



PSA via Singular Value Decomposition

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Outline

AGATA PSA

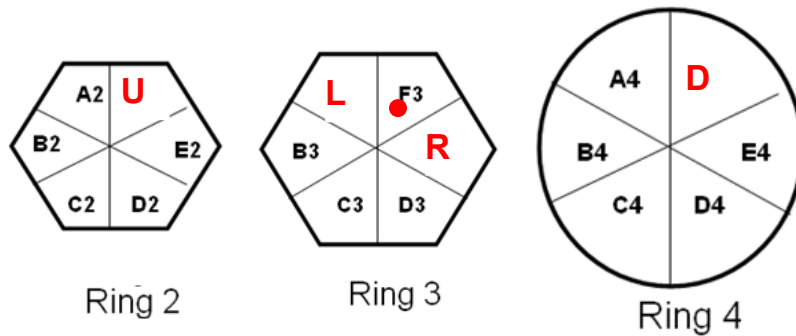
Pulse shape comparison with X^2 threshold

SVD & Principal component analysis

Fast pulse shape comparison

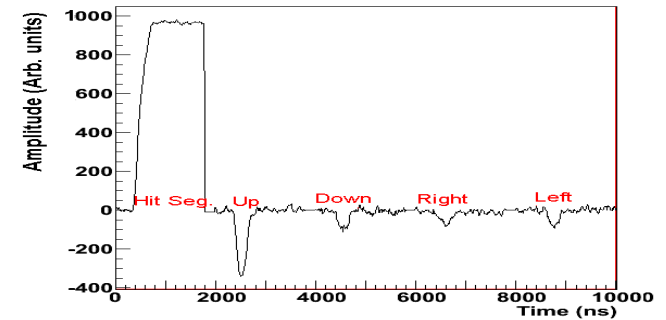
Next steps

PSA



Position of gamma interaction

ADL /
scanner
→
?
←



Pulse shape
(hit segment and direct neighbours)

PSA via reference signals

1. build database of signals for know positions
2. find best fit to reference signal to determine position

→ e.g. Adaptive Grid Search

Core of the program

- Just a sequence of loops with almost no calculations
- Runs fast because all structures fit in memory

```

#define RERRE 24
#define RPHI 25
#define RZETA 60
#define RTIME 70
#define RSEGM 25

int amplitude[RERRE][RPHI][RZETA][RTIME][RSEGM]; //erre phi zeta time segment      252 MByte

chi2min=1000000000;
for(ierre=4;ierre<=RERRE;ierre+=step1A) {
  for(ipher=1;(ipher <=2+ierre) && (ipher<=RPHI) ;ipher+=step1A) {
    for(izeta=zetar[slice][0];izeta<=zetar[slice][1];izeta+=step1A) {
      for(dt=-rt;dt<=rt;dt++){
        for(itime=10; chi2=0; (itime<=RTIME-10) && (chi2<chi2min); itime+=step1B) {
          for(kk=1; (kk<=segments[slice][0]) ;kk++){
            isegm=segments[slice][kk];
            aa=samples[itime-3+dt][isegm-1];
            bb=amplitude[ierre-1][ipher-1][izeta-1][itime-1][isegm-1];
            if (isegm==ref)
              chi2 += (long) (metrica[aa-bb+5000]/weights[mult-1]);
            else
              chi2 += metrica[aa-bb+5000];
          }
        }
        if(chi2<chi2min) {
          chi2min = chi2;
          erreb = ierre;
          phib = ipher;
          zetab = izeta;
          bestdt = dt;
        }
      }
    }
  }
}

```

* R. Venturelli presentations at AGATA weeks (e.g. Liverpool June 2006) available at [http://www-win.gsi.de/agata/](http://www.win.gsi.de/agata/)

** R. Venturelli, et al., LNL Annual Report 2002, INFN-LNL(REP)198/2003, pp. 154–156

Core of the program

- Just a sequence of loops with almost no calculations
- Runs fast because all structures fit in memory

```
#define RERRE 24
#define RPFI 25
#define RZETA 60
#define RTIME 70
#define RSEGM 25
```

```
int amplitude[RERRE][RPFI][RZETA][RTIME][RSEGM]; //erre phi zeta time segment 252 MByte
```

```
chi2min=1000000000;
```

```
for(ierre=4;ierre<=RERRE;ierre+=step1A){
```

```
  for(iphi=1;(iphi <=2+ierre)&&(iphi<=RPFI) :iphi+=step1A){
```

```
    for(izeta=zetar[slice][0];izeta<=zetar[slice][1];izeta+=step1B){
```

```
      for(dt=-rt;dt<=rt;dt++){
```

```
        for(itime=10, chi2=0; (itime<=RTIME-1) && (chi2<chi2min); itime+=step1B){
```

```
          for(kk=1;(kk<=segments[slice][0]);kk++){
```

```
            isegm=segments[slice][kk];
```

```
            aa=samples[itime-3+dt][isegm-1];
```

```
            bb=amplitude[ierre-1][iphi-1][izeta-1][itime-1][isegm-1];
```

```
            if (isegm==ref)
```

```
              chi2 += (long) (metrica[aa-bb+5000]/weights[mult-1]);
```

```
            else
```

```
              chi2 += metrica[aa-bb+5000];
```

```
          }
```

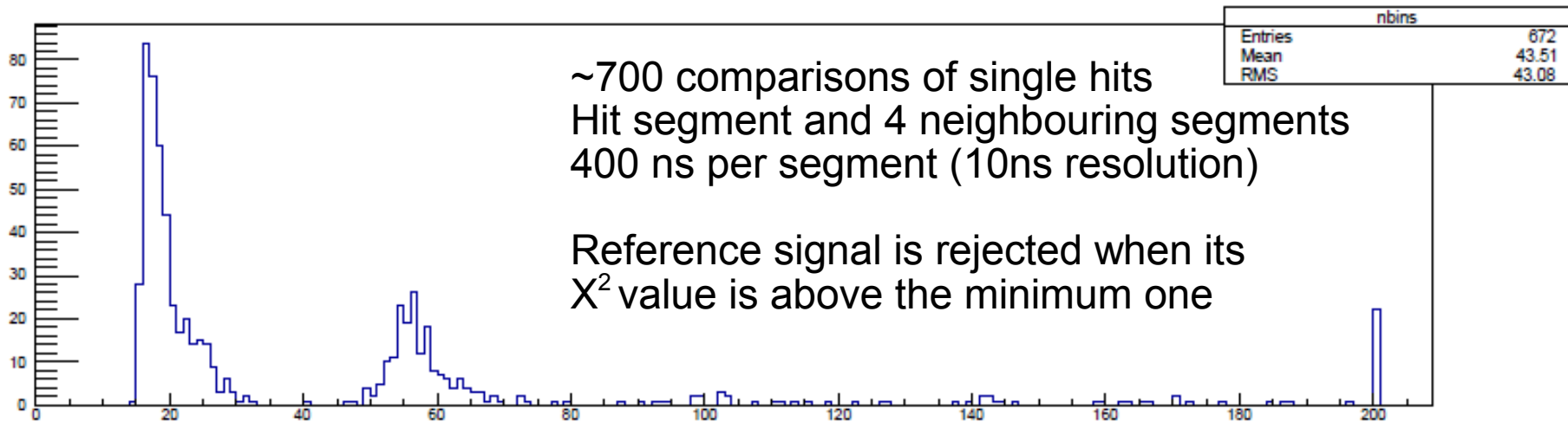
```
        }
      }
    }
  }
}
```

We are not interested in full X^2 distribution, but only in the best fit!
→ Use „adaptive threshold“

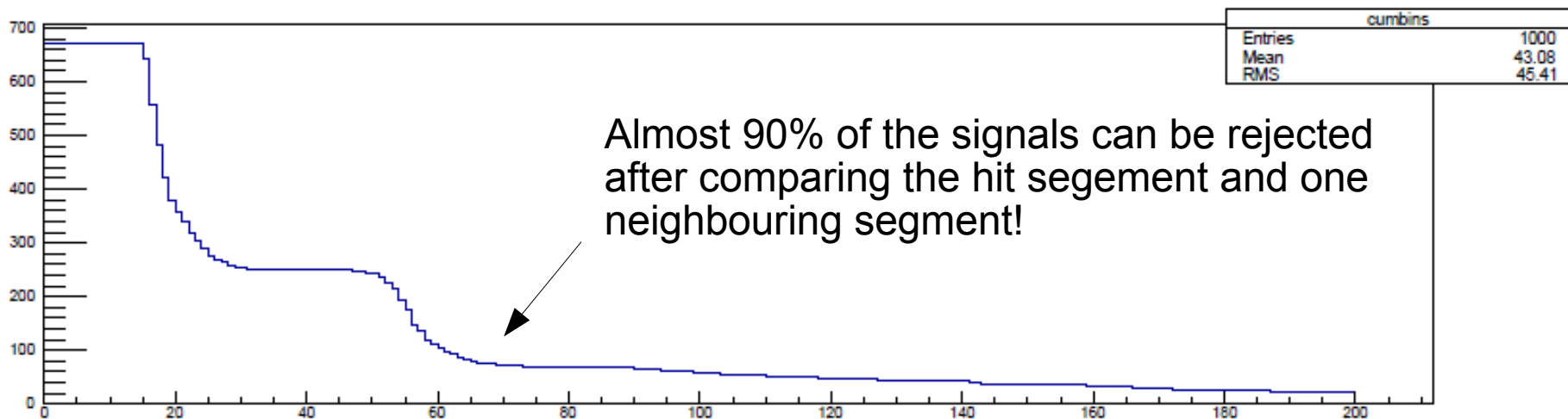
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Pulse Shape Comparison with χ^2 threshold



number of reference signals that can be rejected after comparing x bins



number of best fit candidates after comparing x bins

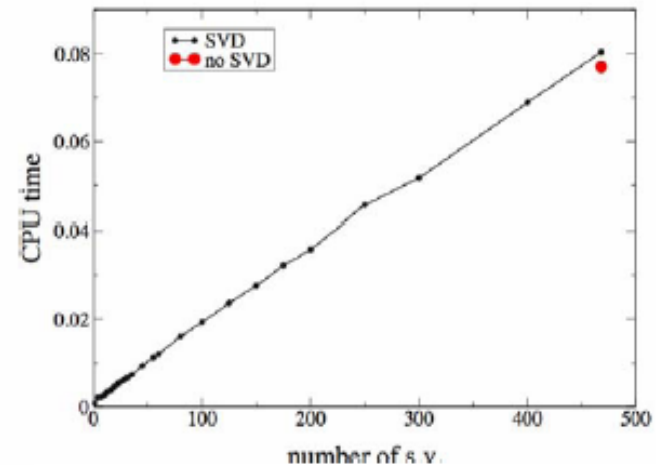
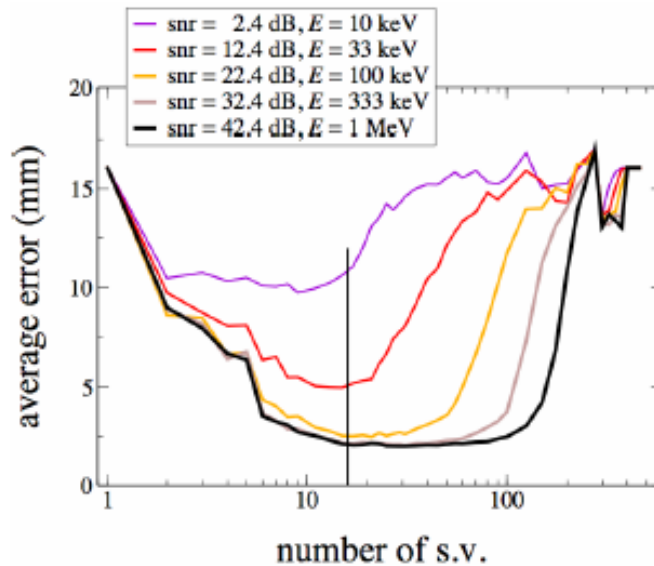
• method

SVD :

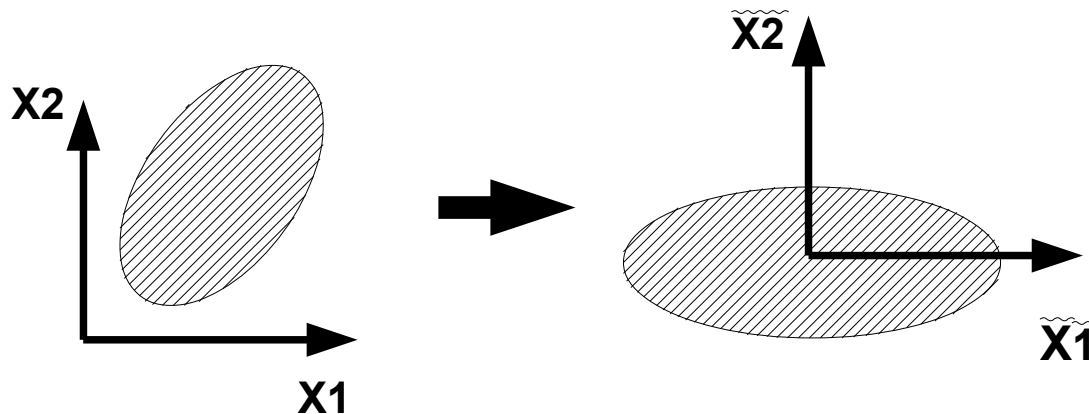
- reduces the size of the signals
- keeping the max. information
- improving the conditioning
- out-of line calculations

- ⇒ reduces the CPU time
- ⇒ no resolution loss
- ⇒ improve resol. and decomposition power

• results



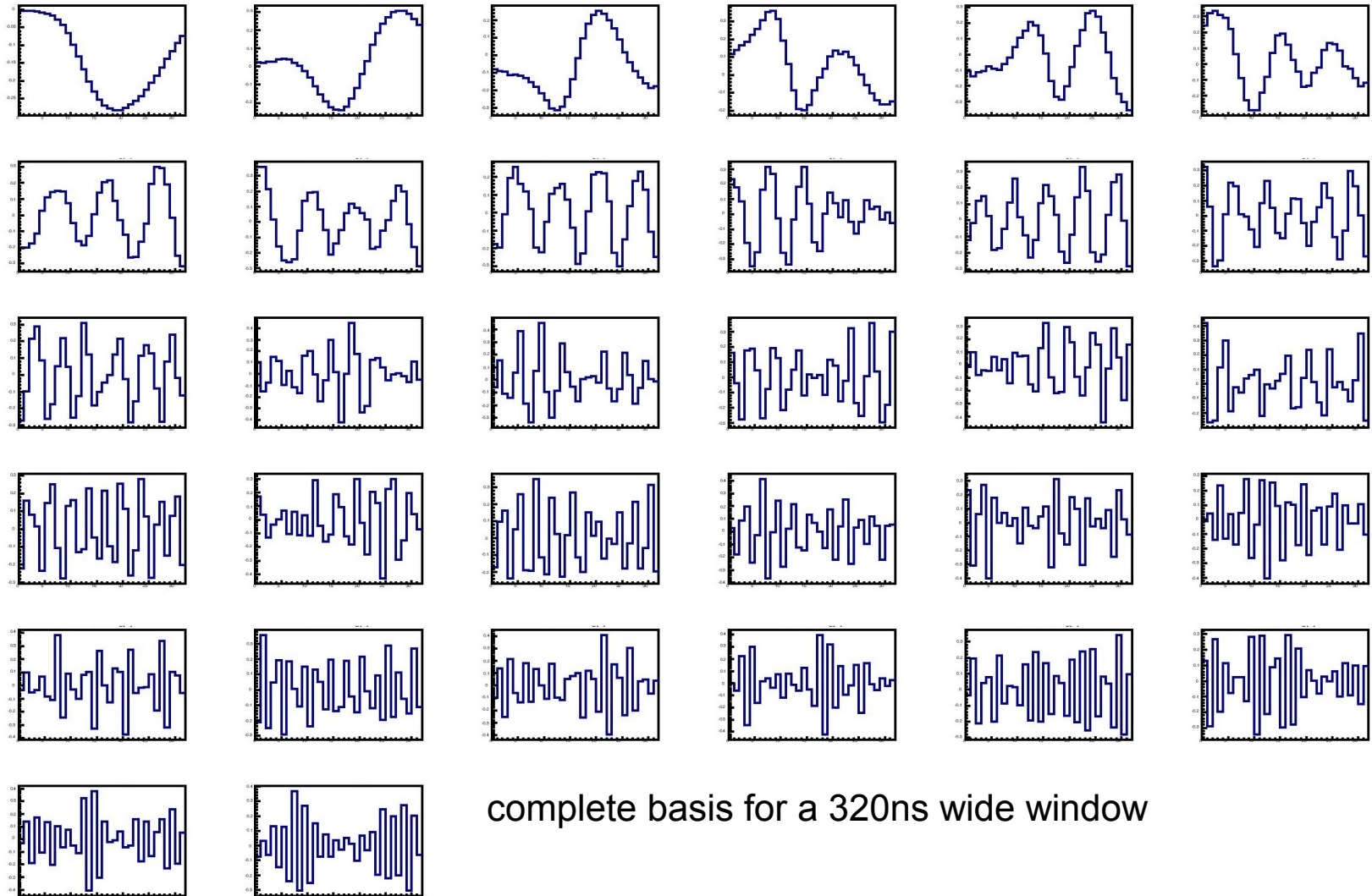
Principal Component Analysis



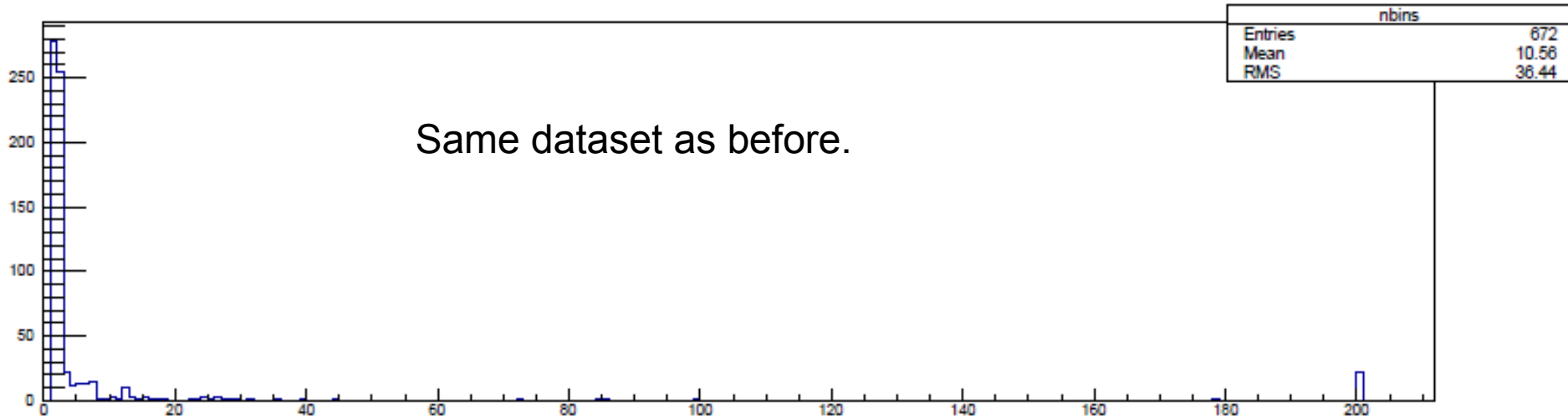
Transform dataset to have maximum information in minimum number of components.

New basis is obtained from singular value decomposition of the set of reference signals (singular vectors).

Singular vectors – Principal components

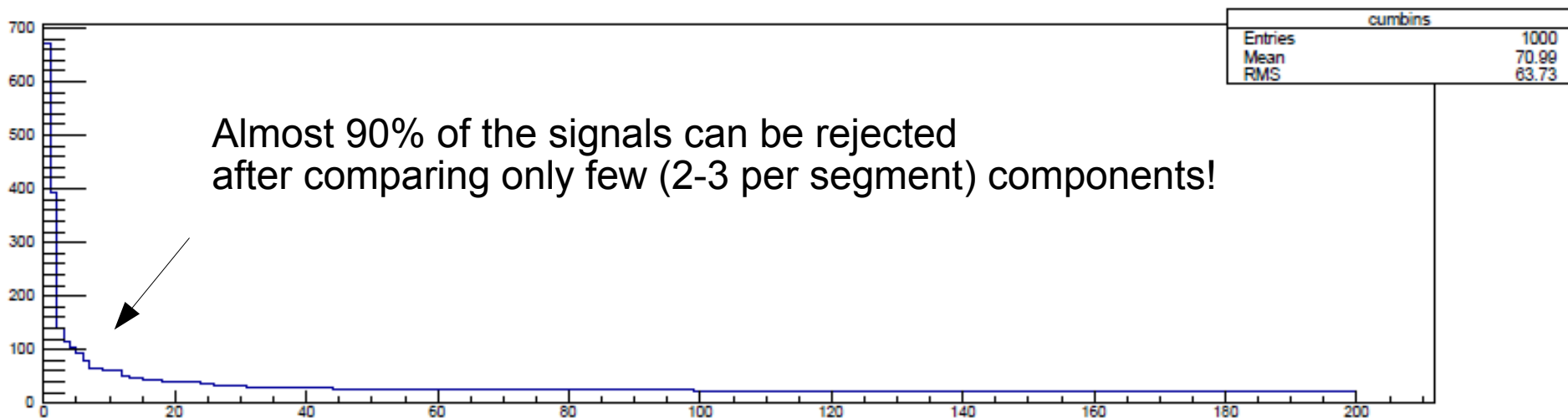


Fast Pulse shape comparison



Same dataset as before.

number of reference signals that can be rejected after comparing x bins



Almost 90% of the signals can be rejected after comparing only few (2-3 per segment) components!

number of best fit candidates after comparing x bins

Procedure

OFFLINE

1. Singular value decomposition of signal basis
→ singular vectors / principal components
2. Transform signal basis

ONLINE

1. Transform measured pulse shape
2. Use Adaptive Grid Search
... but compare the transformed pulse shapes

Procedure

OFFLINE

1. Singular value decomposition of signal basis
→ singular vectors / principal components
2. Transform signal basis

very time
consuming
!!!

ONLINE

1. Transform measured pulse shape
2. Use Adaptive Grid Search
... but compare the transformed pulse shapes

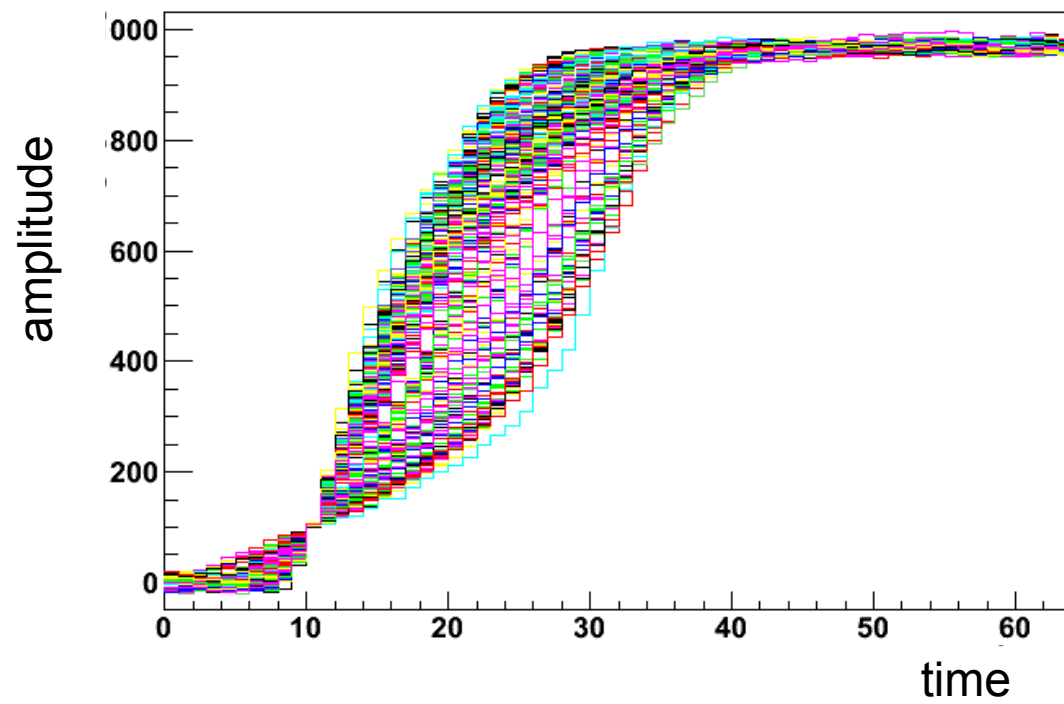
small
overhead

much
faster
!!!

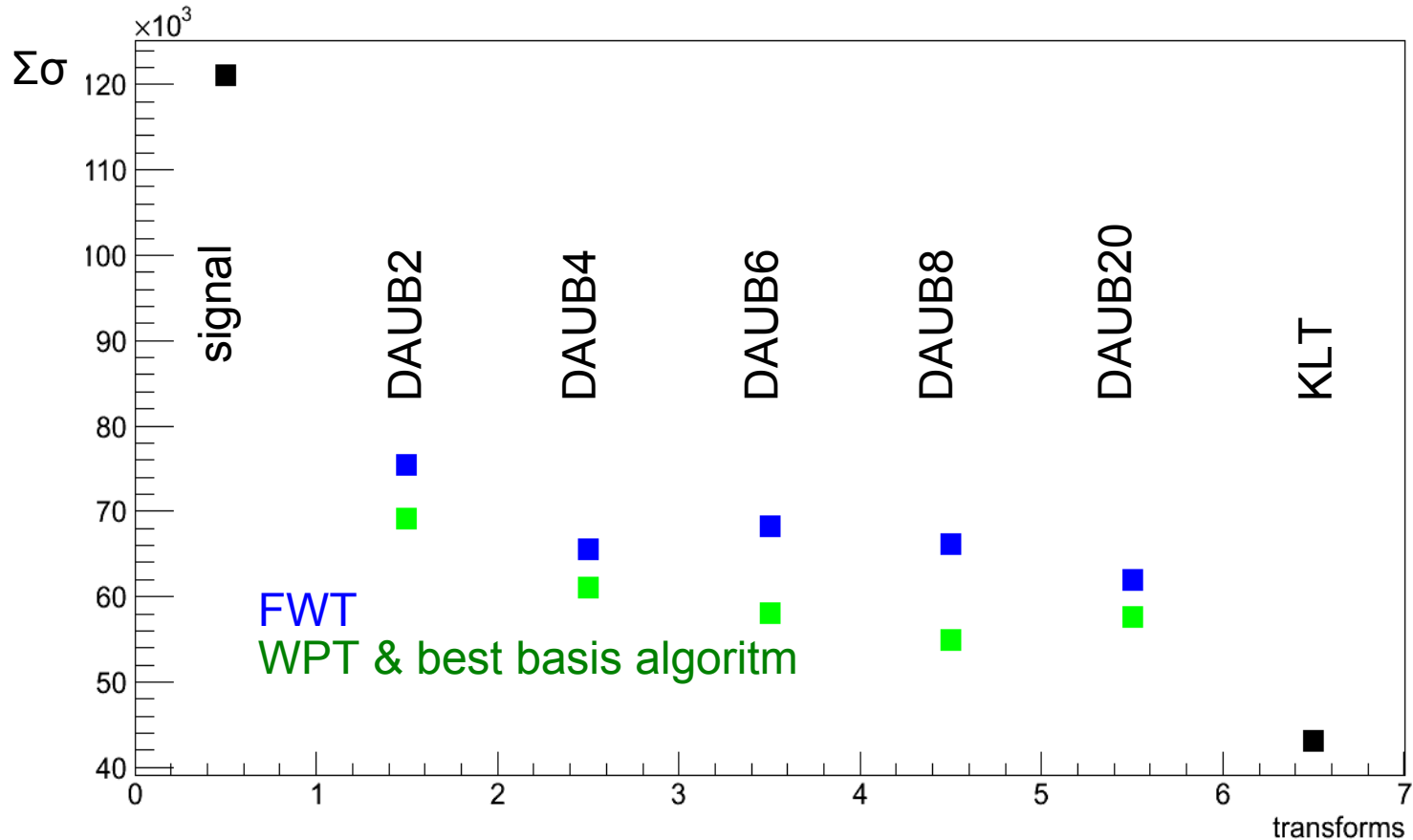
Next steps

- Apply SVD on AGATA signal basis
- Implement Fast Pulse Shape Comparison for NARVAL



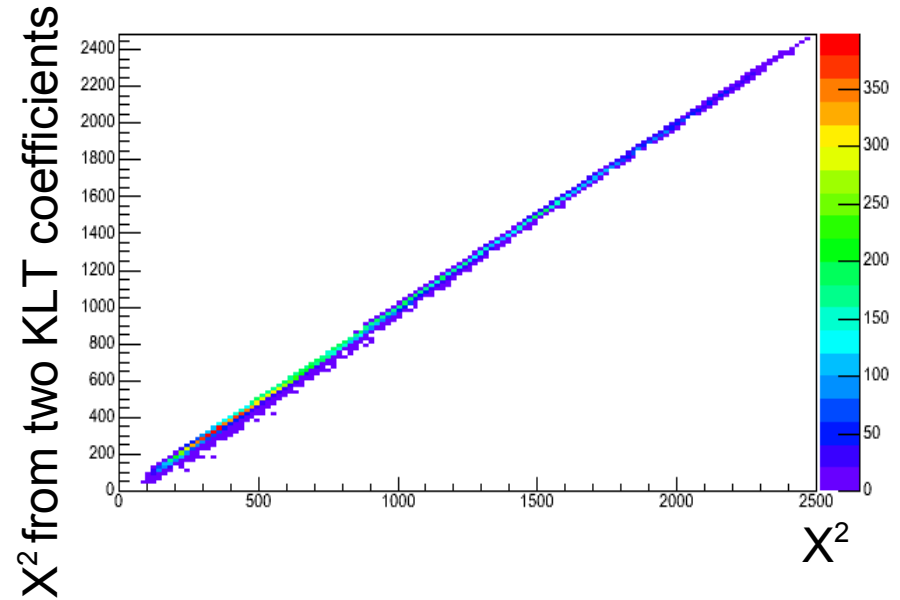
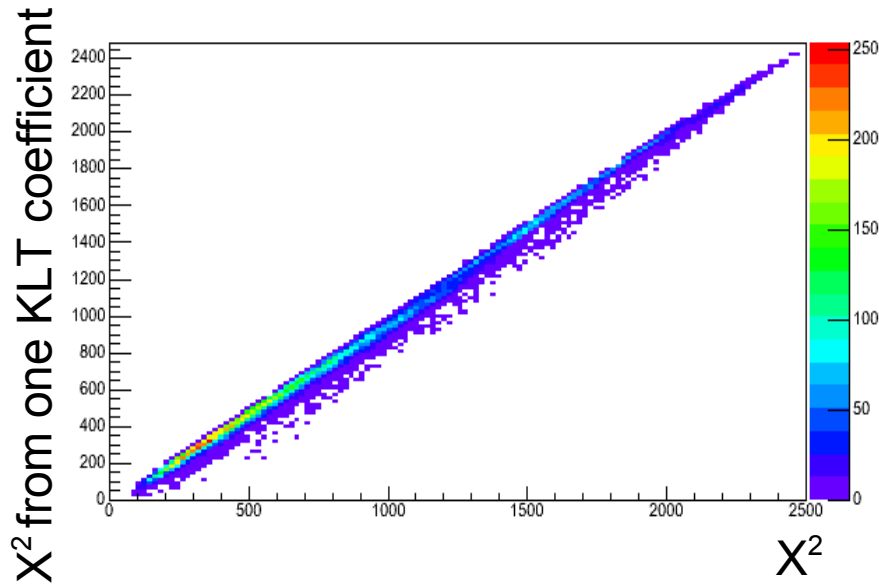


Wavelet vs KLT/Singular value decomposition



→ Karhunen-Loeve-Transform is optimal linear filter for linear approximation

Results



Many basis signals can be rejected already with one coefficient.

Already two coefficients result in a good approximation of x^2 .