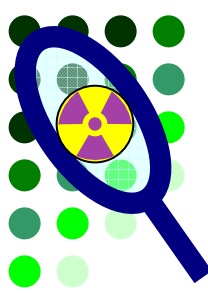
How well do automations work under pressure?





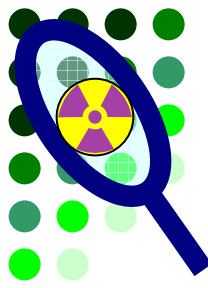
Characteristics of methods

	GADRAS	PCA Classifier	MLG
Type	Estimator	Classifier	Hybrid
Algorithm	MLR	PCA/MSE (dot prod)	Gauss-Newton ML/LBF
Metric	Chi Squared	Poisson Likelihood	AIC
Groups/Trials	5	528	25
Nuclide Output	Composition	1 Library Element	1 Nuclide Group
Reports “No Detection”	Yes	Yes	Yes
Reports “Unknown”	No	Yes	Yes
Additional Output	SNM catagory	Sorted Library List	Fitness of each Isotope Group
Run Speed	1 – 3 seconds	0.5 seconds	0.25 seconds



Evaluation methodology

- Evaluated on two sets
 - Verification – library elements as inputs to algorithm
 - Test – set of similar inputs representing real world inputs
- Ran 100 trials for each using Poisson random draws
- Varied the expected input signal from 50 to 1000 counts for test set
- Background added to sample was Poisson random draw with expected 400 counts
- Reference background was Poisson random draw with expected 120000 counts
- Produced
 - Score based on exact matches or inclusion
 - Cross Correlation Matrix of outputs produced by sample
 - Threat class based assignment

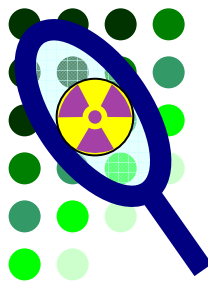


Verification results

- When given library elements and asked figure out which one, we expect nearly perfect performance.
- We didn't get it.

Algorithm	False Negative	False Positive
GADRAS	29%	34%
PCA	10%	10%
MLG	7%	6%

(at 300 cps)

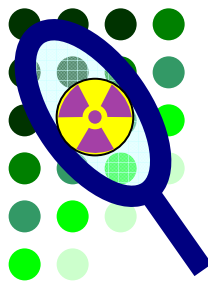


Verification results

- PCA and MLG both performed significantly better than GADRAS (1/3 total number of errors)
- Some items failed to identify or were labeled as mixtures by all methods.

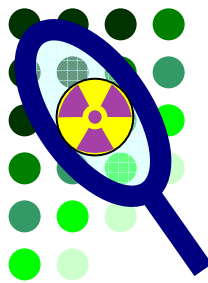


Implies library contains indistinguishable nuclides that cannot be resolved with NaI resolution.



Choice of test cases

- Sources
 - Most commonly detected medical isotopes
 - ^{99m}Tc
 - ^{131}I
 - ^{201}Tl
 - ^{67}Ga
 - Common industrial and potential RDD, ^{137}Cs
 - Natural nuisance sources
 - Bananas
 - Wood
 - Fertilizer w/ potash
 - Weapon surrogates with and without lead shielding
 - HEU
 - Virgin HEU
 - Weapons grade Pu
 - Reactor grade Pu



Correlation Result for Tests

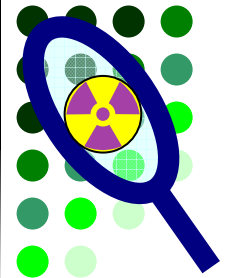
Outputs are 90% identification point with noted errors.
Best is overall performance at all count levels.

Isotope	GADRAS	PCA	MLG	Best
^{99m}Tc	100	100	100	PCA
^{131}I	1000 ^{133}Ba	1000 ^{133}Ba	1000 ^{133}Ba	PCA
^{201}Tl	^{139}Ce	300	200	MLG
^{67}Ga	300	^{237}Np 32%	300	MLG
^{137}Cs	200	100	100	MLG
Bananas	300	500	300	GAD
Wood	300	500	500	GAD
Potash	500	^{152}Eu 70%	^{152}Eu 30%	GAD
HEU	300	200	200	GAD
Virgin HEU	300	200	300	GAD
RG Pu	1000 ^{137}Cs	^{40}K 30%	^{137}Cs 12%	MLG
WG Pu	500	300	500	PCA
HEU 1" Pb	^{232}U 84%	^{232}U 55%	^{232}U 55%	none
Virgin HEU 1" Pb	^{238}U 100%	^{238}U 100%	^{238}U 100%	none
RG Pu 1" Pb	^{232}Th 79%	^{232}Th 79%	^{252}Cf 40%	none
WG Pu 1" Pb	^{137}Cs 70%	^{252}Cf 33%	^{252}Cf 90%	none

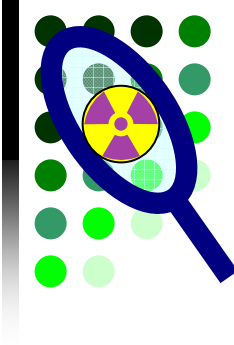


Conclusions

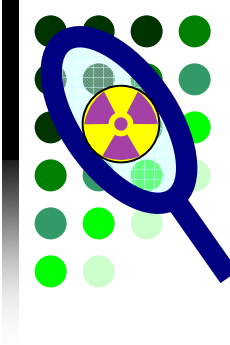
The bottom line



Conclusion



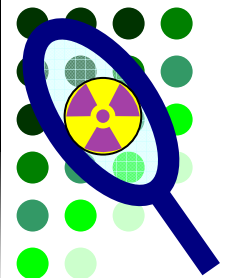
- What are the problems in nuclide identification
 - Ambiguity, Nuisance sources, Inadequate resolution
- What are the problems in a Classifier and an Estimator
 - Classifier – Restricted to ranked list
 - Estimator – Restricted to one solution
- How well do current technologies work
 - Not so swift
- What is to be gained by defining a test bench
 - We can optimize our methods to improve their performance

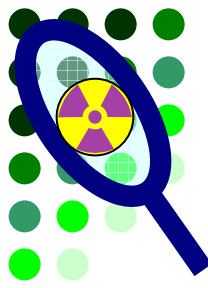




Learning from our mistakes

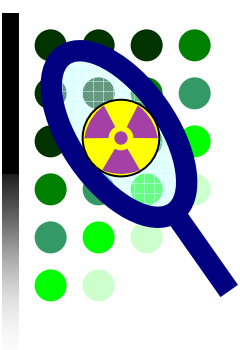
Fool me once shame on you,
Fool me twice shame on me.





LLNL's Template Based Isotopic Identifier

- Produces
 - Optimum fit for single isotope or specified mixtures with arbitrary ratios
 - An “unknown” result in response to novel samples or unexpected mixtures
 - Multiple solutions where more than one possible
 - Significance and Confidence levels for each solution
- Designed for low count spectra (<1000 cts)
- Works with large (2x4x16) NaI detectors
- Uses non-linear Gauss-Newton solver



Verification Result

- Verified by probing algorithm 100 random samples for each template.
 - Samples were drawn to have ~300 counts of source with 334 counts of background
 - Reference 5 minute background provided
- Scored based on false positive and false negative

Un-weighted Results:

False Negative: 134/14300 trials less than 1%.

False Positive: 4217/14300 trials ~30%

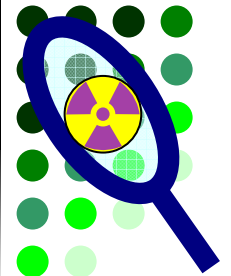
- Require real world frequency data to compute weighted score

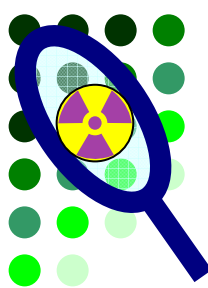




Feature Extraction

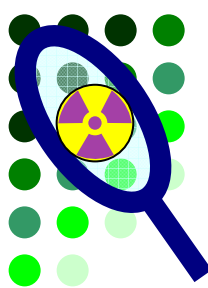
Getting the most out of an observation





Channel Decimation

- Method
 - Adds groups of neighboring channels
- Properties
 - Maintains statistical properties
 - Resulting information loss can cause ambiguity
 - Increases statistical significance of each channel
 - Must strike a balance between information loss and improved significance
 - Radiation spectrum has natural grouping by the square root of the channel energy



Principle Component Analysis

- Method

- Transform input along the basis vector representing the greatest variance (most representative)

- Mathematics

$$R = PP^T = U^T \Lambda V$$

Eigenvalue decomposition

$$W = [U_1 \cdots U_k]$$

Select K largest eigenvalues

$$X = W^T O$$

Transform on this basis

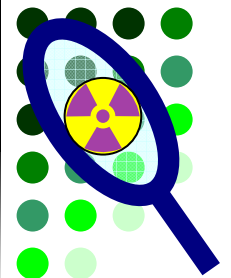
- Properties

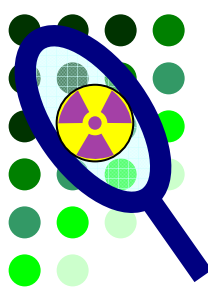
- Does not preserve the statistics of the observation
- Good to reduce high dimensional data sets with minimal loss
- Can reduce noise if it can not be represented in the feature space



Classifiers

What is the best match?





Nearest Neighbor Classifier

Dot Product

- Purpose

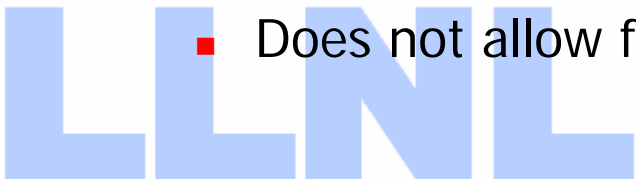
- *Compute the nearest template using L2 norm to the observation.*

- Mathematics

$$\underset{\varepsilon}{\text{minimize}} \quad \varepsilon = \|O - E - kT_i\|_2^2 \quad \longrightarrow \quad \underset{\varepsilon}{\text{maximize}} \quad \varepsilon = (O - E) \cdot \frac{T_i}{\|T_i\|}$$

- Properties

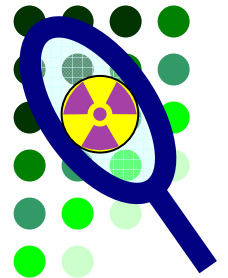
- Shaped based spectral analysis frequently paired with PCA.
- Computationally inexpensive for each template.
- Requires many templates to fully span input domain.
- Does not account for statistical nature of noise.
- Does not allow for fluxations in background conditions.

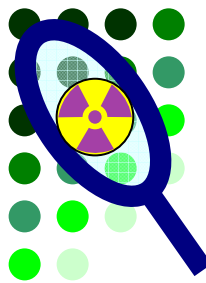




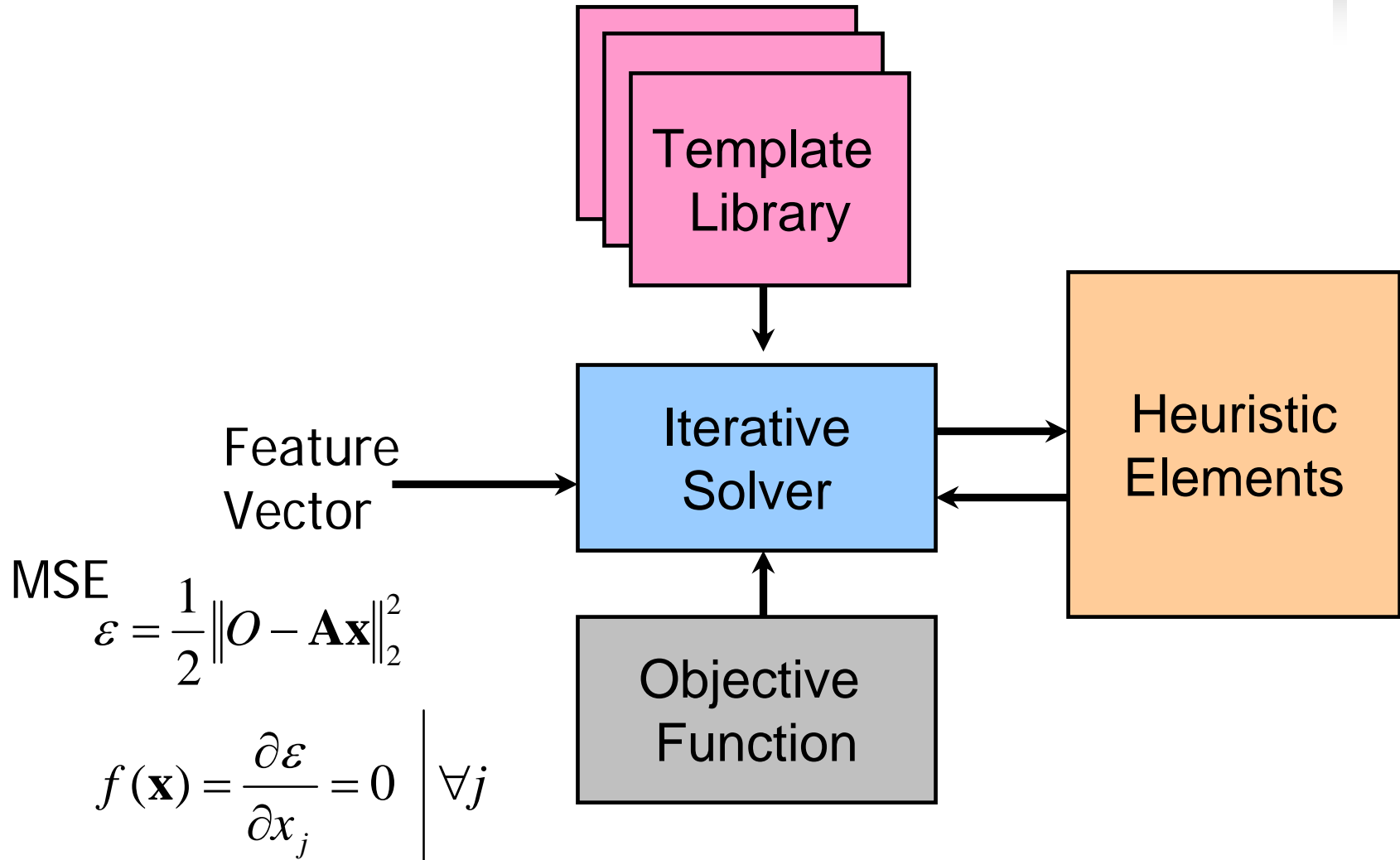
Estimators

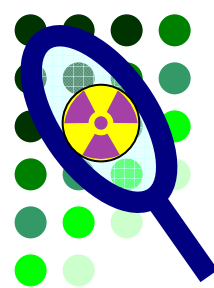
How much of each?





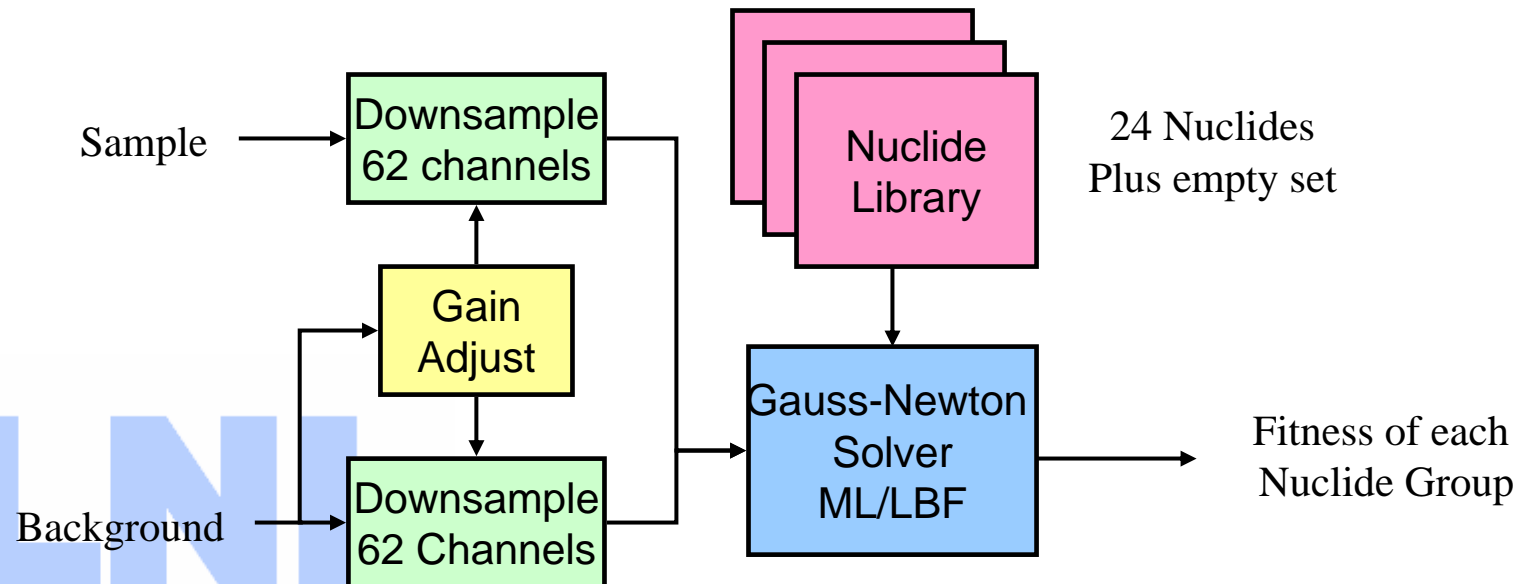
Anatomy of An Estimator

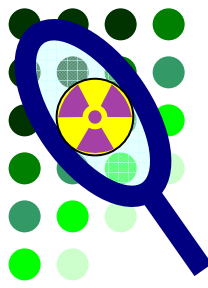




Maximum Likelihood of Subgroups

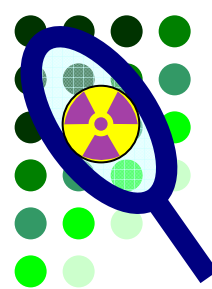
- Estimator using Poisson Statistics and Maximum Likelihood Nonlinear Solver
- Requires Logarithmic Barrier Function (LBF) to impose fully positive constraints
- Computes Likelihood of best fit for Background plus each set of nuclides (all shieldings)



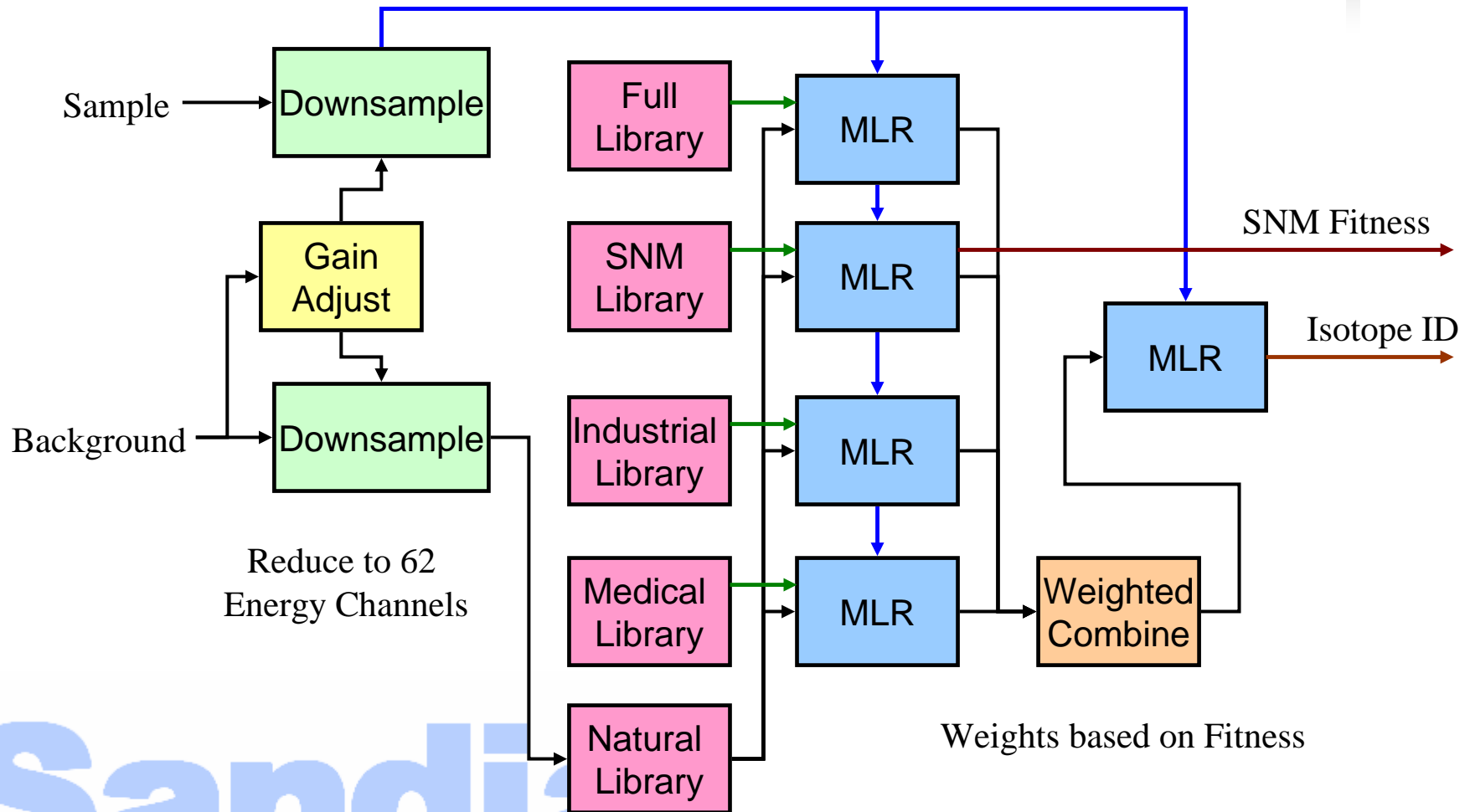


Anatomy of fittodb (GADRAS)

- Core is Heuristic guided MLR (multiple linear regression)
- Objective Function is χ^2 based on 7 variance estimations
- Breaks the library into 4 classes for subgroup analysis
 - Natural
 - Medical
 - Industrial
 - SNM
- Produces
 - Fitness of SNM
 - Nuclide Identification (only reports “significant contributors”)
 - Significance (reduced χ^2)



Anatomy of fittodb (GADRAS)



Reduce to 62
Energy Channels

Weights based on Fitness

143 Isotope/Shield Pairs

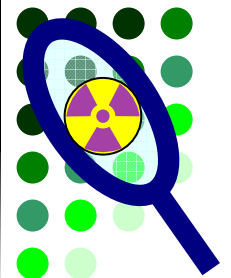
PROPRIETARY

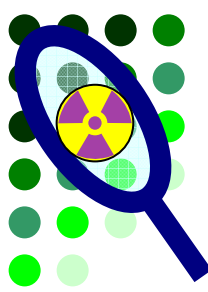
Sandia



Other Techniques

One of these things are not like the others.





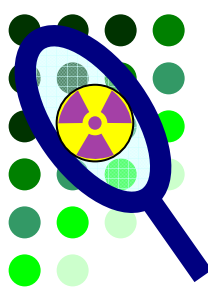
Materials Basis Set

- Purpose
 - Estimate the shielding located between a known nuclide and the detector.

- Mathematically

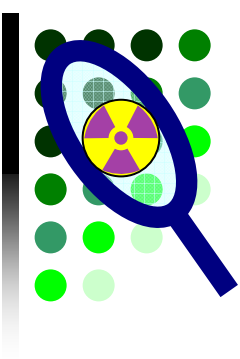
$$b_{z,i} = -\ln\left(\frac{c_{z,i}}{c_{z,ref}}\right) \quad U = \begin{bmatrix} b_{0,ref} & b_{0,z1} & \cdots & 1 \\ b_{1,ref} & & & 1 \\ \vdots & & \ddots & 1 \\ b_{n,ref} & & \cdots & 1 \end{bmatrix} \quad b = U \cdot \rho$$

- Properties
 - Uses Beer's Law ($A=\varepsilon/c$) estimate shielding thickness
 - Works in a log space
 - Assumes that increased shielding decreases the mean counts in a channel



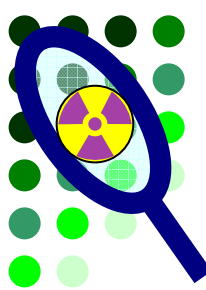
Materials Basis Set

- Primary functions on the photo-peaks
 - Compton scatter breaks assumptions thus is biasing the solution
- Working in a logarithmic space makes mixture analysis difficult.
 - Nuclide signatures add linearly
 - Complexity of estimator will increase as N^2 as additional nuclides are included in the mixture
 - Perturbations from Compton Scattering likely to bias mixture solution
- Assumes χ^2 statistics on reconstructed spectra



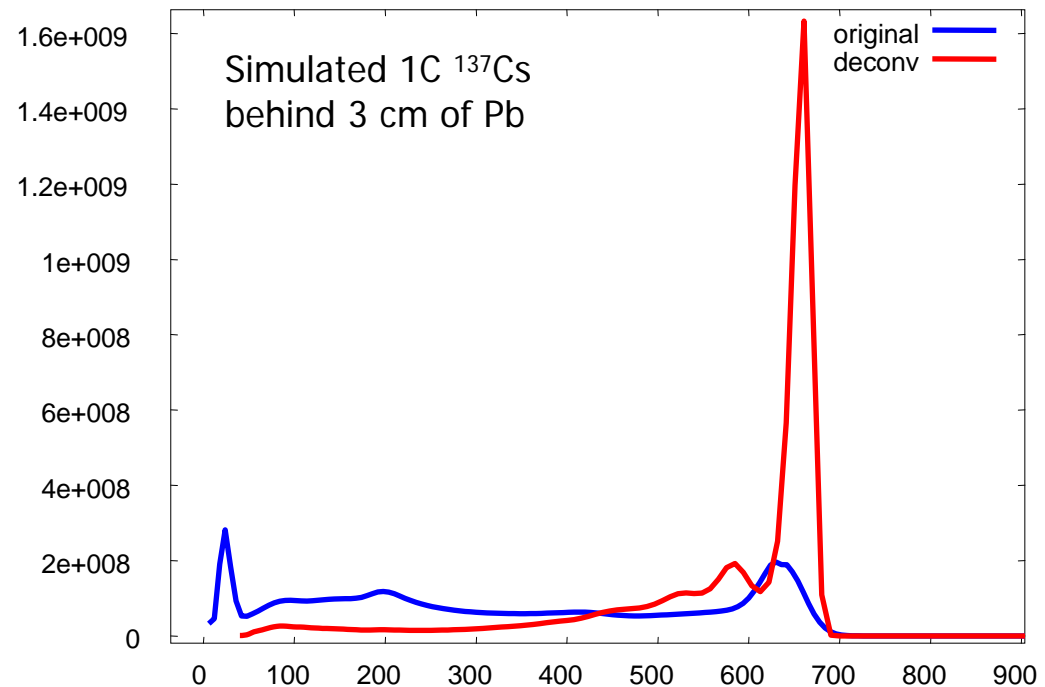
Deconvolution

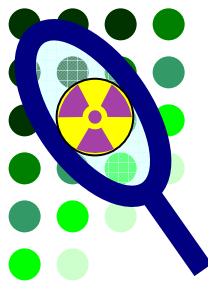
- Purpose
 - Compute the energy flux at the surface of the detector from the observation
- Properties
 - Requires relatively large numbers of counts to produce a quality result (10k)
 - Requires extremely well tuned detector model
 - Removes virtually all detector effects
 - Compton Scatter
 - Backscatter peaks
 - Decreases the FWHM of the detector by a factor of 4
 - Allows isolation of peaks within $\frac{1}{2}$ of the FWHM



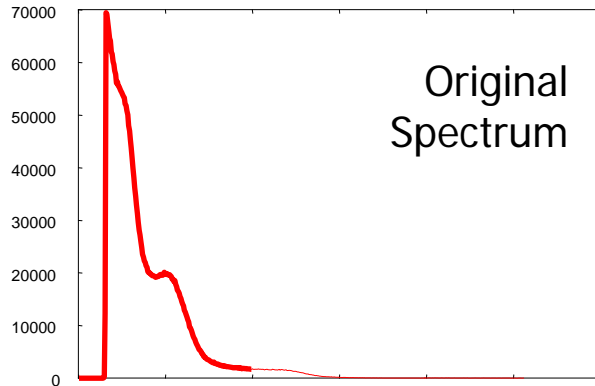
NaI Deconvolution

- Deconvolution using EM algorithm
- Allows direct evaluation of shielding from Compton scattering or branching ratios
- Removed backscatter peak
- Recovered Compton Scatter in the detector to the photo-peak



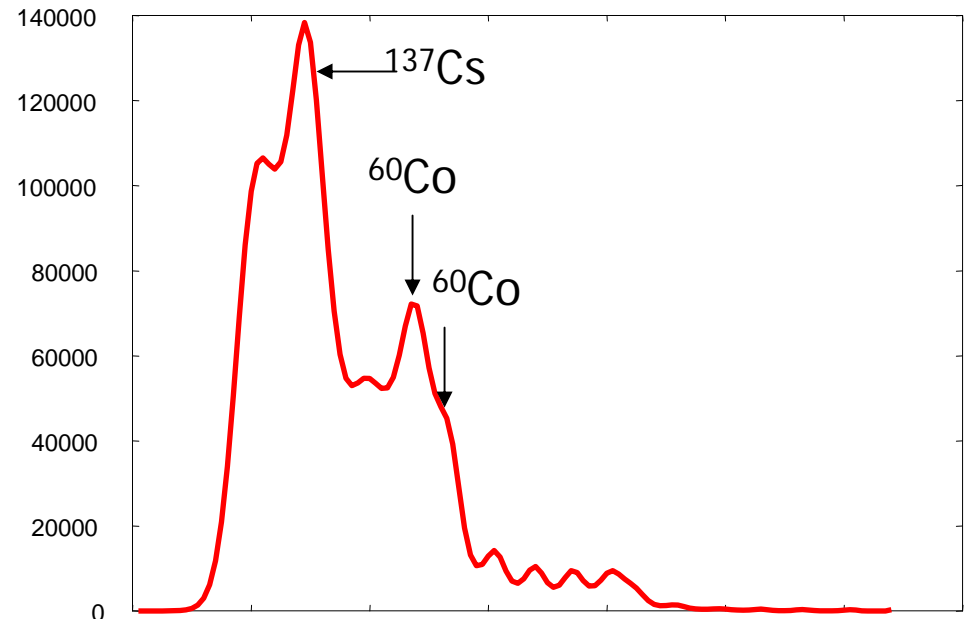


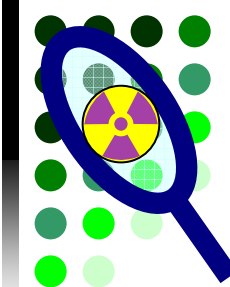
Plastic Deconvolution



- Provides
 - Peaks otherwise not visible can be identified
 - Allows calibration of energy scale

- Deconvolution using EM algorithm
 - 128 bin energy square root energy
 - Gaussian basis

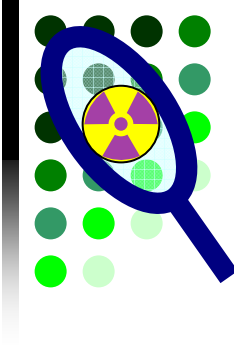




Multi-detector Nuclide Comparison

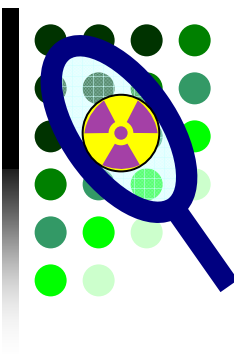
- Purpose
 - Uses multiple detectors to improve identification or match spectra between previous encounters
- Properties
 - Simultaneously solve detectors by information sharing or summing
 - Depending on the number of energy channels used can operate with less counts at a particular encounter than possible otherwise
 - Demonstrated in DTS to improve nuclide identification for high speed traffic by using detections from multiple locations

Asymmetric Detector Nuclide Identification/Deconvolution



- Purpose
 - Using two detectors with different resolutions and efficiencies to improve identification.
- Properties
 - Simultaneously solves identification in both detectors by information sharing
 - Complexity increases at number of detectors squared
 - Pairs small high resolution (HPGe, SiLi, CZT) detector with larger low resolution detector (NaI, Plastic)
 - Creates hybrid detector with resolution and efficiency somewhere between the two

LLNL



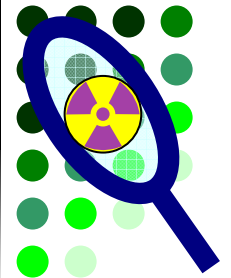
Non-negative Iterative Solvers

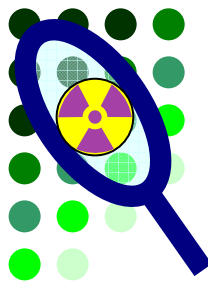
- NNLS - Lewson and Hanson
- Logarithmic Barrier Function – method of generalized inequality constraints
- Expectation Maximization – uses probability density function that are positive by design



Statistics

What do these silly numbers all mean?





Model Confidence (Trust)

- Purpose

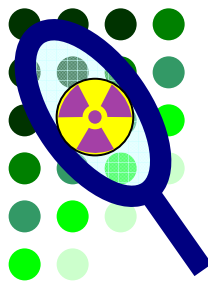
- *Indicates how frequently a specified model is a correct identification.*

- Mathematics

$$C_{M_i} = \frac{P(O | M_i)P(M_i)}{\sum_{M_j} P(O | M_j)P(M_j)}$$

- Properties

- Depends on immeasurable quantity $P(M)$
- Requires the sum of all possible solutions that may not be computable in finite time.
- At best we can compute a empirical estimate of confidence or psuedo-confidence



Nuclide Confidence

- Purpose

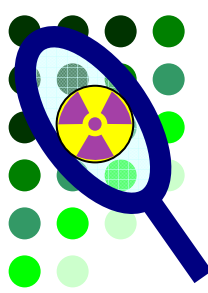
- *Indicate how frequently a specified nuclide is a correct identification.*

- Mathematics

$$C_X = \frac{\sum_{X \subset M_i} P(O | M_i) P(M_i)}{\sum_{M_j} P(O | M_j) P(M_j)}$$

- Properties

- As with Model Confidence, depends on immeasurable quantities. Thus we can at best estimate it.



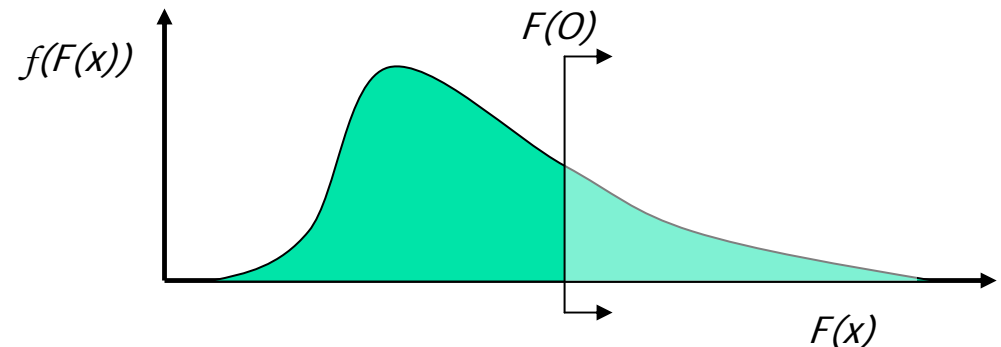
Significance

■ Purpose

- *Indicates how frequently a similar observation may be produced from a model.*

■ Mathematics

$$P(F(x) > F(O) | M)$$



■ Properties

- Evaluated by either a null hypothesis test (χ^2) or by likelihood ratios.
- Used as the objective function in estimators.
- Well defined for radio-nuclide identification problem.
- Controls the rate of Type II errors.