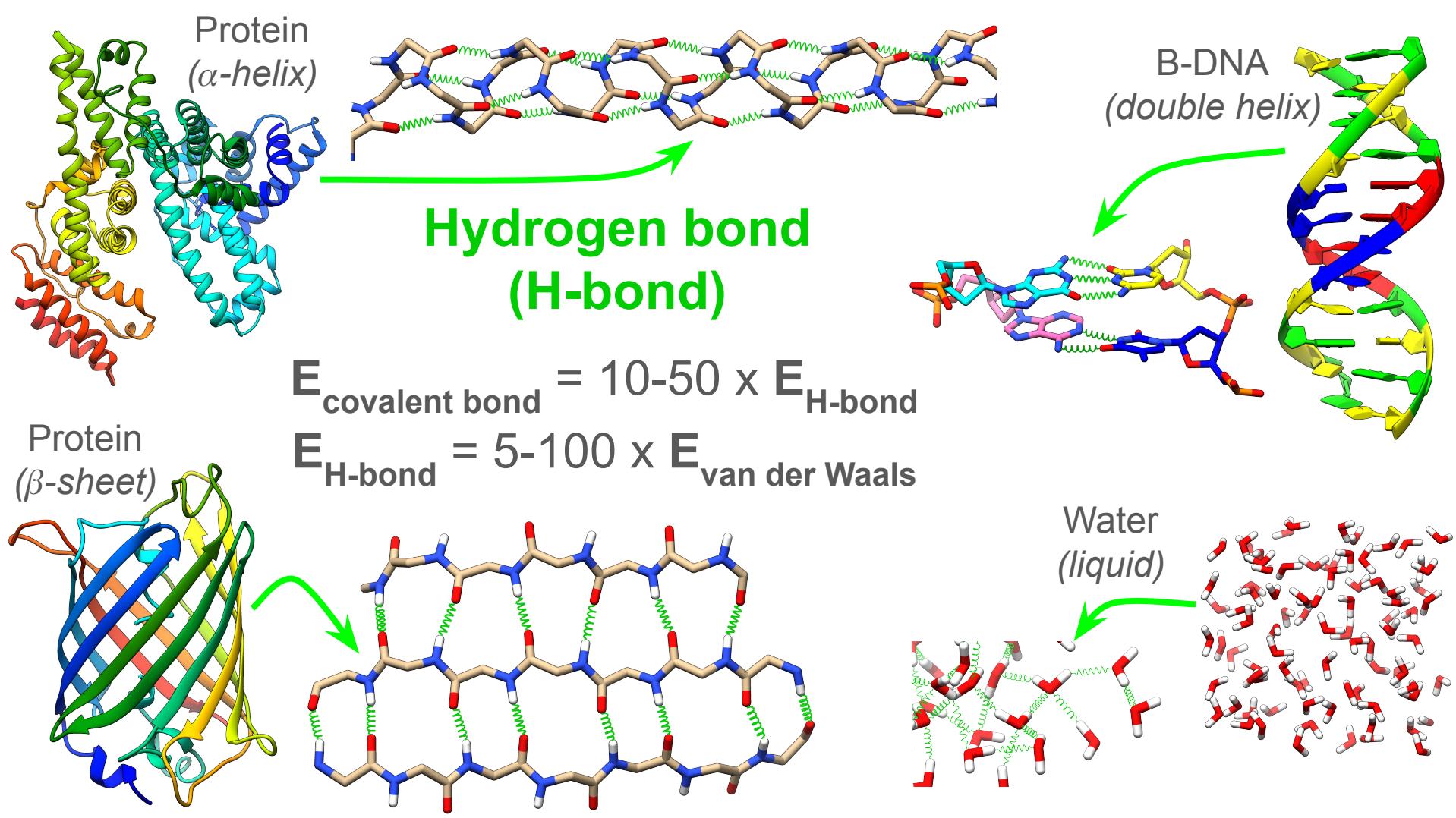


Development and application of parallel CPU/GPU tools to extend the characterization of direct and solvent-mediated H-bond interactions in large structural protein databases

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Rosa Maria Vitale (*ICB-CNR*), Melania Paturzo (*ISASI-CNR*),
Pietro Amodeo (*ICB-CNR*)





Protein Data Bank

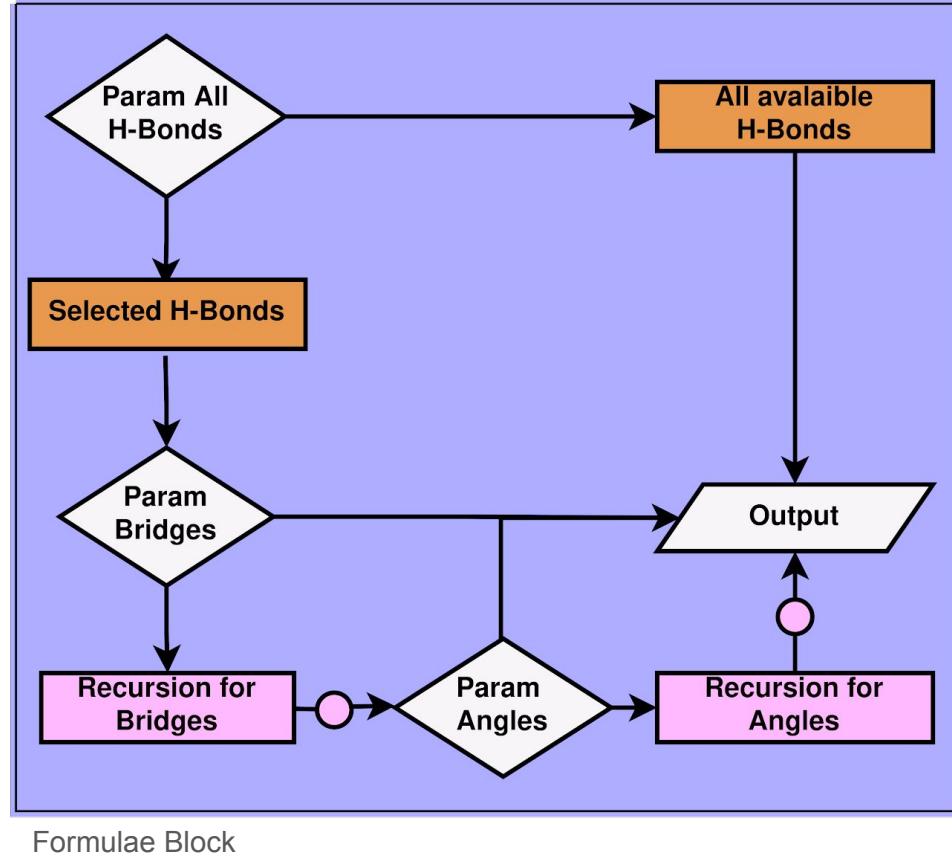
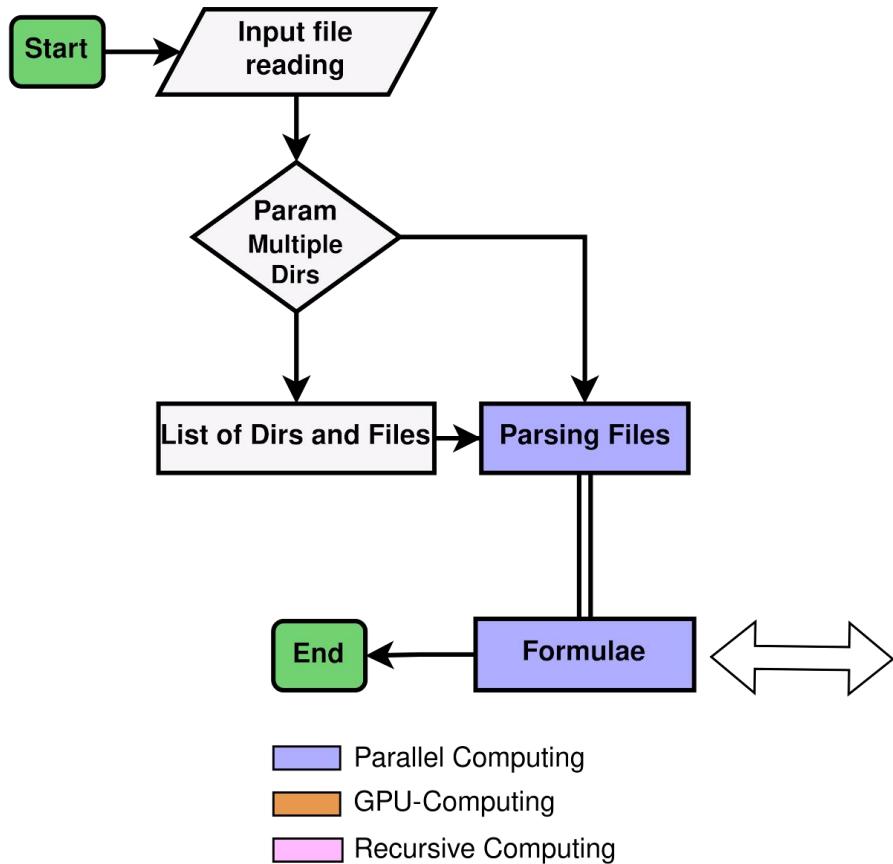
147803 are Protein Complexes

The screenshot shows the RCSB PDB homepage. At the top center, it displays "217,966 Structures from the PDB" with a red box around the number. Below this, there are two other statistics: "1,068,577 Computed Structure Models (CSM)". To the right, there is a link to "3D Structures" with a question mark icon. At the bottom of the header, there are several links: "PDB-101", "www.PDB", "EMDataResource", "NAKB", "WWPDB Foundation", and "PDB-Dev".

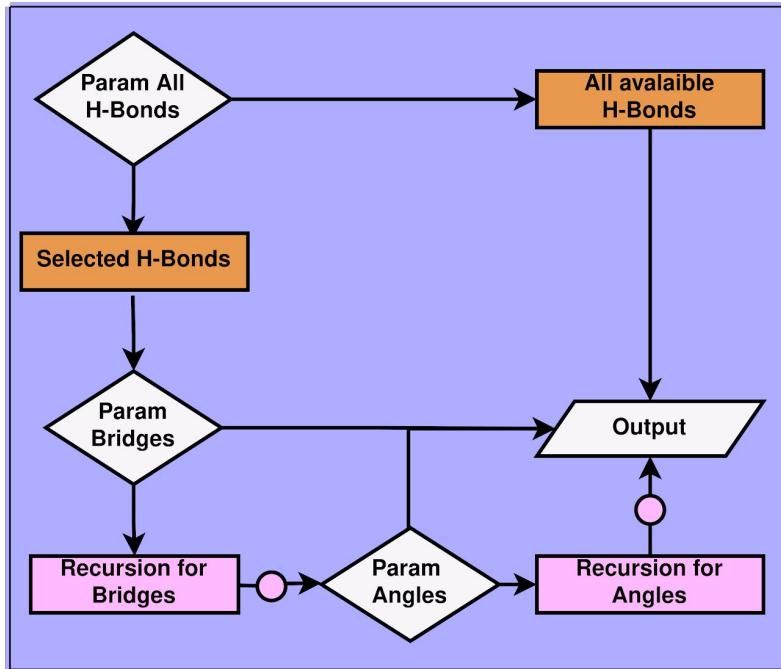
PDB Format Example

ATOM	1	N	GLY	A	6	21.232	-3.592	-12.215	1.00	10.46		N
ANISOU	1	N	GLY	A	6	1537	1300	1136	110	-528	-59	N
ATOM	2	CA	GLY	A	6	21.163	-4.550	-11.066	1.00	10.57		C
ANISOU	2	CA	GLY	A	6	1558	1396	1063	-267	-343	87	C
ATOM	3	C	GLY	A	6	22.447	-4.422	-10.244	1.00	8.93		C
ANISOU	3	C	GLY	A	6	1193	1290	908	-97	-117	152	C
ATOM	4	O	GLY	A	6	23.428	-3.846	-10.715	1.00	10.86		O
ANISOU	4	O	GLY	A	6	1267	1673	1187	-79	-127	346	O
ATOM	5	N	SER	A	7	22.415	-4.967	-9.066	1.00	9.67		N
ANISOU	5	N	SER	A	7	1214	1683	777	-108	-24	51	N
ATOM	6	CA	SER	A	7	23.627	-4.941	-8.213	1.00	10.00		C
ANISOU	6	CA	SER	A	7	1162	1784	855	89	-78	87	C

Algorithm flowchart



Formulae and Input



Parallel Computing
GPU-Computing
Recursive Computing

Input.txt example

```
1 output_directory= /home/user/Desktop
2 output_file_name= IBiSCO_ws
3 output_extension= all
4 input_resolution= 1.5
5 input_directory= /home/user/Dataset
6 input_multi_dir= True
7 input_multi_name= DB_CIF_
8 input_multi_numb= 9
9 molecule_standard_res= ARG(NE)
10 molecule_define_res_1= HOH, 0: don-acceptors
11 calculate_direct_hbonds= ARG
12 calculate_bridges_hbond= True
13 calculate_bridges_set= 3
14 calculate_angles= True
15 calculate_angles_define_1= HOH(0) | ARG(NE), acc
16 calculate_angles_define_2= HOH(0) | ARG(NE), don
17 calculate_angles_define_3= ARG(NE) | acc, HOH(0)
18 calculate_angles_define_4= ARG(NE) | acc
```

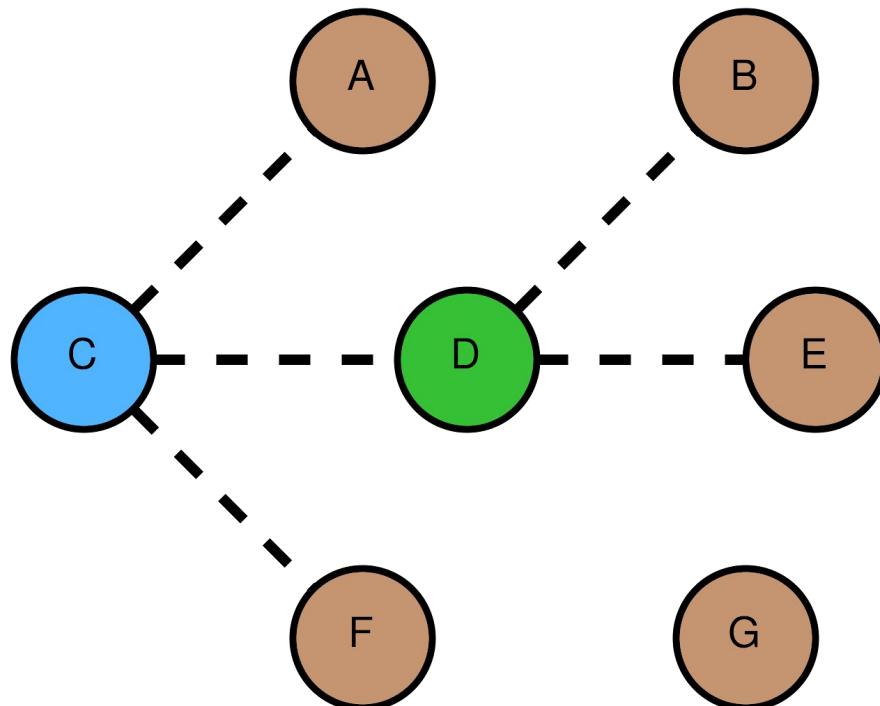
Descriptors highlight

```
molecule_standard_res= ARG(NE)
molecule_define_res_1= HOH, 0: don-acceptors
```

Bridging H-Bonds

C molecule_standard_res=
D molecule_define_res_1=

ARG(NE)
HOH, O: don-acceptors



Paths	Length 5	Length 3
	A - C - D - B - E	A - C - D
	F - C - D - B - E	A - C - F
Length 4	F - C - D	F - C - D
	A - C - D - B	C - D - B
	A - C - D - E	B - D - E
	F - C - D - B	C - D - E
	F - C - D - E	

Selected Atom

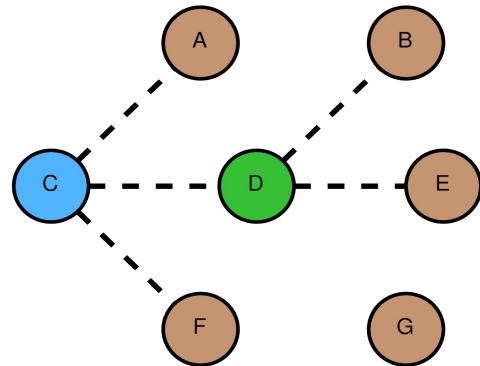
Defined Atcm

Other Donors-Acceptors

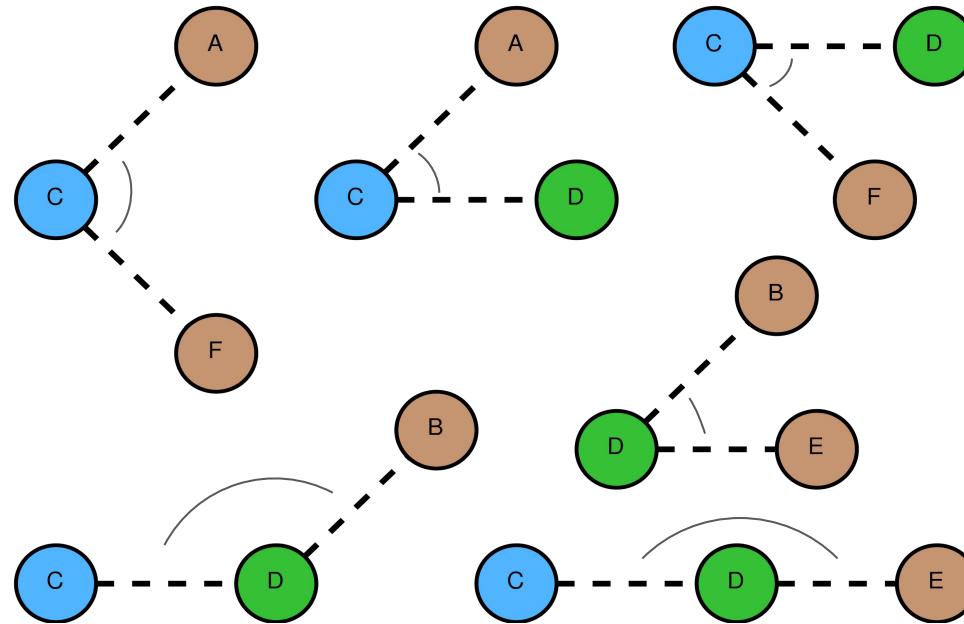
Chem3D Pro 12.0.2

Bridge Angles

```
calculate_angles_define_1= HOH(0) | ARG(NE), acc  
calculate_angles_define_2= HOH(0) | ARG(NE), don  
calculate_angles_define_3= ARG(NE) | acc, HOH(0)  
calculate_angles_define_4= ARG(NE) | acc
```



- Selected Atom
- Defined Atom
- Other Donors-Acceptors



Output Examples

michele@schielhau: ~/HBF													
(base) michele@schielhau:~/HBF\$ cat IBiSCo_ws.txt grep 2A6Z													
2A6Z Direct H-Bonds													
2A6Z ARG NE A 43 (289) ['GLU OE1 A 217 (1682) d. 3.063']	1	2A6Z	Direct H-Bonds										
2A6Z ARG NE A 75 (558) ['GLU OE1 A 217 (1682) d. 2.866']	2	2A6Z	ARG NE A 43 (289)	GLU OE1 A 217 (1682) d. 3.063									
2A6Z ARG NE A 99 (744) ['HOH O B 424 (1970) d. 3.203']	3	2A6Z	ARG NE A 75 (558)	GLU OE1 A 217 (1682) d. 2.866									
2A6Z ARG NE A 128 (971) ['THR OG1 A 126 (951) d. 2.974']	4	2A6Z	ARG NE A 99 (744)	HOH O B 424 (1970) d. 3.203									
2A6Z ARG NE A 165 (1263) ['HOH O B 350 (1896) d. 3.345' , 'SER OG A 7 (10) d. 3.384' , 'GLU OE2 A 71 (518) d. 2.915']	5	2A6Z	ARG NE A 128 (971)	THR OG1 A 126 (951) d. 2.974									
2A6Z ARG NE A 189 (1467) ['ASP O A 172 (1321) d. 2.926']	6	2A6Z	ARG NE A 165 (1263)	HOH O B 350 (1896) d. 3.345	SER OG A 7 (10) d. 3.384	GLU OE2 A 71 (518) d. 2.915							
2A6Z ARG NE A 199 (1552) ['HOH O B 273 (1819) d. 2.859']	7	2A6Z	ARG NE A 189 (1467)	ASP O A 172 (1321) d. 2.926									
2A6Z Other H-Bonds	8	2A6Z	ARG NE A 199 (1552)	HOH O B 273 (1819) d. 2.859									
2A6Z HOH O B 424 (1970) ['ARG NH1 A 99 (746) d. 3.454']	9	2A6Z	Other H-Bonds										
2A6Z THR OG1 A 126 (951) ['HOH O B 229 (1775) d. 2.722' , 'HOH O B 233 (1779) d. 3.086']	10	2A6Z	GLU OE1 A 217 (1682)										
2A6Z HOH O B 350 (1896) ['GLU OE2 A 71 (518) d. 2.89']	11	2A6Z	HOH O B 424 (1970)	ARG NH1 A 99 (746) d. 3.454									
2A6Z SER OG A 7 (10) ['HOH O B 367 (1913) d. 2.743' , 'GLU OE2 A 71 (518) d. 3.384']	12	2A6Z	THR OG1 A 126 (951)	HOH O B 229 (1775) d. 2.722	HOH O B 233 (1779) d. 3.086								
2A6Z GLU OE2 A 71 (518) ['HOH O B 350 (1896) d. 2.89']	13	2A6Z	HOH O B 350 (1896)	GLU OE2 A 71 (518) d. 2.89									
2A6Z ASP O A 172 (1321) ['ASN N A 174 (1337) d. 3.377' , 'LEU N A 175 (1345) d. 3.47']	14	2A6Z	SER OG A 7 (10)	HOH O B 367 (1913) d. 2.743	GLU OE2 A 71 (518) d. 2.67	ARG NH2 A 165 (1266) d. 3.085							
2A6Z HOH O B 273 (1819) ['LEU O A 58 (404) d. 2.711' , 'ARG NH2 A 199 (1555) d. 3.39']	15	2A6Z	GLU OE2 A 71 (518)	HOH O B 350 (1896) d. 2.89									
2A6Z Bridging H-Bonds	16	2A6Z	ASP O A 172 (1321)	ASN N A 174 (1337) d. 3.377	LEU N A 175 (1345) d. 3.47								
2A6Z [ARG NE A 99 (744) , ARG NH1 A 99 (746) , HOH O B 424 (1970)]	17	2A6Z	HOH O B 273 (1819)	LEU O A 58 (404) d. 2.711	ARG NH2 A 199 (1555) d. 3.39	ILE N A 200 (1556) d. 3.127							
2A6Z [GLU OE2 A 71 (518) , ARG NE A 165 (1263) , HOH O B 350 (1896)]	18	2A6Z	Bridging H-Bonds										
2A6Z [ASP O A 172 (1321) , ASN N A 174 (1337) , ARG NE A 189 (1467)]	19	2A6Z	ARG NE A 99 (744)	ARG NH1 A 99 (746)	HOH O B 424 (1970)								
2A6Z [LEU O A 58 (404) , ARG NE A 199 (1552) , HOH O B 273 (1819)]	20	2A6Z	GLU OE2 A 71 (518)	ARG NE A 165 (1263)	HOH O B 350 (1896)								
2A6Z [ARG NE A 199 (1552) , ARG NH2 A 199 (1555) , HOH O B 273 (1819)]	21	2A6Z	ASP O A 172 (1321)	ASN N A 174 (1337)	ARG NE A 189 (1467)								
2A6Z [ARG NE A 199 (1552) , ILE N A 200 (1556) , HOH O B 273 (1819)]	22	2A6Z	ASP O A 172 (1321)	LEU N A 175 (1345)	ARG NE A 189 (1467)								
2A6Z [SER OG A 7 (10) , GLU OE2 A 71 (518) , ARG NE A 165 (1263) , HOH O B 350 (1896)]	23	2A6Z	LEU O A 58 (404)	ARG NE A 199 (1552)	HOH O B 273 (1819)								
2A6Z [SER OG A 7 (10) , GLU OE2 A 71 (518) , ARG NE A 165 (1263) , HOH O B 350 (1896)]	24	2A6Z	ARG NE A 199 (1552)	ARG NH2 A 199 (1555)	HOH O B 273 (1819)								
2A6Z [SER OG A 7 (10) , GLU OE2 A 71 (518) , HOH O B 350 (1896)]	25	2A6Z	ARG NE A 199 (1552)	ILE N A 200 (1556)	HOH O B 273 (1819)								
2A6Z [SER OG A 7 (10) , ARG NE A 165 (1263) , HOH O B 350 (1896)]	26	2A6Z	SER OG A 7 (10)	GLU OE2 A 71 (518)	ARG NE A 165 (1263)	HOH O B 350 (1896)							
2A6Z GLU OE2 A 71 (518) ARG NE A 165 (1263) HOH O B 350 (1896) 54.457 degr	27	2A6Z	SER OG A 7 (10)	GLU OE2 A 71 (518)	ARG NE A 165 (1263)								
2A6Z LEU O A 58 (404) ARG NE A 199 (1552) HOH O B 273 (1819) 38.817 degr	28	2A6Z	SER OG A 7 (10)	GLU OE2 A 71 (518)	HOH O B 350 (1896)								
2A6Z SER OG A 7 (10) ARG NE A 165 (1263) GLU OE2 A 71 (518) 49.533 degr	29	2A6Z	SER OG A 7 (10)	ARG NE A 165 (1263)	HOH O B 350 (1896)								
2A6Z SER OG A 7 (10) ARG NE A 165 (1263) HOH O B 350 (1896) 63.318 degr	30	2A6Z	Bridge Angles										
	31	2A6Z	GLU OE2 A 71 (518)	HOH O B 273 (1819) HOH O B 273 (1819) 55.168 degrees									
	32	2A6Z	LEU O A 58 (404)	HOH O B 273 (1819) HOH O B 273 (1819) 99.796 degrees									
	33	2A6Z	SER OG A 7 (10)	HOH O B 273 (1819) HOH O B 273 (1819) 47.915 degrees									
	34	2A6Z	SER OG A 7 (10)	HOH O B 273 (1819) HOH O B 273 (1819) 58.873 degrees									
	35	2A6Z	ARG NE A 99 (744)	HOH O B 273 (1819) HOH O B 273 (1819) 39.954 degrees									
	36	2A6Z	ARG NE A 199 (1552)	HOH O B 273 (1819) HOH O B 273 (1819) 41.471 degrees									
	37	2A6Z	ARG NE A 199 (1552)	HOH O B 273 (1819) HOH O B 273 (1819) 113.444 degrees									
	38	2A6Z	GLU OE2 A 71 (518)	ARG NE A 165 (1263) ARG NE A 165 (1263) 54.457 degr									
	39	2A6Z	LEU O A 58 (404)	ARG NE A 165 (1263) ARG NE A 165 (1263) 38.817 degr									
	40	2A6Z	SER OG A 7 (10)	ARG NE A 165 (1263) ARG NE A 165 (1263) 49.533 degr									
	41	2A6Z	SER OG A 7 (10)	ARG NE A 165 (1263) ARG NE A 165 (1263) 63.318 degr									

TXT - CSV

Output Examples

```

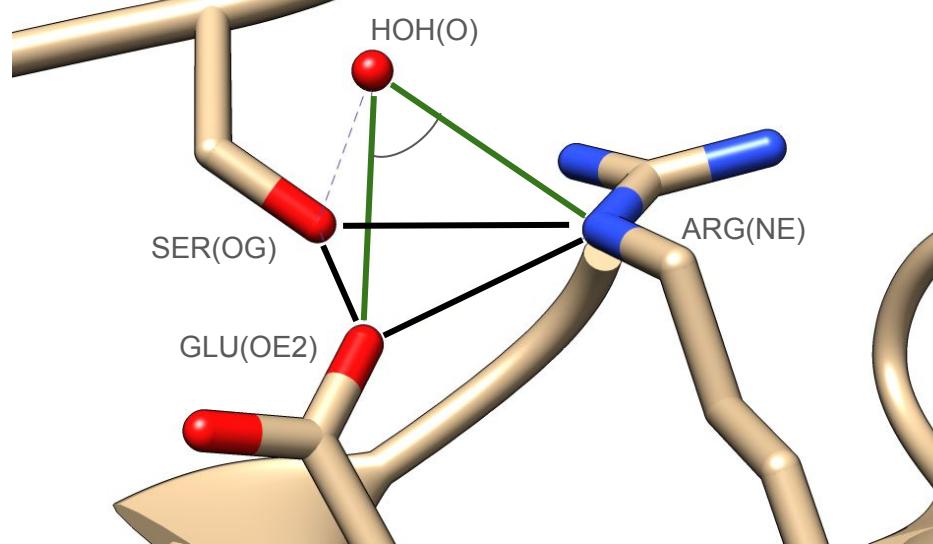
1 [
2   "2A6Z": {
3     "Direct H-Bonds": {
4       "ARG NE A 43 (289)": "[ 'GLU OE1 A 217 (1682) d. 3.063' ]",
5       "ARG NE A 75 (558)": "[ 'GLU OE1 A 217 (1682) d. 2.866' ]",
6       "ARG NE A 99 (744)": "[ 'HOH O B 424 (1970) d. 3.203' ]",
7       "ARG NE A 128 (971)": "[ 'THR OG1 A 126 (951) d. 2.974' ]",
8       "ARG NE A 165 (1263)": "[ 'HOH O B 350 (1896) d. 3.345', 'SER OG A 7 (10) d. 3.384', 'GLU OE2 A 71 (518) d. 2.915' ]",
9       "ARG NE A 189 (1467)": "[ 'ASP O A 172 (1321) d. 2.926' ]",
10      "ARG NE A 199 (1552)": "[ 'HOH O B 273 (1819) d. 2.859' ]"
11    }
12  },
13  "2A6Z": {
14    "Other H-Bonds": {
15      "GLU OE1 A 217 (1682)": "[ ]",
16      "HOH O B 424 (1970)": "[ 'ARG NH1 A 99 (746) d. 3.454' ]",
17      "THR OG1 A 126 (951)": "[ 'HOH O B 229 (1775) d. 2.722', 'HOH O B 233 (1779) d. 3.086' ]",
18      "HOH O B 350 (1896)": "[ 'GLU OE2 A 71 (518) d. 2.89' ]",
19      "SER OG A 7 (10)": "[ 'HOH O B 367 (1913) d. 2.743', 'GLU OE2 A 71 (518) d. 2.673', 'ARG NH2 A 165 (1266) d. 3.085' ]",
20      "GLU OE2 A 71 (518)": "[ 'HOH O B 350 (1896) d. 2.89' ]",
21      "ASP O A 172 (1321)": "[ 'ASN N A 174 (1337) d. 3.377', 'LEU N A 175 (1345) d. 3.47' ]",
22      "HOH O B 273 (1819)": "[ 'LEU O A 58 (404) d. 2.711', 'ARG NH2 A 199 (1555) d. 3.395', 'ILE N A 200 (1556) d. 3.127' ]"
23    }
24  },
25  "2A6Z": {
26    "Bridging H-Bonds": {
27      "1": "[ARG NE A 99 (744), ARG NH1 A 99 (746), HOH O B 424 (1970)]",
28      "2": "[GLU OE2 A 71 (518), ARG NE A 165 (1263), HOH O B 350 (1896)]",
29      "3": "[ASP O A 172 (1321), ASN N A 174 (1337), ARG NE A 189 (1467)]",
30      "4": "[ASP O A 172 (1321), LEU N A 175 (1345), ARG NE A 189 (1467)]",
31      "5": "[LEU O A 58 (404), ARG NE A 199 (1552), HOH O B 273 (1819)]",
32      "6": "[ARG NE A 199 (1552), ARG NH2 A 199 (1555), HOH O B 273 (1819)]",
33      "7": "[ARG NE A 199 (1552), ILE N A 200 (1556), HOH O B 273 (1819)]",
34      "8": "[SER OG A 7 (10), GLU OE2 A 71 (518), ARG NE A 165 (1263), HOH O B 350 (1896)]",
35      "9": "[SER OG A 7 (10), GLU OE2 A 71 (518), ARG NE A 165 (1263), HOH O B 350 (1896)]",
36      "10": "[SER OG A 7 (10), GLU OE2 A 71 (518), ARG NE A 165 (1263), HOH O B 350 (1896)]",
37      "11": "[SER OG A 7 (10), ARG NE A 165 (1263), HOH O B 350 (1896)]"
38    }
39  },
40  "2A6Z": {
41    "Bridge Angles": {
42      "1": "[GLU OE2 A 71 (518) HOH O B 350 (1896)]",
43      "2": "[LEU O A 58 (404) HOH O B 273 (1819)]",
44      "3": "[SER OG A 7 (10) HOH O B 350 (1896)]",
45      "4": "[SER OG A 7 (10) HOH O B 350 (1896)]",
46      "5": "[ARG NE A 99 (744) HOH O B 424 (1970)]",
47      "6": "[ARG NE A 199 (1552) HOH O B 273 (1819)]",
48      "7": "[ARG NE A 199 (1552) HOH O B 273 (1819)]",
49      "8": "[GLU OE2 A 71 (518) ARG NE A 165 (1263)]",
50      "9": "[LEU O A 58 (404) ARG NE A 199 (1552)]",
51      "10": "[SER OG A 7 (10) ARG NE A 165 (1263) GLU OE2 A 71 (518) | 36.811 degrees, HOH O B 350 (1896) | 49.533 degrees]",
52      "11": "[SER OG A 7 (10) ARG NE A 165 (1263) HOH O B 350 (1896) | 63.318 degrees]"
53    }
54  }.
```

JSON - DATABASE

TABELLE		id	Name	Atom	H Bonds
BridgeAngles	11	1 2A6Z	ARG NE A 43 (289)	['GLU OE1 A 217 (1682) d. 3.063']	
BridgingHBonds	1 2	2A6Z	ARG NE A 75 (558)	['GLU OE1 A 217 (1682) d. 2.866']	
DirectHBonds	7 3	2A6Z	ARG NE A 99 (744)	['HOH O B 424 (1970) d. 3.203']	
OtherHBonds	8 4	2A6Z	ARG NE A 128 (971)	['THR OG1 A 126 (951) d. 2.974']	
sqlite_sequence	3 5	2A6Z	ARG NE A 165 (1263)	['HOH O B 350 (1896) d. 3.345', 'SER OG A 7 (10) d. 3.384', 'GLU OE2 A 71 (518) d. 2.915']	
	6 2A6Z		ARG NE A 189 (1467)	['ASP O A 172 (1321) d. 2.926']	
	7 2A6Z		ARG NE A 199 (1552)	['HOH O B 273 (1819) d. 2.859']	

Viewing Results

```
2A6Z ARG NE A 165 (1263) ['HOH O B 350 (1896) d. 3.345', 'SER OG A 7 (10) (1896) d. 3.384', 'GLU OE2 A 71 (518) d. 2.915']  
2A6Z HOH O B 350 (1896) ['GLU OE2 A 71 (518) d. 2.89']  
2A6Z SER OG A 7 (10) ['HOH O B 367 (1913) d. 2.743', 'GLU OE2 A 71 (518) d. 2.673', 'ARG NH2 A 165 (1266) d. 3.085']  
2A6Z GLU OE2 A 71 (518) ['HOH O B 350 (1896) d. 2.89']  
2A6Z [GLU OE2 A 71 (518), ARG NE A 165 (1263), HOH O B 350 (1896)]  
2A6Z [SER OG A 7 (10), GLU OE2 A 71 (518), ARG NE A 165 (1263), HOH O B 350 (1896)]  
2A6Z [SER OG A 7 (10), GLU OE2 A 71 (518), ARG NE A 165 (1263), HOH O B 350 (1896)]  
2A6Z [SER OG A 7 (10), ARG NE A 165 (1263), HOH O B 350 (1896)]  
2A6Z GLU OE2 A 71 (518) HOH O B 350 (1896) ARG NE A 165 (1263) | 55.168 degrees  
2A6Z SER OG A 7 (10) HOH O B 350 (1896) GLU OE2 A 71 (518) | 47.915 degrees  
2A6Z SER OG A 7 (10) HOH O B 350 (1896) ARG NE A 165 (1263) | 58.873 degrees  
2A6Z GLU OE2 A 71 (518) ARG NE A 165 (1263) HOH O B 350 (1896) | 54.457 degrees  
2A6Z SER OG A 7 (10) ARG NE A 165 (1263) GLU OE2 A 71 (518) | 49.533 degrees  
2A6Z SER OG A 7 (10) ARG NE A 165 (1263) HOH O B 350 (1896) | 63.318 degrees
```



Future Developments:

HBF
Platform

- Multi-platform stand-alone program, downloadable, for calculations on local dataset
- Online service with textual input
- Online service with GUI
- Free access to pre-computed community datasets
- Development of further analyses

Thank you for your attention!



Consiglio Nazionale
delle Ricerche



Istituto di
Chimica
Biomolecolare



Institute of Applied Sciences
and Intelligent Systems
"Eduardo Caianiello"

