

Covariance NMR

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Covariance NMR



- Resolution enhancement along indirect dimension
- “Transfer” resolution of direct dimension to indirect dimension
 - Result: Fully symmetric spectrum
- Singular value decomposition (SVD) of matrix square-root
 - Computationally efficient
- Amenable to wide range of solids and liquids applications

Covariance NMR



Principle:

After FT in direct dimension:

- Compare all columns (indirect dimension)
- For columns which are similar:
 - transfer info of frequency in direct dimension to the indirect dimension

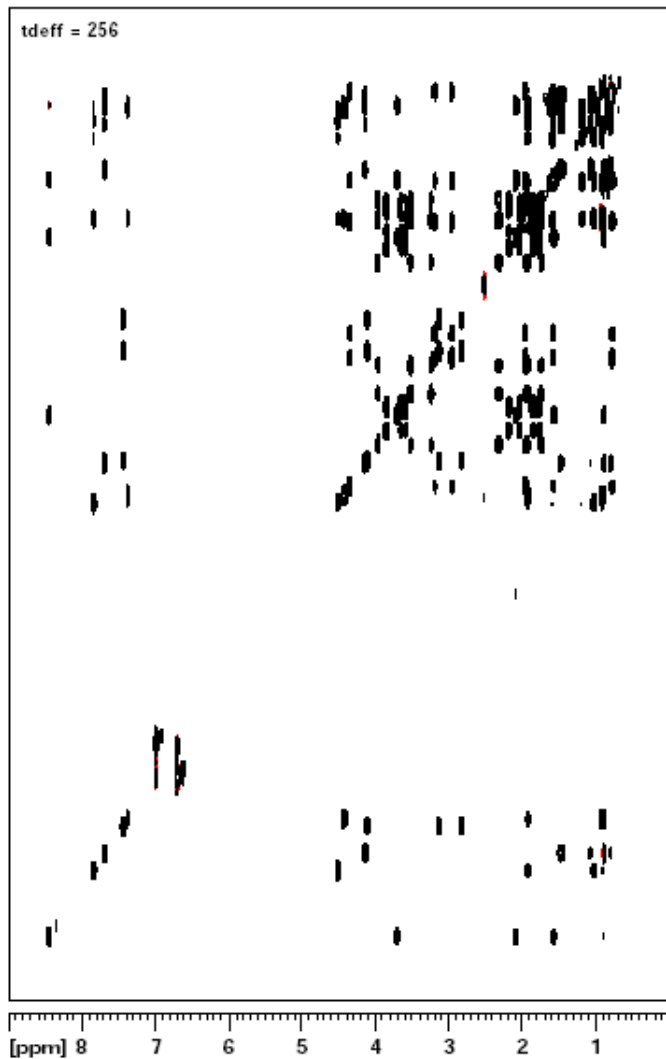
No further FT applied

Advantage: enhanced resolution in indirect dimension

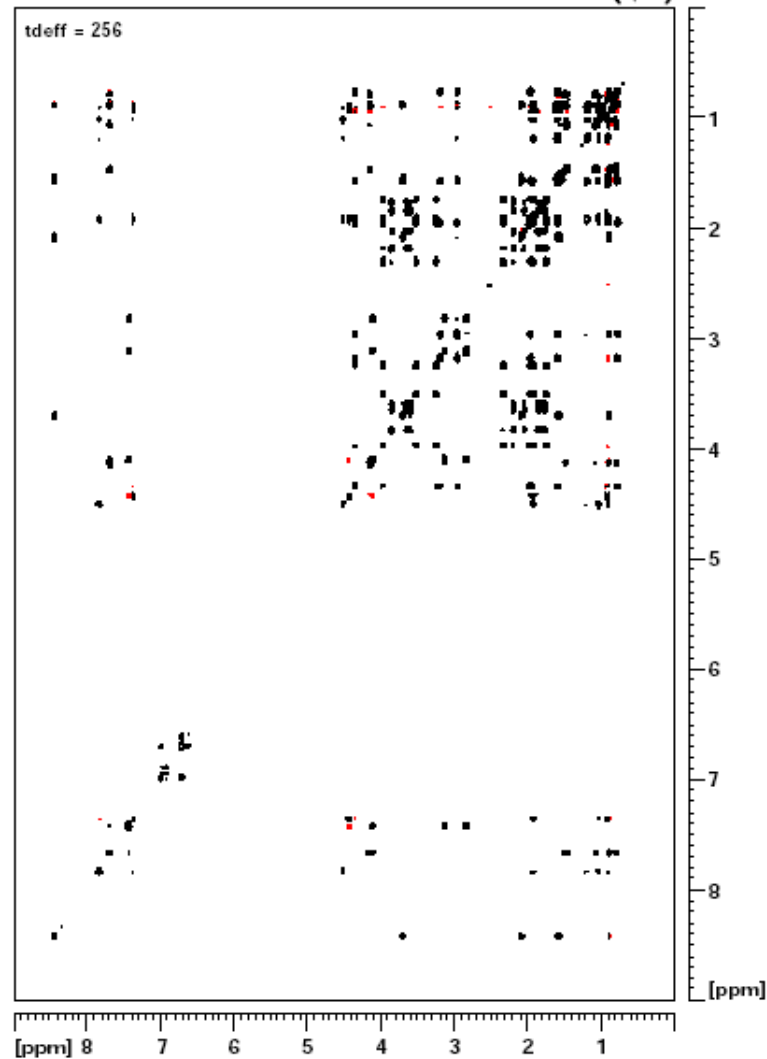
Covariance NMR



DIPS12 xfb



DIPS12 xf2 + covariance (\sqrt{C})



Covariance NMR



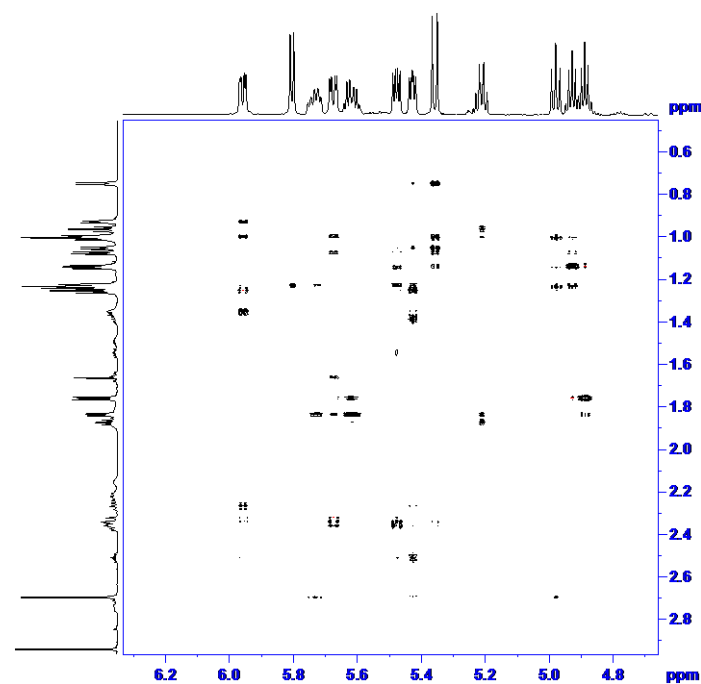
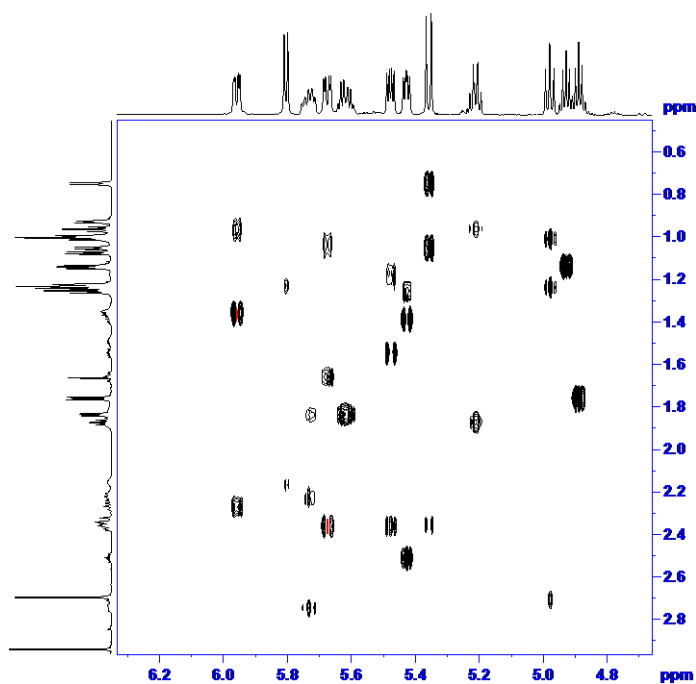
Application: Resolution enhancement

Example: Cyclosporine

TOCSY TD (f1) = 128 \rightarrow SI = 2kx2k

FT

Covariance



Covariance NMR



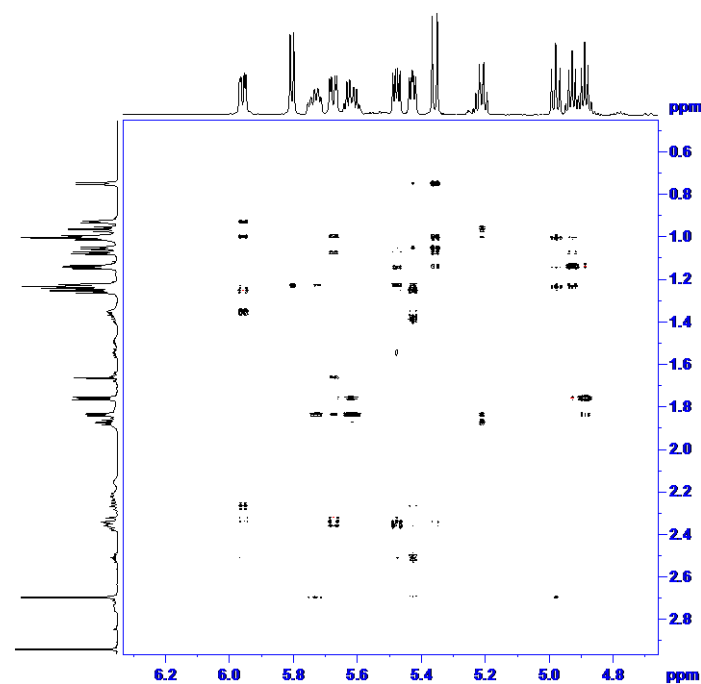
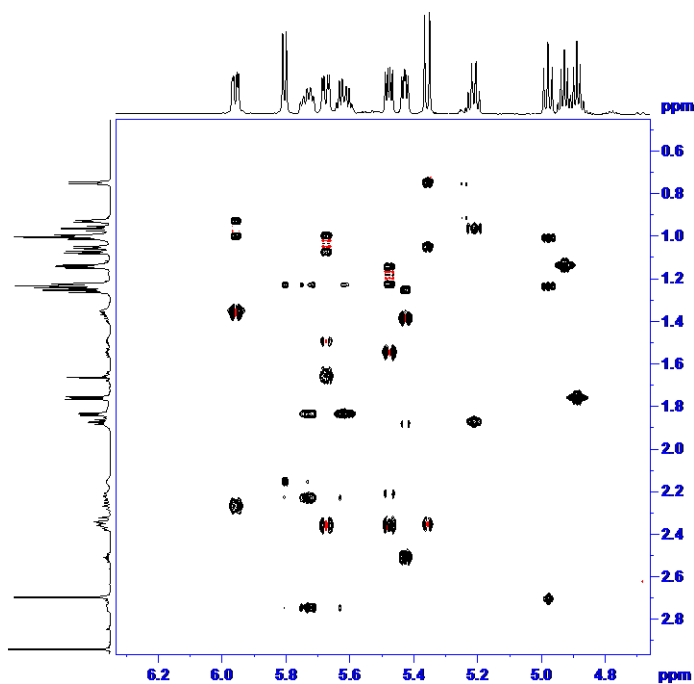
Application: Resolution enhancement

Example: Cyclosporine

TOCSY TD (f1) = 512 \rightarrow SI = 2kx2k

FT

Covariance

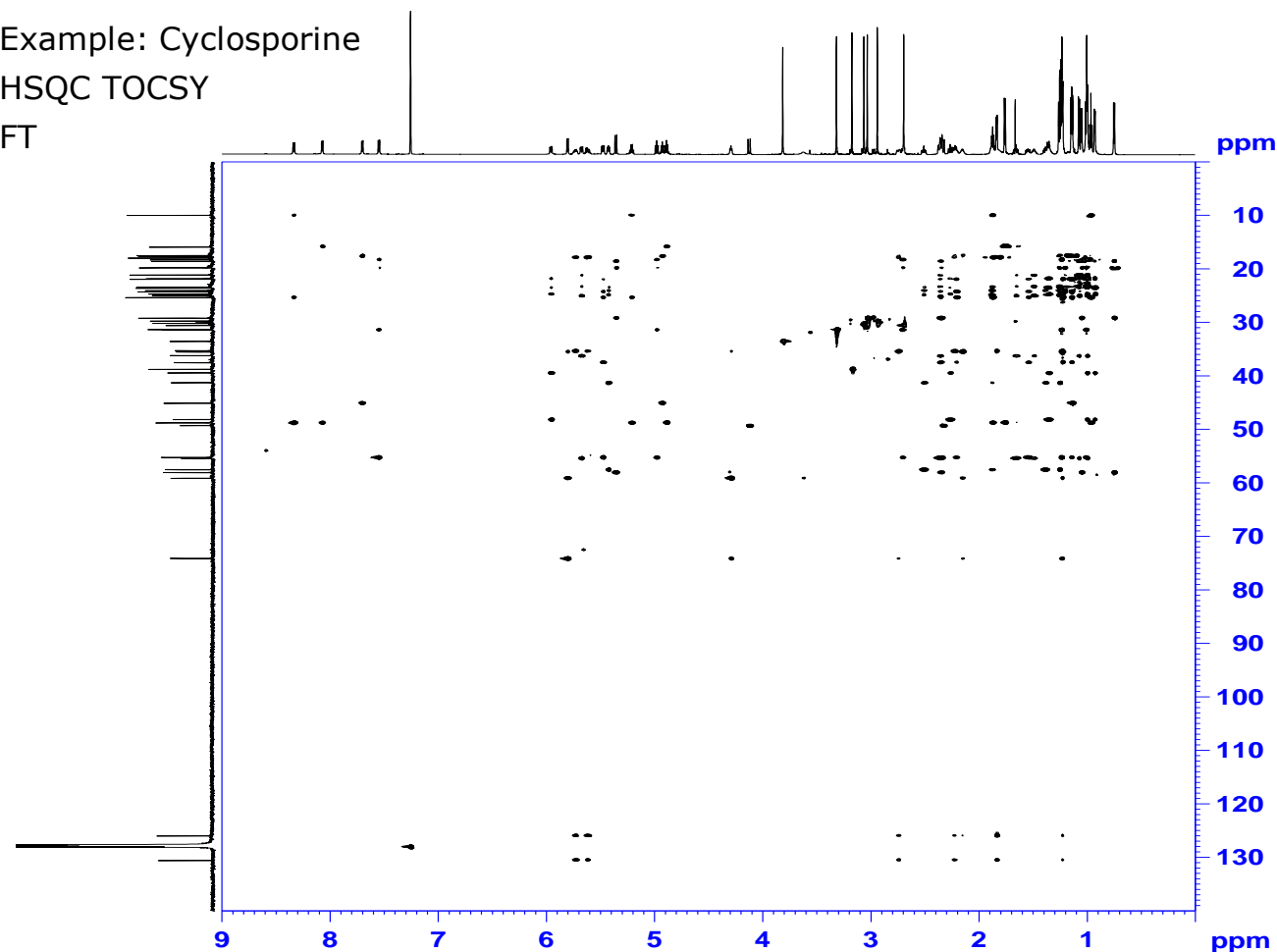


Covariance NMR



Application: Inverse spectroscopy

Example: Cyclosporine
HSQC TOCSY
FT



Covariance NMR

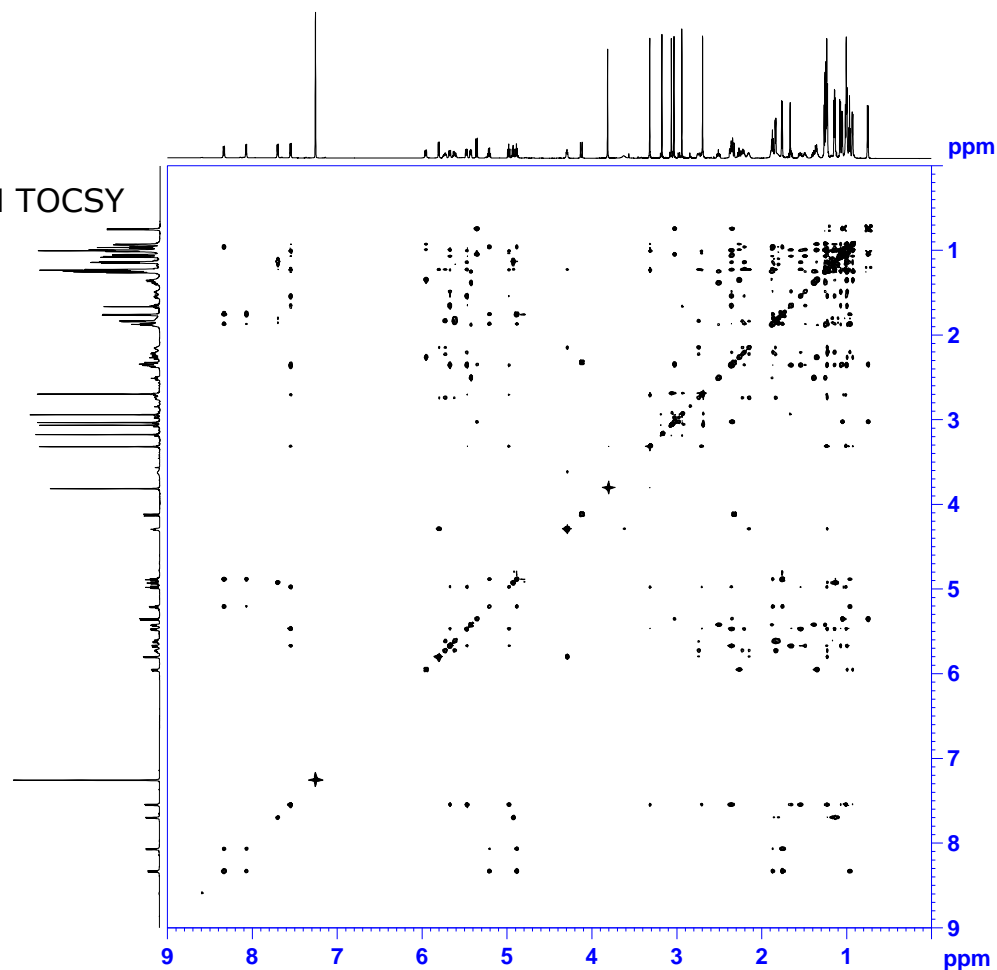


Application: Inverse spectroscopy

Example: Cyclosporine

HSQC TOCSY

Covariance with FT in f2 → H,H TOCSY



Covariance NMR

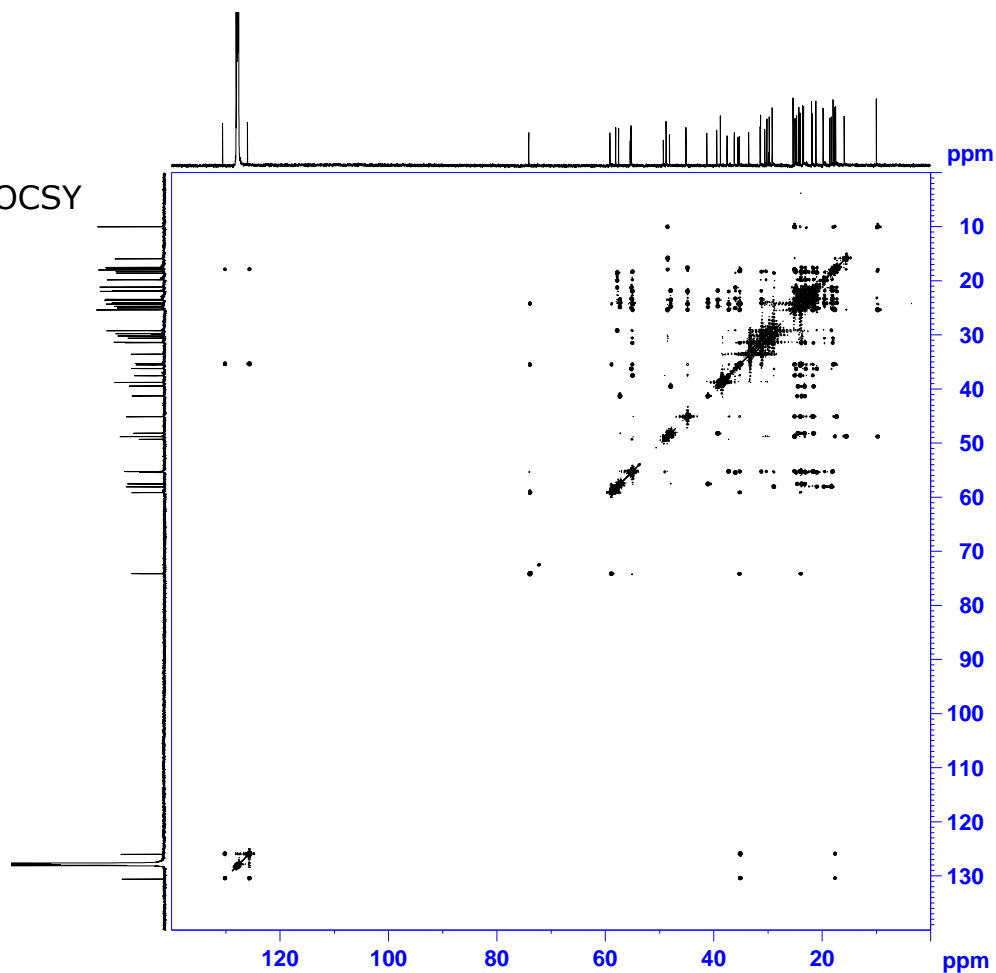


Application: Inverse spectroscopy

Example: Cyclosporine

HSQC TOCSY

Covariance with FT in f1 → C,C TOCSY



Implementation in Topspin



Data Processing

Process with fourier transform and phase correct.

Start AU program:

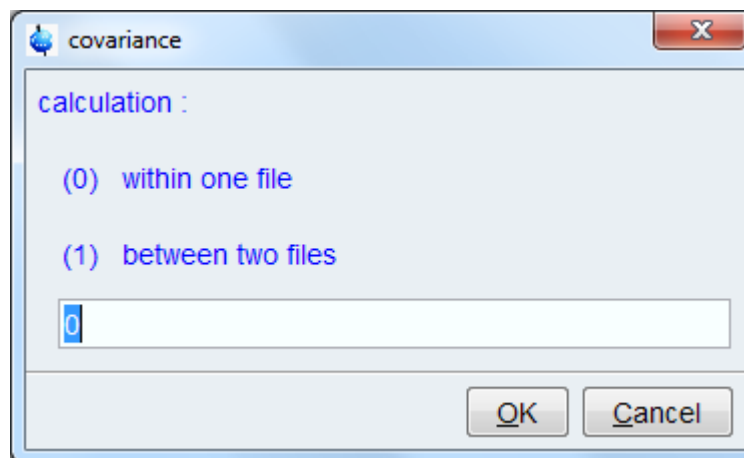
covariance (with no argument for default options)

covariance man (for interactive mode)

Implementation in Topspin



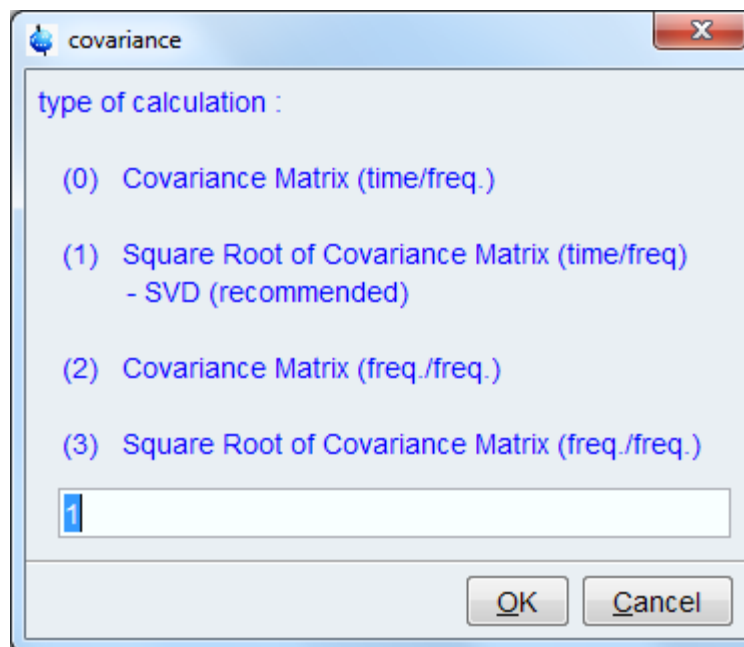
covariance man



Implementation in Topspin



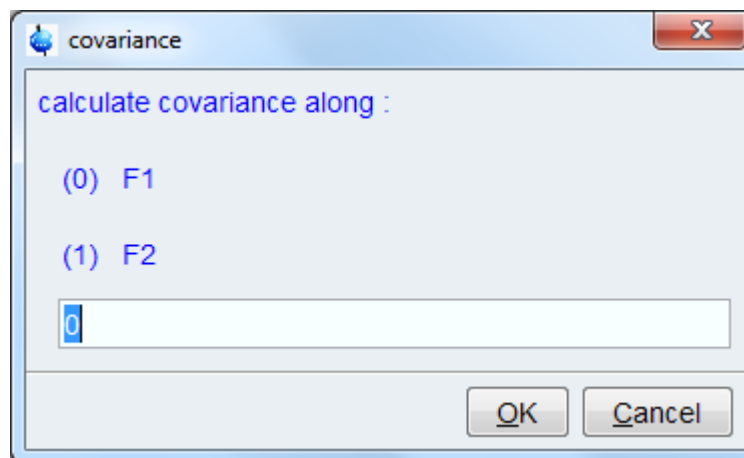
covariance man



Implementation in Topspin



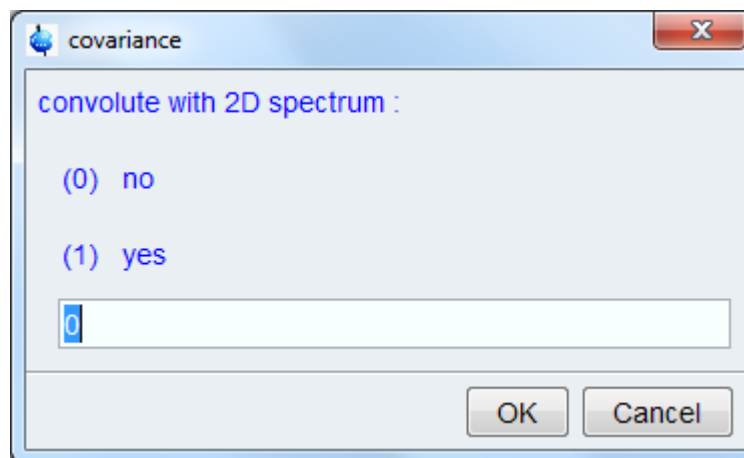
covariance man



Implementation in Topspin



covariance man

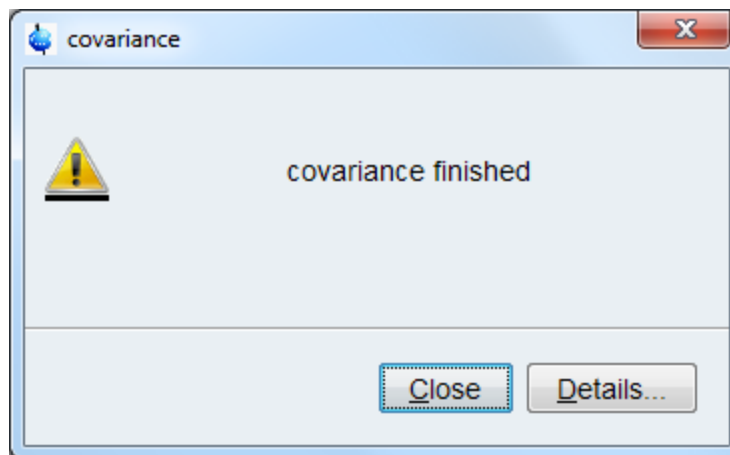


Implementation in Topspin



covariance man

Wait until



Can take seconds or minutes, depending on processor speed and data set size.

Availability



AU program is included as part of Topspin3.0 and newer

For Topspin2:

<http://spin.magnet.fsu.edu/software/covNMR/covNMR.html>

Covariance NMR

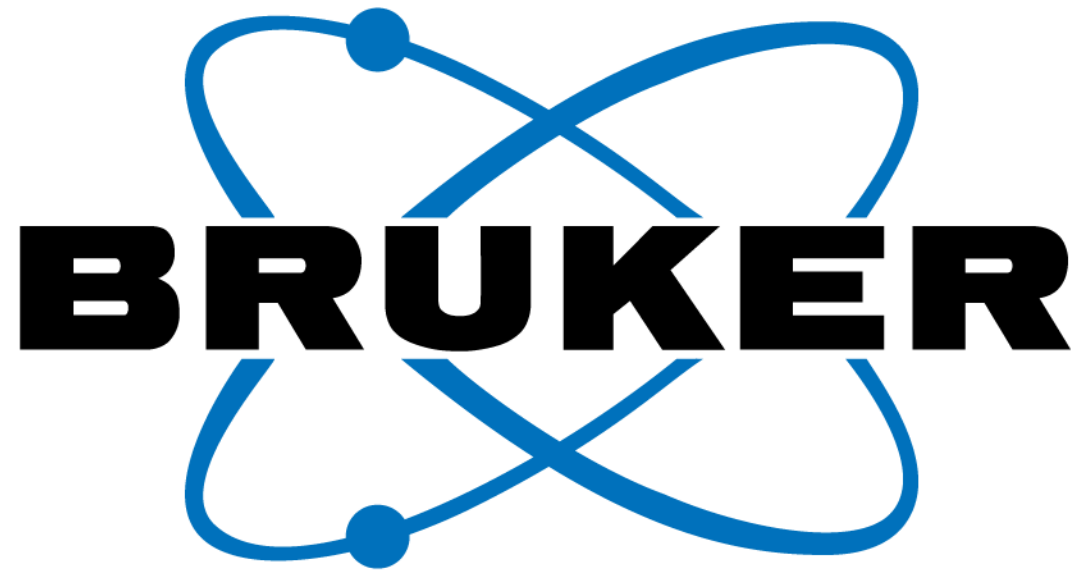


Literature

- [1] R. Bruschiweiler and F. Zhang, Covariance nuclear magnetic resonance spectroscopy
J. Chem. Phys. 120, 5253-5260 (2004)

- [2] N. Trbovic, S. Smirnov, F. Zhang, and R. Bruschiweiler, Covariance NMR spectroscopy by singular value decomposition
J. Magn. Reson. 171, 277-283 (2004)

- [3] F. Zhang and R. Bruschiweiler, Indirect covariance NMR spectroscopy
J. Am. Chem. Soc. 126, 13180-13181 (2004)



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