

# New for APSY



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What is new:

😊 Finally, it is back 😊

1. Supported since Topspin 3.2pl2
2. Support for Topspin 3.0 & 3.1: on request, contact DMO
3. Matched projection angles for optimum resolution in projections
4. Easy setup using a flowbar interface
5. No license required



## Motivation for using APSY:

1. APSY is a method based on projection spectroscopy
2. 2D-projections are used to describe a nD space
3. Rapid acquisition method
4. Simple experiment setup and analysis

## What do we get from APSY:

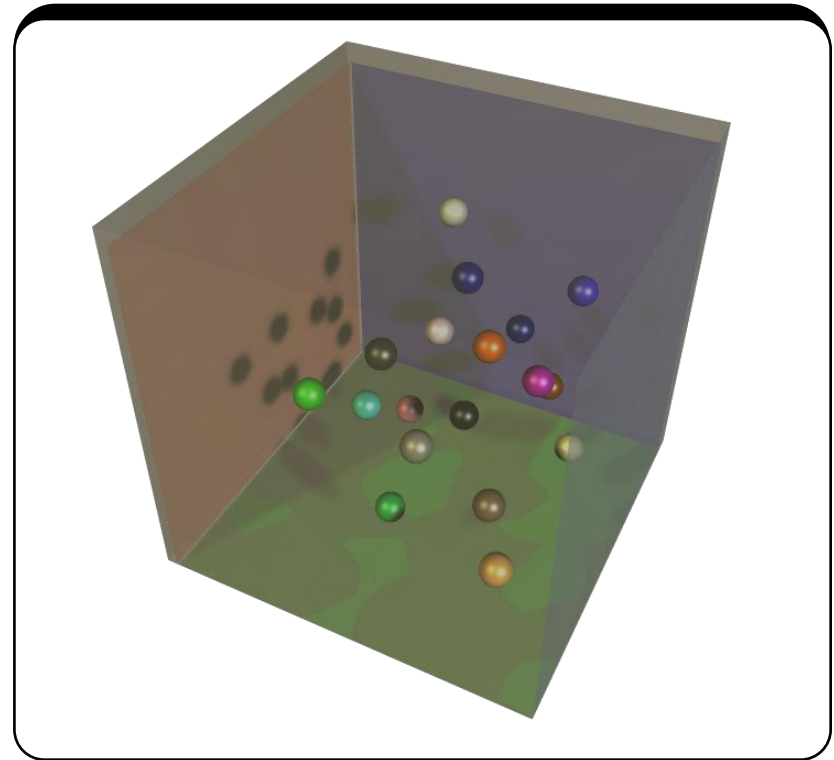
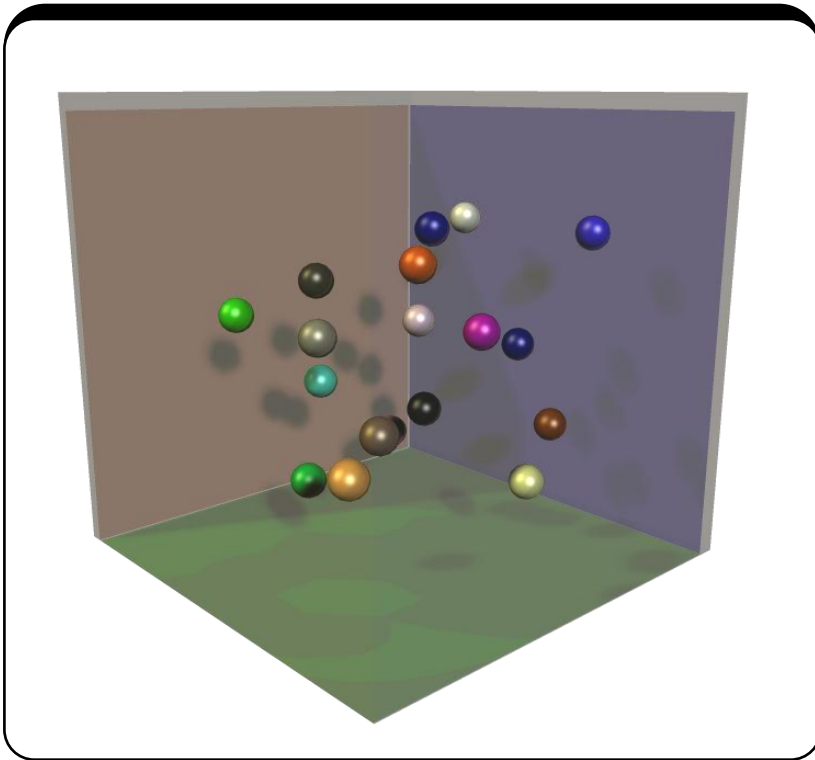
1. A nD peak list, not a spectrum
2. Peak list of high precision
3. Strength of APSY: high precision peak lists

# Projection Spectroscopy



What is 'projection spectroscopy'?

1. **Projections** use different viewpoints to evaluate content of a n-D space

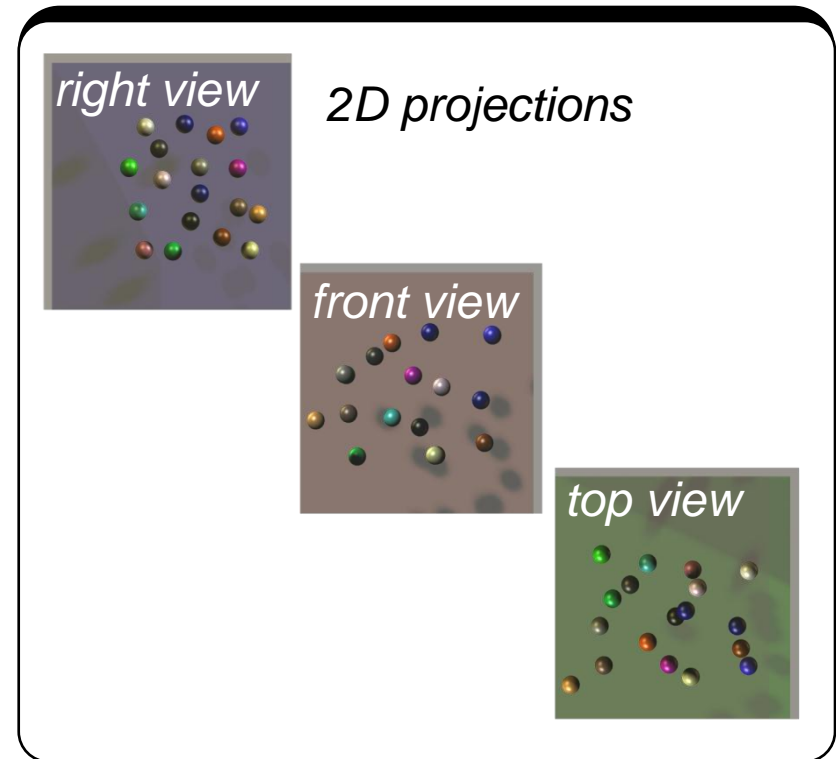
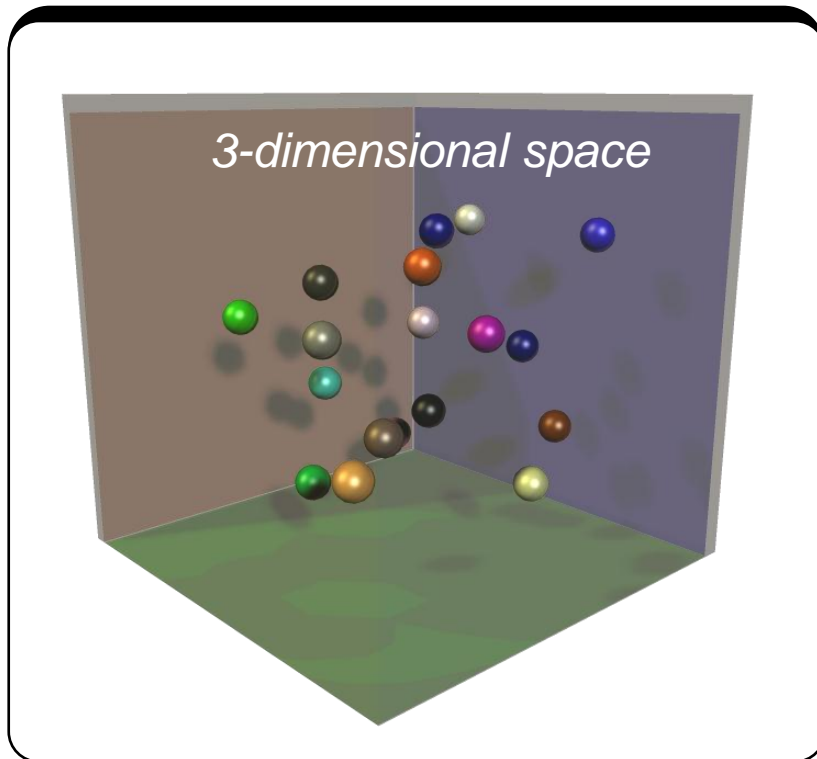


# Projection Spectroscopy



What is 'projection spectroscopy'?

1. **Projections** use different viewpoints to evaluate content of a n-D space
2. **Reduction of dimensionality:** example: 2D-projections for description of a n-D space.

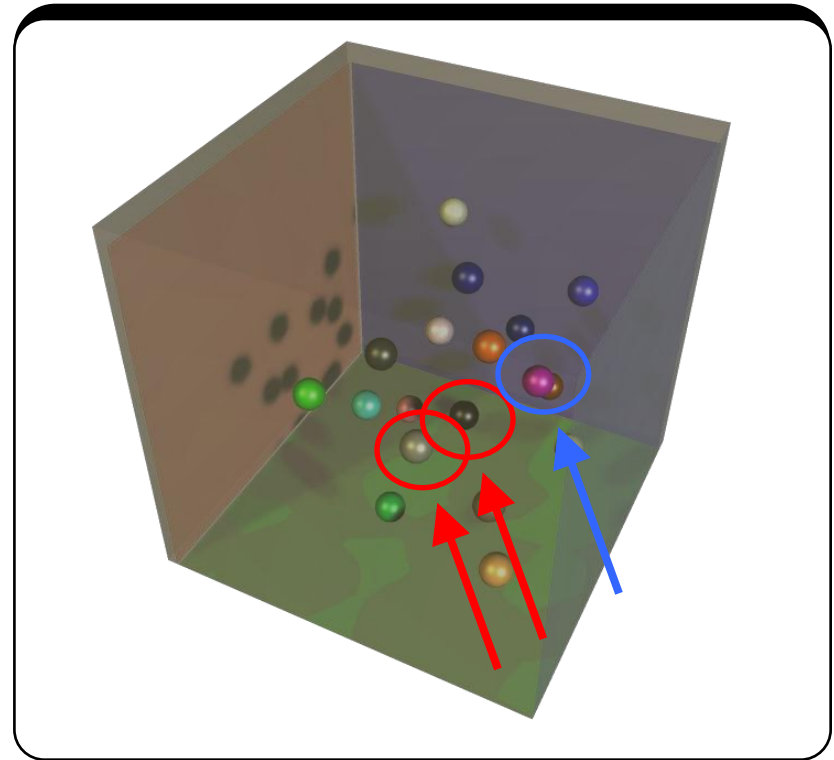
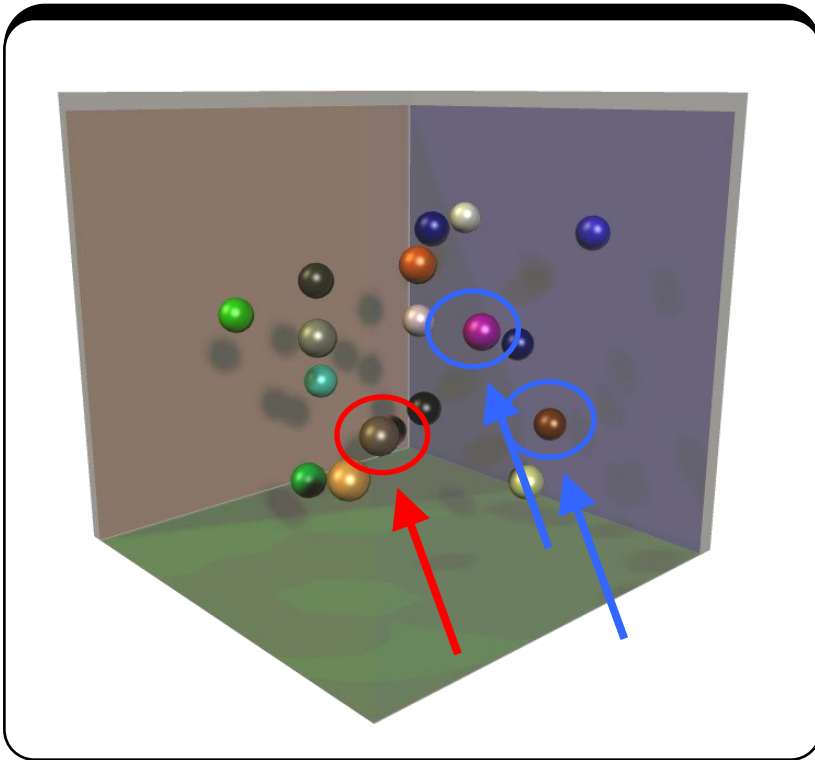


# Projection Spectroscopy



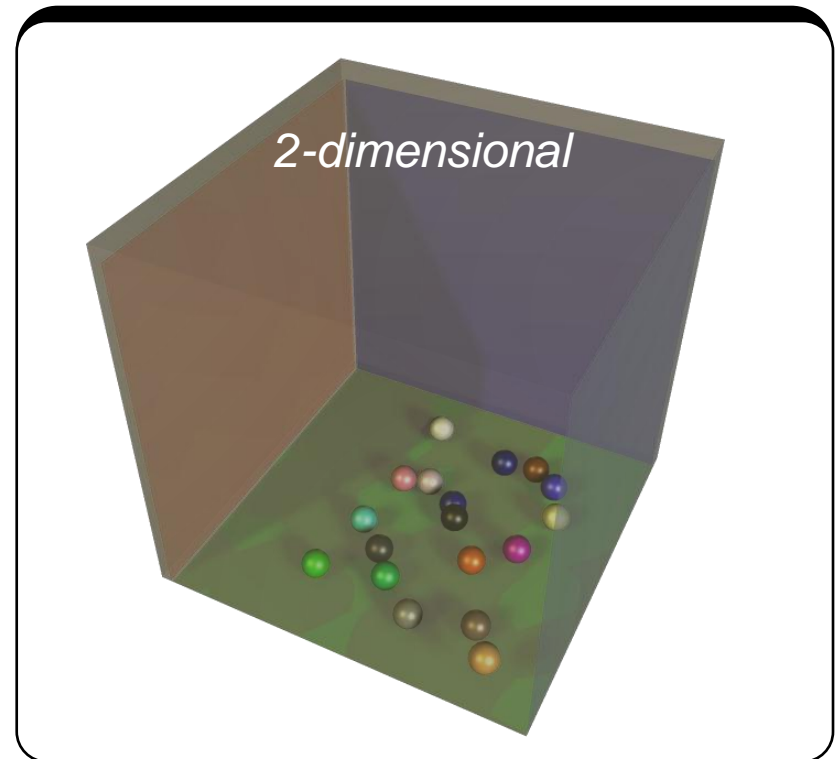
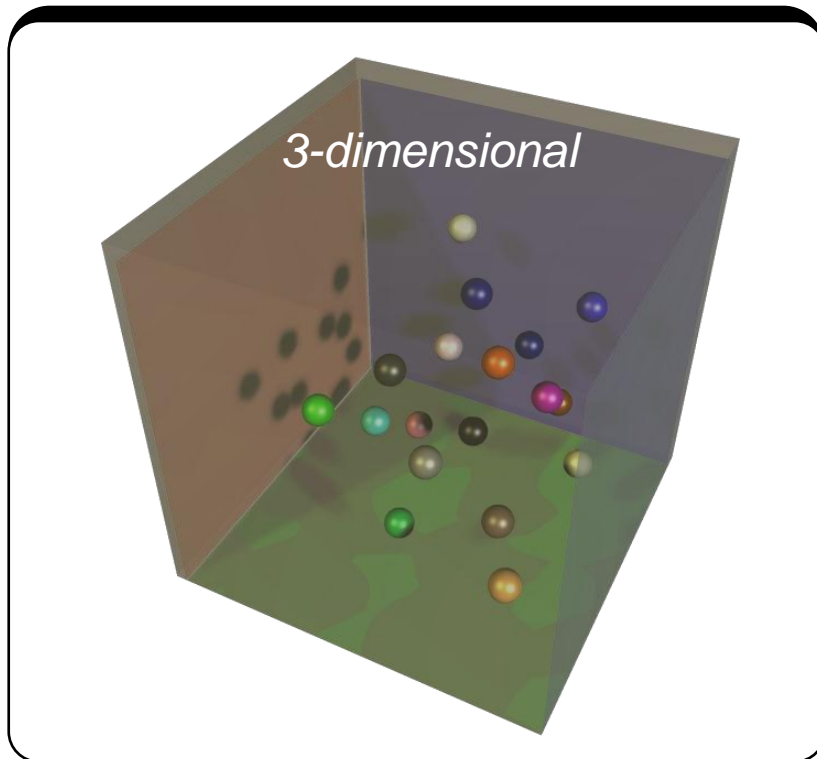
What is 'projection spectroscopy'?

**Projection angles** Multiple viewpoints are required



## **Reduction of dimensionality:**

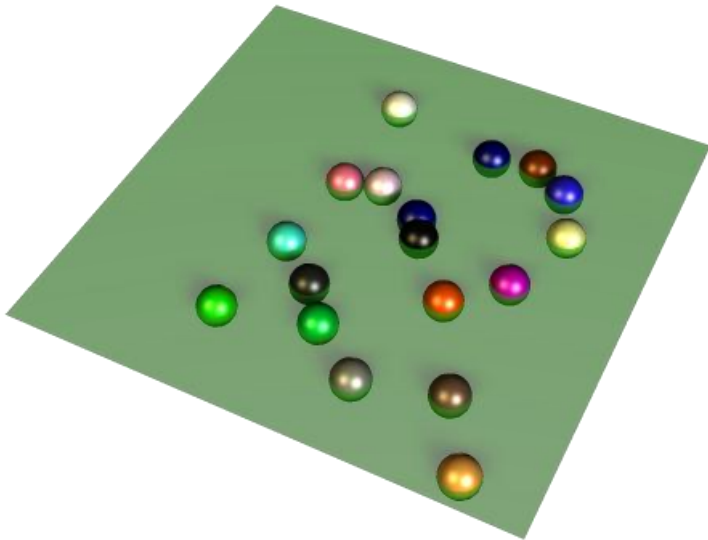
Consequence: Information of additional dimensions is lost



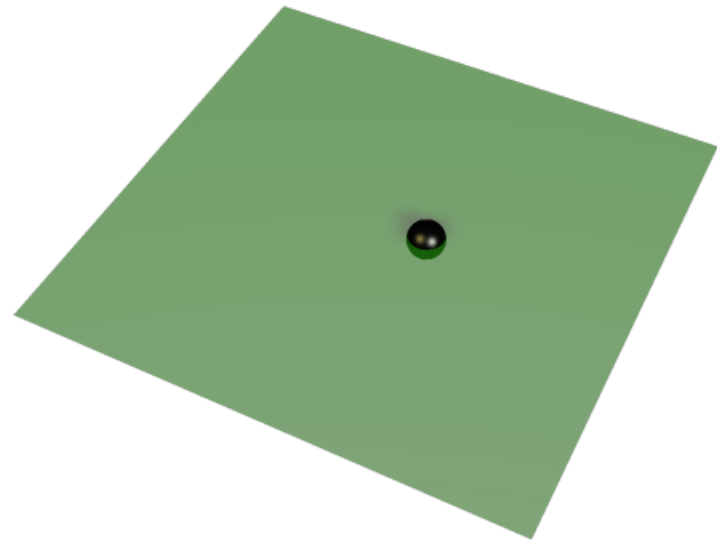
## **Reduction of dimensionality:**

Consequence: Information of additional dimensions is lost

*2-dimensional, all peaks*



*2-dimensional, one selected peak*

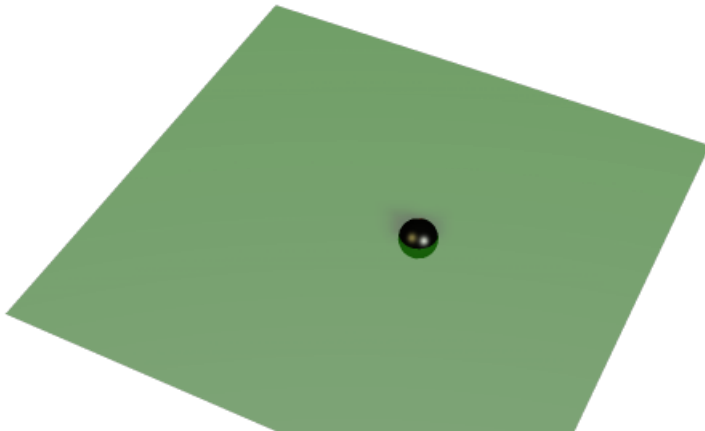




## **Reduction of dimensionality:**

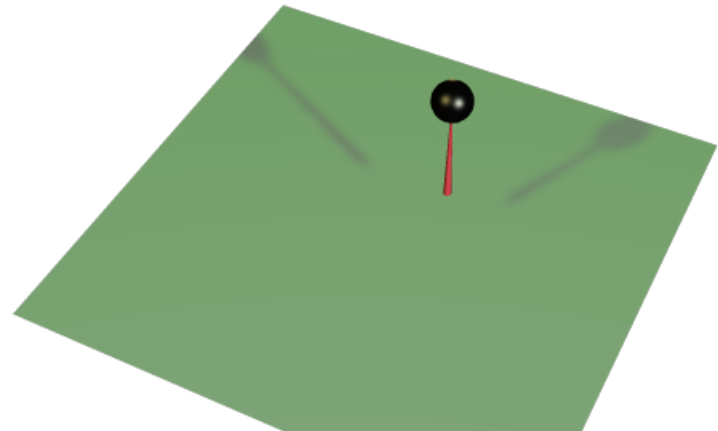
Consequence: Information of additional dimensions is lost

*2-dimensional, one selected peak  
No information about shift in Z*



dimensionality:  
reduced

*2-dimensional, one selected peak  
With information about shift in Z*



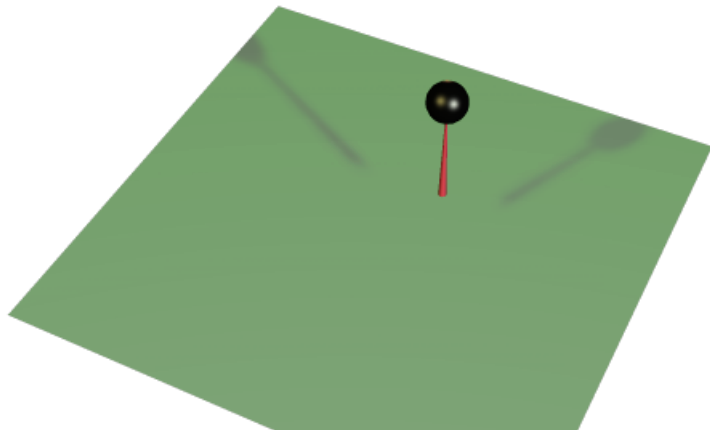
dimensionality:  
full

Consequence of **Reduction of dimensionality**:

Shift information of reduced dimensions is lost, but:

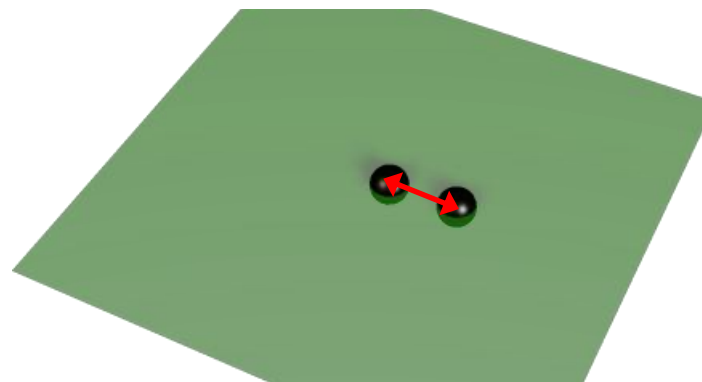
- Shift information is coded as a distance
- By additional splitting of single peaks

*2-dimensional, one selected peak  
With information about shift in Z*



dimensionality:  
full

*2-dimensional, one selected peak  
Information about shift in Z coded  
in a **distance***



dimensionality:  
reduced

# Recording of projection spectra



## Example: 3D HNCO experiment

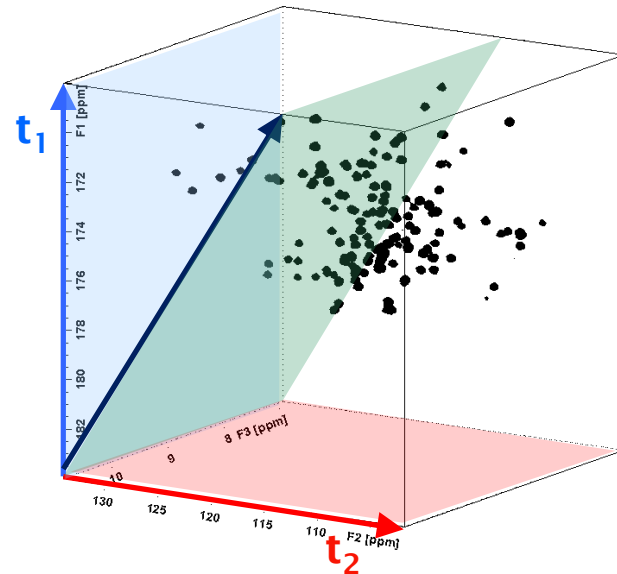
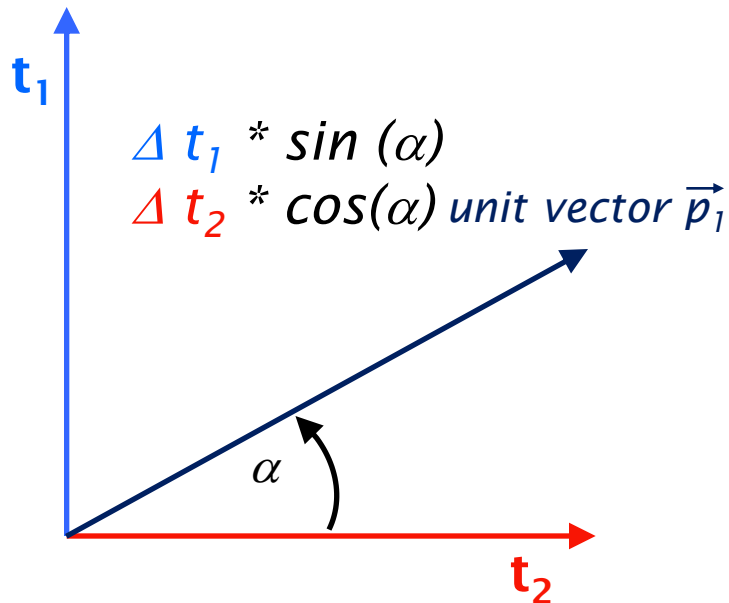
single evolution during  $t_1$  only:

2D H,C plane ( $\alpha = 90^\circ$ )

single evolution during  $t_2$  only:

2D H,N plane ( $\alpha = 0^\circ$ )

simultaneous evolution during  $t_1$  and  $t_2$ : 2D H,NC plane ( $\alpha = \alpha^\circ$ )





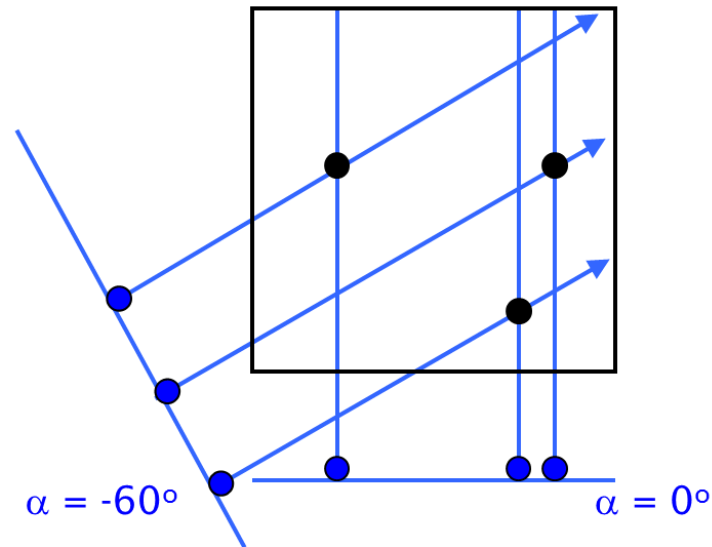
What can be done with the projections?

1. Reconstruct a n-dimensional spectrum:
  - *projection reconstruction*
2. Reconstruct a n-dimensional peak list:
  - *APSY*

# Principle of GAPRO



1. step: two projections are measured and selected arbitrarily:  
e.g.  $0^\circ$  and  $-60^\circ$

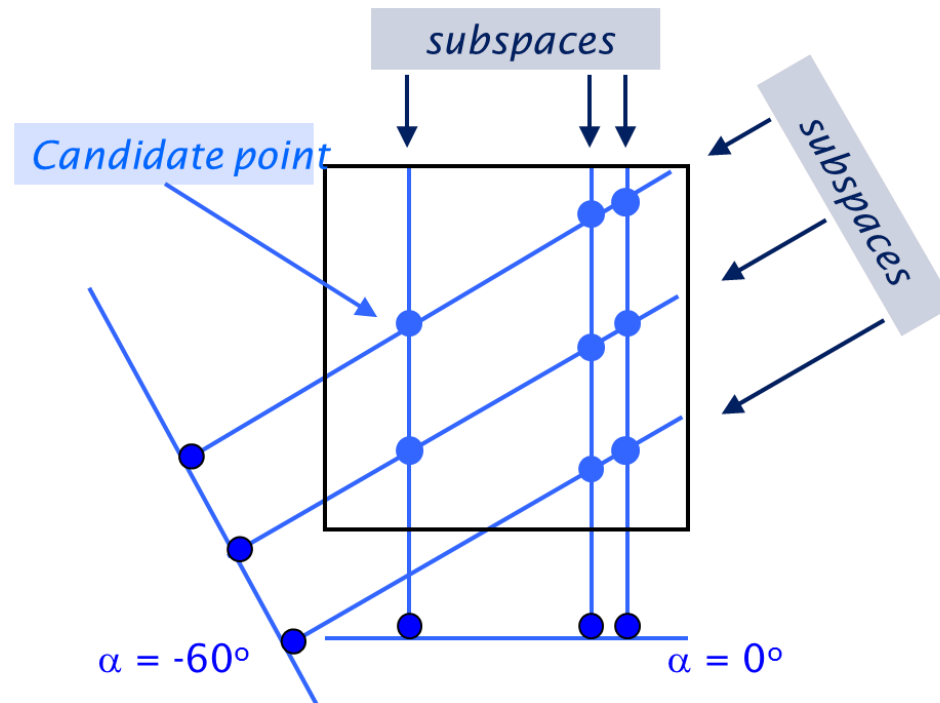


# Principle of GAPRO



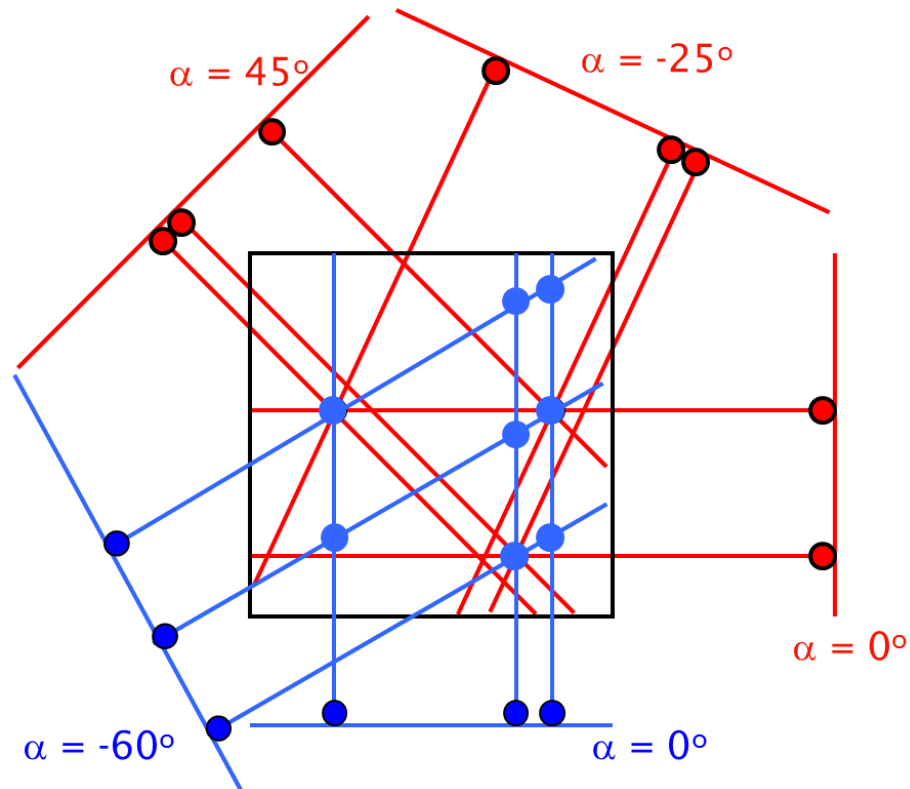
1. step: two projections are selected arbitrarily:  
e.g.  $0^\circ$  and  $-60^\circ$

Intersection of subspaces creates **candidate points**

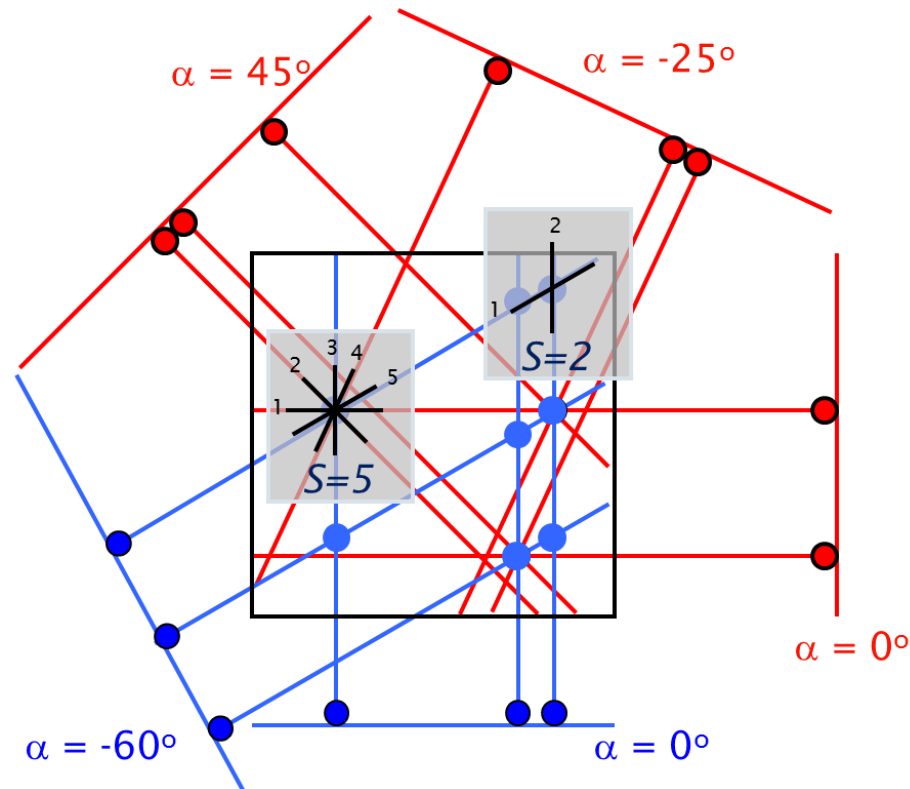


2. step: additional projections included: e.g.  $0^\circ$ ,  $-25^\circ$  and  $45^\circ$

Calculate additional **intersections/subspaces**

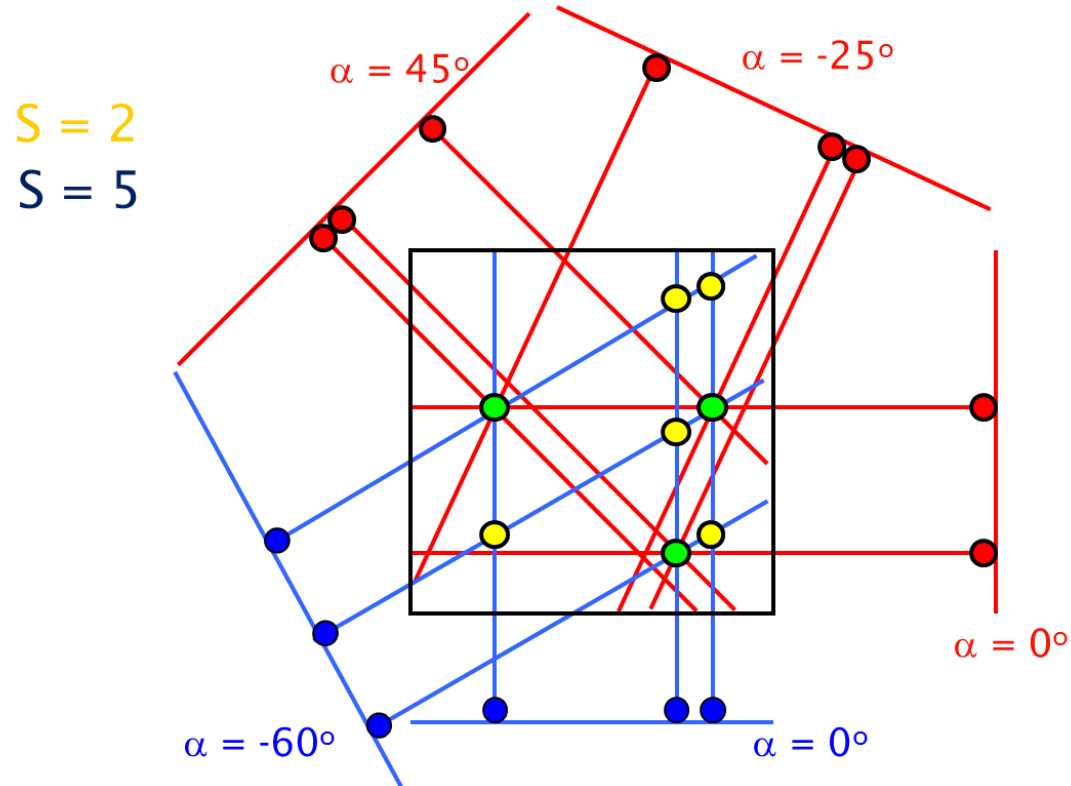


3. step: number of intersecting subspaces (*support S*) is calculated for each candidate point





3. step: *support S*: high values result in high ranking





## APSY as rapid acquisition method

### Conventional

48 and 128 increments  
for dimensions  $t_1$  and  $t_2$   
32 increments for  $t_3$   
16 for  $t_4$ ,  $t_5$  and  $t_6$

### APSY

96 increments for  
all projection angles

### Time saving

| Dimensionality<br>APSY | exp. time [h] | #projections | exp. time [h] | factor conventional : |
|------------------------|---------------|--------------|---------------|-----------------------|
| 3D                     | <b>2</b>      | 20           | <b>0.5</b>    | 4                     |
| 4D                     | <b>54</b>     | 40           | <b>1.0</b>    | 54                    |
| 5D                     | <b>864</b>    | 60           | <b>1.5</b>    | 576                   |
| 6D                     | <b>13824</b>  | 80           | <b>2.0</b>    | 6912                  |
| 7D                     | <b>221184</b> | 100          | <b>2.5</b>    | 88474                 |



## Acquisition parameters for APSY 6,2-HNCOCANH

Bruker TopSpin 3.2.b.29 on B-CH as dmo

Start Acquire Process Analyse Publish View Manage

Back Angles GAPRO Setup Exp-Time Run Stop Re-Process Re-Evaluate Results Help

Del-Analysis

1 APSY\_62\_HNCOCANH 10 1 C:\data\APSY

Spectrum ProcPars AcqPars Title PulseProg Peaks Integrals Sample Structure Plot Fid Acqu

Probe: 5 mm CPTCI 1H/2H-13C/15N Z-GRD Z12345/1

|             | F6                                 | F5         | F4        | F3        | F2         | F1        | Frequency axis                   |
|-------------|------------------------------------|------------|-----------|-----------|------------|-----------|----------------------------------|
| Experiment  |                                    |            |           |           |            |           |                                  |
| PULPROG     | rd_hncocanh_62                     |            |           |           |            |           | Current pulse program            |
| AQ_mod      | DQD                                |            |           |           |            |           | Acquisition mode                 |
| FnMODE      | States States States States States |            |           |           |            |           | Acquisition mode for 2D, 3D etc. |
| FnTYPE      | traditional(planes)                |            |           |           |            |           | nD acquisition mode for 3D etc.  |
| TD          | 2048                               | 48         | 128       | 40        | 48         | 48        | Size of fid                      |
| DS          | 16                                 |            |           |           |            |           | Number of dummy scans            |
| NS          | 16                                 |            |           |           |            |           | Number of scans                  |
| TD0         | 1                                  |            |           |           |            |           | Loop count for 'td'              |
| Width       |                                    |            |           |           |            |           |                                  |
| SW [ppm]    | 16.0182                            | 35.5073    | 32.9916   | 11.1297   | 35.5073    | 3.3984    | Spectral width                   |
| SWH [Hz]    | 8012.820                           | 1800.000   | 4150.000  | 1400.000  | 1800.000   | 1700.000  | Spectral width                   |
| IN_F [µsec] | 555.56                             |            |           |           |            |           | Increment for delay              |
| AQ [sec]    | 0.1277952                          | 0.01333333 | 0.0154217 | 0.0142857 | 0.01333333 | 0.0141176 | Acquisition time                 |
| FIDRES [Hz] | 7.825020                           | 75.000000  | 64.843750 | 70.000000 | 75.000000  | 70.833336 | Fid resolution                   |
| FW [Hz]     | 125000.000                         |            |           |           |            |           | Filter width                     |
| Receiver    |                                    |            |           |           |            |           |                                  |
| RG          | 144                                |            |           |           |            |           | Receiver gain                    |
| DW [µsec]   | 62.400                             |            |           |           |            |           | Dwell time                       |

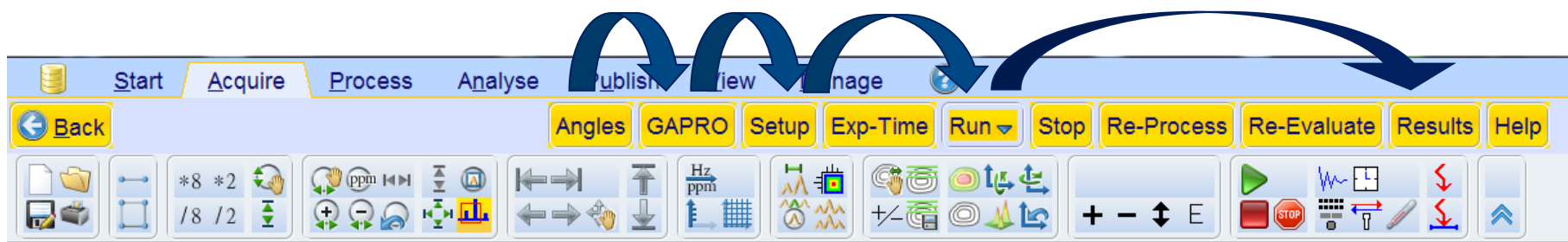
# APSY experiments and software



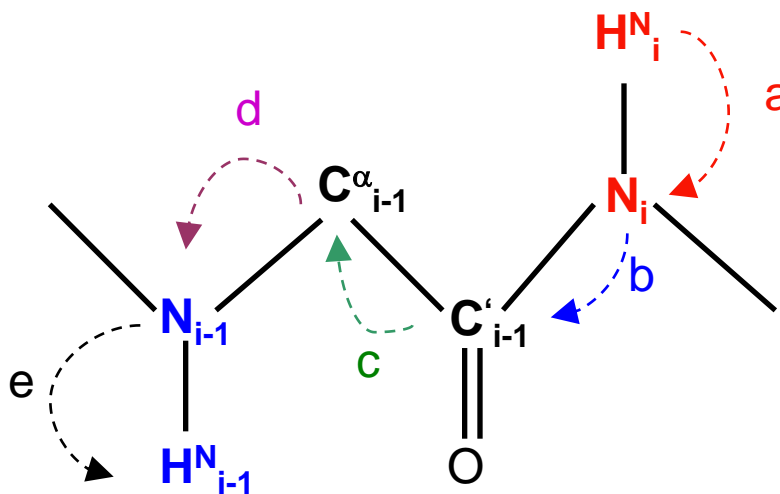
## APSY flow bar

- Define projection angles
- Define GAPRO processing parameters
- Run APSY experiment
- Reprocessing of projections
- Reevaluation of peak lists

1. Set projection angles
2. Set GAPRO parameters
3. Acquire
4. Result



6D sequential  $H_i-N_i-CO_{i-1}-CA_{i-1}-N_{i-1}-H_{i-1}$



# APSY: the peak list

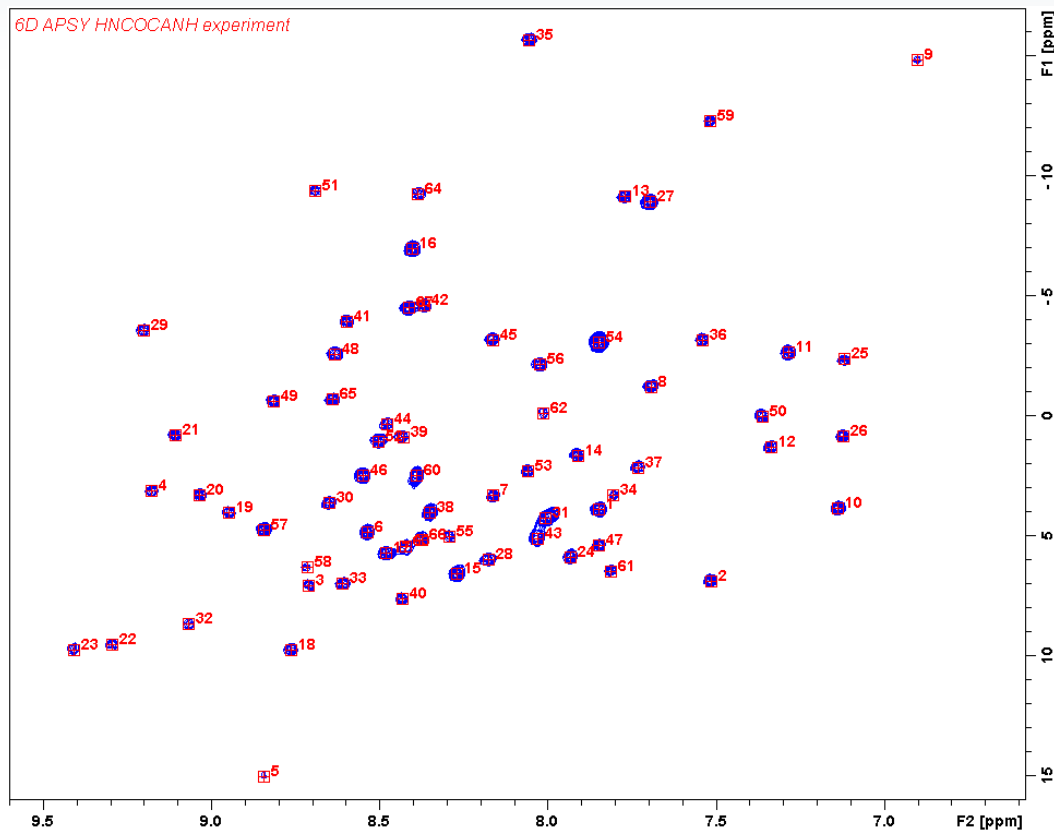


- High Precision: peak lists of high precision from an optimum number of projections.

| $H_{i-1}$     | $N_{i-1}$       | CO       | CA      | $N_i$           | $H_i$         | $\delta(N_{i-1}-N_i)$ |
|---------------|-----------------|----------|---------|-----------------|---------------|-----------------------|
| ppm           | ppm             | ppm      | ppm     | ppm             | ppm           | Hz                    |
| 8.0613        | 102.4332        | 177.1224 | 52.3724 | <b>133.0131</b> | <b>8.8405</b> | -4.635                |
| <b>8.8578</b> | <b>133.1058</b> | 174.3129 | 56.454  | <b>125.1379</b> | <b>8.7061</b> | 1.415                 |
| <b>8.7125</b> | <b>125.1096</b> | 175.6223 | 58.7179 | <b>122.0751</b> | <b>8.9428</b> | -1.24                 |
| <b>8.9591</b> | <b>122.0999</b> | 174.9824 | 52.7983 | <b>124.366</b>  | <b>8.7145</b> | -4.17                 |
| <b>8.716</b>  | <b>124.4494</b> | 173.6779 | 54.8708 | <b>123.1657</b> | <b>8.378</b>  | ---                   |

Sequential assignment of  $[^{13}\text{C}, ^{15}\text{N}]$ -ubiquitin using the peak list from a 6,2-APSY-HNCOANH experiment.

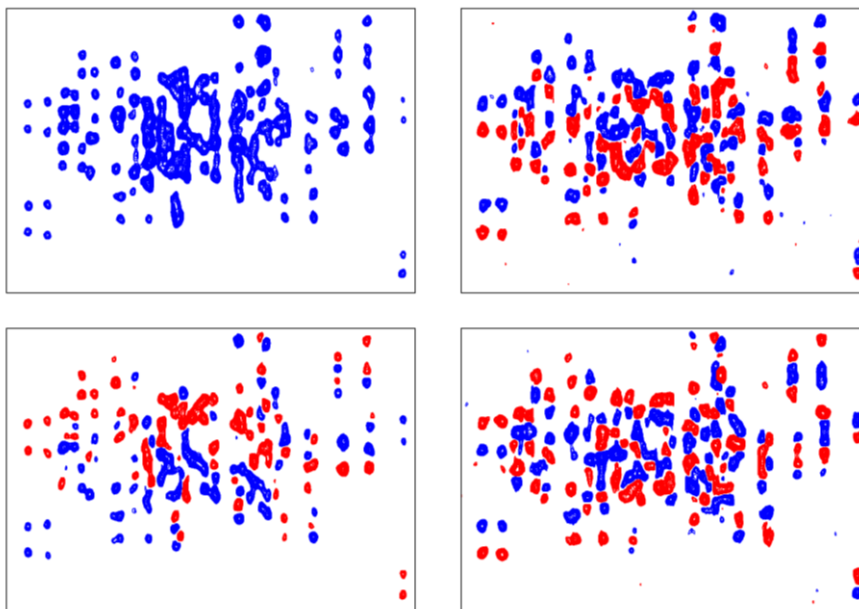
# APSY: the peak list



*2D projection of a 6,2-APSY-HNCOCANH experiment. The displayed peak list was calculated from the 6D peak list and displayed on the projection*

## Challenge / limitations:

- Signal loss for experiments of high dimensionality: relaxation
- Increased complexity of spectra due to simultaneous incrementation and multiple quadrature



4,2-HNCOCA  
projection angles:  
 $\alpha, \beta \neq 0^\circ, 90^\circ$



**Challenge:** APSY peak lists with high precision even if recorded on different:

- Spectrometers
- Date

## Solution

- NMR-Thermometer!

## Example

- 4D HNCOCA APSY
- Recorded @ 600 and 800MHz
- Compare peak lists
- Sample: 0.4mM  $^{13}\text{C}$ ,  $^{15}\text{N}$  ubiquitin in 90%  $\text{H}_2\text{O}$ , 10%  $\text{D}_2\text{O}$ , NMR Thermometer compound: 10mM TSP-d<sub>13</sub>

# APSY and the NMR-Thermometer



## Comparison of 4,2 HNCOCA APSY peak lists, 600 & 800MHz

0.4mM  $^{13}\text{C}$ ,  $^{15}\text{N}$  ubiquitin in 90%  $\text{H}_2\text{O}$ , 10%  $\text{D}_2\text{O}$ , NMR-Thermometer compound: 10mM TSP- $\text{d}_{13}$

| $\Delta[\text{ppm}]$ 800-600MHz |                   |                   |                   | $\Delta[\text{Hz @800MHz}]$ 800-600MHz |                |                   |                   |                   |                   |
|---------------------------------|-------------------|-------------------|-------------------|--|----------------|-------------------|-------------------|-------------------|-------------------|
| N                               | CA                | Ca                | H                 | N                                      | CA             | Ca                | H                 |                   |                   |
| 0.0034                          | -0.0112           | 0.0158            | 0                 | 0.27574                                | -2.2512        | 3.1758            | 0                 |                   |                   |
| -0.0058                         | -0.0024           | 0.0098            | 0.0014            | -0.47038                               | -0.4824        | 1.9698            | 1.120182          |                   |                   |
| 0.0038                          | -0.0047           | 0.0061            | 0.0002            | 0.30818                                | -0.9447        | 1.2261            | 0.160026          |                   |                   |
| -0.0026                         | -0.0112           | -0.0067           | -0.001            | -0.21086                               | -2.2512        | -1.3467           | -0.80013          |                   |                   |
| -0.0132                         | -0.0166           | -0.003            | 0.0005            | -1.07052                               | -3.3366        | -0.603            | 0.400065          |                   |                   |
| -0.0032                         | 0.0012            | -0.0442           | 1E-04             | -0.25952                               | 0.2412         | -8.8842           | 0.080013          |                   |                   |
| -0.0103                         | -0.0048           | 0.0165            | -0.0005           | -0.83533                               | -0.9648        | 3.3165            | -0.400065         |                   |                   |
| 0.0019                          | -0.0116           | 0.0313            | -0.0006           | 0.15409                                | -2.3316        | 6.2913            | -0.480078         |                   |                   |
| -0.0153                         | -0.0051           | 0.0069            | 0.0005            | -1.24083                               | -1.0251        | 1.3869            | 0.400065          |                   |                   |
| -0.0189                         | -0.0051           | -0.0045           | -0.0008           | -1.53279                               | -1.0251        | -0.9045           | -0.640104         |                   |                   |
| -0.0033                         | -0.0102           | 0.0141            | 0.0001            | -0.26763                               | -2.0502        | 2.8341            | 0.080013          |                   |                   |
| 0.002                           | -0.0224           | -0.0122           | -0.0008           | 0.1622                                 | -4.5024        | -2.4522           | -0.640104         |                   |                   |
| 0.0095                          | -0.0146           | -0.0085           | 0.0006            | 0.77045                                | -2.9346        | -1.7085           | 0.480078          |                   |                   |
| -0.0017                         | -0.0128           | 0.0328            | -0.0004           | -0.13787                               | -2.5728        | 6.5928            | -0.320052         |                   |                   |
| -0.0005                         | -0.0141           | 0.009             | -0.0003           | -0.04055                               | -2.8341        | 1.809             | -0.240039         |                   |                   |
| -0.0092                         | -0.0078           | 0.0056            | 0.0007            | -0.74612                               | -1.5678        | 1.1256            | 0.560091          |                   |                   |
| -0.0253                         | -0.0136           | 0.0315            | 0                 | -2.05183                               | -2.7336        | 6.3315            | 0                 |                   |                   |
| <b>STD/ppm:</b>                 | <b>0.03086873</b> | <b>0.00776172</b> | <b>0.02262312</b> | <b>0.00065478</b>                      | <b>STD/Hz:</b> | <b>2.50345434</b> | <b>1.56010552</b> | <b>4.54724794</b> | <b>0.52391188</b> |

Precision is in the order of the Hz/Pt resolution used for processing:

$SI[F2] = 8\text{K}$ ;  $SI[F1] = 1\text{K}$ ;

$\text{Hz/Pt}[F2]=0.78\text{Hz}$ ;  $\text{Hz/Pt}[F1]=1.4 - 3.5\text{Hz}$ ;



**GAPRO:** peak picking, geometrical analysis, matched projection angles

1. Calculation of projection angles will be transferred to **manageapsy**

**manageapsy:** main program controlling setup, run, analysis & report

1. Calculation of matched projection angles: optimum resolution for indirect dimensions

**Pulse programs:**

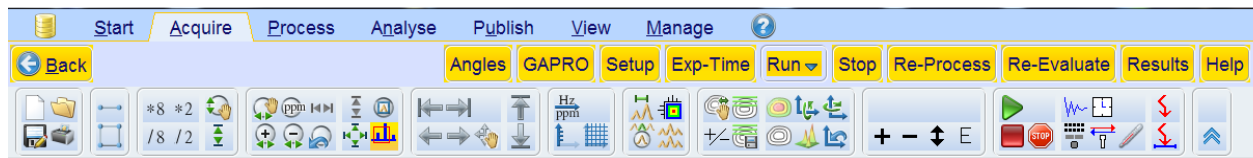
1. Implementation of missing experiments

**All:** adaption to future Topspin versions

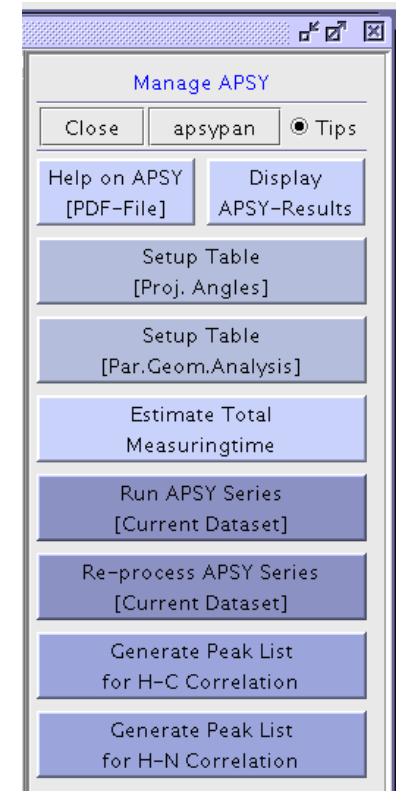


Before I forget...How to start APSY:

“apsy”



“bpan apsypan”



## Acknowledgement

- Gerhard Wider, Barbara Krähenbühl: ETH Zürich
- Sebastian Hiller: Biozentrum, University Basel
- Werner Mausshard, Wolfgang Bermel, Frank Schumann