

Biological halogen bonds in protein-ligand complexes: a combined QTAIM and NCIPLOT study in four representative cases

Antonio Frontera^{a,*} and Antonio Bauzá^{a,*}

*^aDepartment of Chemistry, Universitat de les Illes Balears
Ctra. de Valldemossa km 7.5, 07122 Palma de Mallorca (Spain)
E-mail: toni.frontera@uib.es; antonio.bauza@uib.es*

Table S1
Cartesian coordinates of the PDB models

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Table S1. List of X-ray structures exhibiting biological HaBs with indication of the Ligand and Protein ID, the HaB donor and acceptor counterparts (Interaction), HaB distance (d in Å) and angle (A in °).

Ligand ID	PDB ID	Interaction	d	A
I7T	2Y6Q	I···S (MET375)	2.831	176.6
IMM	1PGF	I···OC(LEU384)	3.748	178.9
IRF	2J08	I···OC(GLY342)	3.351	166.8
ISF	1PGE	I···OC(LEU384)	3.151	177.5
K8Q	6RLW	I···OC(GLY312)	3.431	176.2
IZG	3PP1	I···OC(VAL127)	3.298	178.1
KON	6RZM	I···OC(GLY182)	3.249	165.9
LCJ	6NYB	I···OC(VAL127)	3.607	178.4
LSG	3MBL	I···OC(VAL127)	3.252	176.2
LUG	3EQB	I···OC(VAL127)	3.081	176.7
M15	4LB3	I···O(THR113)	2.947	167.1
M3K	4ARK	I···OC(VAL127)	3.439	178.0
MEK	3DV3	I···OC(VAL127)	3.147	177.2
P96	4AGQ	I···OC(LEU145)	2.978	176.0
P74	4AGO	I···OC(LEU145)	3.000	174.7
P51	4AGP	I···OC(LEU145)	2.989	175.2
NXG	4AGN	I···OC(LEU52)	2.989	174.4
PIH	3DNA	I···SeC(MSE102) I···OC(LEU84)	3.383 3.396	170.5 164.2
PYZ	7NQX	I···OC(THR42)	3.642	170.1
Q2J	6U99	I···OC(GLY135)	3.746	164.7
Q2A	6U9A	I···OC(GLY135)	3.691	161.4
Q2A	6U9B	I···OC(GLY135)	3.818	165.6
Q2D	6U98	I···OC(GLY135)	3.617	164.5
QOA	7M0W	I···OC(VAL127)	3.561	177.1
R22	2BE2	I···OC(TYR188)	3.372	168.5
V04	3V04	I···OC(VAL127)	3.311	176.3
T3W	7B94	I···OC(VAL127)	3.480	175.2
VKD	7JUS	I···OC(VAL127)	3.644	168.9
VRA	3E8N	I···OC(VAL127)	3.309	176.2
YJ8	2YJ8	I···OC(GLY61)	3.121	176.8
YQY	4ANB	I···OC(VAL127)	3.430	173.9
0BT	4L6C	I···OC(ARG163)	3.172	171.4
136	1GJD	I···OC(VAL41)	3.418	169.9
1CX	3DY7	I···OC(VAL127)	3.354	178.6
1WW	4LAZ	I···OC(THR113)	3.031	165.6
21F	4LYJ	I···OC(THR58)	3.276	178.3
21F	4LYH	I···OC(THR58)	3.477	174.8
24I	2C6C	I···OC(ILE456)	3.767	165.4
2P7	4AN9	I···OC(VAL127)	3.777	172.2
33E	4TKI	I···OC(GLY1043)	2.907	179.2
3AC	2B5J	I···OC(TYR188)	3.567	169.4
3BM	3EQC	I···OC(VAL127)	3.113	178.4
3OR	3ORN	I···OC(VAL127)	3.592	179.1
3V0	3V01	I···OC(VAL127)	3.281	173.9
3WH	4X21	I···SC(MET146)	3.051	173.1
42F	4XP5	I···OC(ASP121)	3.273	165.9

4BM	7JUJ	I...OC(VAL127)	3.543	175.4
4BM	7JV0	I...OC(VAL127)	3.747	169.4
4BM	7M0X	I...OC(VAL127)	3.173	177.4
4BM	3EQG	I...OC(VAL127)	3.177	176.5
4BM	3VVH	I...OC(VAL127)	3.198	177.2
4O9	4LBR	I...OC(THR113)	2.933	165.6
573	4MNE	I...OC(VAL127)	3.423	174.0
5EA	1S9I	I...OC(VAL131)	3.165	175.6
5I5	2I6A	I...OC(ILE39)	3.282	162.1
5Y0	4AN3	I...OC(VAL127)	3.436	175.1
61V	5HK2	I...CC(TYR206)	3.572	165.2
77D	3SLS	I...OC(VAL127)	3.130	174.8
7FO	5TMV	I...OC(GLU419)	3.175	170.8
7OV	5U00	I...OC(TYR827)	3.062	176.1
9GH	5O1I	I...OC(LEU145)	2.887	175.0
9GK	5O1G	I...OC(LEU145)	2.919	177.1
9GN	5O1H	I...OC(LEU145)	2.921	177.4
9GO	5O1F	I...OC(LEU145)	2.929	177.6
9GT	5O1E	I...OC(LEU145)	2.913	177.5
9GW	5O1D	I...OC(LEU145)	2.944	177.1
9GZ	5O1C	I...OC(LEU145)	2.873	176.5
BBM	1S9J	I...OC(VAL127)	3.134	176.1
CZI	1UHI	I...OC(THR166)	3.119	174.2
DIZ	1T4E	I...OC(GLN72)	3.085	163.9
ELB	6G3Y	I...OC(GLY312)	3.353	173.4
EUI	7JUY	I...OC(VAL127)	3.861	170.8
FIB	3DN3	I...SC(MET102)	2.867	172.1
FIB	3DN8	I...SeC(MSE102)	2.989	173.9
G9J	6DFN	I...OC(VAL533)	3.089	173.5
9H2	5O1B	I...OC(LEU145)	2.895	176.8
9H5	5O1A	I...OC(LEU145)	2.867	176.9
DIU	2BXL	I...OC(ARG257)	3.469	173.3
DIU	4LUH	I...OC(ASP450)	3.499	170.2
DIU	5OSW	I...OC(GLU449)	3.814	166.2
P84	4AGL	I...OC(LEU145)	2.980	175.3
P86	4AGM	I...OC(LEU145)	2.967	174.4
T2B	3WGW	I...OC(GLY127)	3.069	174.1
Y0V	5AOJ	I...OC(LEU145)	2.914	172.4
9ZZ	5OOH	I...OC(THR177)	3.390	175.1
A0Z	6SPW	I...OC(ASP237)	2.888	175.1
K8X	3KXN	I...OC(VAL116)	2.937	168.1
T4A	1KGI	I...OC(ALA709)	3.074	164.6
T44	4LNX	I...OC(PHE218)	3.060	176.0
T44	1SN0	I...OC(LEU109)	3.295	173.7
T44	1HK1	I...OC(ALA406)	3.516	170.0
T44	1F86	I...OC(SER117)	3.660	176.3
T44	1ICT	I...OC(SER117)	3.302	171.5
T44	1HK2	I...OC(ALA406)	3.720	169.2
T44	1HK3	I...OC(ALA406)	3.532	169.4
T44	1ETB	I...OC(THR109)	3.227	168.5
T44	2CEO	I...OC(GLU377)	3.793	162.1
T44	1IE4	I...OC(ALA109)	3.071	167.0
T44	5MHE	I...OC(MET111)	3.700	164.8

4HY	2QPY	I···CC(TYR834)	3.631	174.7
4HY	3D57	I···OC(PHE272)	3.347	173.4
4HY	3JZC	I···OC(PHE272)	3.237	176.1
4HY	3JZB	I···OC(PHE218)	3.215	173.7
4HY	2PIN	I···OC(PHE272)	3.279	176.4
4HY	2PKL	I···CC(TYR834)	3.476	175.8
4HY	2PIU	I···CC(TYR834)	3.594	175.1
4HY	2PIT	I···CC(TYR834)	3.671	175.1
4HY	1NQ0	I···OC(PHE272)	3.329	175.8
4HY	1NUO	I···OC(PHE272)	3.691	176.1
4HY	1NQ2	I···OC(PHE272)	3.074	174.4
4QL	5B34	I···OC(ILE222)	3.110	177.3
B3I	1BKE	I···OC(ARG257)	3.707	175.8
I3C	3E3S	I···OC(GLY20)	3.486	176.2
I3C	3E79	I···OC(ALA334)	3.040	161.1
I3C	3E3D	I···OC(TYR20)	3.004	162.3
I3C	3E3T	I···OC(LEU141)	2.829	169.4
I3C	4NC7	I···OC(GLU34)	2.110	167.7
I3C	4BVX	I···OC(LEU171)	3.071	168.1
I3C	3Q7S	I···OC(LEU23)	2.791	177.2
I3C	5U84	I···OC(ARG193)	2.901	168.5
I3C	6EG7	I···OC(ASP146)	2.879	174.4
I3C	6MRY	I···OC(GLN3)	3.122	165.2
I3C	6NQZ	I···OC(THR74)	3.400	170.3
I3C	6QE4	I···OC(LEU292)	3.336	166.4
I3C	7JSY	I···OC(SER312)	3.309	173.8
T3	2H77	I···OC(PHE218)	3.118	176.6
T3	4NLW	I···OC(PHE218)	3.057	175.3
T3	3VKX	I···OC(GLY127)	3.068	171.1
T3	4LNX	I···OC(PHE218)	3.060	176.0
T3	1BSX	I···OC(PHE272)	3.325	175.8
T3	3GWS	I···OC(PHE272)	3.101	174.5
T3	4ZO1	I···OC(GLY344)	3.107	162.1

Cartesian coordinates of the PDB models

3DNA@MSE102

C	4.1787552	-0.8318630	2.4378021
C	3.5257552	0.4721370	1.9648021
O	3.8027552	1.5461370	2.5028021
N	2.6517552	0.3701370	0.9678021
C	1.9667552	1.5451370	0.4348021
C	1.0867552	2.2161370	1.4878021
O	1.1217552	3.4381370	1.6468021
C	1.1557552	1.1791370	-0.8091979
C	2.0227552	0.9051370	-2.0231979
Se	0.9867552	0.7171370	-3.6481979
C	0.5617552	2.5801370	-3.9891979
N	0.3197552	1.4121370	2.2188021
C	-0.5632448	1.9301370	3.2678021
C	-3.4772448	-1.8308630	-0.1601979
C	-4.3512448	-2.6668630	0.5358021
C	-4.8272448	-3.8318630	-0.0721979
C	-4.4292448	-4.1548630	-1.3711979
C	-3.5542448	-3.3138630	-2.0671979
C	-3.0762448	-2.1498630	-1.4591979
I	-1.7582448	-0.8768630	-2.4781979
H	3.9029897	-0.9927505	3.4987171
H	-0.5402563	3.0326664	3.1935926
H	2.7194545	2.3239439	0.1929050
H	0.4776694	0.3236565	-0.6026147
H	0.4887488	2.0404298	-1.0228342
H	2.5924994	-0.0431694	-1.9325065
H	2.7701876	1.7107972	-2.1783501
H	0.9999209	3.2163070	-3.1956375
H	-0.5345378	2.7272104	-3.9928038
H	0.9819466	2.8871936	-4.9656326
H	-3.1076213	-0.9096816	0.3169060
H	-4.6629007	-2.4027683	1.5580956
H	-5.5175889	-4.4945900	0.4722953
H	-4.8028117	-5.0704689	-1.8551864
H	-3.2460247	-3.5704564	-3.0921680
H	3.9039249	-1.7336014	1.8537916
H	5.2774618	-0.7001381	2.4066119
H	-1.6021143	1.5671126	3.1276481
H	-0.2100731	1.6327934	4.2777114
H	0.4235798	0.4012679	2.1144865
H	2.4908989	-0.5343602	0.5211292

3DNA@LEU84

C	-5.2245734	0.2486797	-0.5622114
C	-3.8505734	-0.4093203	-0.6672114
O	-3.5805734	-1.1363203	-1.6252114
N	-2.9965734	-0.1603203	0.3247886

C	-1.6205734	-0.6603203	0.3087886
C	-1.5035734	-2.1353203	0.6897886
O	-0.6675734	-2.8533203	0.1367886
C	-0.7185734	0.1846797	1.2197886
C	-0.5695734	1.6816797	0.9207886
C	0.4784266	2.2966797	1.8387886
C	-0.2235734	1.9296797	-0.5472114
N	-2.3345734	-2.5763203	1.6317886
C	-2.2595734	-3.9463203	2.1517886
C	4.5644266	-0.6023203	-2.5492114
C	5.4994266	0.3416797	-2.9792114
C	5.7134266	1.4986797	-2.2322114
C	4.9944266	1.7136797	-1.0572114
C	4.0574266	0.7746797	-0.6272114
C	3.8424266	-0.3833203	-1.3752114
I	2.4394266	-1.7933203	-0.7312114
H	-5.9931096	-0.5489286	-0.5783003
H	-1.0527768	0.0481420	2.2749313
H	0.2883604	-0.2843136	1.1665743
H	-1.5444100	2.1848272	1.1305816
H	-0.0566738	3.0083103	-0.7446643
H	0.7059022	1.3877750	-0.8280049
H	-1.0262507	1.5870364	-1.2311159
H	0.2468270	2.1235437	2.9107101
H	0.5623677	3.3916616	1.6834282
H	1.4784194	1.8530591	1.6418021
H	-1.2574095	-0.6284703	-0.7380780
H	-1.3623262	-4.4133294	1.7061111
H	4.3971476	-1.5161773	-3.1384448
H	6.0637804	0.1638336	-3.9074284
H	6.4516571	2.2431259	-2.5684549
H	5.1604494	2.6249787	-0.4620106
H	3.4914281	0.9470214	0.3007830
H	-3.1514741	-4.5411374	1.8625522
H	-2.1719155	-3.9442692	3.2576718
H	-5.3843668	0.8668435	-1.4670700
H	-5.3783463	0.8796114	0.3370006
H	-3.2412128	0.5375650	1.0297191
H	-3.0782525	-1.9542698	1.9534689

4x21

C	0.0811085	-3.3264682	5.8894747
C	-0.9528915	-2.2284682	5.7494747
O	-1.3898915	-1.6334682	6.7354747
N	-1.3128915	-1.9584682	4.5194747
C	-2.2318915	-0.8924682	4.2014747
C	-3.3438915	-1.3844682	3.2794747
O	-3.2478915	-2.4404682	2.6844747
C	-1.4798915	0.2685318	3.5384747
C	-0.2828915	0.7855318	4.2984747
S	0.9791085	1.5055318	3.2874747

C	2.1611085	0.2255318	3.2154747
N	-4.3698915	-0.5694682	3.1264747
C	-5.4278915	-0.9184682	2.2024747
C	-1.3188915	0.5095318	-3.9685253
C	-1.2018915	0.1675318	-6.3755253
N	-1.8708915	0.2305318	-5.1275253
N	-0.0468915	0.7905318	-3.9855253
C	0.5241085	1.0435318	-2.8055253
C	-0.2398915	1.0335318	-1.6395253
I	0.4061085	1.4105318	0.2924747
C	-1.5978915	0.7275318	-1.7565253
N	-2.1298915	0.4605318	-2.9215253
C	1.9351085	1.3585318	-2.8095253
C	2.6681085	1.9285318	-1.8215253
N	3.9661085	2.0555318	-2.1875253
C	4.0971085	1.5795318	-3.4385253
C	2.8621085	1.1175318	-3.8775253
C	2.7411085	0.5675318	-5.1515253
C	3.8651085	0.4805318	-5.9665253
C	5.0951085	0.9505318	-5.4995253
C	5.1931085	1.5045318	-4.2435253
H	0.2272227	-3.9374024	4.9760613
H	-2.1800779	1.0984176	3.3068589
H	-1.1430150	-0.0998405	2.5436110
H	-0.5863591	1.5413453	5.0525482
H	0.1890787	-0.0305870	4.8832135
H	1.7548936	-0.7063154	2.7650500
H	2.5874790	-0.0123745	4.2134341
H	2.9871706	0.5714929	2.5627478
H	-2.6541251	-0.5639771	5.1774724
H	-6.0645225	-1.7412663	2.5936429
H	-0.4708992	-0.6732918	-6.4299496
H	-2.2538532	0.6966625	-0.8664977
H	2.3597550	2.3494943	-0.8666861
H	4.6954598	2.5002559	-1.6344985
H	1.7550044	0.2247044	-5.4874295
H	3.7830675	0.0457376	-6.9743434
H	5.9892539	0.8927089	-6.1389537
H	6.1687308	1.8764824	-3.8866167
H	-1.9417563	0.0289082	-7.1865486
H	-0.6269405	1.0962739	-6.5817042
H	-4.9914804	-1.2682791	1.2467147
H	-6.0648902	-0.0344012	2.0116838
H	-0.2060516	-3.9845180	6.7310957
H	1.0528036	-2.8643807	6.1593108
H	-1.1067324	-2.5994585	3.7478094
H	-4.4576699	0.2678565	3.7023528
H	-2.8607740	-0.0112111	-5.0787009

5HK2

C	-5.5883863	1.1225453	-0.3926241
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C	-4.7113863	0.2645453	0.5133759
O	-5.0433863	-0.8854547	0.7993759
N	-3.5953863	0.8265453	0.9663759
C	-2.7123863	0.1235453	1.8883759
C	-3.4203863	-0.0864547	3.2213759
O	-3.3483863	-1.1654547	3.8123759
C	-1.4113863	0.8995453	2.0973759
C	-0.3233863	0.1065453	2.7903759
C	-0.3433863	-1.2834547	2.7953759
C	0.7206137	0.7465453	3.4473759
C	0.6496137	-2.0134547	3.4263759
C	1.7166137	0.0245453	4.0823759
C	1.6756137	-1.3544547	4.0683759
O	2.6626137	-2.0764547	4.6973759
N	-4.1083863	0.9565453	3.6783759
C	-4.8313863	0.9265453	4.9463759
C	5.5476137	-0.2584547	-7.5746241
C	3.8236137	0.1025453	-5.8936241
C	3.3916137	-0.1114547	-4.5936241
C	2.0976137	0.2645453	-4.2586241
C	1.6216137	0.0635453	-2.9716241
C	2.4446137	-0.5164547	-2.0186241
C	3.7406137	-0.8974547	-2.3506241
C	4.2186137	-0.6964547	-3.6416241
N	5.0236137	-0.4074547	-6.2106241
O	3.1116137	0.7185453	-6.6866241
I	1.7156137	-0.8114547	-0.0856241
H	-6.5584034	1.2855052	0.1183158
H	-1.6193201	1.8355404	2.6606495
H	-1.0314951	1.2143308	1.0951845
H	-1.1689792	-1.8398489	2.3363532
H	0.7564535	1.8480917	3.4670482
H	0.6279832	-3.1123845	3.4373551
H	2.5378423	0.5492670	4.5982954
H	3.2931281	-1.4476017	5.0929284
H	-2.5374033	-0.8882818	1.4783534
H	-4.5569103	-0.0163180	5.4554107
H	4.8642154	0.4245214	-8.1117557
H	1.4742528	0.7218703	-5.0389664
H	0.5947294	0.3623216	-2.7132412
H	4.3933301	-1.3490817	-1.5885001
H	5.2647670	-0.9669911	-3.8372187
H	-5.1606641	2.1082057	-0.6677436
H	-5.8040165	0.5499255	-1.3149974
H	-4.5507752	1.7888065	5.5857823
H	-5.9310092	0.9368388	4.7902431
H	6.5710861	0.1705512	-7.5640367
H	5.5746149	-1.2310764	-8.1107782
H	-3.3029193	1.7367940	0.6055792
H	-4.2322219	1.7605381	3.0609993
H	5.5000399	-1.0533808	-5.5873484

3PP1

C	-2.2994255	-4.6424371	-2.3078033
C	-1.8304255	-4.6964371	-0.8718033
O	-2.4734255	-5.3364371	-0.0348033
N	-0.7064255	-4.0364371	-0.5918033
C	-0.1844255	-3.9164371	0.7721967
C	-1.2204255	-3.2664371	1.6921967
O	-1.8654255	-2.2724371	1.3371967
N	-1.3804255	-3.8594371	2.8681967
C	-2.2964255	-3.3534371	3.8731967
C	2.4665745	6.7955629	-2.1048033
N	2.7435745	5.8415629	-1.1388033
C	2.0235745	4.6815629	-0.8258033
C	0.8865745	4.2695629	-1.5238033
C	0.2145745	3.1015629	-1.1608033
C	0.6445745	2.3205629	-0.0808033
I	-0.4014255	0.5535629	0.4741967
C	1.7825745	2.7445629	0.6151967
C	2.4735745	3.9085629	0.2511967
F	3.5735745	4.2915629	0.9481967
H	-1.5680117	-4.2187625	-3.0254322
H	0.1254186	-4.9162077	1.1456833
H	-1.8351405	-3.3929561	4.8806649
H	-3.2458109	-3.9309104	3.8894643
H	0.5209700	4.8691915	-2.3692009
H	-0.6766044	2.8023875	-1.7333398
H	2.1718261	2.1765762	1.4739771
H	-2.5379399	-2.3059660	3.6143834
H	-2.5661051	-5.6687278	-2.6237856
H	-3.2275792	-4.0375019	-2.3509973
H	-0.3210591	-3.4020918	-1.2911868
H	-0.9556625	-4.7756794	3.0167521
H	0.7109617	-3.2648634	0.7410726
H	3.5208467	6.0181558	-0.5054149
H	2.4560089	6.3838065	-3.1450675
H	1.4799488	7.3087160	-1.9671342
H	3.2409590	7.5865750	-2.0886292