

Hydrogen bond co-ordination in organic crystal structures: statistics, prediction and application

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Table S1. Fitting statistics for logistic regression models to predict donor acceptor co-ordination derived in this work. The values represent how much better the optimised model equation (residual) fits the sample data, compared with the independent model (null). "D. O. F." is independent degrees of freedom per sample, "P value" is the probability that one could obtain a given result randomly assuming the null hypothesis is correct. The acceptance criterion was $P < 0.05$. "AIC" is Akaike's Information Criterion and is another estimate of the residual deviance associated with lost information as a result of applying the model as a predictor of the complete data. "Area under ROC (receiver operator characteristic) curve" is a measure of classification of predictions. It is desirable to achieve a value as close to unity as possible, whereas 0.5 indicates a random predictor prestigious.

Model name	Residual deviance	D.O.F	Null deviance	D.O.F	log likelihood	P value (log likelihood test)	AIC	Area Under Roc curve
Br_Br_a_1	66.6885	574	66.6885	574	-33.3442	0	68.6885	1
Br_ion_Br_a_1	22.2611	119	28.1081	120	-11.1306	0.00886732	26.2611	0.861582
Br_ion_Br_a_2	47.2159	119	160.723	120	-23.6079	8.43E-27	51.2159	0.971304
Br_ion_Br_a_3	38.639	119	117.695	120	-19.3195	3.06E-19	42.639	0.976043
Br_ion_Br_a_4	26.5183	119	58.9206	120	-13.2592	6.45E-09	30.5183	0.971239
Cl_Cl_a_1	282.326	1751	295.455	1753	-141.163	0.000704751	288.326	0.696132
Cl_ion_Cl_a_1	2.51E-09	432	0	432	-1.26E-09	0	2	1
Cl_ion_Cl_a_2	153.58	431	470.395	432	-76.7902	3.59E-71	157.58	0.968985
Cl_ion_Cl_a_3	273.751	431	585.771	432	-136.876	3.98E-70	277.751	0.943216
Cl_ion_Cl_a_4	198.314	431	392.857	432	-99.1568	1.63E-44	202.314	0.942884
Cl_ion_Cl_a_5	60.2204	431	109.724	432	-30.1102	1.01E-12	64.2204	0.939628
I_a_1	8.08019	146	12.0012	148	-4.0401	0.070395	14.0802	0.966216
I_ion_I_a_1	41.1898	57	50.3971	58	-20.5949	0.00131675	45.1898	0.782222
I_ion_I_a_2	34.9963	57	46.8321	58	-17.4982	0.000312031	38.9963	0.860294
I_ion_I_a_3	3.42E-10	58	0	58	-1.71E-10	0	2	1
T1NH0_a_1	31.2866	48	50.9134	51	-15.6433	9.67E-05	39.2866	0.902381
T1NH0_a_2	15.5893	48	28.2037	51	-7.79465	0.00258322	23.5893	0.921875
T1OH0_a_1	791.218	733	882.139	738	-395.609	2.08E-18	803.218	0.725646
T1OH0_a_2	447.532	732	876.536	738	-223.766	8.01E-90	461.532	0.931777
T2NH0_N.2_a_1	1262.81	1507	1620.21	1513	-631.405	1.96E-74	1276.81	0.822924
T2NH0_N.2_a_2	77.3623	1510	89.24	1513	-38.6811	0.00362295	85.3623	0.804342
T2OH_O.3_a_1	335.013	340	376.044	344	-167.507	1.26E-08	345.013	0.725963
T2OH_O.3_a_2	56.1167	342	83.3969	344	-28.0584	5.96E-07	62.1167	0.89881
T2OH_O.3_d_1	110.961	340	230.827	344	-55.4803	2.80E-25	120.961	0.860752
T2OH_O.3_d_2	98.2693	341	123.403	344	-49.1346	6.97E-06	106.269	0.836768
T2OH0_a_1	337.047	616	429.306	621	-168.524	1.09E-18	349.047	0.814743
T3CS_S.2_a_1	600.738	535	743.963	539	-300.369	2.84E-30	610.738	0.804428
T3CS_S.2_a_2	252.064	534	380.879	539	-126.032	2.07E-26	264.064	0.903727
T3NH1_d_1	819.604	1351	1310.95	1355	-409.802	2.49E-105	829.604	0.87555
T3NH1_d_2	653.236	1349	806.332	1355	-326.618	8.35E-31	667.236	0.815805
T3NH2_a_1	91.8939	249	228.17	254	-45.947	5.41E-28	103.894	0.785125
T3NH2_d_1	56.8518	254	56.8518	254	-28.4259	0	58.8518	1
T3NH2_d_2	209.98	250	260.653	254	-104.99	1.26E-10	219.98	0.798804
T3NH2_d_3	190.001	250	211.197	254	-95.0005	0.000132318	200.001	0.6758
T4NH2_d_1	14.5676	257	32.7839	261	-7.28378	0.000504397	24.5676	0.981982
T4NH2_d_2	111.333	259	169.463	261	-55.6666	1.19E-13	117.333	0.885756
T4NH2_d_3	228.221	258	286.721	261	-114.111	6.05E-13	236.221	0.796935
T4NH2_d_4	36.6635	259	49.493	261	-18.3318	0.000818619	42.6635	0.897276
T4PO_O.2_a_1	70.1498	157	124.402	159	-35.0749	8.28E-13	76.1498	0.90579
T4PO_O.2_a_2	60.0967	157	160.129	159	-30.0483	9.49E-23	66.0967	0.96582
acyclic_n_N.2_a_1	106.294	398	332.706	404	-53.1472	2.19E-46	120.294	0.978734
acyclic_n_N.2_a_2	2.75E-06	403	14.0053	404	-1.37E-06	9.70E-05	4	1
al_oh_O.3_a_1	1948.74	2072	2604.59	2078	-974.372	1.03E-138	1962.74	0.828376

al_oh_O.3_a_2	228.845	2074	338.92	2078	-114.423	3.44E-23	238.845	0.907758
al_oh_O.3_d_1	938.002	2073	1643.19	2078	-469.001	1.85E-150	950.002	0.870978
al_oh_O.3_d_2	546.692	2075	571.546	2078	-273.346	7.97E-06	554.692	0.63995
ar_n_N.ar_a_1	1452.03	1415	1967.86	1421	-726.016	1.62E-108	1466.03	0.816017
ar_n_N.ar_a_2	286.442	1415	371.599	1421	-143.221	1.46E-16	300.442	0.876901
ar_oh_O.3_a_1	1103.37	1459	1468.5	1463	-551.686	4.72E-78	1113.37	0.853326
ar_oh_O.3_a_2	131.87	1458	211.453	1463	-65.935	4.94E-16	143.87	0.939889
ar_oh_O.3_d_1	588.882	1460	2026.38	1463	-294.441	1.0767E-311	596.882	0.970421
ar_oh_O.3_d_2	287.546	1460	330.507	1463	-143.773	1.23E-09	295.546	0.806458
carbamoyl_N.am_d_1	39.1272	287	58.4125	289	-19.5636	3.25E-05	45.1272	0.893192
carbamoyl_N.am_d_2	198.048	285	290.233	289	-99.0241	2.21E-19	208.048	0.856124
carbamoyl_N.am_d_3	56.5818	285	73.2243	289	-28.2909	0.001012225	66.5818	0.845301
carbamoyl_O.2_a_1	79.8799	287	99.9358	289	-39.94	2.21E-05	85.8799	0.858213
carbamoyl_O.2_a_2	239.478	283	401.529	289	-119.739	1.06E-32	253.478	0.888418
carbamoyl_O.2_a_3	32.6189	284	58.4125	289	-16.3094	4.37E-05	44.6189	0.97007
carbonyl_O.2_a_1	3883.08	3606	4984.24	3612	-1941.54	5.82E-235	3897.08	0.815624
carbonyl_O.2_a_2	701.414	3606	1086.62	3612	-350.707	2.09E-80	715.414	0.900188
carboxylate_O.co2_a_1	416.463	1334	685.944	1339	-208.232	1.79E-56	428.463	0.915544
carboxylate_O.co2_a_2	1480.11	1336	1853.76	1339	-740.053	5.60E-81	1488.11	0.789845
carboxylate_O.co2_a_3	411.642	1335	537.894	1339	-205.821	1.21E-26	421.642	0.846201
cooh_O.2_a_1	773.26	788	1096.54	792	-386.63	5.10E-69	783.26	0.852655
cooh_O.2_a_2	308.047	787	404.802	792	-154.024	1.24E-19	320.047	0.872601
cooh_O.3_a_1	285.063	791	373.549	795	-142.531	1.35E-18	295.063	0.833941
cooh_O.3_d_1	64.5855	793	492.476	795	-32.2927	6.08E-94	70.5855	0.979262
cooh_O.3_d_2	159.794	792	179.479	795	-79.8969	9.41E-05	167.794	0.763192
cyclic_amide_N.am_d_1	279.069	1374	455.791	1378	-139.534	1.86E-37	289.069	0.886709
cyclic_amide_N.am_d_2	376.433	1373	436.414	1378	-188.216	5.84E-12	388.433	0.772502
cyclic_amide_O.2_a_1	1363.72	1488	1662.45	1491	-681.862	9.35E-65	1371.72	0.788887
cyclic_amide_O.2_a_2	783.738	1487	1212.47	1491	-391.869	8.55E-92	793.738	0.898681
cyclic_n_N.2_a_1	631.376	734	1015.54	740	-315.688	3.50E-80	645.376	0.874271
cyclic_n_N.2_a_2	128.043	738	161.95	740	-64.0217	2.17E-08	134.043	0.85676
cyclic_nn_N.2_a_1	183.392	176	253.03	182	-91.6958	2.29E-13	197.392	0.82654
cyclic_nn_N.2_a_2	1.06E-09	182	0	182	-5.31E-10	0	2	1
ether_O.3_a_1	1924.69	3705	2108.62	3711	-962.347	2.43E-37	1938.69	0.749569
nh3_N.4_d_1	1.24E-07	583	14.7449	585	-6.22E-08	0.000314158	6	1
nh3_N.4_d_2	25.1114	581	47.8688	585	-12.5557	6.51E-05	35.1114	0.947595
nh3_N.4_d_3	99.4905	582	117.074	585	-49.7452	0.000254184	107.49	0.836963
nh3_N.4_d_4	754.143	581	793.088	585	-377.072	3.40E-08	764.143	0.636067
nh3_N.4_d_5	219.002	579	265.125	585	-109.501	1.28E-08	233.002	0.807623
nitrile_N.1_a_1	624.196	495	676.138	497	-312.098	2.63E-12	630.196	0.752096
nitrile_N.1_a_2	66.8244	494	81.9693	497	-33.4122	0.000798674	74.8244	0.848214
nitro_O.2_a_1	1159.14	1329	1539.61	1333	-579.569	2.29E-81	1169.14	0.821934
nitro_O.2_a_2	248.734	1328	495.545	1333	-124.367	1.31E-51	260.734	0.95212
oxime_N.2_a_1	272.297	274	389.46	280	-136.148	3.10E-23	286.297	0.866373
oxime_O.3_a_1	124.569	276	154.313	280	-62.2845	2.59E-06	134.569	0.753422
oxime_O.3_d_1	1.63E-09	279	23.7666	280	-8.15E-10	5.65E-07	4	1
prim_amine_a_1	126.991	1159	400.081	1163	-63.4954	3.42E-58	136.991	0.985196
prim_amine_d_1	522.085	1160	618.635	1163	-261.043	4.24E-21	530.085	0.809198
prim_amine_d_2	1170.32	1158	1567.9	1163	-585.159	4.89E-84	1182.32	0.823883
prim_amine_d_3	533.372	1159	577.56	1163	-266.686	2.81E-09	543.372	0.722639
sec_amine_a_1	288.975	2114	687.75	2119	-144.487	2.70E-84	300.975	0.966093
sec_amine_d_1	2016.27	2114	2787.08	2119	-1008.14	1.19E-164	2028.27	0.836787
sec_amine_d_2	530.589	2115	635.289	2119	-265.295	4.82E-22	540.589	0.816189
sulfinyl_O.2_a_1	1.03E-07	31	32.7996	36	-5.16E-08	1.88E-06	12	1
sulfinyl_O.2_a_2	27.865	35	32.7996	36	-13.9325	0.0152315	31.865	0.72043
sulfonyl_O.2_a_1	1659.29	1552	2148.31	1557	-829.647	9.33E-104	1671.29	0.804994
sulfonyl_O.2_a_2	542.256	1553	923.04	1557	-271.128	1.96E-81	552.256	0.915467
tert_amine_N.3_a_1	507.52	576	750.648	580	-253.76	9.75E-52	517.52	0.863894
tert_amine_N.3_a_2	25.1526	578	37.5813	580	-12.5763	0.00100027	31.1526	0.934833
tert_ammonia_N.4_d_1	5.87981	228	171.402	233	-2.9399	3.23E-34	17.8798	0.999653
tert_ammonia_N.4_d_2	107.518	231	154.759	233	-53.759	2.76E-11	113.518	0.864087
thioether_S.3_a_1	144.323	898	144.323	898	-72.1616	0	146.323	1
transamide_N.am_d_1	1328.65	2237	2246.36	2243	-664.327	2.78E-195	1342.65	0.886814
transamide_N.am_d_2	306.515	2238	335.559	2243	-153.258	1.03E-05	318.515	0.749887
transamide_O.2_a_1	2001.01	2158	2800.02	2164	-1000.51	1.25E-169	2015.01	0.85048
transamide_O.2_a_2	901.113	2159	1378.83	2164	-450.557	2.56E-101	913.113	0.88946
water_O.3_a_1	524.097	501	555.031	502	-262.049	1.38E-08	528.097	0.68727
water_O.3_a_2	426.141	500	466.431	502	-213.071	8.91E-10	432.141	0.701862
water_O.3_d_1	60.3828	501	65.0741	502	-30.1914	0.0176423	64.3828	0.594567
water_O.3_d_2	220.591	501	232.818	502	-110.296	0.000252439	224.591	0.681862
water_O.3_d_3	299.315	501	303.102	502	-149.657	0.0308631	303.315	0.59049