## CSD product datasheet 2022



Academic users		advancing structural science
	CSD-Community	CSD-Enterprise
Data		
CSD deposited and curated data	√۱	$\checkmark$
CSD teaching resources	$\checkmark$	$\checkmark$
Proprietary CSD extension		$\checkmark$
Deposit		
Guided data deposition	$\checkmark$	$\checkmark$
CIF syntax check	$\checkmark$	$\checkmark$
Reduced cell check	$\checkmark$	$\checkmark$
Data validation	$\checkmark$	$\checkmark$
CSD DOI and curated CSD entry on publication	$\checkmark$	$\checkmark$
Direct publication through CSD Communications	$\checkmark$	$\checkmark$
Enhanced data discoverability	$\checkmark$	$\checkmark$
Persistent, free storage of your data	$\checkmark$	$\checkmark$
Deposition portal allowing you to access, edit and share your deposits	$\checkmark$	$\checkmark$
Access		
Retrieve via bibliographic info, DOI, CSD identifiers or compound name	$\checkmark$	V
Link from published articles and repositories	$\checkmark$	$\checkmark$
Publisher referee services	$\checkmark$	$\checkmark$
Search		
Search by chemical formula, cell parameters, 2D/3D substructure, similarity, and more		$\checkmark$
Protein-ligand binding sites		$\checkmark$
Visualise		
3D display and manipulation	$\checkmark$	$\checkmark$
High resolution graphics and movie generation	$\checkmark$	$\checkmark$
3D printing file output	$\checkmark$	$\checkmark$
PXRD pattern simulation	$\checkmark$	$\checkmark$
PXRD pattern comparison		$\checkmark$
Molecule and structure editing	$\checkmark$	$\checkmark$
2D diagram generation		$\checkmark$
Analyse		
Plotting and charting		$\checkmark$
Descriptive statistics		×
Interactive visualisation		√ ,
Filtering and categorisation		$\checkmark$
Reporting		$\checkmark$
Integrations		
CSD Pipeline Pilot Component Collection		$\checkmark$
CSD KNIME Component Collection		$\checkmark$

## CSD product datasheet 2022

## Academic users



	CSD-Community	CSD-Enterprise
Conformations		
Bond length assessment		$\checkmark$
Valence angle assessment		$\checkmark$
Torsion angle assessment		$\checkmark$
Ring geometry assessment		$\checkmark$
Conformer generation		$\checkmark$
Interactions		
Fragment interaction maps (IsoStar)		$\checkmark$
Protein interaction maps (SuperStar)		$\checkmark$
Full interaction maps		$\checkmark$
Ligand-based drug discovery		
Ligand overlay		$\checkmark$
Field-based ligand screener		$\checkmark$
Scaffold hopping		$\checkmark$
Structure-based drug discovery		
Protein-ligand docking		✓
Ensemble docking		$\checkmark$
Pose analysis		$\checkmark$
Proprietary structures		$\checkmark$
Cavity similarity searching		$\checkmark$
Solid form analysis		
Motif searching		$\checkmark$
Packing feature searching		$\checkmark$
Crystal packing similarity		$\checkmark$
Calculations		$\checkmark$
Hydrogen bond propensity and coordination assessment		$\checkmark$
Hydrate analysis		$\checkmark$
Solvate analysis		$\checkmark$
Aromatic analysis		$\checkmark$
Co-crystal design		$\checkmark$
Hydrogen bond statistics assessment		$\checkmark$
Particle analysis		
Predict particle facets		$\checkmark$
Visualize surface chemistry and charge		$\checkmark$
Identify slip planes		$\checkmark$
Surface interactions analysis		$\checkmark$
Full interaction maps on surface		$\checkmark$
Determine H-bond dimensionality		$\checkmark$
Visualize surface topology		$\checkmark$
Quantify surface chemistry and topology		$\checkmark$
Python API particle analysis		 ■恕?

<sup>1</sup>Essential curated information for each entry is freely available through Access Structures on the CCDC website. More enhanced curated information is available through CSD-Core, CSD-Discovery, CSD-Materials and CSD-Enterprise

