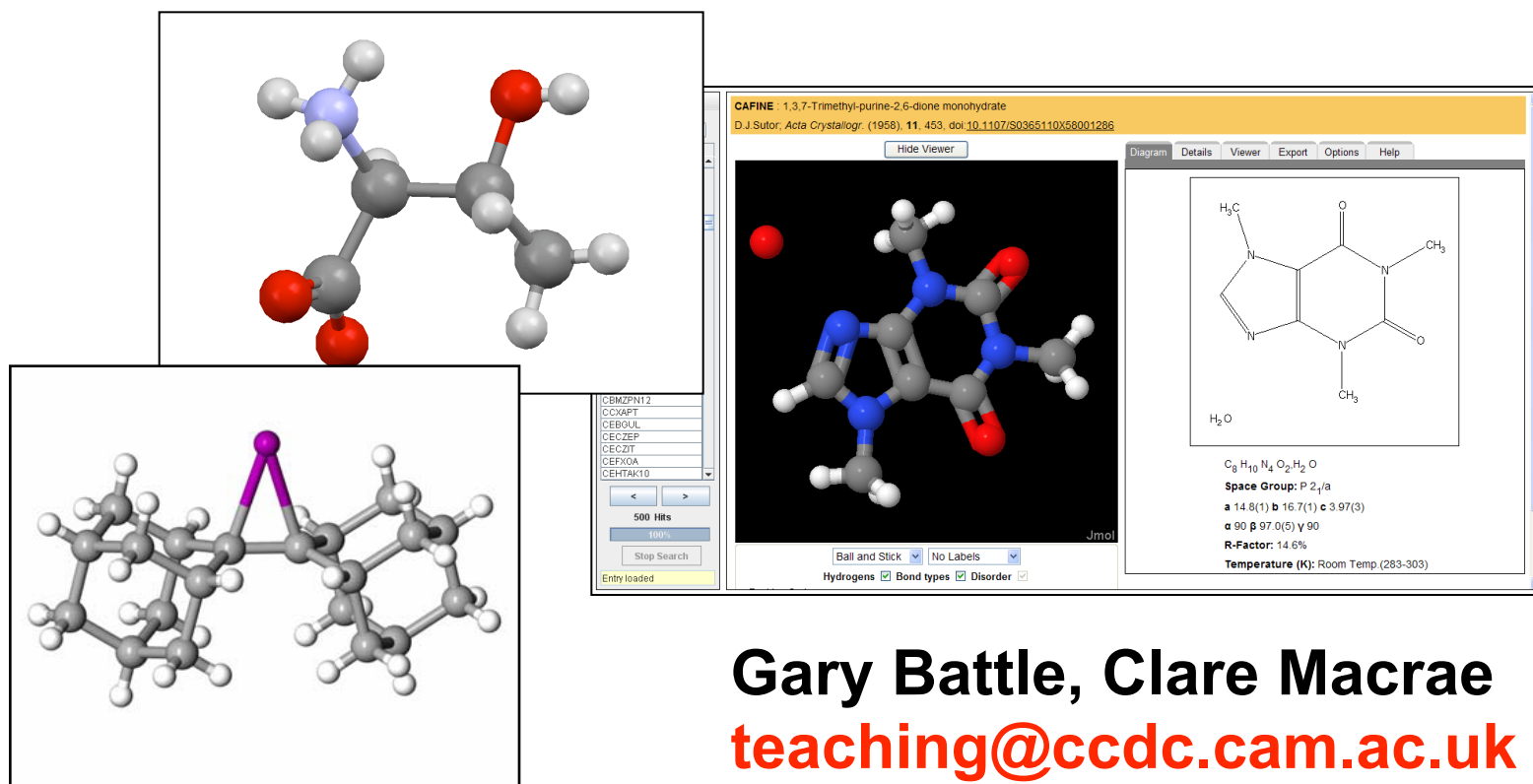




Imperial College Cambridge Structural Database Workshop



Gary Battle, Clare Macrae
teaching@ccdc.cam.ac.uk



Schedule...

- Introduction: CSDS content and coverage, use in teaching structural chemistry
- Web-based access: software tools and example uses
- Advanced desktop applications: software tools and examples
- IsoStar library of intermolecular interactions: overview and demo
- SuperStar: predicting interactions



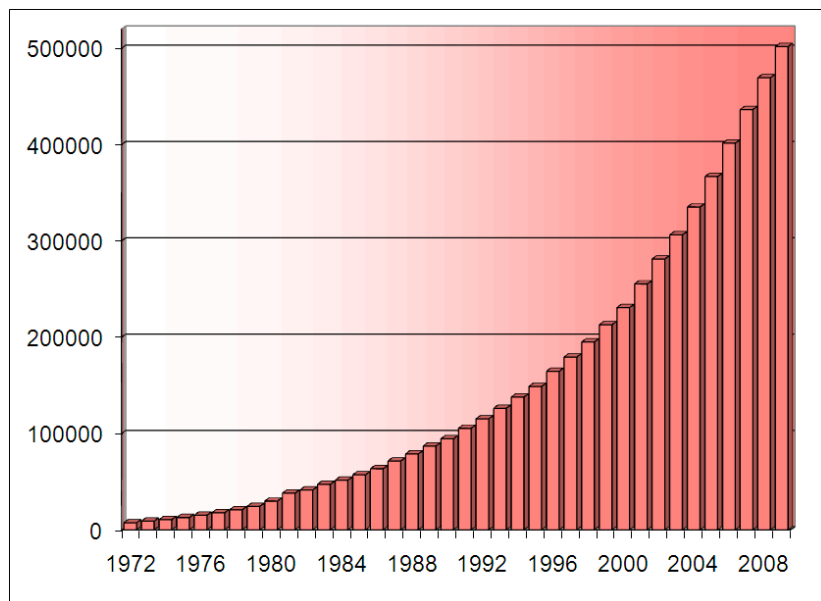
Crystal Structures are Uniquely Suited for Teaching Structural Chemistry

- X-ray analysis method of choice for 3D structure characterization
- Very precise: standard uncertainties <0.005 Å on bond lengths and <0.5 on valence and torsion angles
- Provide remarkable richness of structural information: both the 3D geometric structures of molecules and also the nature and geometry of their interactions
- $> 750,000$ crystal structures published in past 60 years, and modern instrumentation adding $> 60,000$ novel structures each year



Cambridge Structural Database

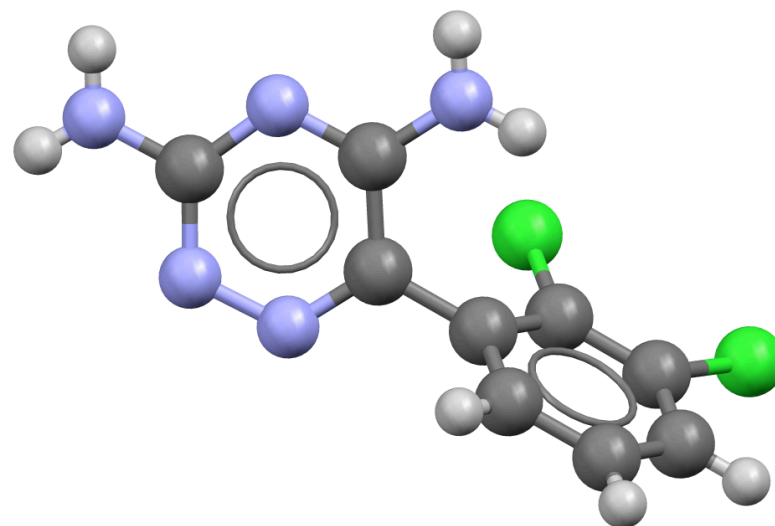
Worldwide repository of validated small-molecule organic & metal-organic crystal structures



Lamotrigine

Acta Cryst., Sect. C: Cryst. Struct. Commun. (2009), **65**, o460

Refcode: **EFEMUX01**



CSD Growth 1970-2010

Dec 09 – 500,000th structure milestone reached



Pedagogical Value

- 3D visualisations enhance students conceptual understanding and spatial abilities
 - Williamson, V. M. *J. Chem. Educ.*, **2008**, 85, 718-723
 - Bodner, G. M.; Guay, R. B. "The Purdue Visualization of Rotations Test" *The Chemical Educator*
- Use of experimentally measured data is of great pedagogical value, and has been shown to enhance student learning
 - DeHaan, R. L. *J. Sci. Educ. Technol.*, **2005**, 14, 253-269
 - Handelsman, J. *Science* **2004**, 304, 521-522
 - Prince, M. *J. Eng. Educ.*, **2004**, 89, 1-9



Using the Cambridge Structural Database to Introduce Important Inorganic Concepts



Tiana V. Davis, M. Shahzad Zaveer, and Marc Zimmer*

Chemistry Department, Connecticut College, New London, CT 06320; *mzimmer@conncoll.edu

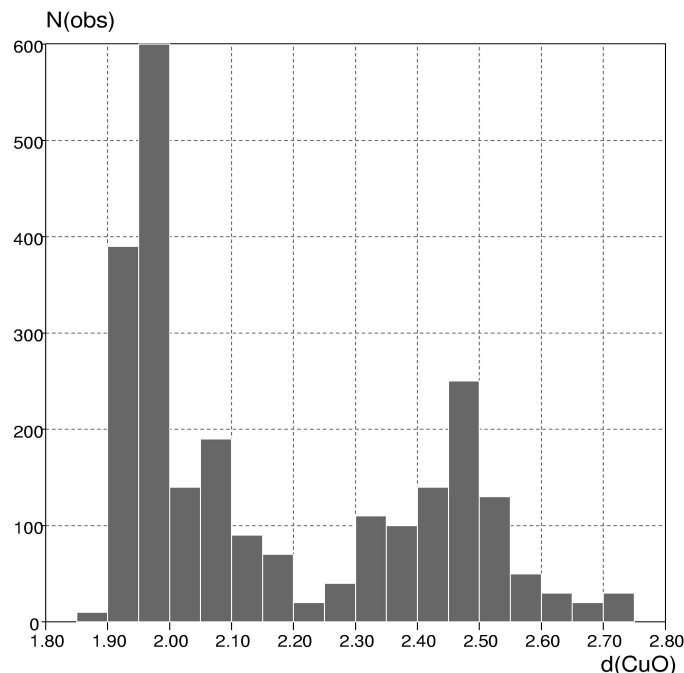
Data and structure correlation analysis is an increasingly important area in science (1). This is particularly true in biology where genomic and proteomic studies are generating vast amounts of data. In order to expose our students to chemoinformatics and introduce them to an important resource in chemistry, the Cambridge Structural Database, we have devised a simple series of inorganic exercises that can be done in introductory inorganic classes.

The latest version of the Cambridge Structural Database (CSD), version 5.22, contains the CSD database with 245,392 structures, ConQuest (an interface to CSD), Mercury (a visualization program), Isostar (software for superimposing molecular fragments) and Vista (a statistics package designed for use with the CSD). CSD has 245,392 X-ray and neutron diffraction structures of organocarbon compounds (2, 3).¹ All the compounds in the CSD have less than 1,000 atoms. Peptides with up to 24 residues are covered (4), while larger peptides and proteins are in the Protein Database.² The CSD has 109,349 structures that contain one or more transition metal ions. Classroom ConQuest comes with a reduced database of 11,300 entries. Aqueous with at least one normal

of congeneric families are useful as they can reveal the different backbone conformations the structure can adopt in the different crystals environments. These analyses, in turn, can provide information about conformations available to the backbone, interconversions of the conformers, and environmental factors responsible for certain conformations (6).

The release of a new user-friendly graphics interface called ConQuest, easy to use tutorials, a Windows PC and Linux version, and a classroom edition have prompted us to develop some inorganic laboratory exercises utilizing the database. The purpose of these exercises is to expose the students to database analyses, and to demonstrate inorganic structural properties and structure correlation. Four example exercises are outlined below; all the structures used in the exercises are shown in Table 1. The exercises can be presented as open-ended discovery assignments or as more traditional problems. As the tutorials are complete, easy to understand and instructive, the students need no assistance in doing ConQuest searches. The resultant data can easily be exported to Isostar, Vista, or Excel.

Backbonding
Eighteen-Electron Rule
High-Spin vs. Low-Spin
Jahn-Teller Effect





In the Classroom

Teaching with Technology

edited by
Gabriela C. Weaver
Purdue University
West Lafayette, IN 47907

Using the Cambridge Structural Database To Teach Molecular Geometry Concepts in Organic Chemistry

Jay Wm. Wackerly, Philip A. Janowicz, Joshua A. Ritchey, Mary M. Caruso, and Erin L. Elliott

Department of Chemistry, University of Illinois at Urbana-Champaign, Urbana, IL 61801

Jeffrey S. Moore*

Departments of Chemistry, Materials Science and Engineering, and the Beckman Institute, University of Illinois at Urbana-Champaign, Urbana, IL 61801; *jsmoore@illinois.edu

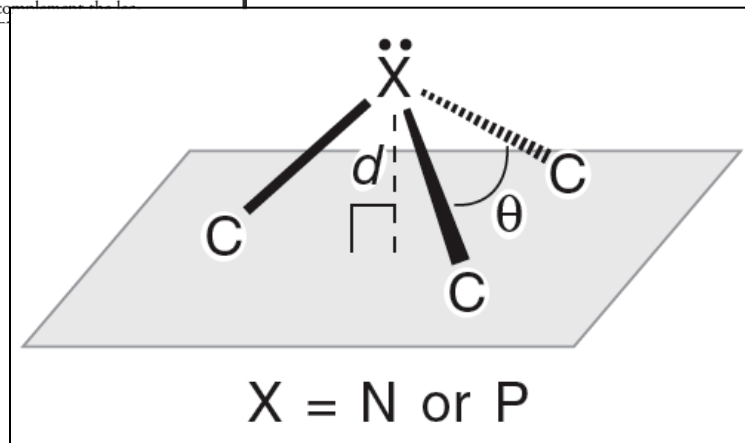
A large lecture classroom may not seem to give students a chance to move beyond their textbook structures and analyze data from actual molecular structures. However with continuous technological advances applied to education, individualized software programs are more commonly available to enhance students' learning (1). It has been shown that students take a greater interest in their coursework when they have the opportunity to analyze "real-world" data (2). Given society's growing dependence on information technology and the need to understand the basic concepts of data mining and parameter correlation, the opportunity to introduce these ideas in an organic chemistry classroom full of future professionals is appealing (3). Herein

The CSD is a database of small organic and organometallic structures that have been elucidated by X-ray or neutron diffraction techniques (5). The database currently has 400,977 structures, and 43% of these are organic compounds. In order to give all of the students convenient access to the database, Classroom ConQuest was used.¹ Classroom ConQuest is provided by the Cambridge Crystallographic Data Center free of charge² although it contains a reduced number of structures (11,300). While only a subset of the full CSD, meaningful searches using Classroom ConQuest for simple structural trends can still be obtained.

These assignments were designed to complement the

J. Chem. Educ., **2009**, 86
(4), p 460
DOI: 10.1021/ed086p460

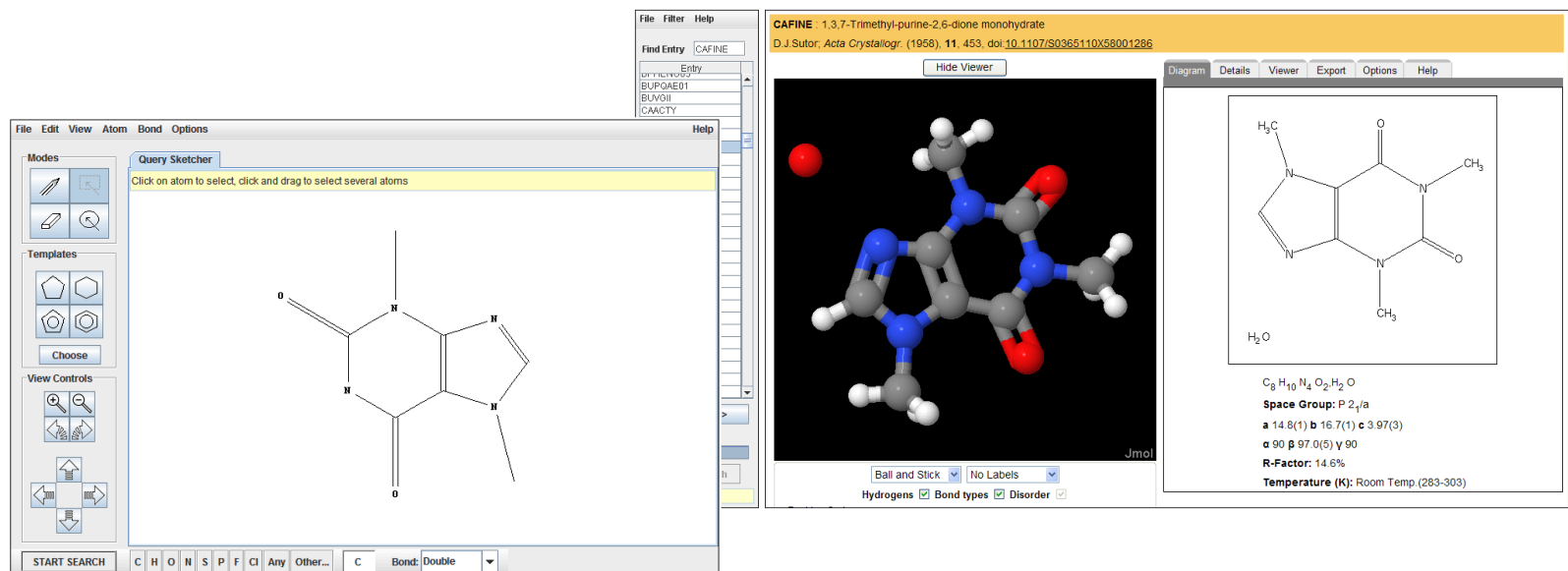
"...gives students a chance to move beyond their textbook structures and analyze data from actual molecular structures"





Improving CSD Accessibility

- **WebCSD**: online search interface to the CSD
- **University-wide access**: not required to download, install or register software on individual machines
- **Ease of use**: search and visualise structures using a standard web browser





Search functionality

- Text and numeric
- Reduced cell
- 2D substructure
- 2D structure similarity
- View specific refcode or browse the CSD

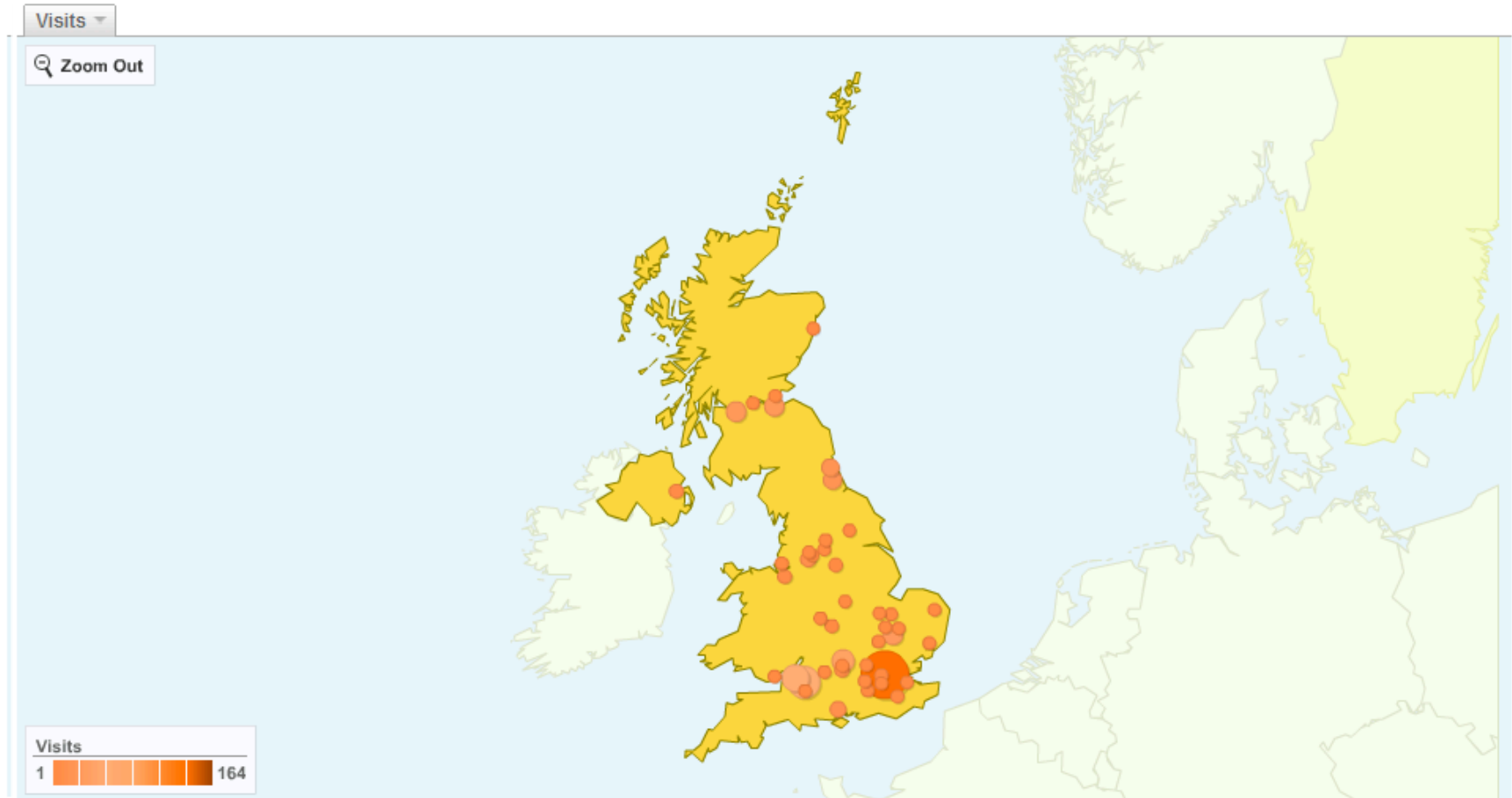
The screenshot shows the Cambridge Structural Database (CSD) search results for the entry CAFINE. The interface includes a search bar on the left, a list of search results, and a detailed view of the selected entry.

CAFINE 1,3,7-Trimethyl-purine-2,6-dione monohydrate
D.J.Sutor, *Acta Crystallogr.* (1958), **11**, 453. doi:10.1107/S0365110X58001286

Identifier CAFINE
Author(s) D.J.Sutor
Reference *Acta Crystallogr.* (1958), **11**, 453. doi:10.1107/S0365110X58001286
Formula C₈H₁₀N₄O₂·H₂O
Compound 1,3,7-Trimethyl-purine-2,6-dione monohydrate
SMILES CN1C=NC2=C1C(=O)N(C)C(=O)N2C.O
Synonym Caffeine monohydrate
Space Group P 2₁/a
Cell Lengths a 14.8(1) b 16.7(1) c 3.97(3)
Cell Angles α 90 β 97.0(5) γ 90
Cell Volume 973.911
Z, Z' Z 4 Z' 1
R-Factor (%) 14.6
Reduced Cell Lengths a 3.97 b 14.8 c 16.7
Reduced Cell Angles α 90 β 90 γ 97
Reduced Cell Volume 973.911
Temperature (K) Room Temp.(283-303)
Density (CCDC) 1.447
Average Sigma (C-C) 0.01 < sigma <= 0.03
Radiation Probe x-ray
Bioactivity stimulant agent which increases CNS activity



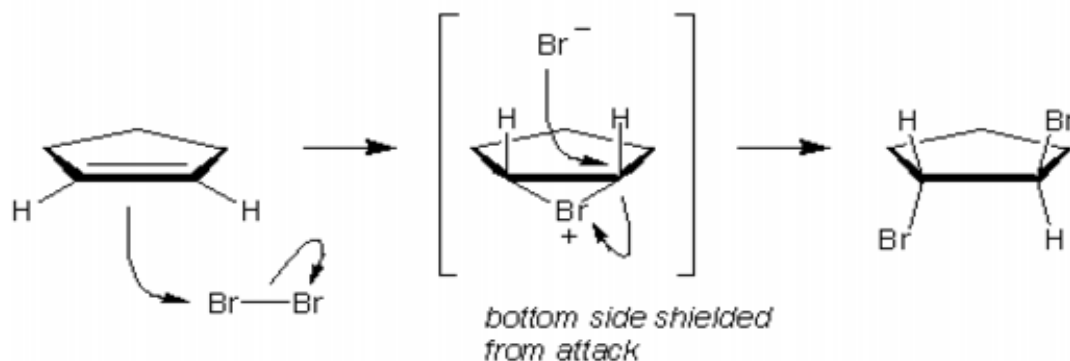
WebCSD Usage Jan - Mar 2011



'This country/territory sent 617 visits via 45 cities

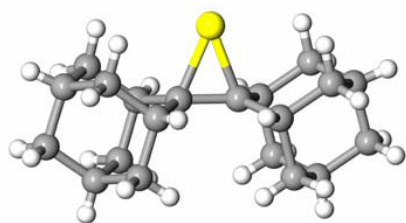


Example 1: Exploring Reaction Mechanisms using Crystallographic Data

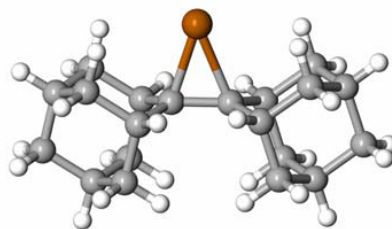


Learning Goals:

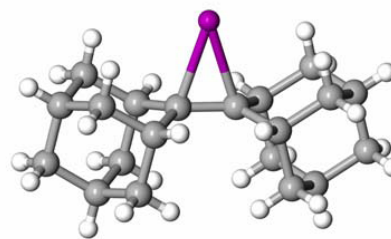
- Stereoselectivity
- Evidence for cyclic halonium ions
- Stabilisation



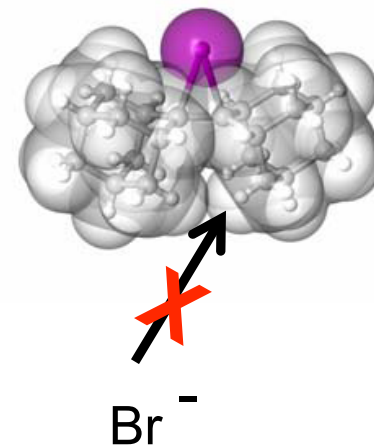
Cl - UGIDP



Br - DAKVUG



I - WEVPOC

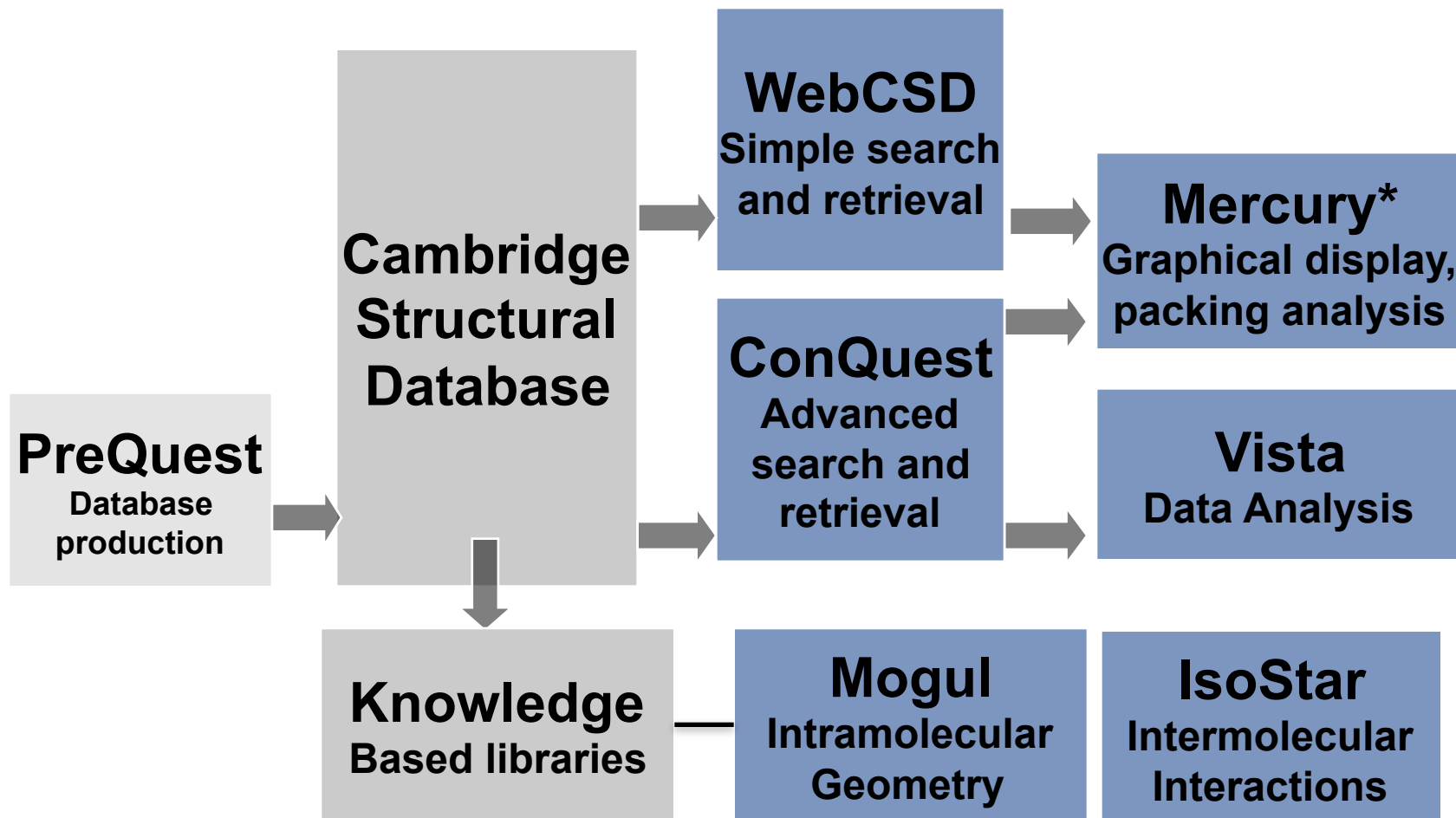


http://www.ccdc.cam.ac.uk/free_services/teaching/



Database plus Access Software

Acta Cryst., B58, 380-388 & 389-397, 2002



*free downloads <http://www.ccdc.cam.ac.uk/>



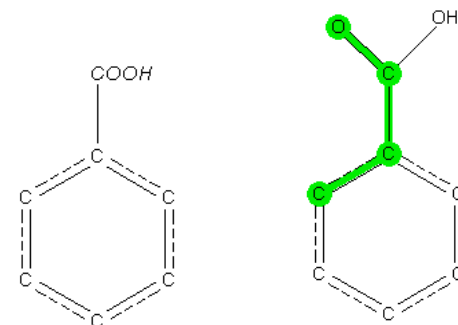
Searching the CSD: ConQuest

- Searches:

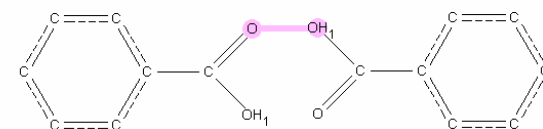
- Text & numerical data

Compound name, author name,
Chemical formula, melting point,
keyword.....

- Chemical substructures in 2D/3D



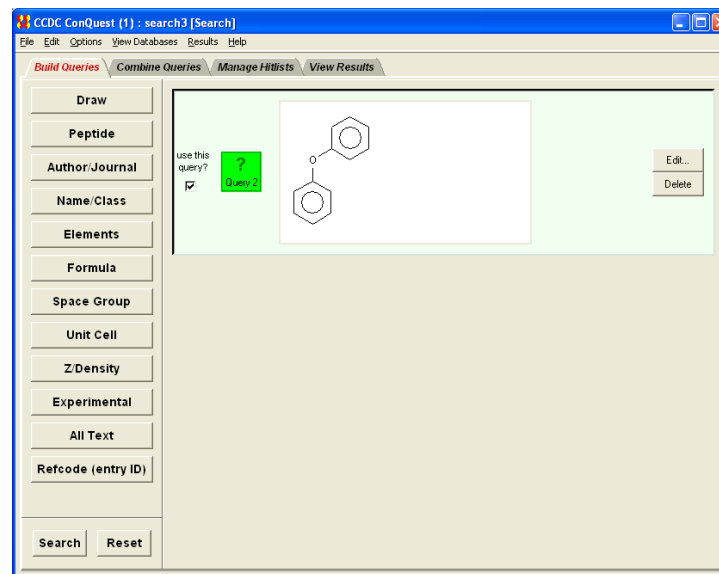
- Intermolecular non-bonded contacts





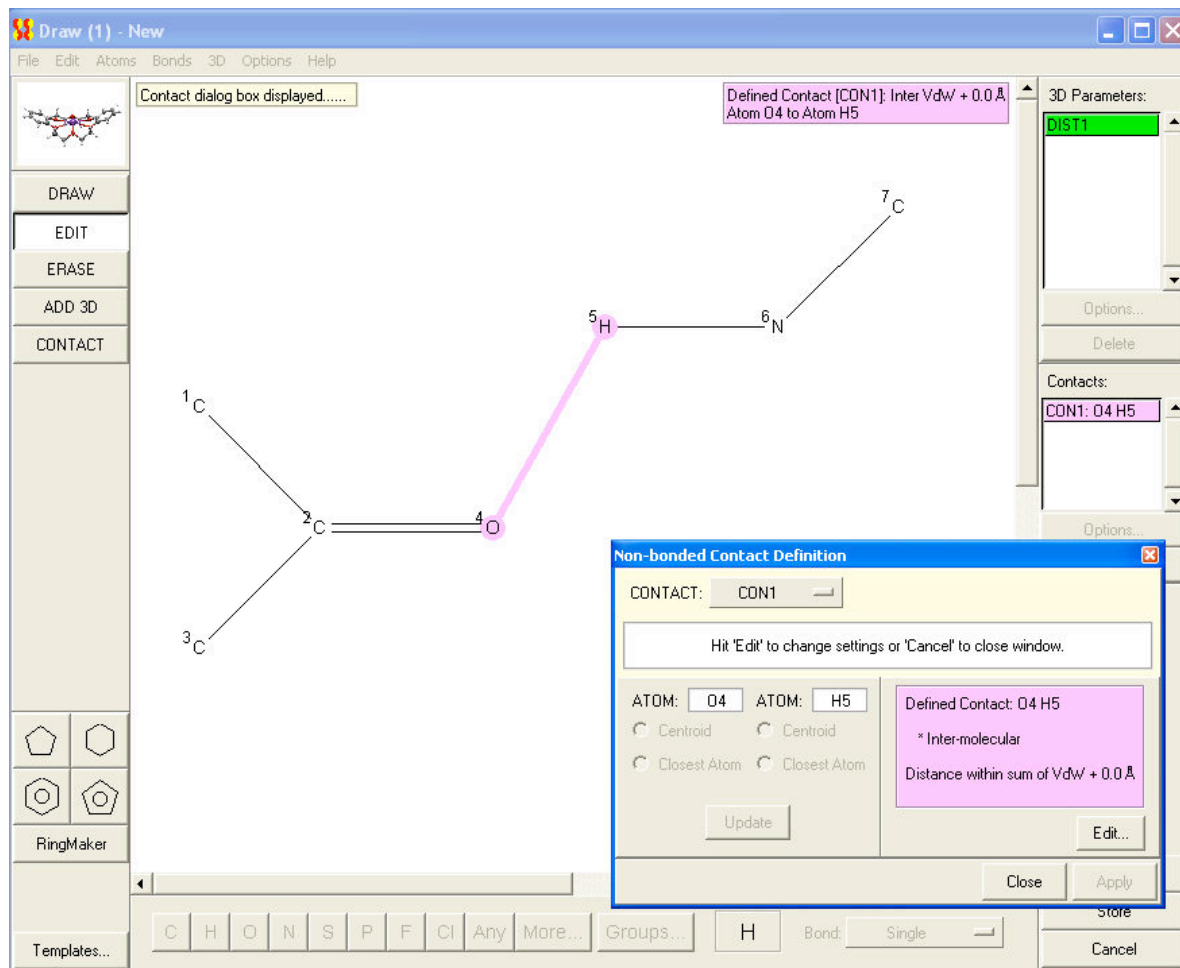
ConQuest

- Searches:
 - Access CSD and in-house data at the same time
 - Large range of filtering options based on experimental or chemical options
 - Combine searches and manage hitlists
 - Export results to Vista (numerical analysis), or Mercury (visualisation)





ConQuest Search: N-H...O=C(amide) H-Bonds



Search criteria:

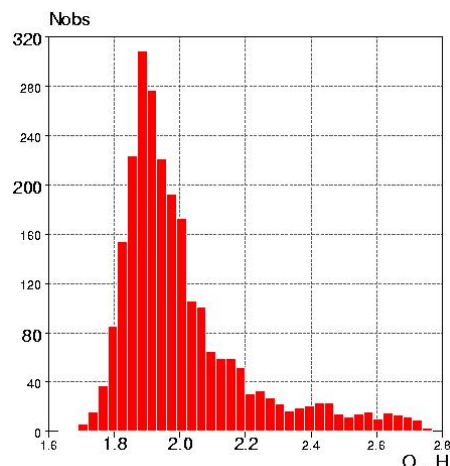
- C=O...H vdW+0.4
- Cryst. R-factor <5%

Calculate geometry:

- O...H distance
- O...H-N angle
- C=O...H angle
- Angle between N-H vector and amide plane



Analysing Geometrical Data: Vista



X-Vista: quest@tern

VISTA v.2.1 TABLE SPREADSHEET

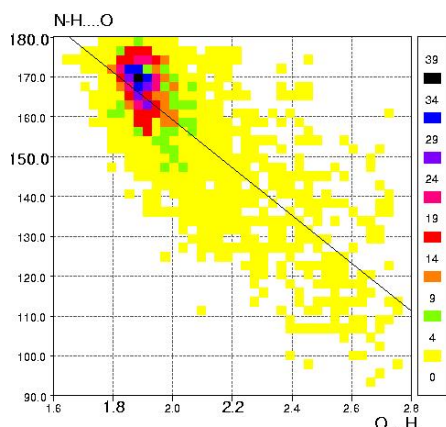
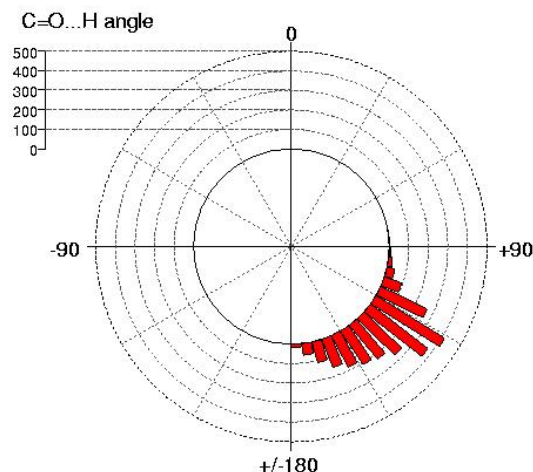
Quest File: cohord

Test: 1 of 1

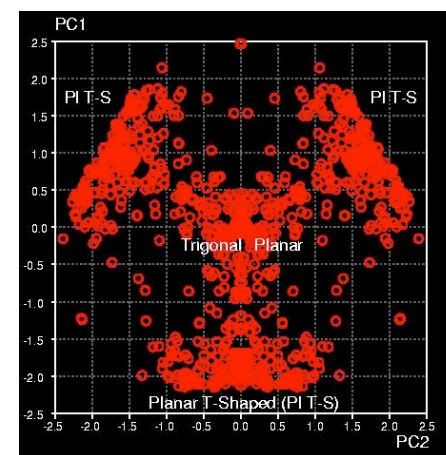
Parameters: 6
Refcodes: 653
Fragments: 794

Total: 0
Selected: n/a
Suppressed: 0

REFCODE	PARAMS	1	2	3	4	5	6	7
1	ACDQUR	1.8567	116.912	155.439	-26.2061			
2	ACDQUR	2.3301	171.215	115.452	-13.2601			
3	ACDQUR	1.7947	146.647	167.055	-52.3011			
4	ACDQUR	1.7417	127.061	169.4051	-62.1468			
5	ACDQUR	1.8574	125.933	177.725	-115.611			
6	ACDQUR	2.3231	103.352	110.413	143.6371			
7	ACDQUR	2.3631	118.948	142.937	-111.5011			
8	ACDQUR	1.9631	123.670	160.209	-67.428			
9	ACDQUR	2.4617	110.068	116.064	177.6201			
10	ACDQUR	1.7621	130.741	150.713	-60.9201			
11	ACDQUR	2.4971	131.504	140.258	102.6801			
12	ACDQUR	2.1611	167.268	130.275	-56.2751			
13	ACDQUR	2.1227	116.327	136.341	-47.5061			
14	ACDQUR	2.3041	167.728	165.363	59.711			
15	ACDQUR	2.0661	139.360	161.999	65.8401			
16	ACDQUR	1.7501	117.302	155.537	146.9891			
17	ACDQUR	1.7571	131.320	151.320	151.3201			
18	ACDQUR	1.7561	133.009	156.941	-153.3001			
19	ACDQUR	1.7081	139.498	176.617	8.5601			
20	ACDQUR	2.0721	138.408	136.167	-74.0561			
21	ACDQUR	2.0181	144.230	163.707	-3.9641			

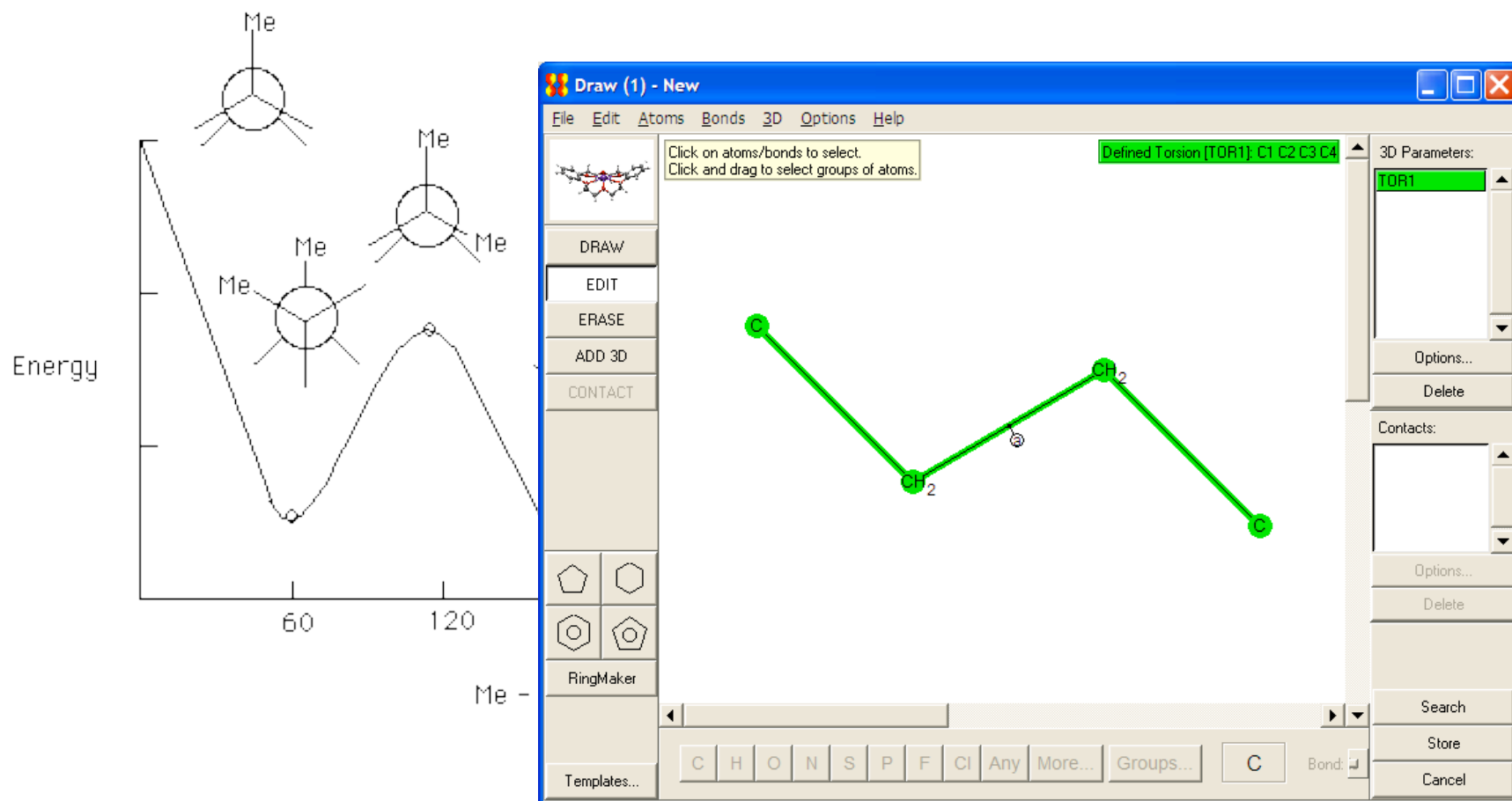


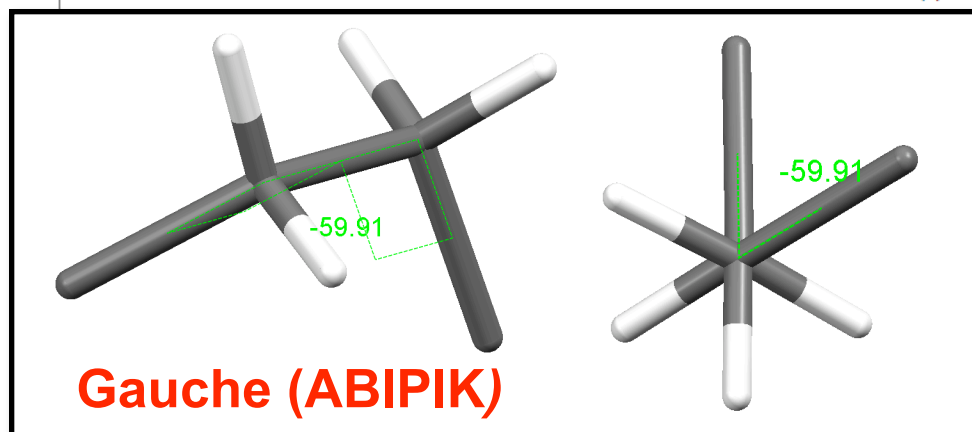
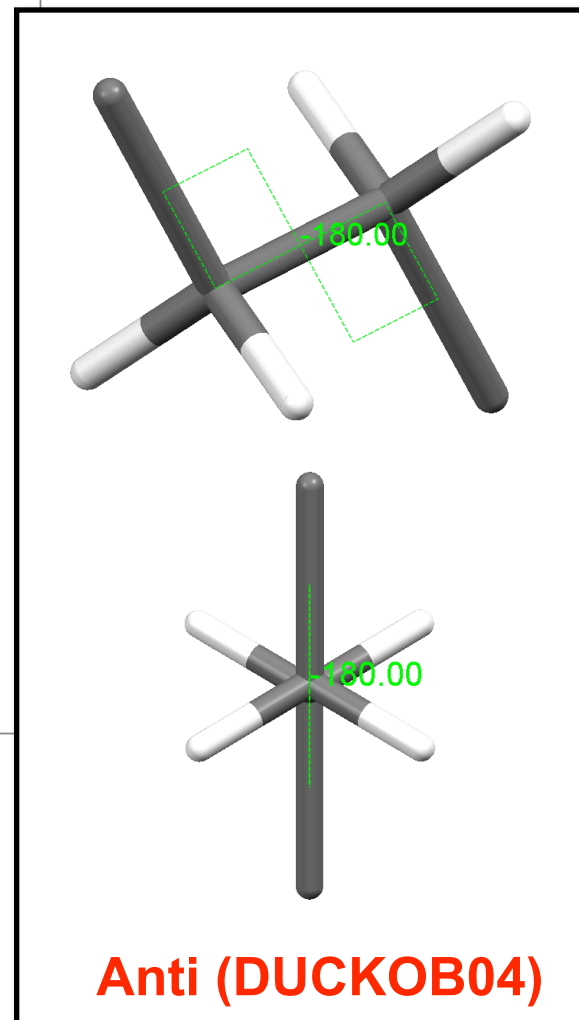
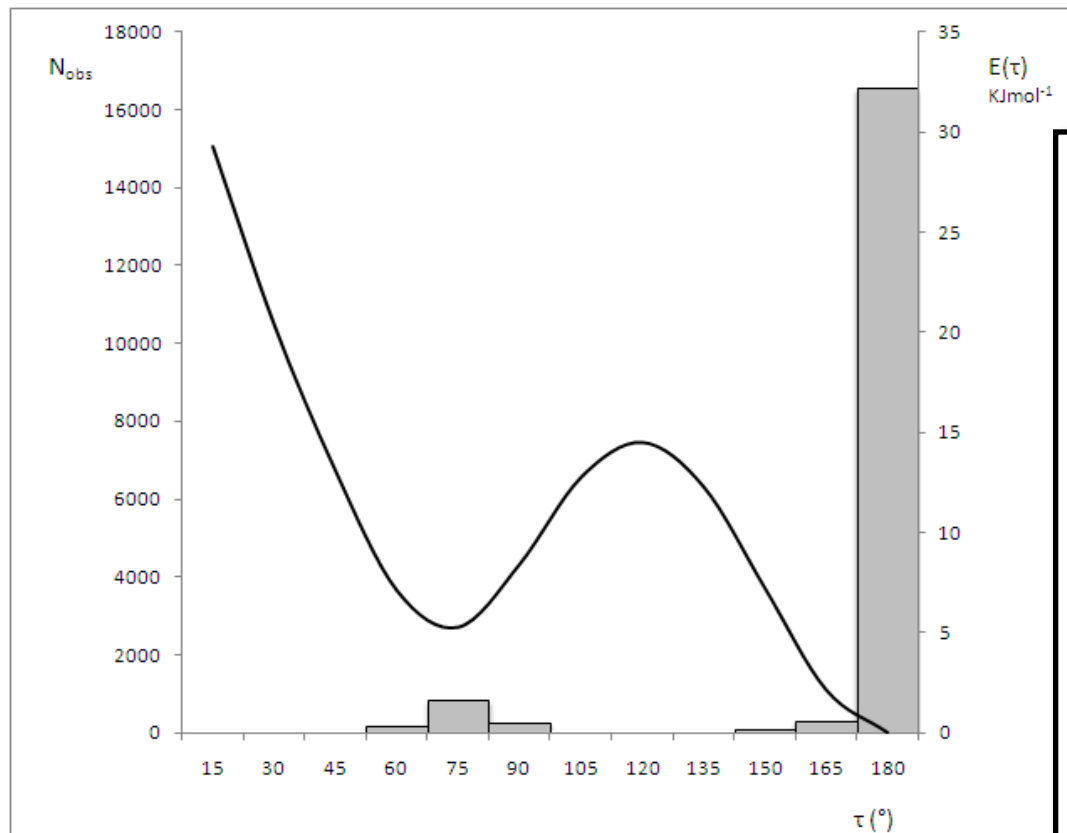
- Histograms (Cartⁿ, Polar)
- Scatterplots (Cartⁿ, Polar)
- Apply mathematical functions to spreadsheet parameters
- Statistical analysis:
 - Means, medians, esd's etc. for parameter distributions
 - Regression
 - Principal components (PCA)





Example 2: Conformational Analysis

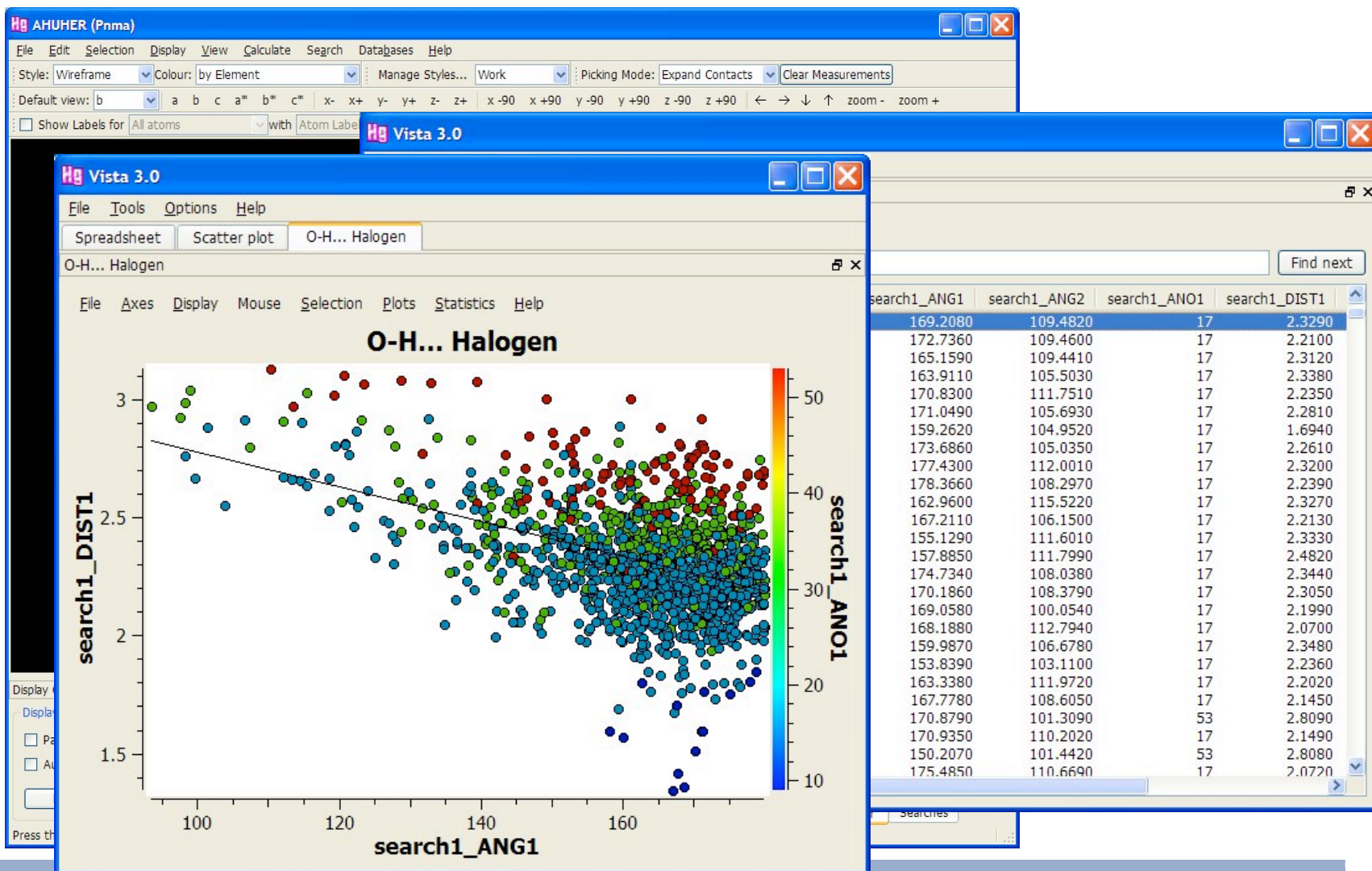






Replacing Vista

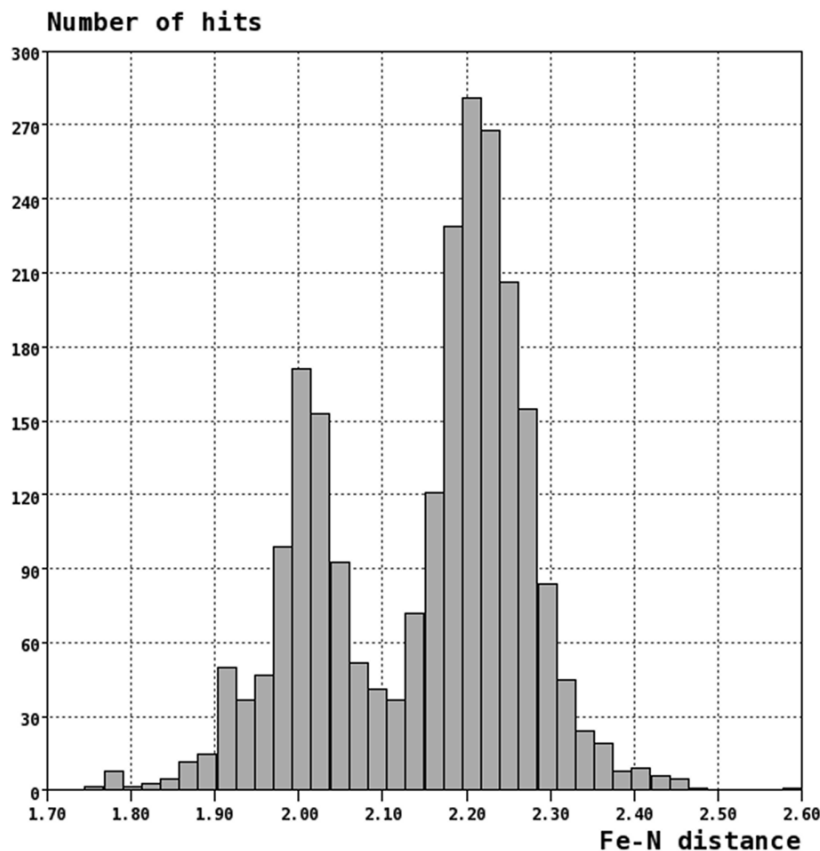
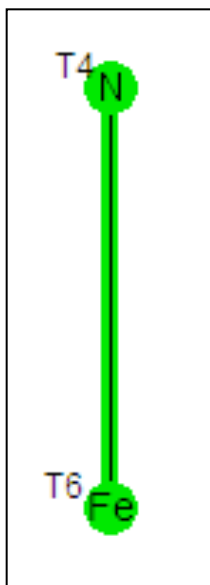
Interactive display and analysis of numeric CSD search data





Example 3:

High spin vs. low spin complexes

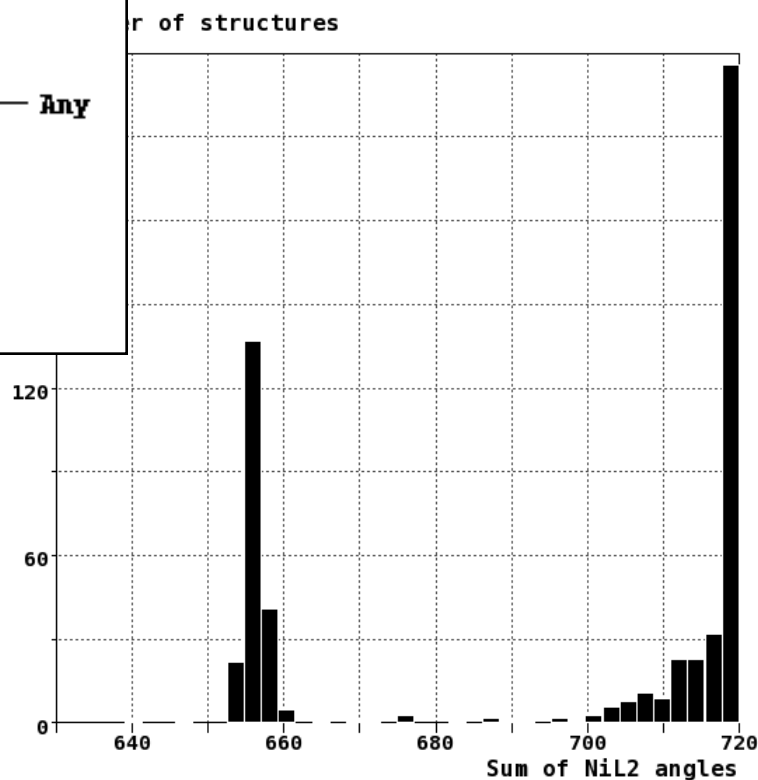
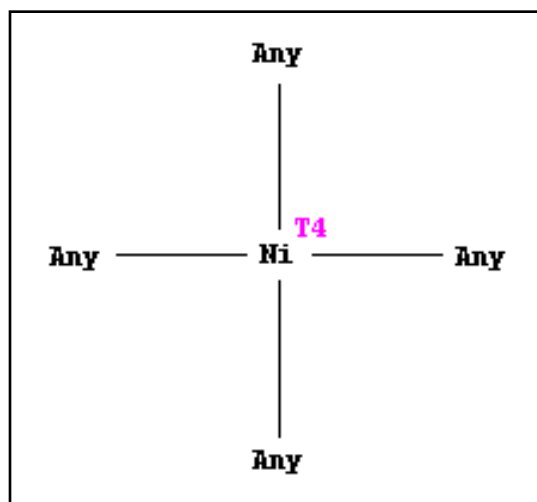


**High spin and low
spin forms of
octahedral Fe(III)**

cf. octahedral Ni(II)



Example 3: Ni coordination geometries



Sum of L-Ni-L
angles showing
square planar and
tetrahedral
geometries



Further Information

- Teaching 3D Structural Chemistry Using Crystal Structure Databases
1. An Interactive Web Accessible teaching Subset of the Cambridge Structural Database
Gary M. Battle and Frank H. Allen, Gregory M. Ferrence
***J. Chem. Educ.*, 2010, 87 (8), pp 809–812**
- Teaching 3D Structural Chemistry Using Crystal Structure Databases
2. Teaching Units that Utilize an Interactive Web Accessible teaching Subset of the Cambridge Structural Database
Gary M. Battle and Frank H. Allen, Gregory M. Ferrence
***J. Chem. Educ.*, 2010, 87 (8), pp 813–818**
- Applications of the Cambridge Structural Database in Chemical Education
Gary M. Battle and Frank H. Allen, Gregory M. Ferrence
***J. Appl. Cryst. Special Teaching Edition*, 2010, 43, 1208-1223**



Free Educational Resources

- Teaching subset of 500 CSD entries chosen to illustrate a wide range of 3D structural issues
- Web-based interface for browsing the teaching subset
- Tutorials, class exercises and worksheets

http://www.ccdc.cam.ac.uk/free_services/teaching/

NSF Discovery Corps Fellowship Grant No. 0725294





Part #2

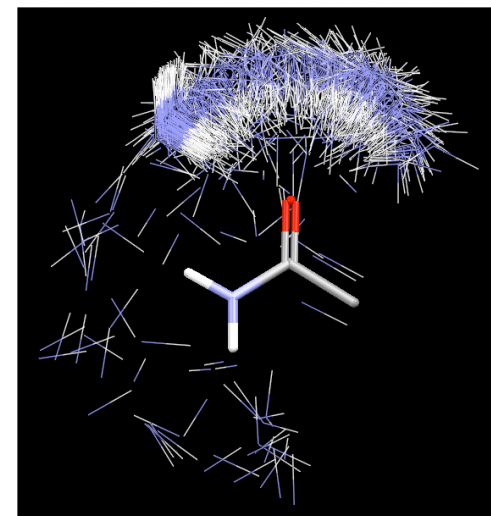
- IsoStar and SuperStar
 - IsoStar: Assessing non-bonded interactions
 - SuperStar: using IsoStar data to predicting binding site interactions



IsoStar

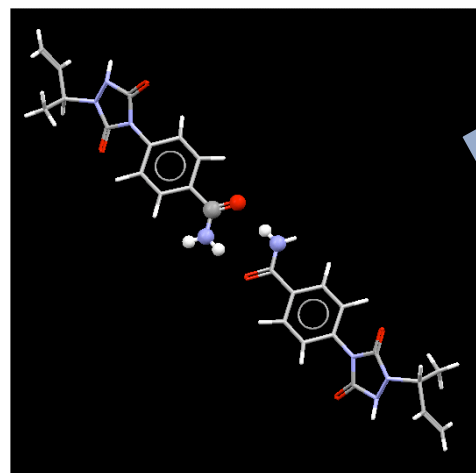
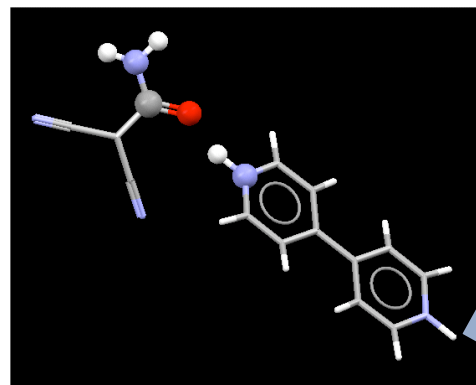
A Knowledge Base of Intermolecular Interactions

- Experimental data taken from:
 - Cambridge Structural Database
 - Protein Data Bank (protein-ligand complexes only)
- Interaction distributions displayed as scatterplots or contour surfaces



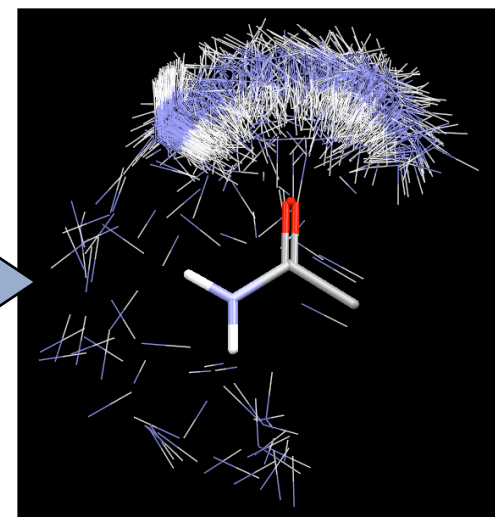
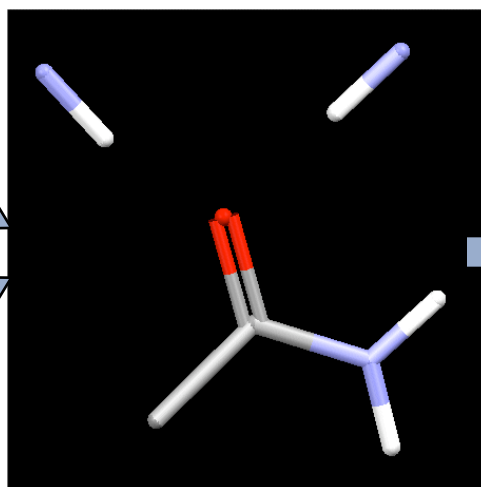


IsoStar Methodology



central group: -CONH_2

contact group: NH

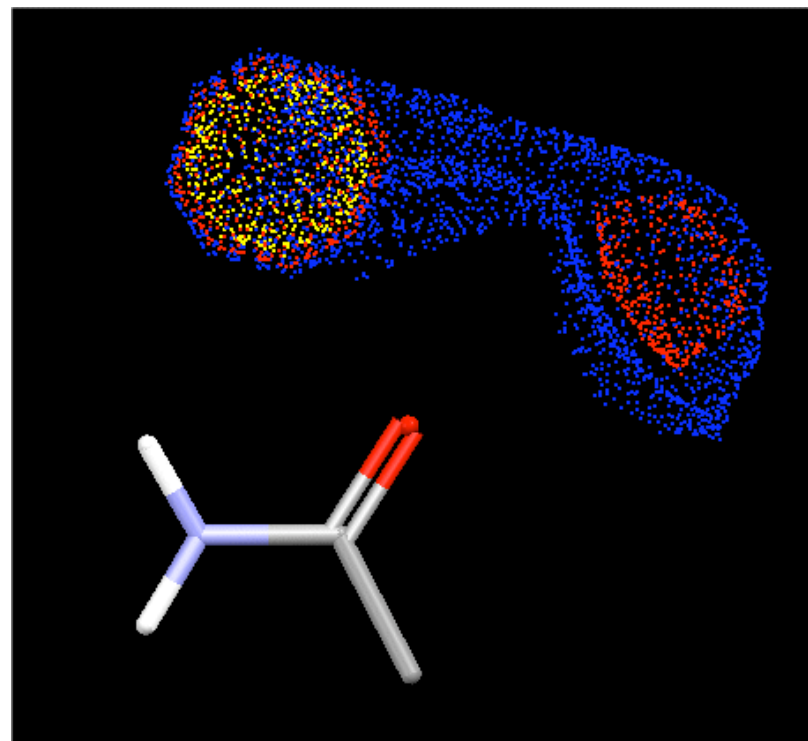
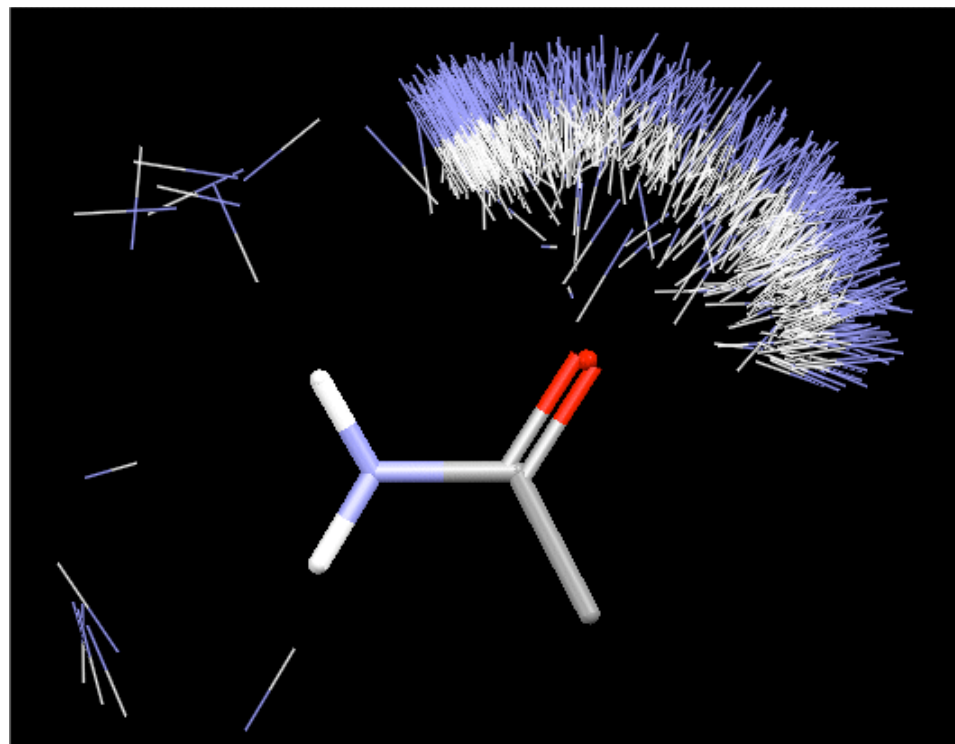


Search CSD or PDB for structures containing desired contact

Superimpose hits and display as scatterplots



Density Maps



Can also represent distribution as density maps



Coverage of Groups

J.Comp.-Aided Mol. Des., 11, 525-537, 1997

- Experimental data from CSD and PDB protein-ligand complexes
- >300 central groups, >50 contact groups
- >22,000 CSD scatterplots
- >7,400 PDB scatterplots
- 1,550 theoretical energy minima from DMA/IMPT

Generate your own scatterplots using IsoGen



Typical Uses of IsoStar

Probability of an interaction occurring

Will fluorine act as an H-bond acceptor?

Is the N or the O of oxazole more likely to H-bond?

Preferred geometries

Do H-bonds to ether O lie along lone pairs?

Do aromatic rings stack or form T-shape arrangements?

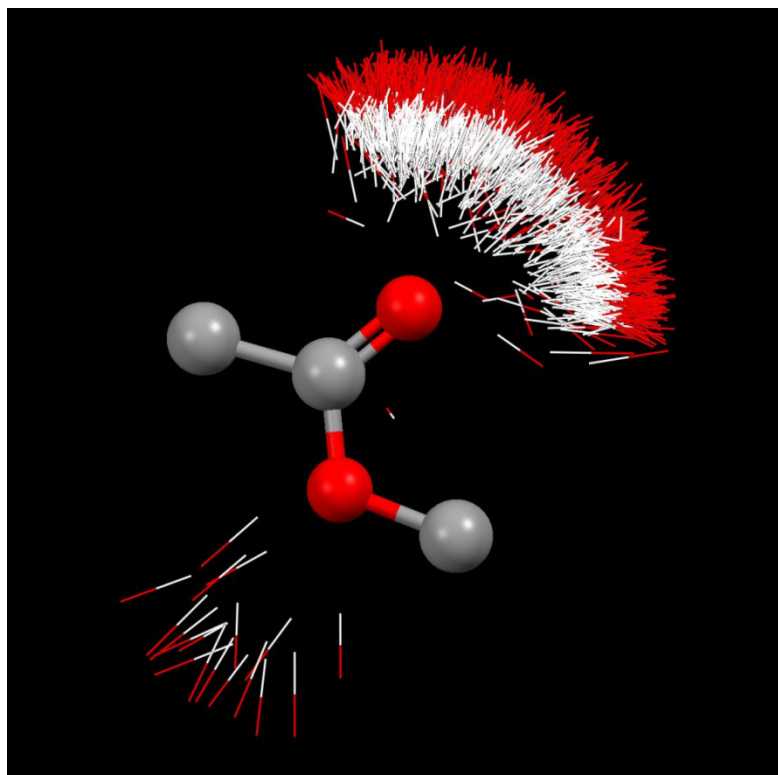
Design strategies

Do thiazole and oxazole form similar interactions?

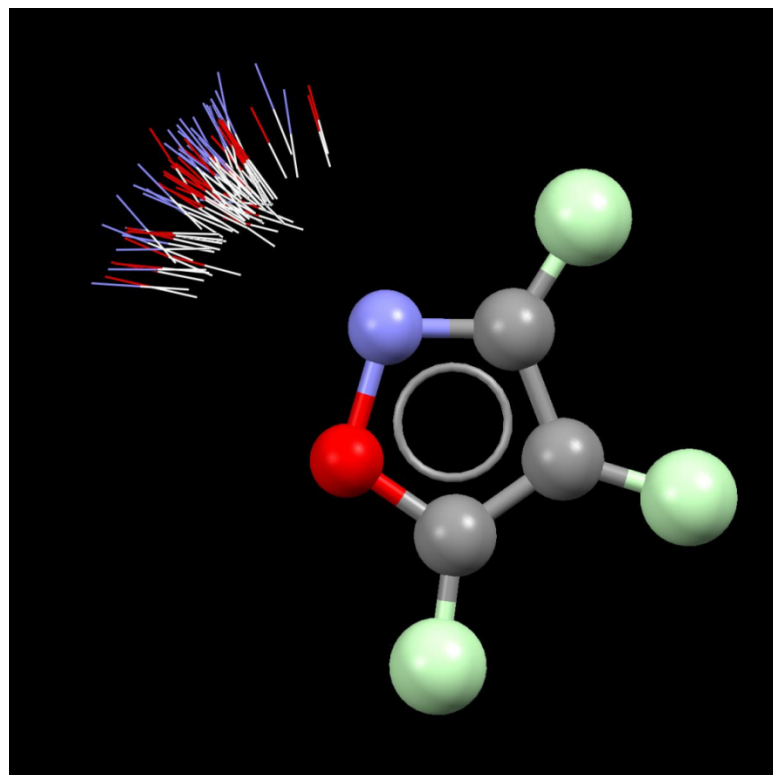
How can we bind to a Trypsin residue?



Strong acceptors compete for donor-H groups



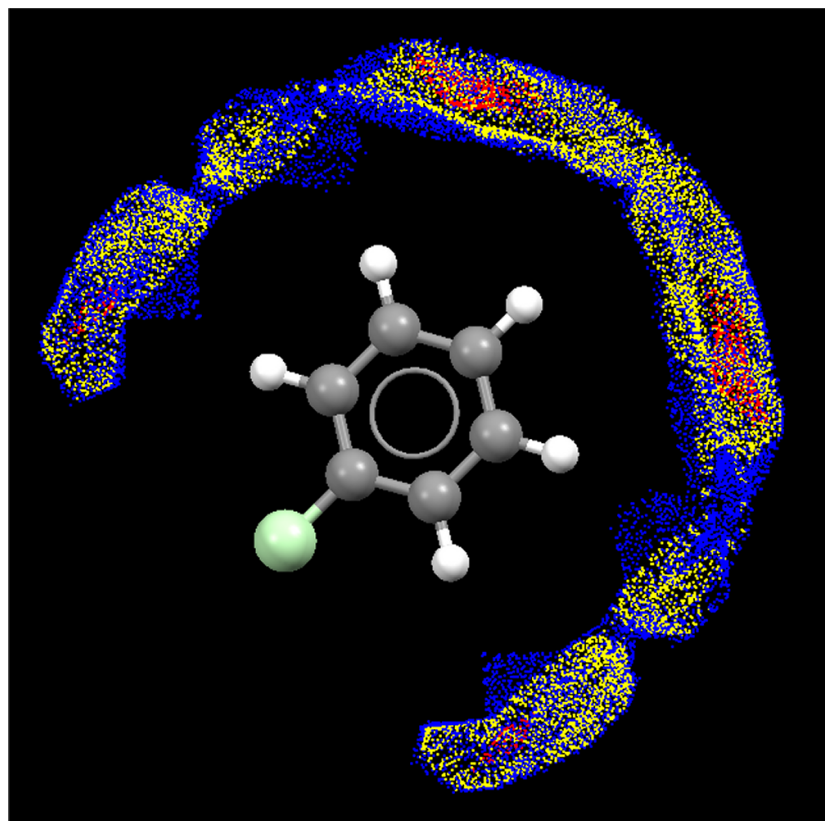
Scatterplot of O-H...O H-bonds in esters



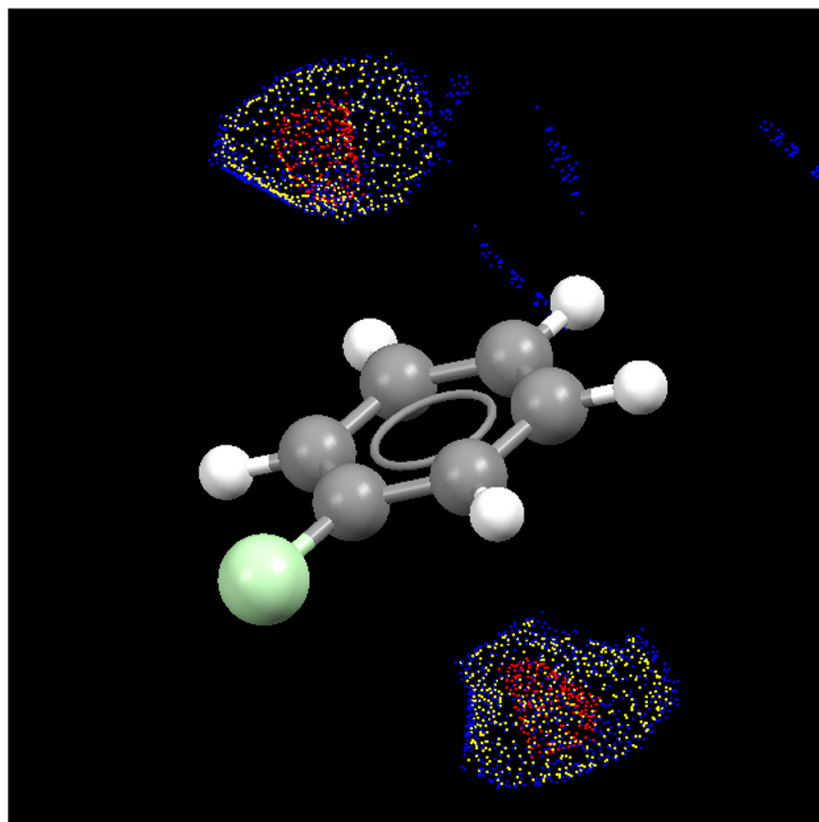
O-H and N-H H-bonds to N- and O-acceptors in isoxazole rings



Electronegativity and electropositivity



carbonyl ($-\text{C}=\text{O}^{\delta-}$) groups around a phenyl ($-\text{C}_6\text{H}_5$) central group



alkyl C-H around a phenyl ($-\text{C}_6\text{H}_5$) central group



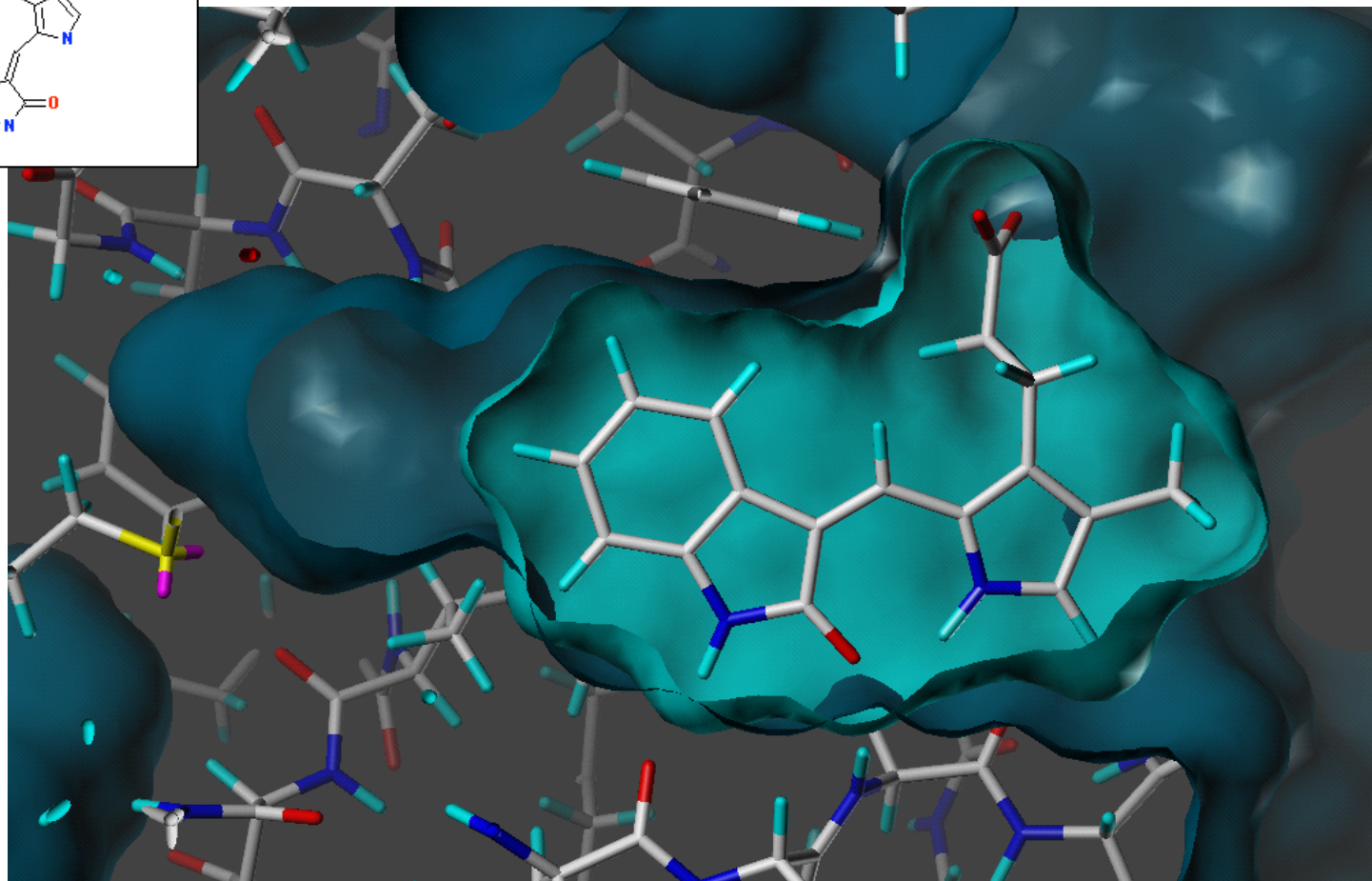
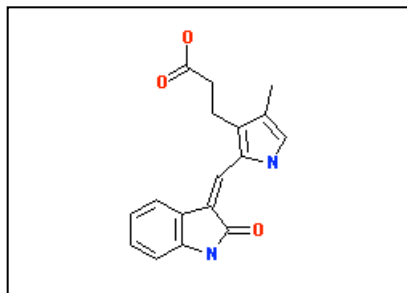
SuperStar

Knowledge-base can provide input for other software

- SuperStar is a program for identifying interaction sites in proteins or around small molecules
- Predicts where a given functional group ('probe') is likely to interact at binding site
- Based exclusively on experimental data in IsoStar
- Validated successfully on >300 protein-ligand complexes from PDB
- Uses include pharmacophore generation

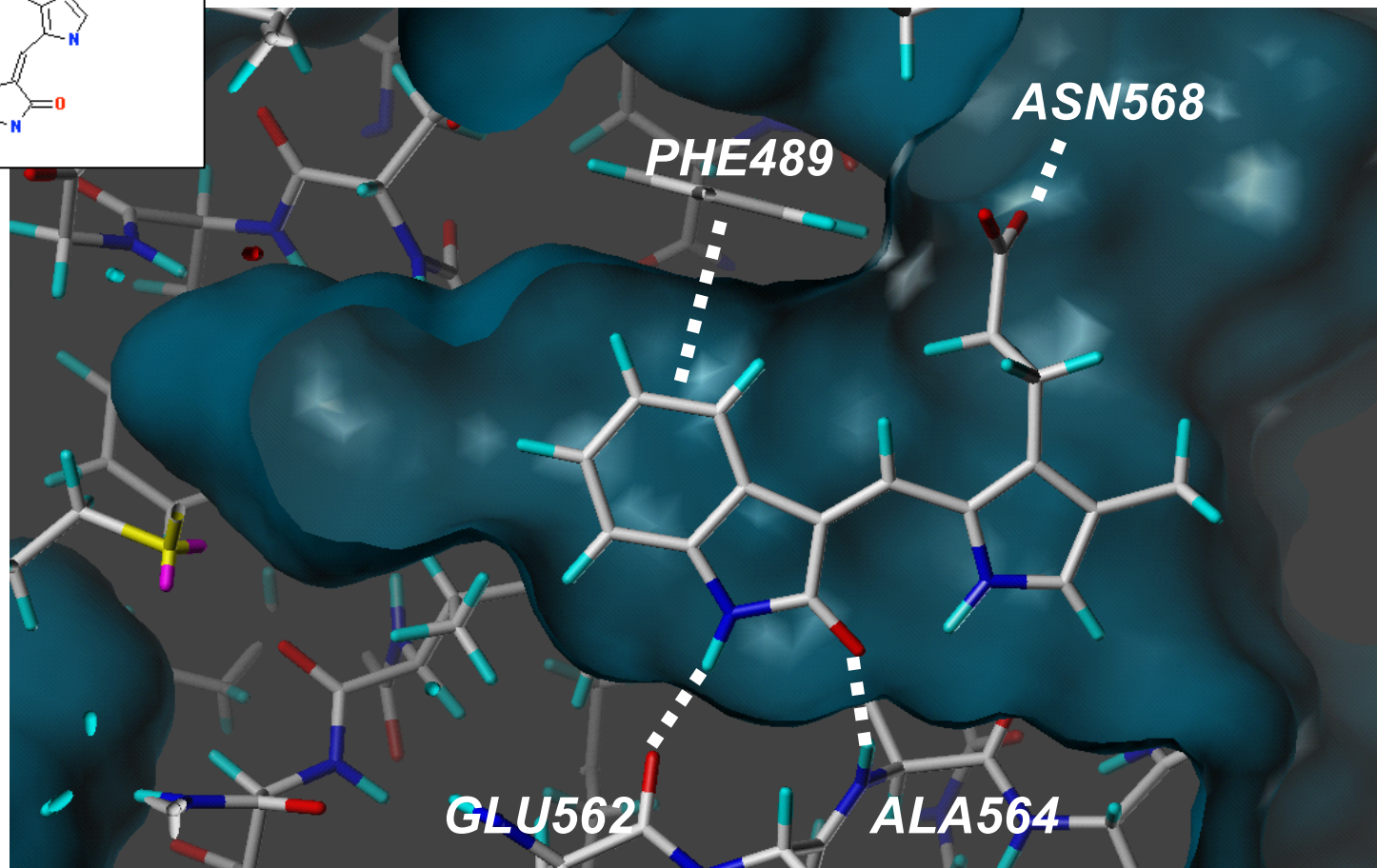
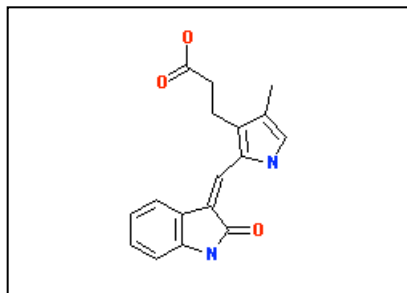


Tyrosine kinase (*1fgi*)



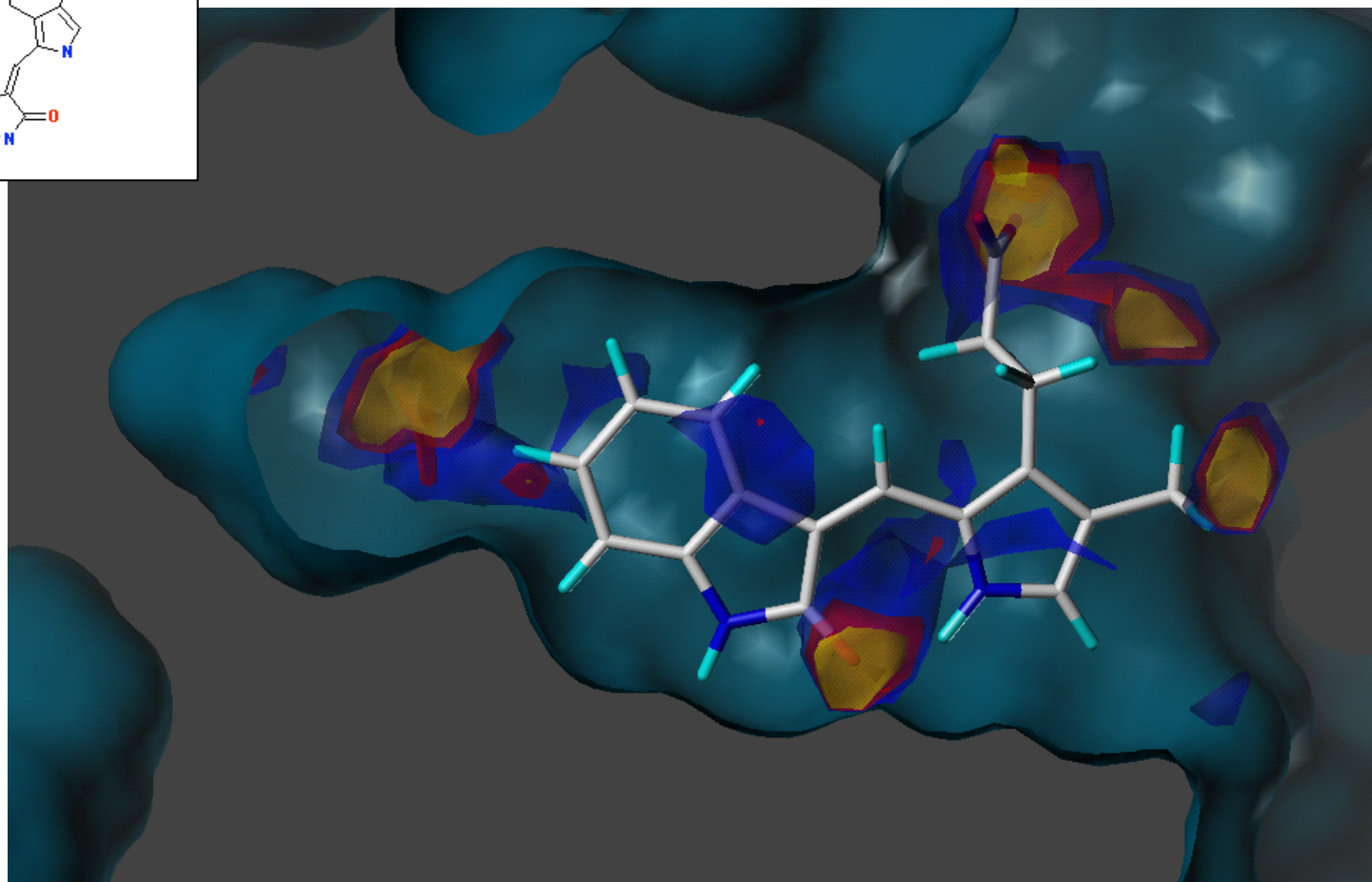
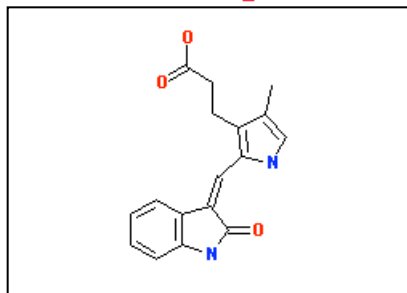


Tyrosine kinase (*1fgi*)



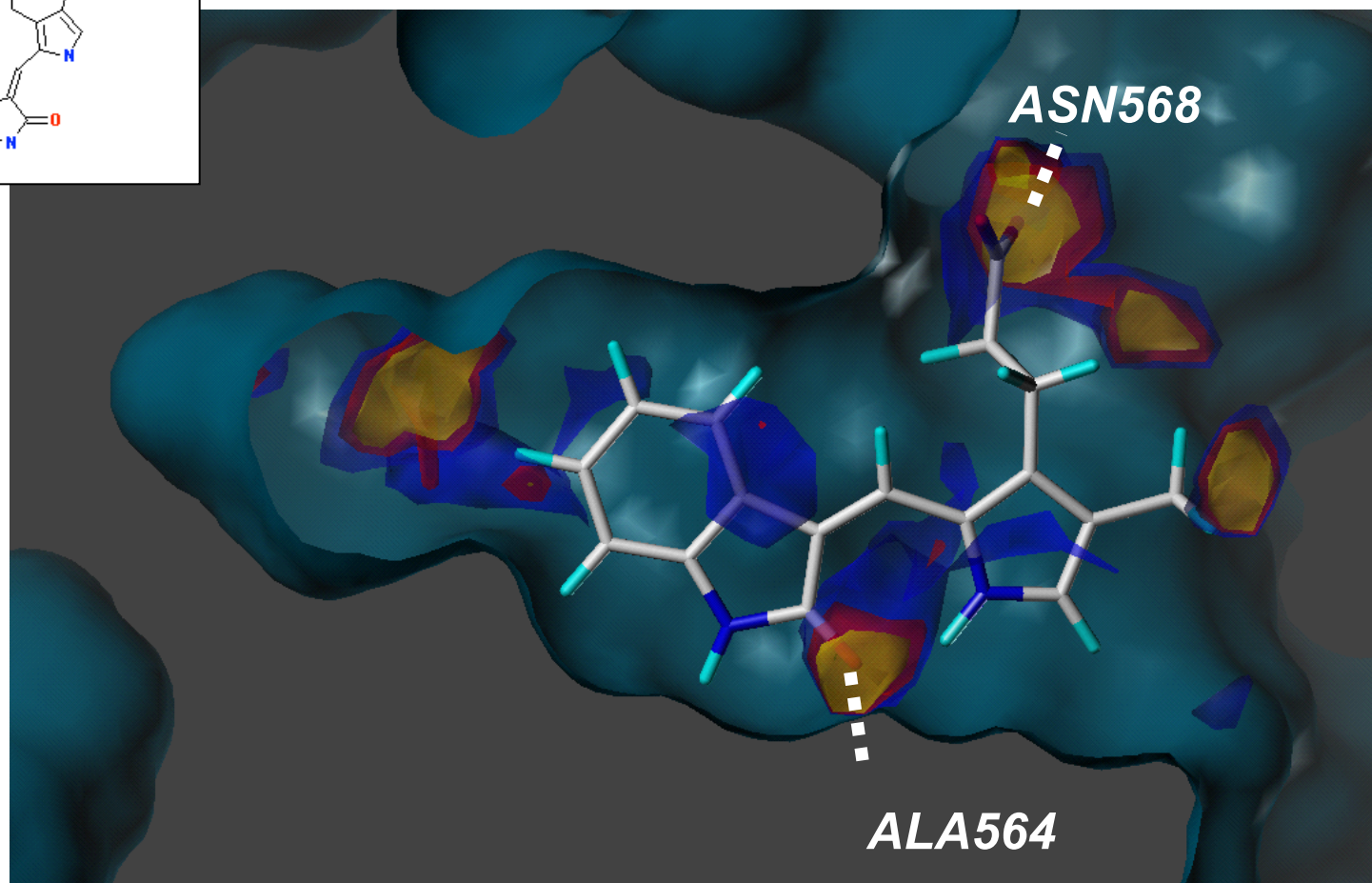
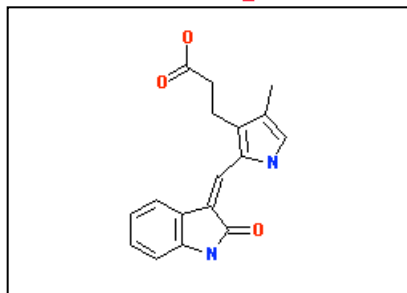


Map for CO Oxygen Probe



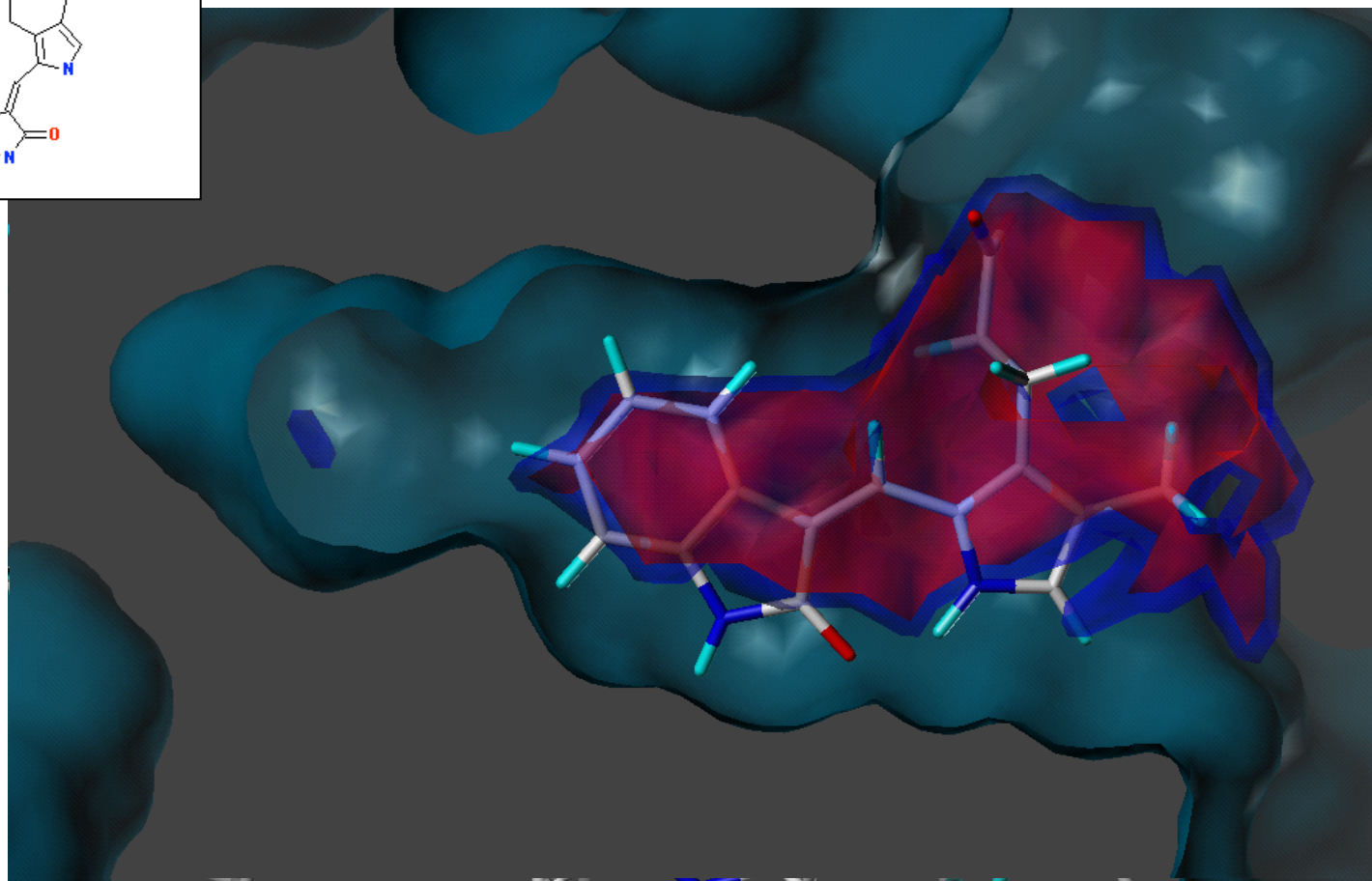
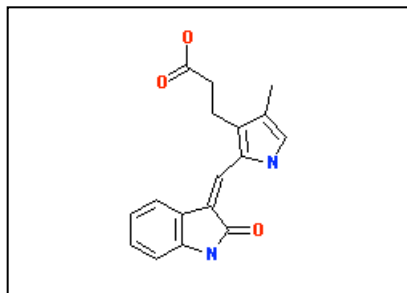


Map for CO Oxygen Probe





Map for aromatic CH carbon probe





Thank you!

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