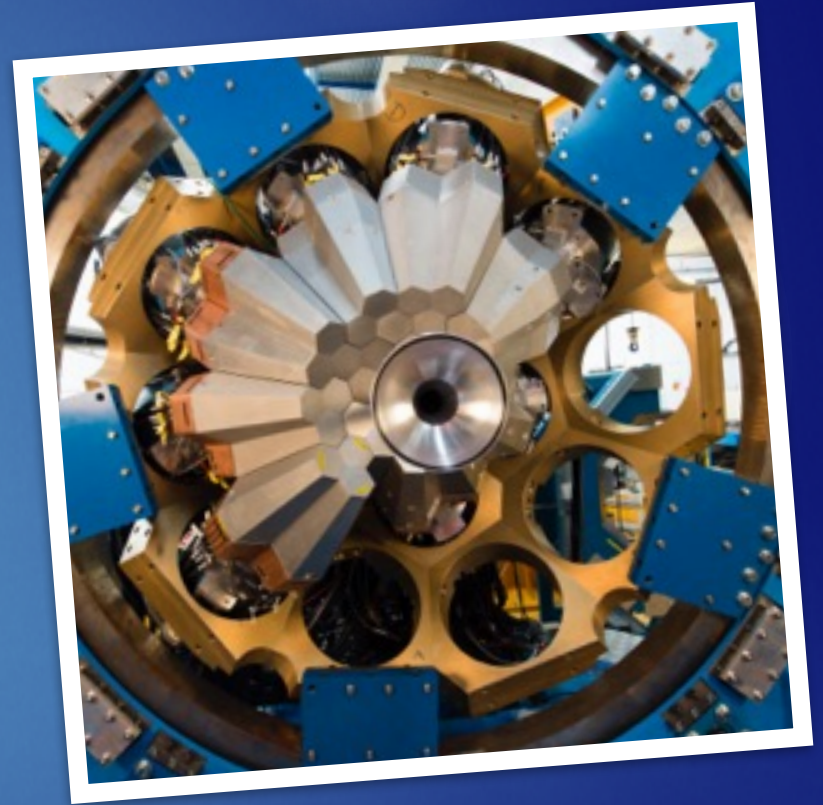


Machine Learning and Topological Data Analysis for Pulse Shape Analysis

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Facilities Council



AGATA
ADVANCED GAMMA
TRACKING ARRAY

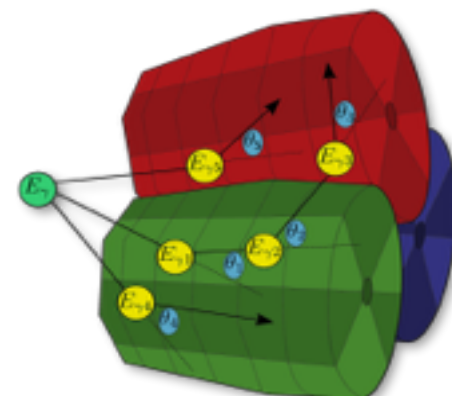


- ▶ γ -ray tracking requires positions at resolution $\sim 5\text{mm}$ FWHM at $\sim 5\text{kHz}/\text{CPU}$.
- ▶ Positions must be inferred from electrical response (PSA).
- ▶ Complex detector response makes parametric methods insufficient.
- ▶ Instead we simulate the detector response in ADL.
- ▶ Interaction locations are then determined by optimisation metrics:

$$\text{Figure of Merit} = \sum_j \sum_{t_i} |A_m^j[t_i] - A_s^j[t_i]|^p$$

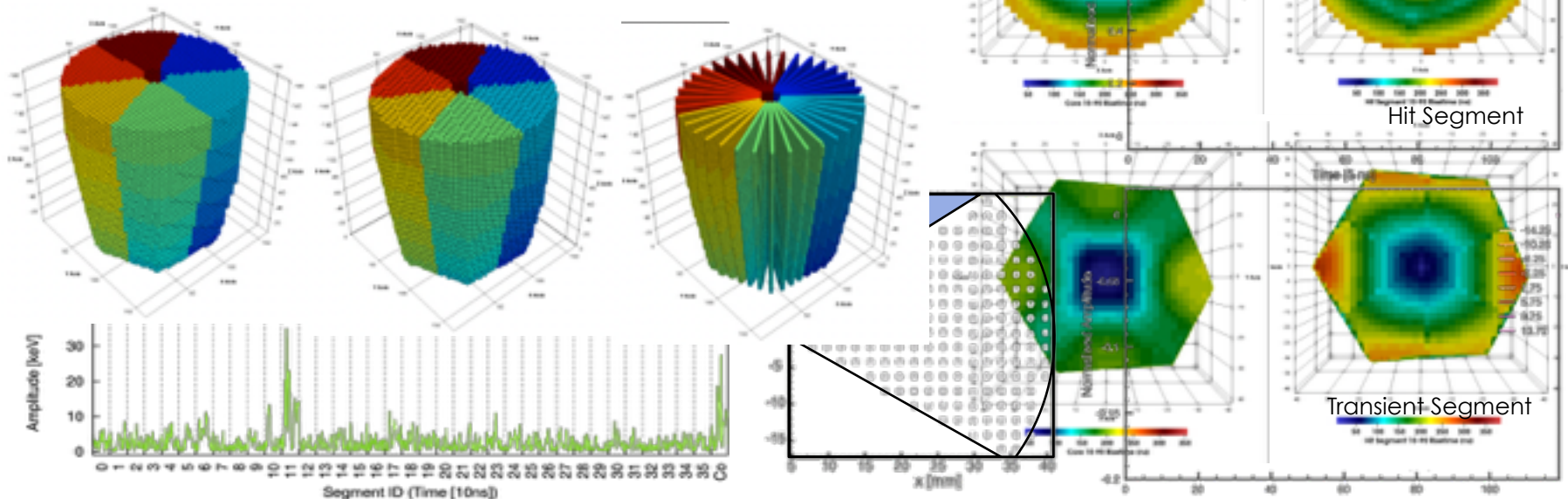
For signals of segment j at time step t_i with p typically $=2$

- ▶ Other metrics can be used to highlight different sensitivities.
 - ▶ Different exponents, weighting for segments.
 - ▶ Time shifting via Dynamic Time-Warping.
- ▶ My work is on developing Novel PSA techniques for AGATA.



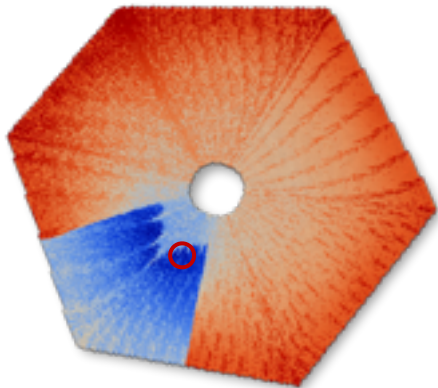


- ▶ Simulated data looks reasonable as expected.
- ▶ Parametric trends are seen in the data, useful for clustering
 - ▶ T_{10-90} , charge asymmetry, knee-point, skewness etc.
 - ▶ These parameters are continuous but break down at high fold.
- ▶ 6-fold symmetric, polar and tetrahedral basis sets simulated.
- ▶ High resolution (0.5mm) basis set generated too.
- ▶ Option for dynamic resolution basis sets.

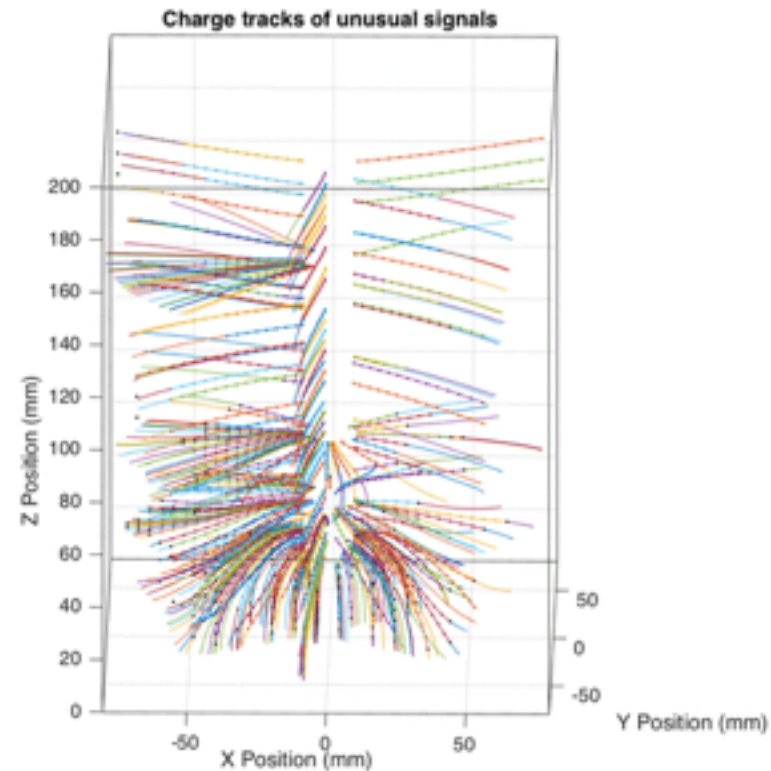
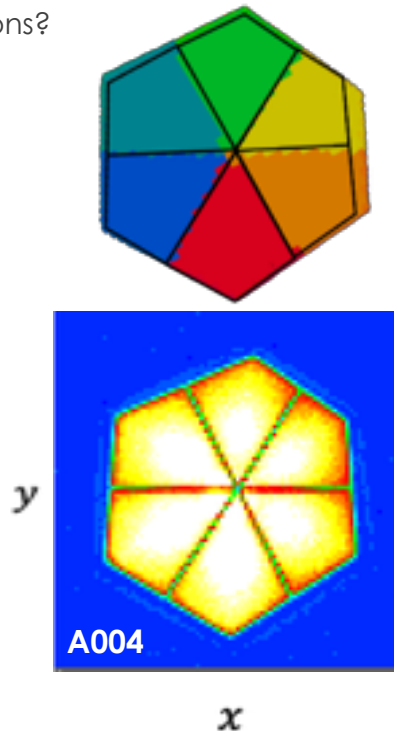




- ▶ Field simulation limited to 1mm spacing, ADL is done at 2mm for a reason.
- ▶ SIMION segmentation is wrong on face of crystals.
- ▶ Odd effects seen at segment boundaries & high resolution:
 - ▶ Unexplained 'charge sharing' between segments.
 - ▶ Sharp discontinuities at edge changes.
 - ▶ Overlap of SIMION definitions?



0.5mm FoM Plot showing odd effects
Optimum Circled





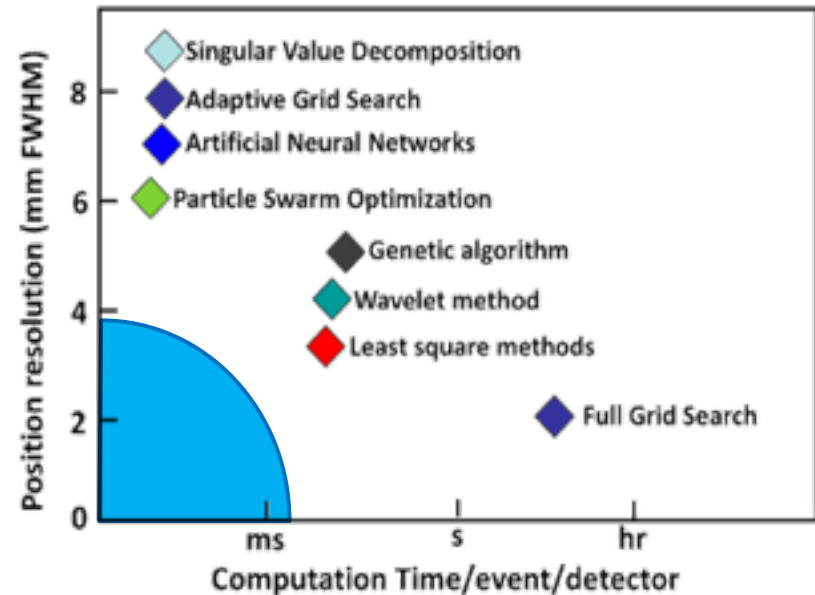
Several PSA algorithms have been tried for AGATA. Time limits for online PSA mean only ~5% of the basis can be searched using current CPU methods.

There are three different ways to solve this issue:

- ▶ Hyper-parallelize the search (GPU acceleration).
- ▶ Use more efficient search methods (**TDA**).
- ▶ Don't search at all, instead infer locations (**ML**).

Moving beyond the basis simulation this becomes a computer science problem, existing techniques can be applied.

- ▶ ∴ Plenty of established fields to learn from.



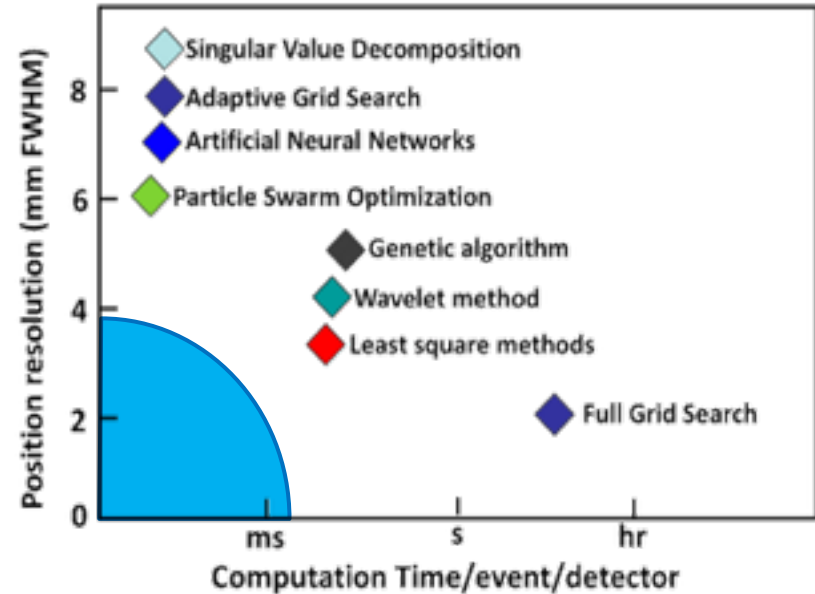


Topological Data Analysis (TDA) techniques try to organize data and form efficient search spaces.

- ▶ Search spaces are Non-Euclidean
- ▶ Generally *kd*-ball or cover trees used.
- ▶ Less prone to local minima.
- ▶ Search algorithms aren't naïve.
- ▶ Each step made moves search closer to optimum.
- ▶ Searching n points can be $\mathcal{O} \log(n)$.

Machine Learning (ML) uses the simulated basis to learn trends via feature extraction.

- ▶ No searching is performed whatsoever.
- ▶ Simulated basis only needed for training.
- ▶ Needs an appropriate model & good data.



Novel Algorithm Development

Tree-based search approaches:

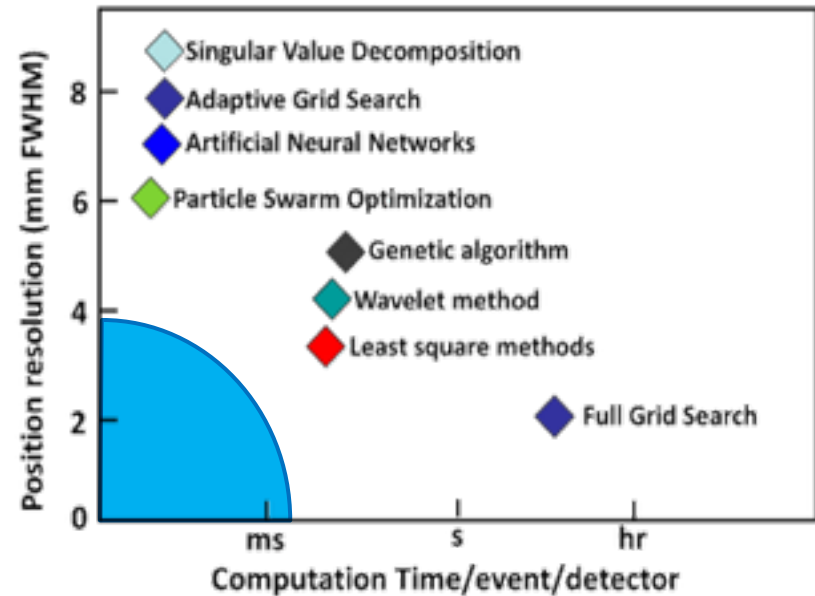
- ▶ k NN - k -dimensional Nearest Neighbors.
- ▶ LSH – Locality-based Sensitivity Hashing.
- ▶ ST/DT MKS – Maximum Kernel Search.

Machine Learning options:

- ▶ Signal Classification.
- ▶ Regression (CNN).
- ▶ Autoencoding/Fingerprinting (β -VAE).

Other options:

- ▶ GPU Acceleration.
- ▶ **All Algorithms have been tested with Gaussian Noise, experimental noise to be determined.**
 - ▶ Performance is likely to decrease.
 - ▶ Will know more when scanning table is operational.



- ▶ GPUs have advanced significantly (10x) since the last AGATA investigation.
- ▶ GPU acceleration can be used on embarrassingly parallel problems:
 - ▶ Exhaustive search.
 - ▶ Adaptive Grid search (two step).
 - ▶ Matrix manipulations.
 - ▶ Figure of merit (although matrix sum $\mathcal{O}(\log_2(n))$)
- ▶ Shared memory makes things complicated.
- ▶ Multiple languages can use GPU accelerated code:
 - ▶ C, C++ (NVCC).
 - ▶ Python (with Numba).
- ▶ Programs can be compiled to use NVBLAS:
 - ▶ MLPACK (Armadillo).
- ▶ GPUs are **very** powerful for ML approaches.

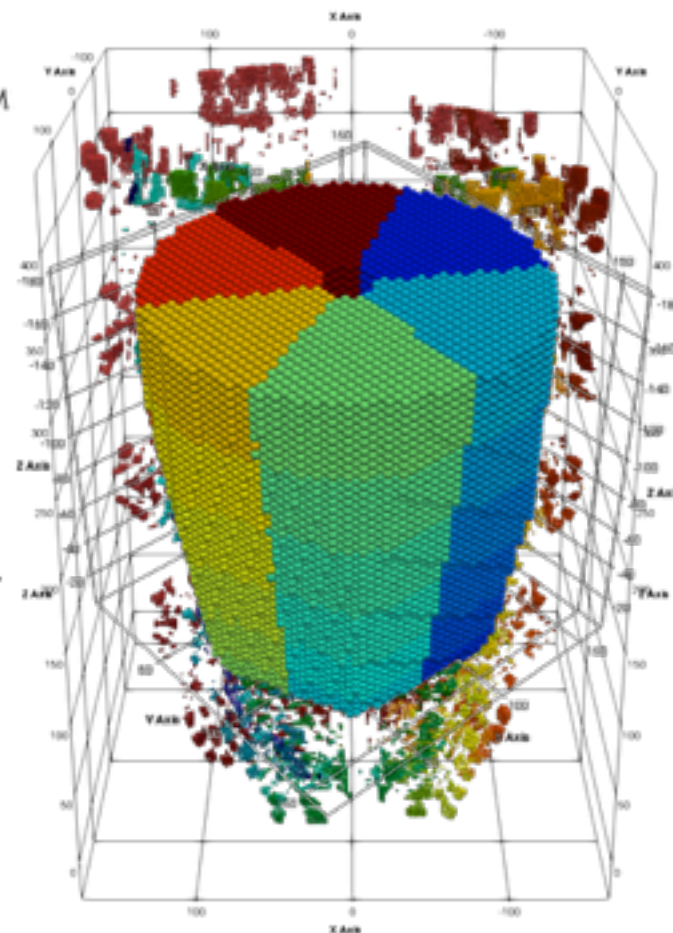


Nvidia P5000
(277 GFLOPS)

Routine	Types	Operation
GEMM		Multiplication of 2 matrices.
SYRK		
TRSM		Triangular solve (right angled)
TRMM		Triangular matrix-matrix multiply
SYMM		Symmetric matrix-matrix multiply

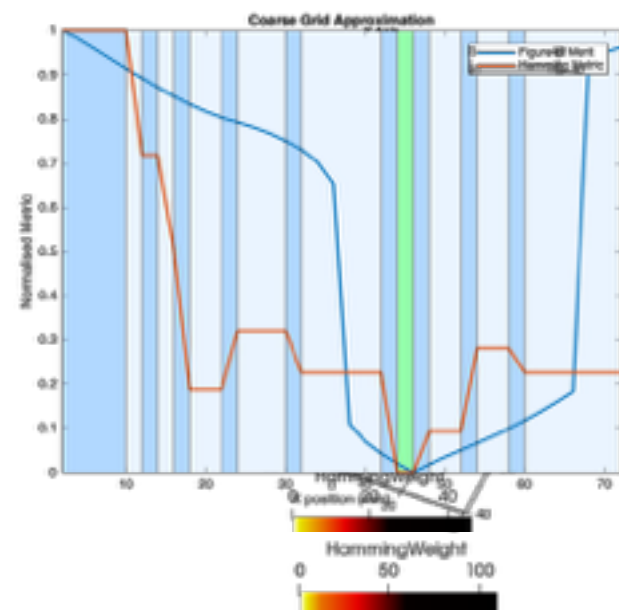
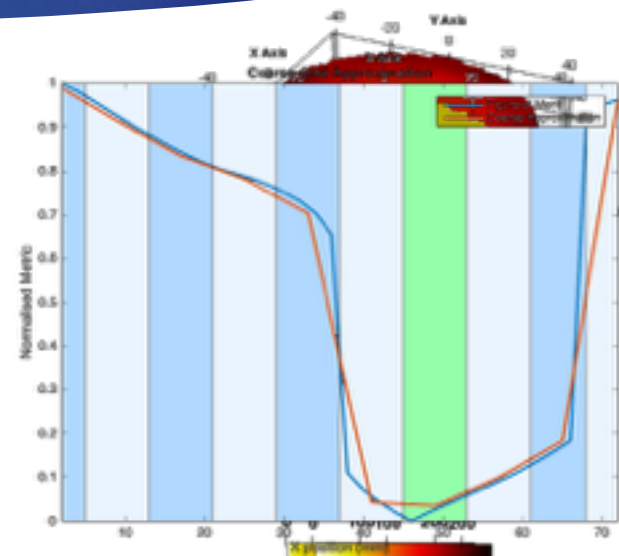


- ▶ Initial investigations were made into optimizing the clustering used in AGS.
- ▶ Instead of using Euclidean splitting the basis was split parametrically:
 - ▶ Segment # $\rightarrow T_{10-90} \rightarrow$ Charge asymmetry \rightarrow Transient Signal Fingerprint \rightarrow FoM
- ▶ This allows for hierarchical ordering of basis & bespoke optimizations.
- ▶ Resolution of metrics inversely related to execution time.
 - ▶ Faster metrics narrow down solution \rightarrow FoM test applied on final cluster.
- ▶ Low resolution metrics mitigate overfitting.
- ▶ Sensitivity of the detector is accounted for.
- ▶ Ultimately parametric clustering difficult (impossible) at high fold.
 - ▶ Accurate fold-invariant metrics difficult to make (might be possible with ML).
- ▶ Method will likely be revisited in the future.
 - ▶ Framework written in C \therefore can be compiled into MTSORT.
- ▶ Made somewhat obsolete by LSH.





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- ▶ Established C++ Library MLPACK used for KNN & MKS operations.
- ▶ GPU acceleration possible using NVBLAS.
- ▶ Additional Python API & Command line interfaces available.
- ▶ Modular design allows for custom Figures of Merit, segment handling.
- ▶ Prefers smooth & convex search spaces.
 - ▶ Doesn't like searching multiple segments.
 - ▶ Metric penalizes segments far from interaction.
- ▶ *Should* work for multiple interactions within the same segment.
 - ▶ Combinations need to be precomputed.
 - ▶ Outrageous memory costs if implemented.
- ▶ Currently 3 techniques look applicable to Fold-1 searches:
 - ▶ *k*NN
 - ▶ LSH
 - ▶ **MKS**



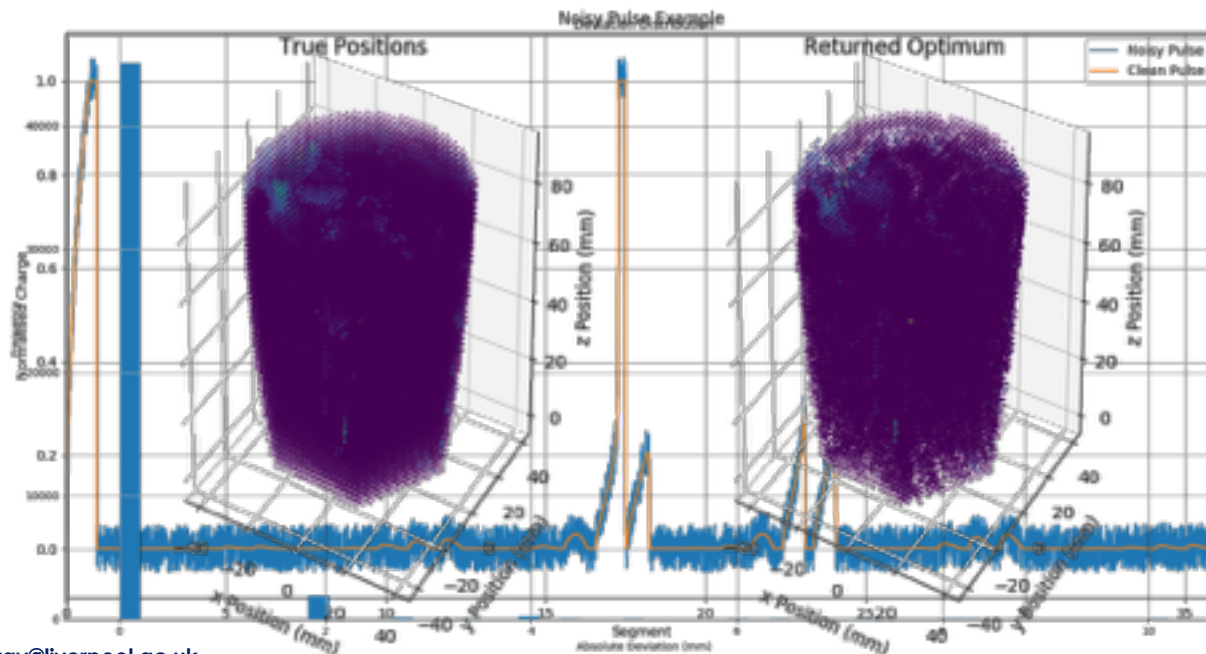


- ▶ Fast Maximum Kernel Search uses two trees to search an ordered data structure.
- ▶ First tree is used to convert reference set into structured data.
- ▶ Second tree is then dynamically built using query set.

- ▶ Efficient comparisons mean that the space can be searched quickly.
- ▶ Mercer Kernels allow for modifications of phase space, improve separations.
 - ▶ More complex kernels have execution penalty.



- ▶ 10% Gaussian noise added to simulated database for preliminary validation.
- ▶ MKS with Gaussian kernel used to return top 5 solutions of kernel search with confidences.
- ▶ 95% of fold-1 events identified at input location.
- ▶ 99% of fold-1 events within 2mm.
- ▶ Discrete distances due to finite grid size.
- ▶ Currently clustering of deviations are not well understood, needs further analysis.

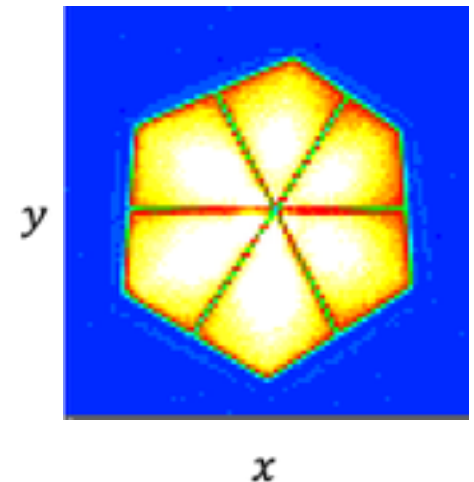




- ▶ Main motivation of this method was to identify interesting sections of the interaction.
 - ▶ Possible groundwork for software-based trigger.
 - ▶ Because of this these networks need to be fast (and likely simple).
- ▶ Position gated pulses used to generate database of hit, transient & noise samples.
- ▶ Various networks trained to predict category.
- ▶ Ultimately the cut is arbitrary, open to interpretation.
- ▶ Doesn't offer much above traditional methods.
 - ▶ However if we want to look for something specific it's pretty useful.

Method	Agreement with Midas Label	Score
Multi-Level Perceptron	~68%	9
Binary Perceptron	~87%	9
Neural Network	~94%	22
Convolution Neural Network	~97.6%	26

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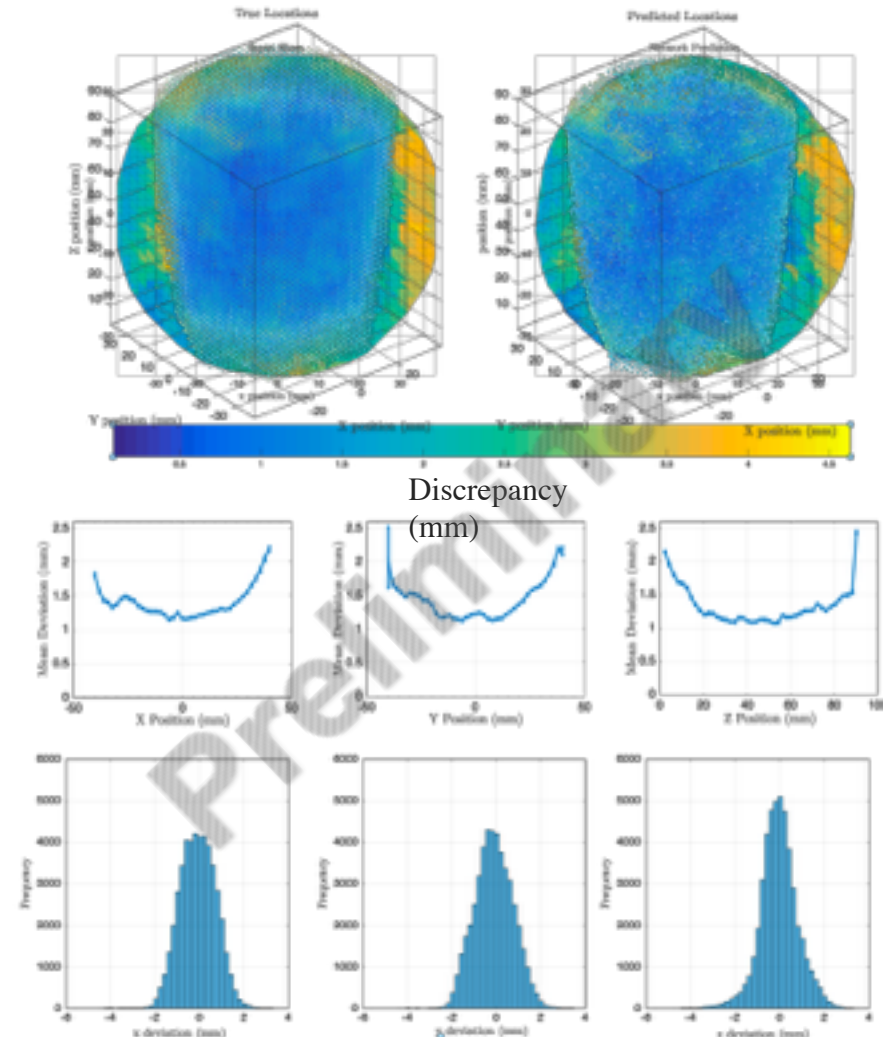
- ▶ Similar setup as before, input data is either core electrode or superpulse.
- ▶ Multiplicity to simulate taken from expected distribution.
- ▶ Two scenarios simulated:
 - ▶ Multiple hits in the same segment.
 - ▶ Multiple hits in the same crystal.
- ▶ Output of network still treated as categorical
 - ▶ Likelihood of fold reported, pick the most likely

Initial results look promising however simulation was heavily idealized.

Issues with this method:

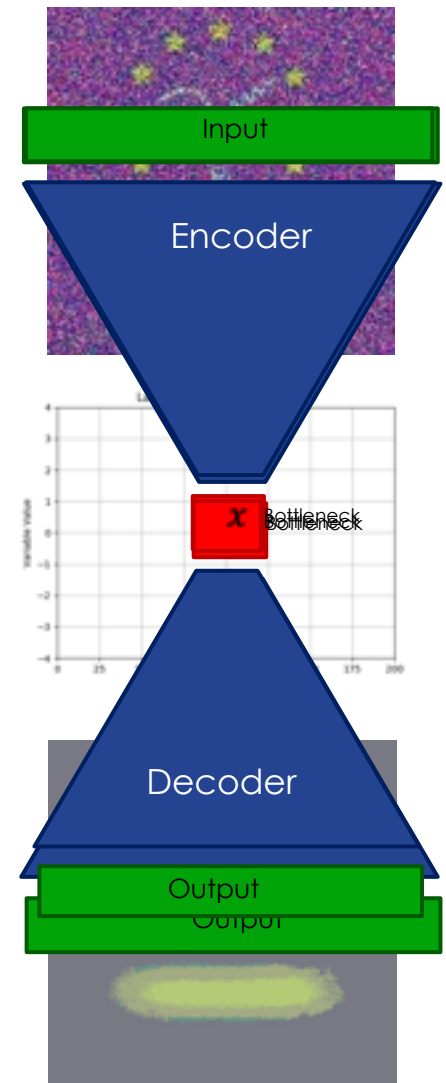
- ▶ Interaction locations & energies picked at random, should use GEANT4 instead.
- ▶ Realistic noise floor needed.

- ▶ CNN used to return continuous outputs.
- ▶ Trained on 6x8x120 tensor (core contact excluded).
 - ▶ Column repeats used for CNN windows.
- ▶ ResNet architecture used for robustness.
- ▶ Gaussian noise & Dropouts used for reliability.
 - ▶ **Should use experimental noise instead.**
- ▶ Works well on detectors with high connectivity.
- ▶ Currently only implemented for fold-1 events.
 - ▶ Training on multi-fold requires separate networks.
 - ▶ This isn't difficult, I'm just waiting for an accurate simulation of multiple fold events.
- ▶ Reasonable execution time $\sim 300\mu\text{s}$.
- ▶ Variable FWHM, performs worse at boundaries.
 - ▶ Will likely decrease with realistic data.



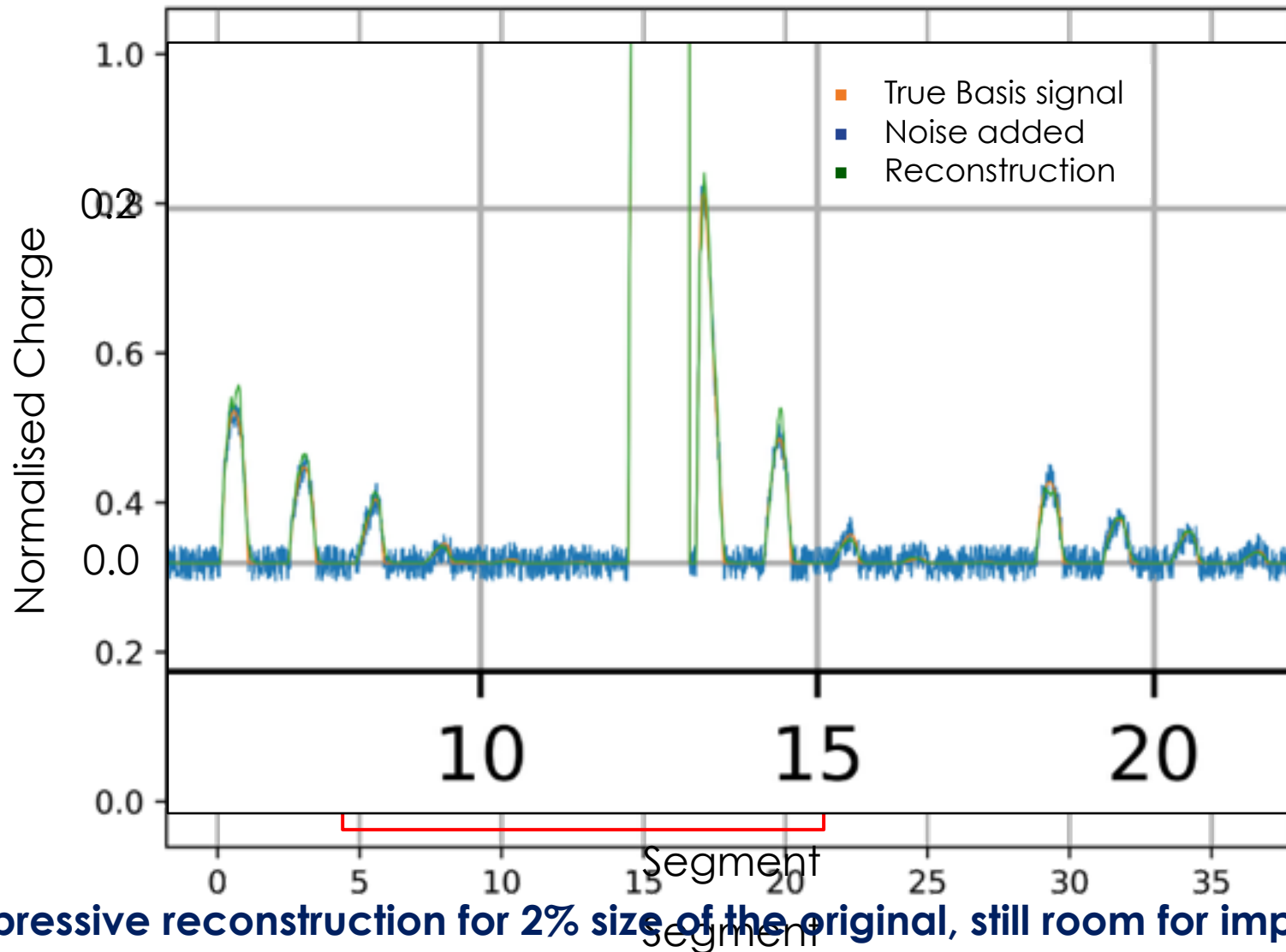


- ▶ Autoencoders combine two separate networks to function:
 - ▶ Encoder: converts input to a learned latent space via feature extraction.
 - ▶ Decoder: converts latent space into a reconstructed output.
- ▶ Autoencoders are **incredibly** efficient however can be lossy.
- ▶ As a whole the network replicates a denoised input.
 - ▶ Signal is intelligently denoised, small transients are unaffected.
 - ▶ Network doesn't see noise as useful information.
- ▶ Current Execution time $\sim 56\mu\text{s}$ however will likely change.
- ▶ Autoencoders become more useful when split into parts:
 - ▶ The Encoder and Decoder compress data far better than traditional methods.
 - ▶ The latent representation can be used to express parametric trends.
 - ▶ This requires disentangling the latent space (difficult)
 - ▶ Can this be used for tagging?
- ▶ Compression isn't necessarily bad, oddly the reconstructed pulses could end up being better than the inputs due to denoising.



Example Reconstructions, ~44x Compression

18



Impressive reconstruction for 2% size of the original, still room for improvement.

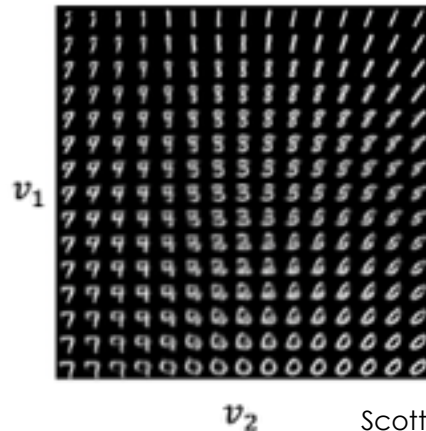
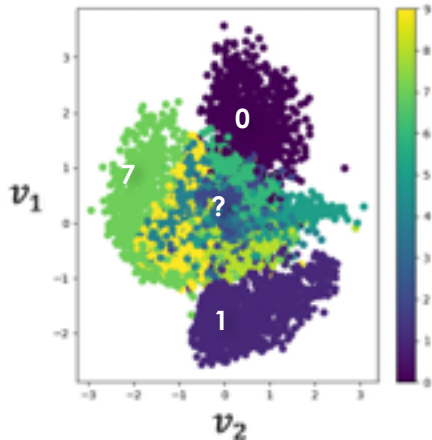
Disentangled Autoencoders

19

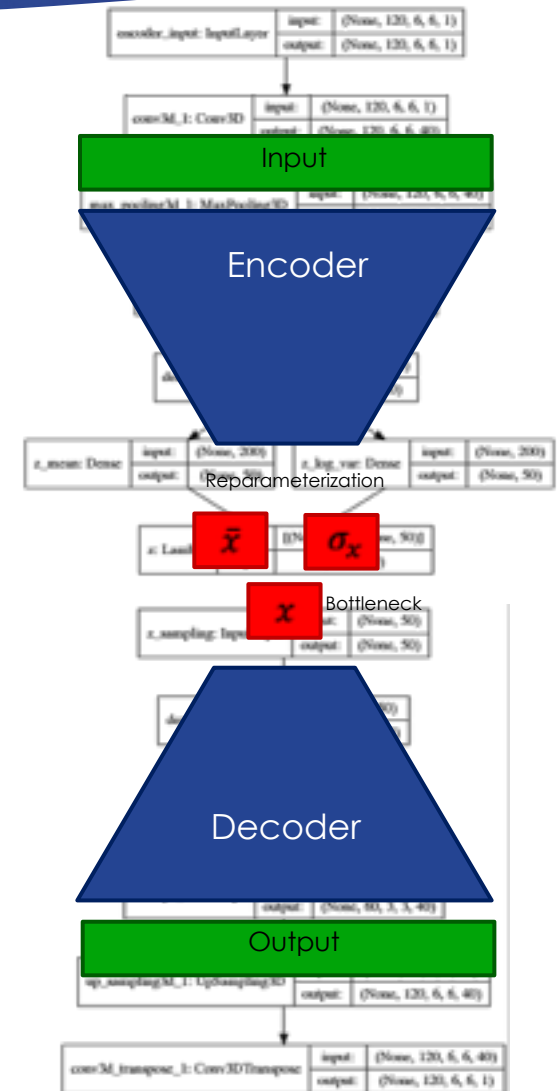


- ▶ Typical AE bottlenecks are impossible to interpret.
- ▶ Optimum bottleneck size is unknown, how many variables contribute?
- ▶ DAE attempt to maximize the usefulness of the latent representation.
- ▶ This is done by making each latent variable strongly independent.
- ▶ Each latent variable should represent a different parametric trend.
 - ▶ Latent space should be separable.
- ▶ Latent representation should be fold-invariant.
- ▶ Perform MKS on latent representation.

MNIST set example:

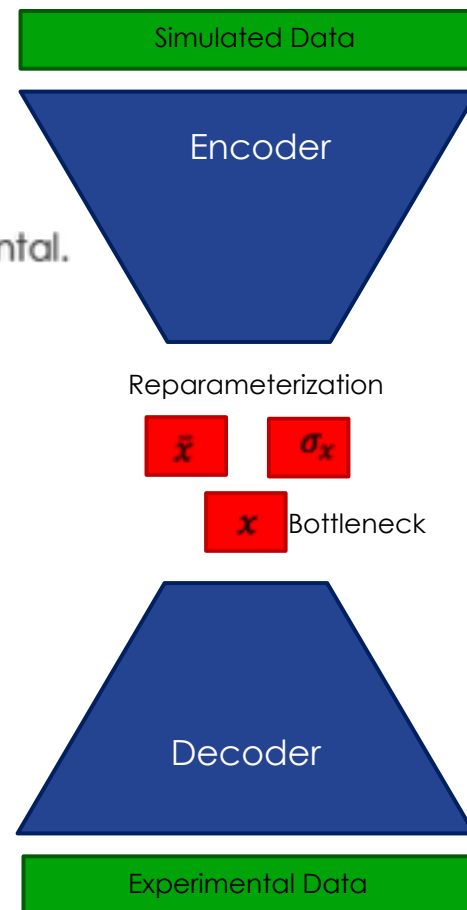


Scott Freitas (CSE 591, 2018)

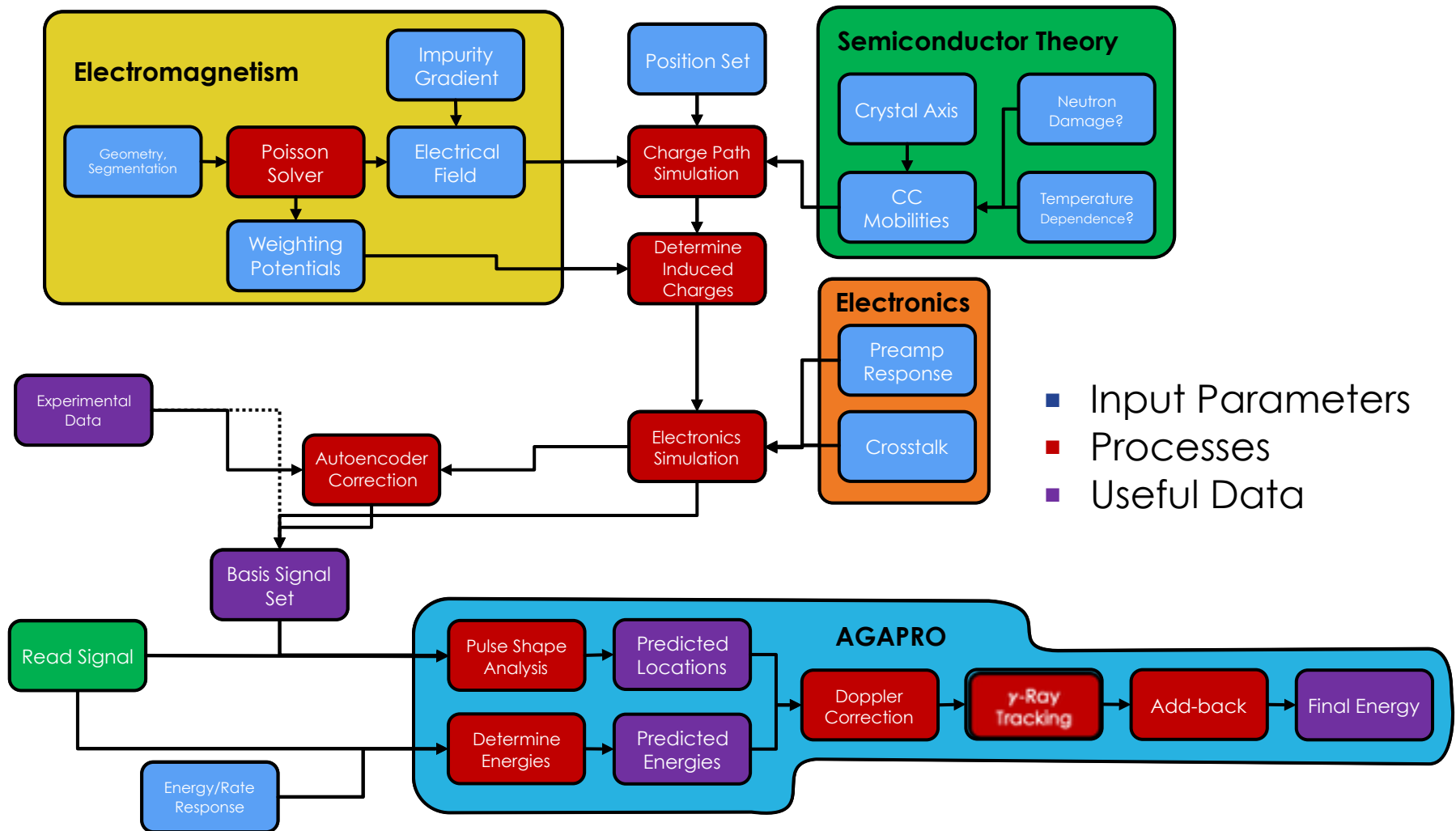




- ▶ PSA and GRT perform differently when given real & simulated data.
- ▶ Therefore there's likely some form of discrepancy between the two.
- ▶ How about using ML to transform simulated into real data?
- ▶ Simulation reduced to latent space & then reconstructed to experimental.
- ▶ This approach requires very good experimental data:
 - ▶ Full x, y, z characterisation of the crystal.
 - ▶ No guarantee that trained model can be adapted to different crystals.
- ▶ Validation data for A005 will be taken anyways.
 - ▶ May as well test the feasibility of this method.
- ▶ Transform of preamplifier response also possible.
 - ▶ **Way easier**

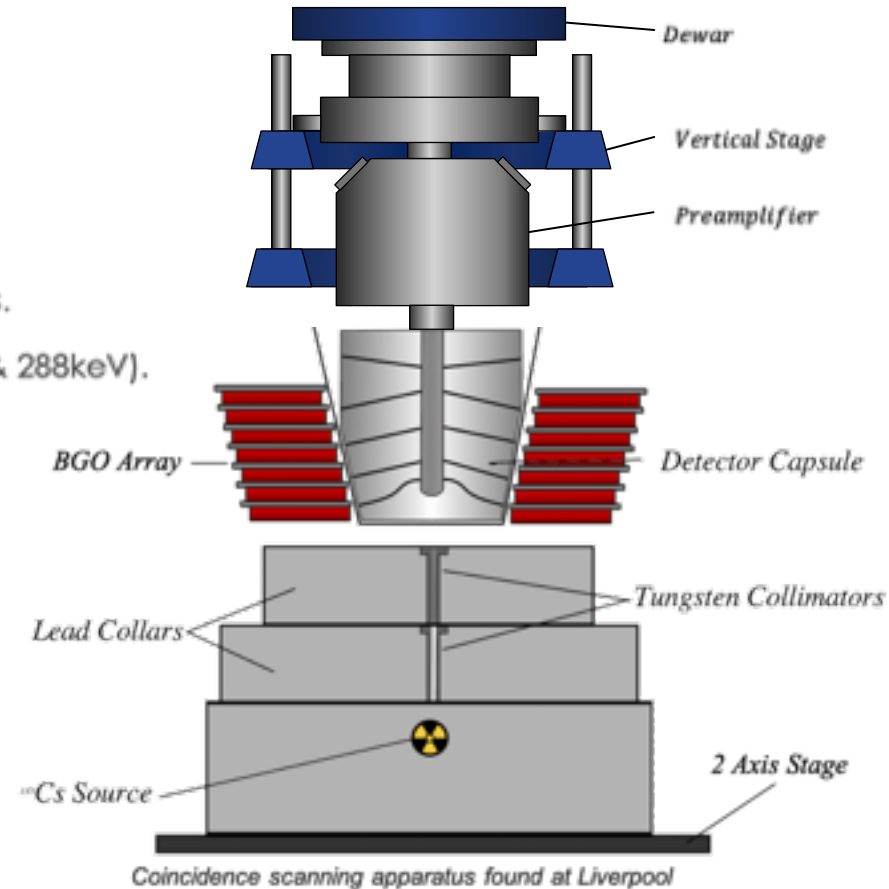


AGATA Pipeline





- ▶ Coincidence scanning will be used to validate simulations, ML efforts and PSCS method (IPHC, Strasbourg).
- ▶ Will provide a definitive & time aligned basis for Geant4.
 - ▶ Allows for proper simulations of high-fold events.
 - ▶ Currently using Caen 1724s, may switch to AGATA digitizers.
- ▶ 1GBq ^{137}Cs source collimated to 1mm on x, y stage.
- ▶ Vertical stage added to apparatus for quick z movements.
- ▶ 90° scatter gating using BGO array & energy gating (374 & 288keV).
- ▶ I'm currently writing the MTSort code for acquisition.
- ▶ Typical validation measurements will be taken:
 - ▶ ^{241}Am surface scan for alignment.
 - ▶ Gated cross & circle measurements for CAO.
 - ▶ Gated coarse cubic grid using vertical stage.
 - ▶ High-resolution pencil beam of front segmentation.
 - ▶ (Time permitting) Automated High-resolution scan.





- ▶ GPUs have advanced significantly over the last decade, likely to continue in the future.
 - ▶ Definitely should be revisited considering future projections.
- ▶ Tree-based search methods are incredibly efficient but difficult to adapt to high fold.
 - ▶ Use fold-invariant search space instead?
 - ▶ Very applicable for Fold-1 regardless.
- ▶ ML approaches offer good learned relationships but need adaptations to high fold.
 - ▶ Realistic high fold dataset necessary.
- ▶ We have a good standing for more ambitious ML techniques.
 - ▶ Discrimination
 - ▶ Regression
 - ▶ Auto-tagging / Fingerprinting
 - ▶ Compression
 - ▶ Basis Correction
- ▶ Variational Autoencoders may simplify pulse storage whilst helping with PSA.
- ▶ I can't take all these methods to completion, future work will involve whittling down algorithms.

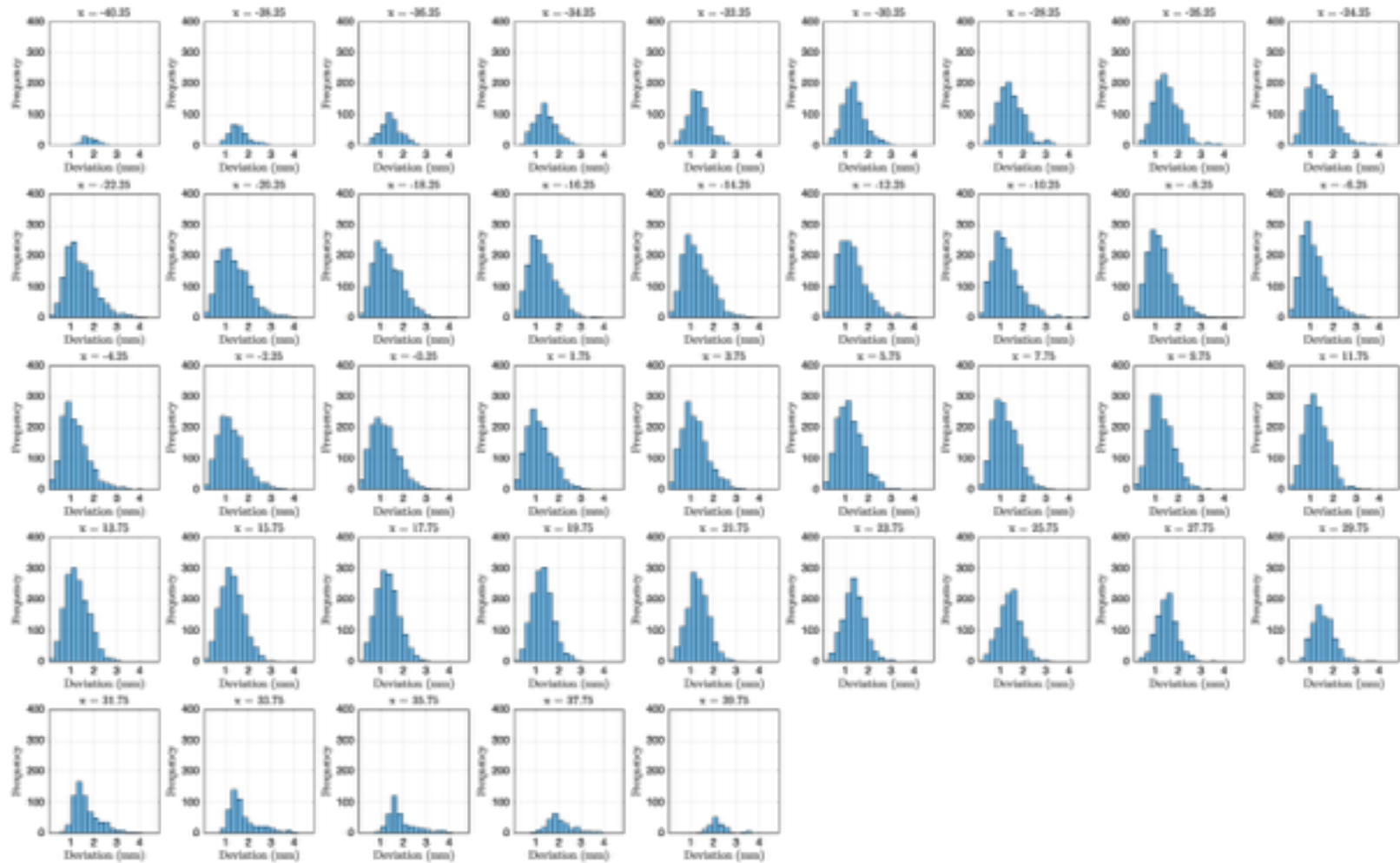
Thanks for Listening

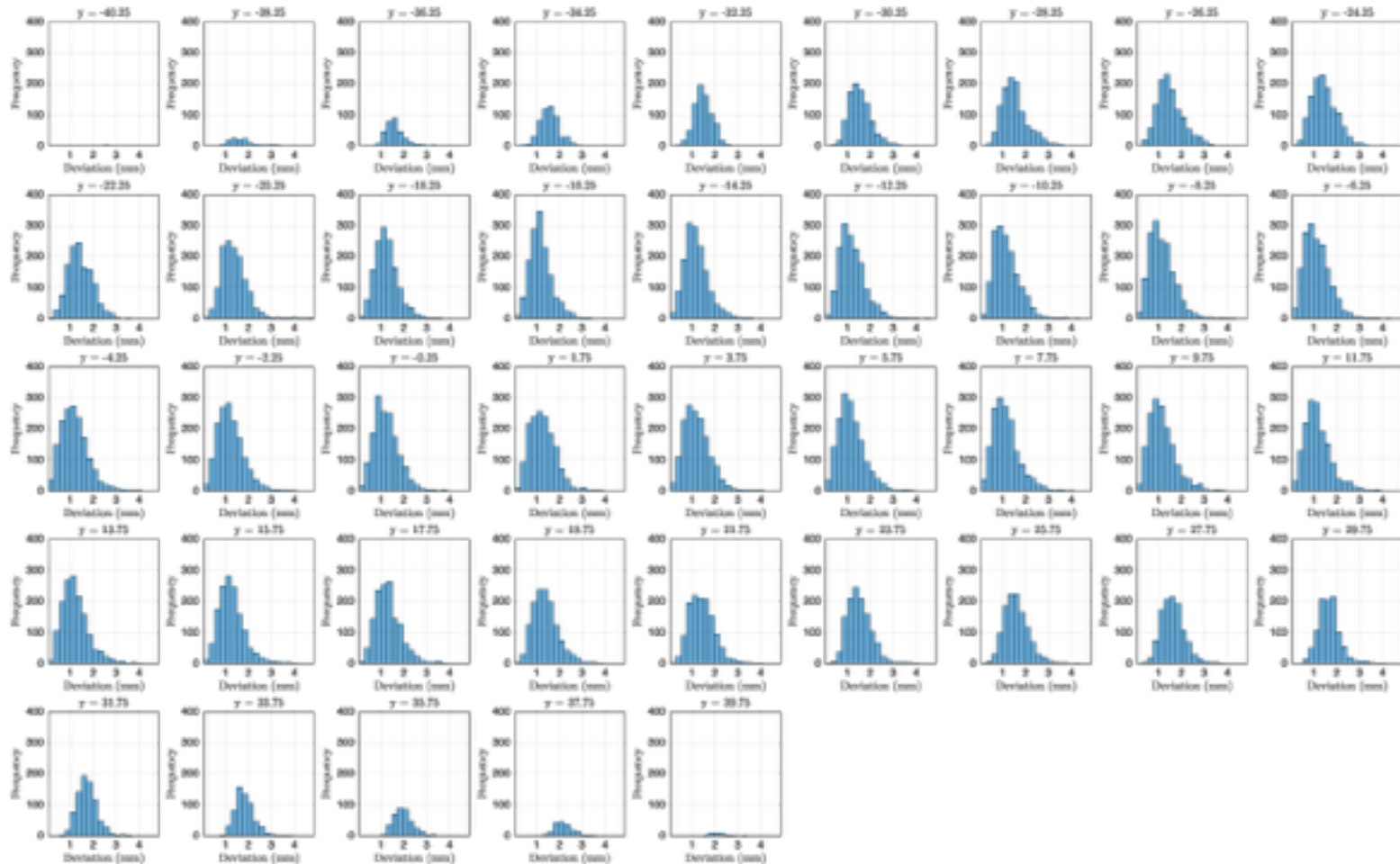
Any Questions?

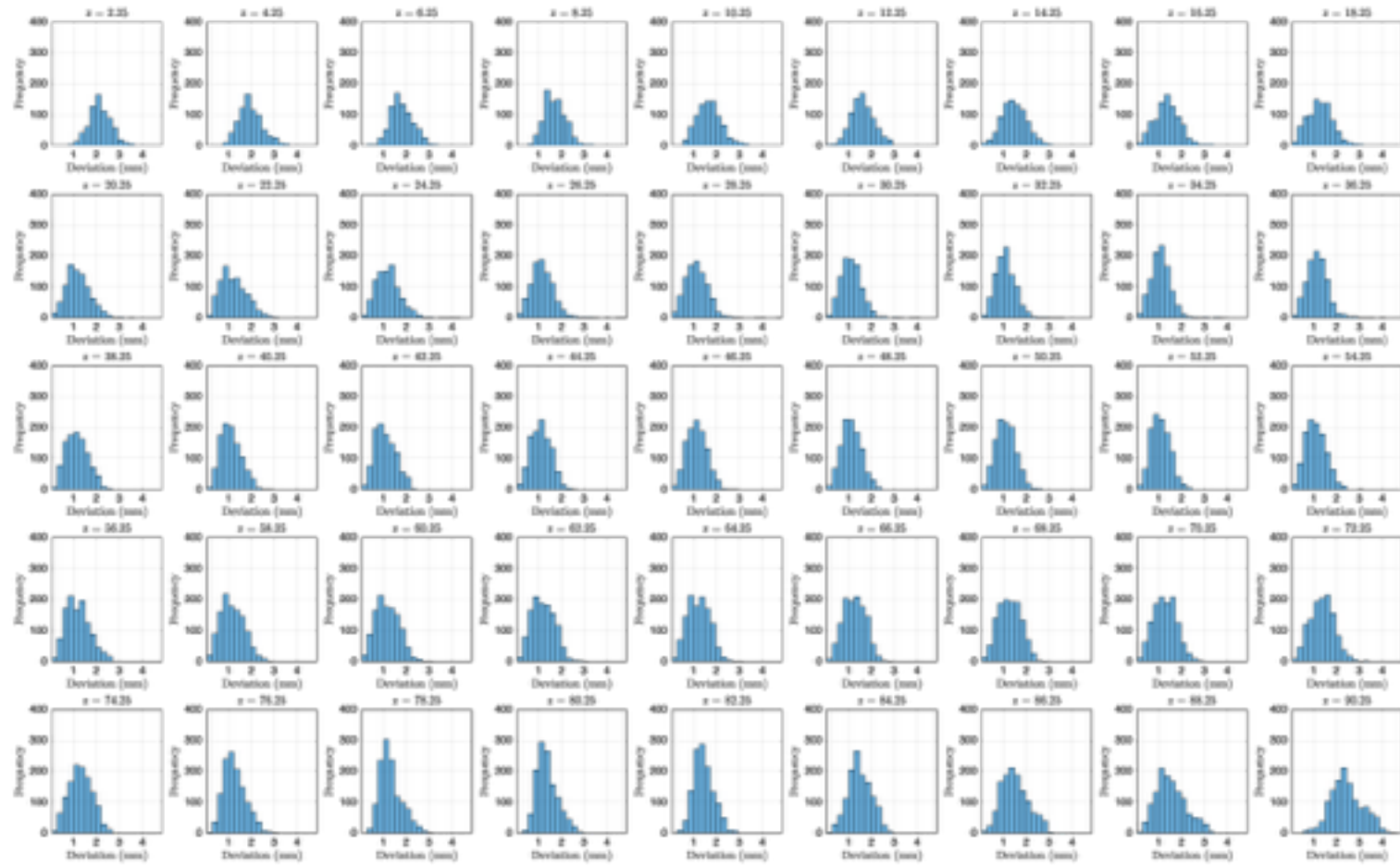


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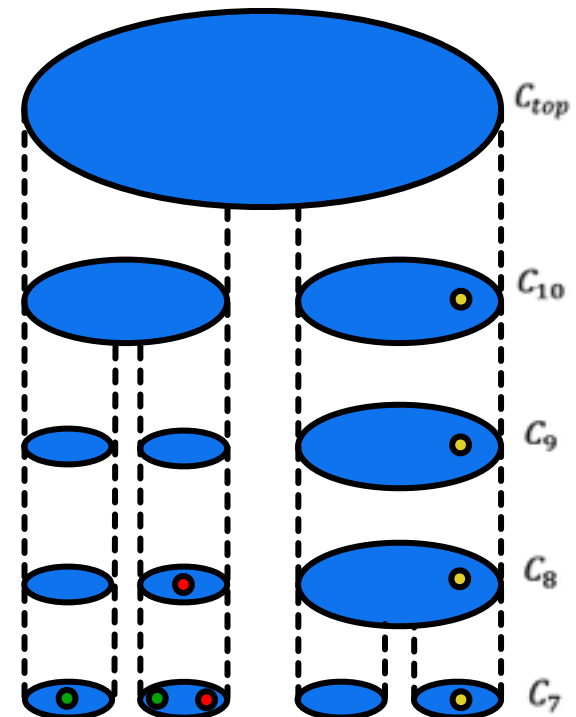




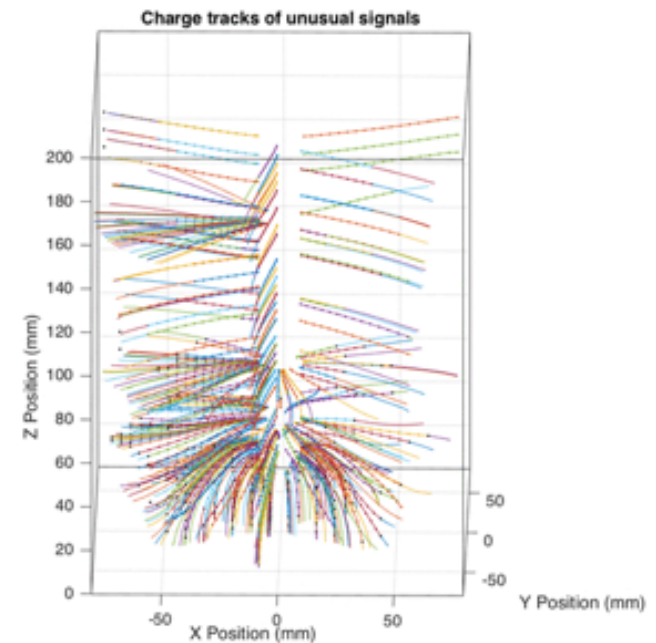
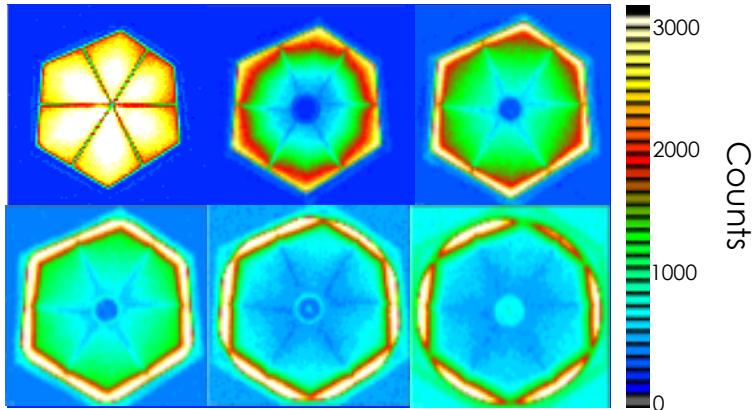




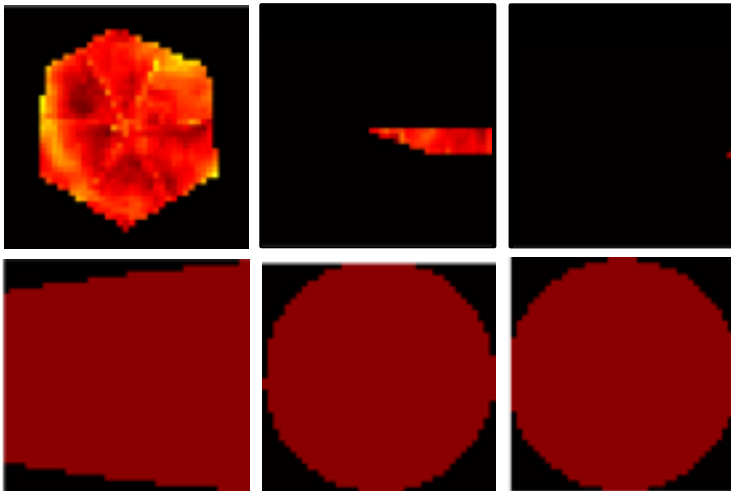
- ▶ For a collection of points C_i on level i of T which represent a subset of points in S the following rules must be enforced:
 - ▶ $C_i \subset C_{i-1}$ - Nesting: any point $p \in S$ that exists in C_i must have an associated node in all lower levels.
 - ▶ $\forall p \in C_{i-1}$ - Covering: for every $p \in C_{i-1}$ there exists one $q \in C_i$ such that $d(p, q) \leq 2^i$ where the node for q is the sole parent of the node for p .
 - ▶ $\forall p, r \in C_i, d(p, r) > 2^i$ - Separation: For all $p, r \in C_i$ then $d(p, r) > 2^i$



- ▶ Several algorithms have been developed for fold-1
- ▶ Adaptions for multiplicity are hard
- ▶ Database needs to be validated experimentally
- ▶ Odd effects in basis set need to be investigated



- ▶ Training set taken from ADL simulated pulses, Gaussian noise added
- ▶ CNN attempts to predict interaction location from superpulse
- ▶ Currently limited to fold-1 events, may be mitigated by using windows



CNN Prediction
Discrepancy

