

# Resource Summary Report

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## CCPN Analysis

RRID:SCR\_016984

Type: Tool

### Proper Citation

CCPN Analysis (RRID:SCR\_016984)

### Resource Information

**URL:** <https://www.ccpn.ac.uk/v2-software/software/analysis>

**Proper Citation:** CCPN Analysis (RRID:SCR\_016984)

**Description:** Software package for interactive NMR spectrum visualization, resonance assignment and data analysis. Graphical elements allow to enter information and to view status of data and library functions manipulate the CCPN data model objects to record the scientific information. Software is cross platform and works on Linux, Mac OSX, Windows and Unix.

**Abbreviations:** CcpNmr Analysis

**Synonyms:** CcpNmr Analysis, CCPN Analysis v2, CCPN Analysis v3

**Resource Type:** data analysis software, data processing software, software application, data visualization software, software resource

**Defining Citation:** [PMID:21953355](#), [PMID:15815974](#)

**Keywords:** interactive, NMR, spectrum, visualization, resonance, data, analysis

**Funding:** Deutsche Forschungsgemeinschaft ;  
Biotechnology and Biological Sciences Research Council (UK)

**Availability:** Public, Available for download, Free of charge for non profit institutions, Tutorial available

**Resource Name:** CCPN Analysis

**Resource ID:** SCR\_016984

**License:** CCPN licence

**Record Creation Time:** 20220129T080333+0000

## Ratings and Alerts

No rating or validation information has been found for CCPN Analysis.

No alerts have been found for CCPN Analysis.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 48 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [SPARC SAWG](#) .

Murphy NP, et al. (2026) Cyclized Peptide Inhibitors of the Small G Protein Cdc42 Mimic Binding of Effector Proteins. *Biochemistry*, 65(3), 297.

Skobelkina A, et al. (2025) KIF2C condensation concentrates PLK1 and phosphorylated BRCA2 on kinetochore microtubules in mitosis. *Nucleic acids research*, 53(11).

?ervený J, et al. (2025) Human Milk Oligosaccharides Multivalently Presented on Defined Synthetic Neo-Glycoproteins Are Nanomolar Ligands of Tandem-Repeat Galectins. *Biomacromolecules*, 26(8), 4995.

Bugge K, et al. (2025) Role of charges in a dynamic disordered complex between an IDP and a folded domain. *Nature communications*, 16(1), 3242.

Almeida T, et al. (2025) Dock-and-lock binding of SxIP ligands is required for stable and selective EB1 interactions. *eLife*, 13.

Yan X, et al. (2025) Intra-condensate demixing of TDP-43 inside stress granules generates pathological aggregates. *Cell*, 188(15), 4123.

Rua AJ, et al. (2024) Perturbations in mitochondrial metabolism associated with defective cardiolipin biosynthesis: An in-organello real-time NMR study. *bioRxiv : the preprint server for biology*.

Soto J, et al. (2024) Dynamics and thermal stability of the bypass polymerase, DinB homolog (Dbh). *Frontiers in molecular biosciences*, 11, 1364068.

Dawes S, et al. (2024) Chaperone BiP controls ER stress sensor Ire1 through interactions with its oligomers. *Life science alliance*, 7(10).

Rampratap P, et al. (2024) Resolving Atomic-Level Dynamics and Interactions of High-Molecular-Weight Hyaluronic Acid by Multidimensional Solid-State NMR. *ACS applied materials & interfaces*, 16(33), 43317.

Rua AJ, et al. (2024) Perturbations in mitochondrial metabolism associated with defective cardiolipin biosynthesis: An in-organello real-time NMR study. *The Journal of biological chemistry*, 300(10), 107746.

Mahana Y, et al. (2024) Structural evidence for protein-protein interaction between the non-canonical methyl-CpG-binding domain of SETDB proteins and C11orf46. *Structure* (London, England : 1993), 32(3), 304.

Lete MG, et al. (2023) NMR Investigation of Protein-Carbohydrate Interactions: The Recognition of Glycans by Galectins Engineered with Fluorotryptophan Residues. *Chemistry* (Weinheim an der Bergstrasse, Germany), 29(5), e202202208.

Aretz J, et al. (2023) Talin and kindlin use integrin tail allostery and direct binding to activate integrins. *Nature structural & molecular biology*, 30(12), 1913.

Gubensäk N, et al. (2023) *Vibrio cholerae*'s ToxRS bile sensing system. *eLife*, 12.

Lin CC, et al. (2023) The combined action of the intracellular regions regulates FGFR2 kinase activity. *Communications biology*, 6(1), 728.

Masone A, et al. (2023) A tetracationic porphyrin with dual anti-prion activity. *iScience*, 26(9), 107480.

Johansen NT, et al. (2022) Mg<sup>2+</sup>-dependent conformational equilibria in CorA and an integrated view on transport regulation. *eLife*, 11.

Martinsen JH, et al. (2022) Structure, dynamics, and stability of the globular domain of human linker histone H1.0 and the role of positive charges. *Protein science : a publication of the Protein Society*, 31(4), 918.

Jara KA, et al. (2022) Multivalency, autoinhibition, and protein disorder in the regulation of interactions of dynein intermediate chain with dynactin and the nuclear distribution protein. *eLife*, 11.