

Contribution of C-H Hydrogen Bonds on anion binding

Yusun Choi, Taehoon Kim, Soonmin Jang*, Jongmin Kang*

Department of Chemistry, Sejong University, Seoul, 143-747, South Korea

kangjm@sejong.ac.kr

1. Fig S1. ^1H NMR spectrum of compound 1 -----	3
2. Fig S2. ^{13}C NMR spectrum of compound 1 -----	4
3. Fig S3. HRMS(FAB) spectrum of compound 1 -----	5
4. Fig S4. ^1H NMR spectrum of compound 2 -----	6
5. Fig S5. ^{13}C NMR spectrum of compound 2 -----	7
6. Fig S6. HRMS(FAB) spectrum of compound 2 -----	8
7. Fig S7. ^1H NMR spectrum of compound 3 -----	9
8. Fig S8. HRMS(FAB) spectrum of compound 3 -----	10
9. Fig S9. ^1H NMR spectrum of compound 4 -----	11
10. Fig S10. ^1H NMR spectrum of compound 5 -----	12
11. Fig S11. ^{13}C NMR spectrum of compound 5 -----	13
12. Fig S12. HRMS(EI) spectrum of compound 5 -----	14
13. Fig S13. A family of UV-vis spectra recorded over the course of titrating a 100 μM DMSO solution of receptor 1 with increased amounts of tetrabutylammonium bromide-----	15
14. Fig S14. A family of UV-vis spectra recorded over the course of titrating a 100 μM DMSO solution of receptor 1 with increased amounts of tetrabutylammonium hydrogen sulfate -----	16
14. Fig S14. ^1H NMR spectra of 2 mM of receptor 1 containing increasing amounts of tetrabutylammonium chloride (0 –44 equiv.) in DMSO- d_6 -----	17
15. Fig S15. ^1H NMR spectra of 2 mM of receptor 1 containing increasing amounts of tetrabutylammonium bromide (0 –86 equiv.) in DMSO- d_6 -----	18
16. Fig S16. ^1H NMR spectra of 2 mM of receptor 1 containing increasing amounts of	

tetrabutylammonium hydrogen sulfate (0–115 equiv.) in DMSO-d₆ -----19

17. Table 1. PDB file of receptor 1 in complex with H₂PO₄⁻ anion. -----20

18. Table 2. PDB file of receptor 1 in complex with Cl⁻ anion.-----24

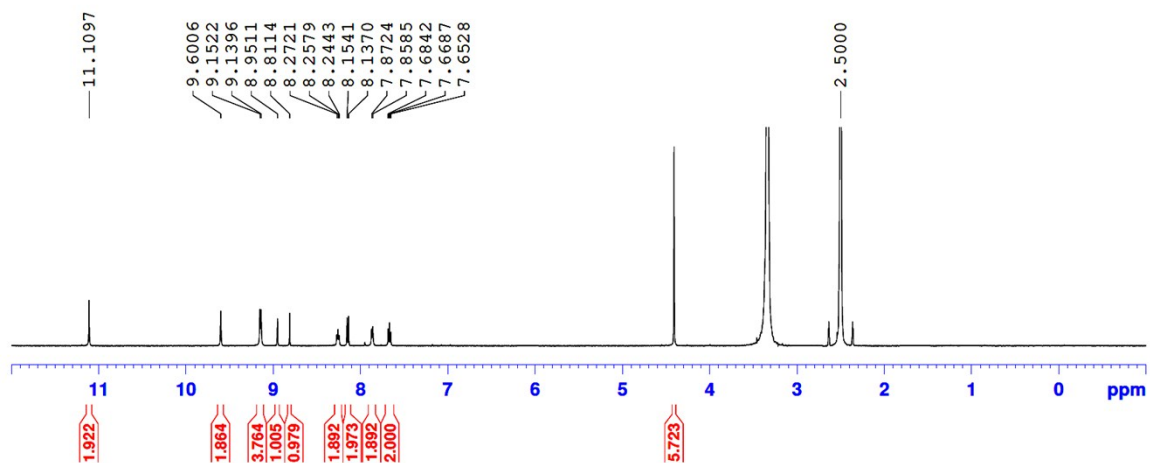


Fig S1. ¹H NMR spectrum of compound **1**

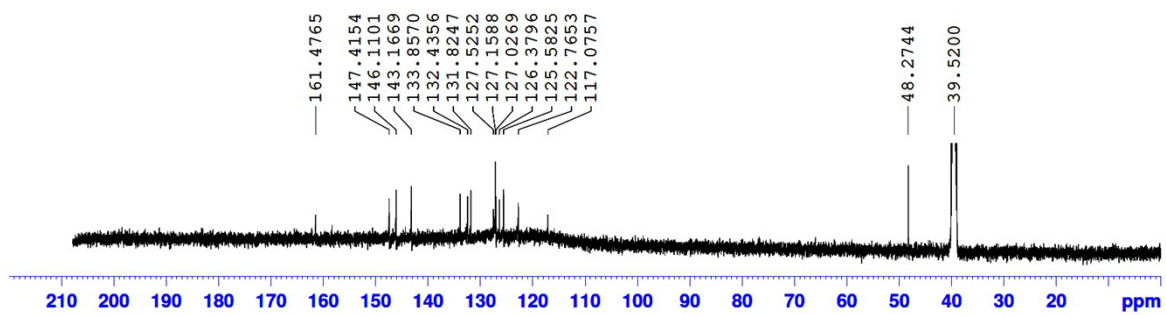


Fig S2. ¹³C NMR spectrum of compound 1

141023_COMPOUND1_H_001-c1 #103-142 RT: 1.55-2.14 Av: 40 NL: 8.61E4
T: + c FAB Full ms [669.50-820.50]

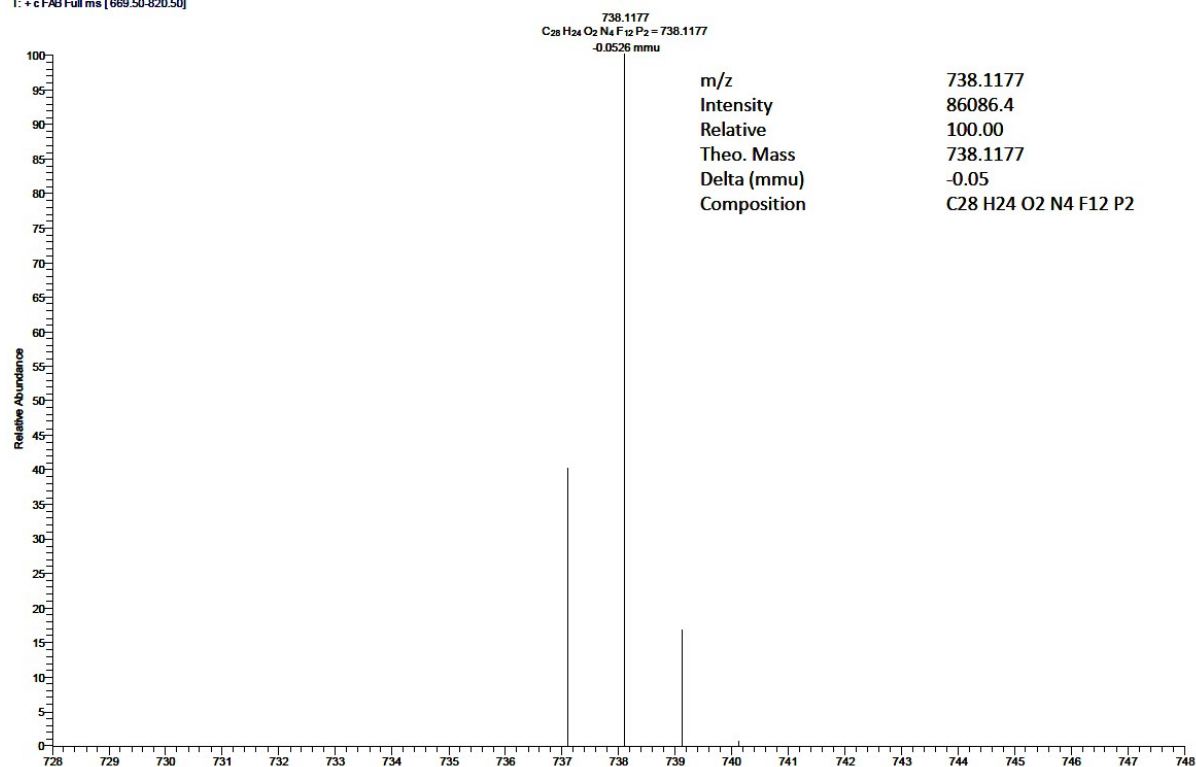


Fig S3. HRMS(FAB) spectrum of compound 1

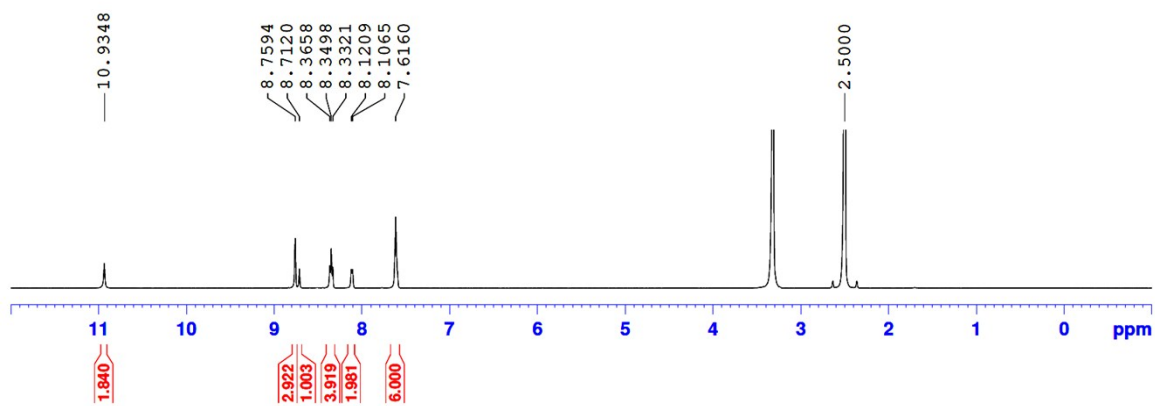


Fig S4. ¹H NMR spectrum of compound 2

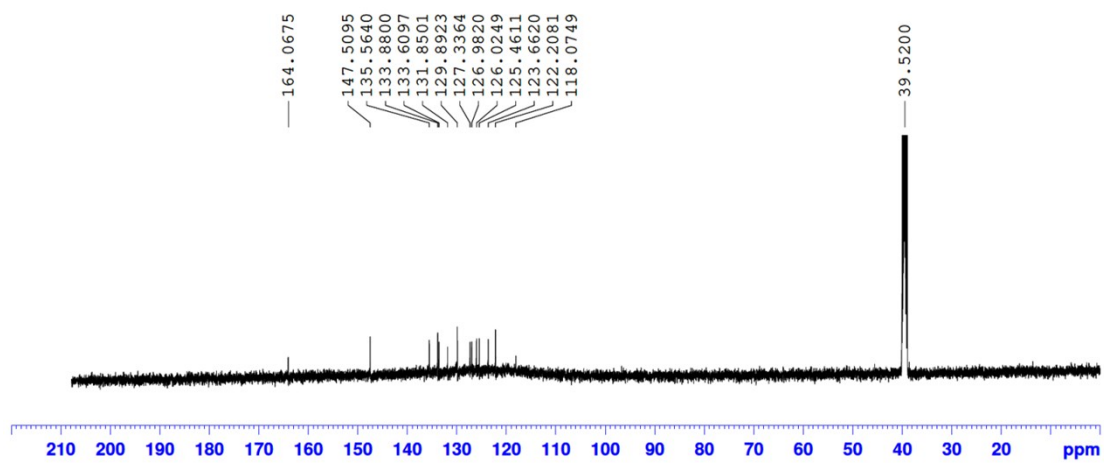


Fig S5. ^{13}C NMR spectrum of compound 2

141023_COMPOUND2_H_002-c1 #71-127 RT: 1.26-2.27 Av: 57 NL: 1.42E4
T: + c FAB Full ms [449.50-600.50]

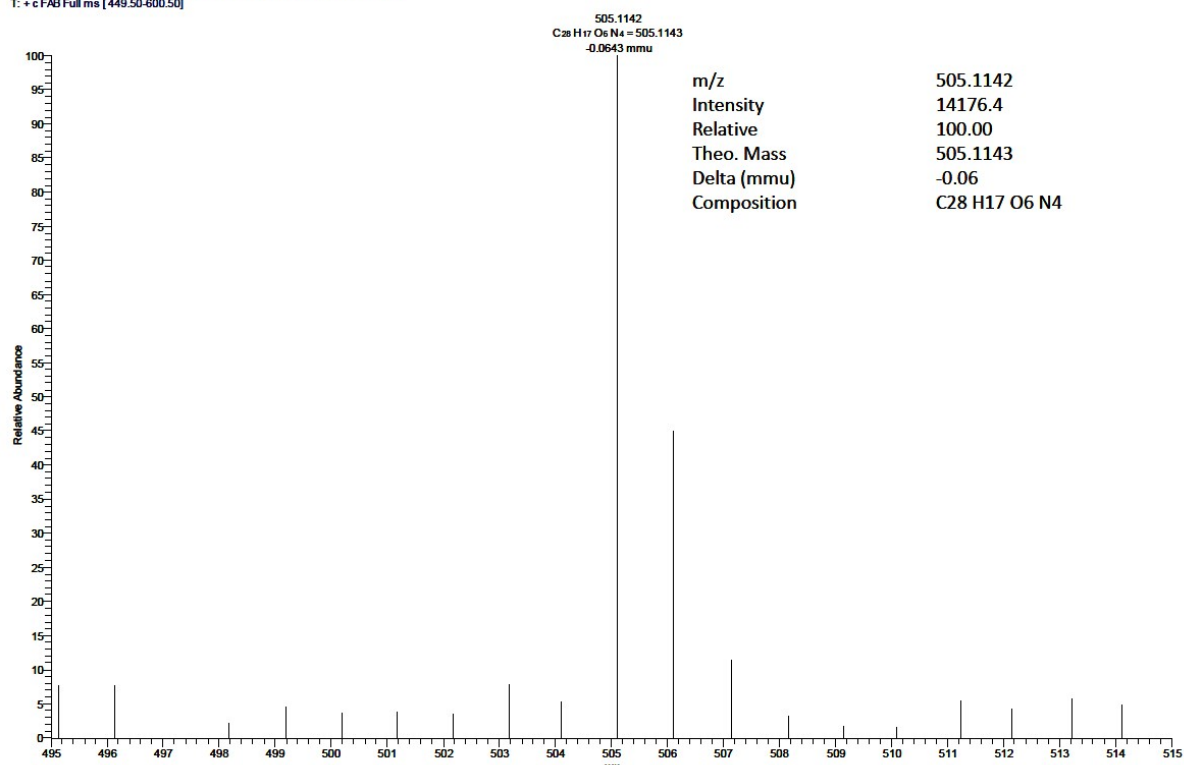


Fig S6. HRMS(FAB) spectrum of compound 2

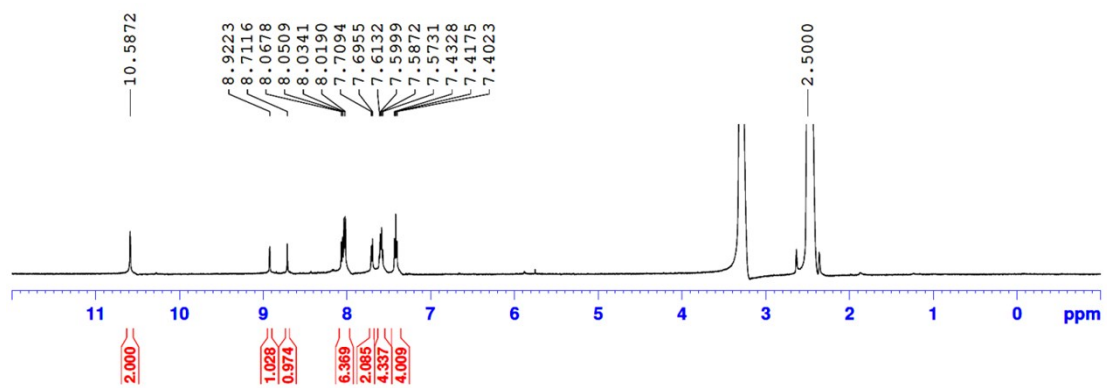


Fig S7 ¹H NMR spectrum of compound 3

141023_COMPOUND3_H_002-e1 #32-56 RT: 0.56-0.99 AV: 25 NL: 1.07E5
T: + cFAB Full ms [314.50-470.50]

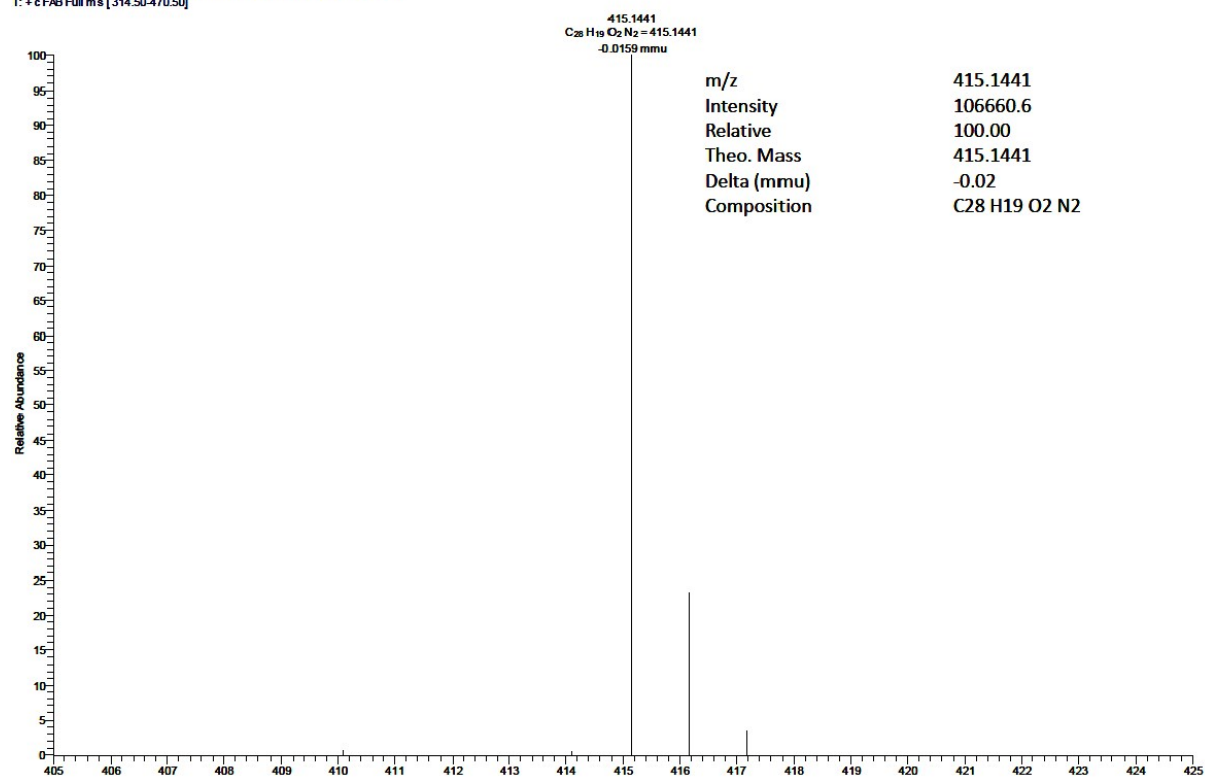


Fig S8. HRMS(FAB) spectrum of compound 3

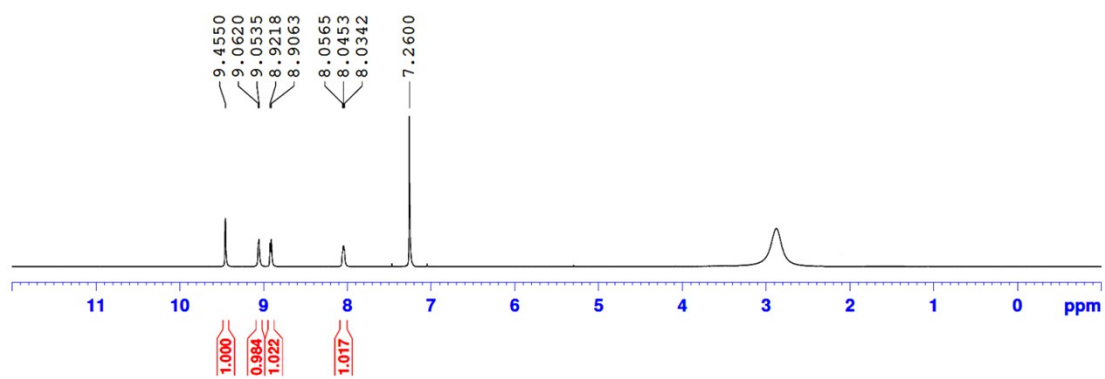


Fig S9 ^1H NMR spectrum of compound 4

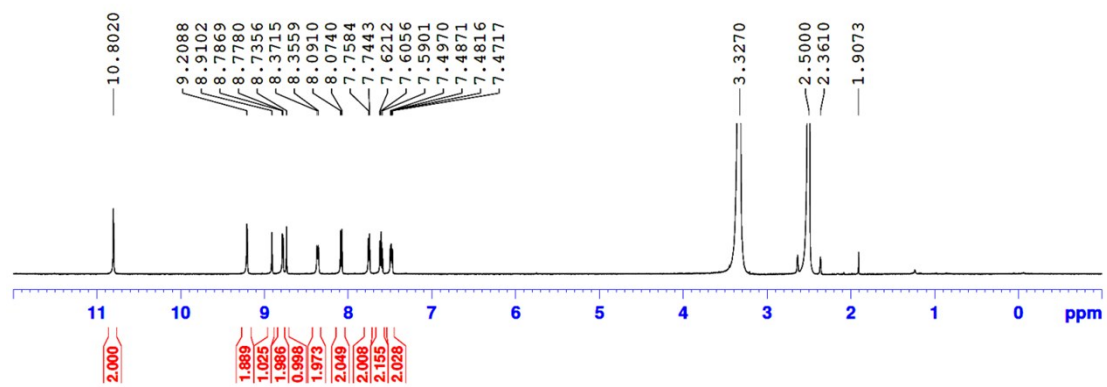
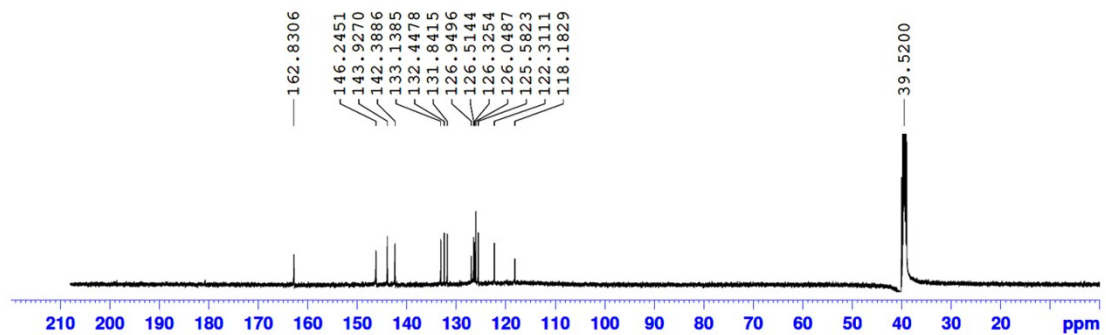


Fig S10. ¹H NMR spectrum of compound 5



11. Fig S11. ^{13}C NMR spectrum of compound 5

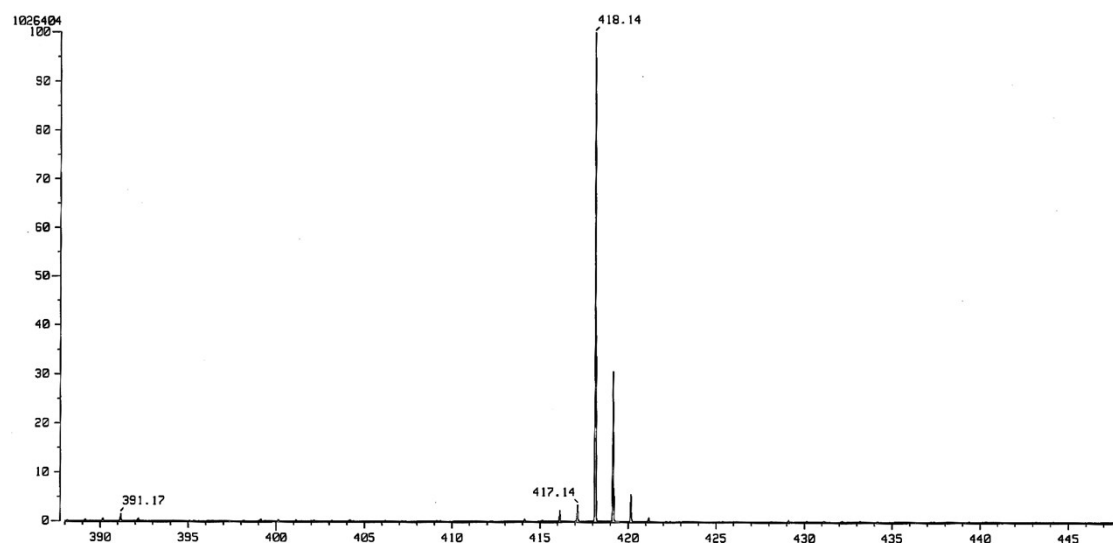


Fig S12. HRMS(EI) spectrum of compound 5

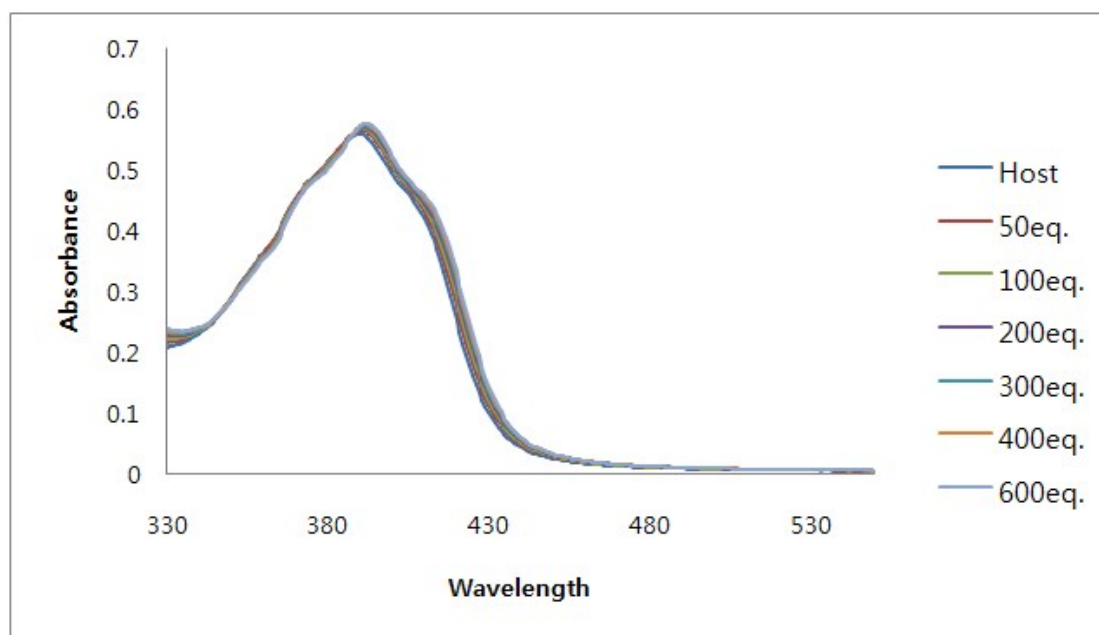


Fig S13. A family of UV-vis spectra recorded over the course of titrating a 100 μM DMSO solution of receptor **1** with increased amounts of tetrabutylammonium bromide

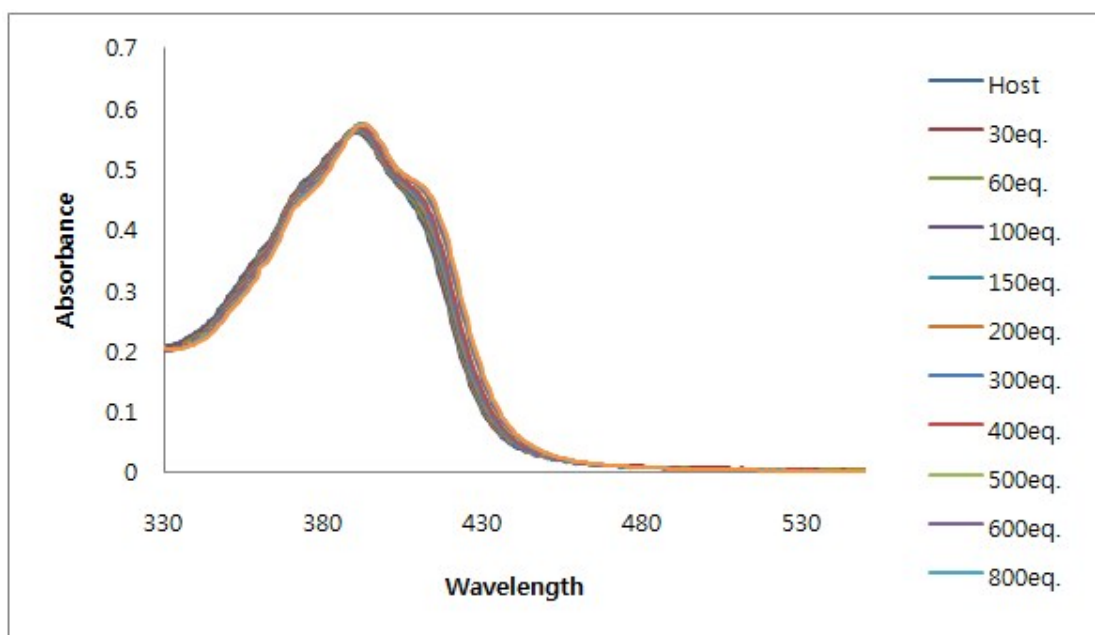


Fig S14. A family of UV-vis spectra recorded over the course of titrating a 100 μM DMSO solution of receptor **1** with increased amounts of tetrabutylammonium hydrogen sulfate

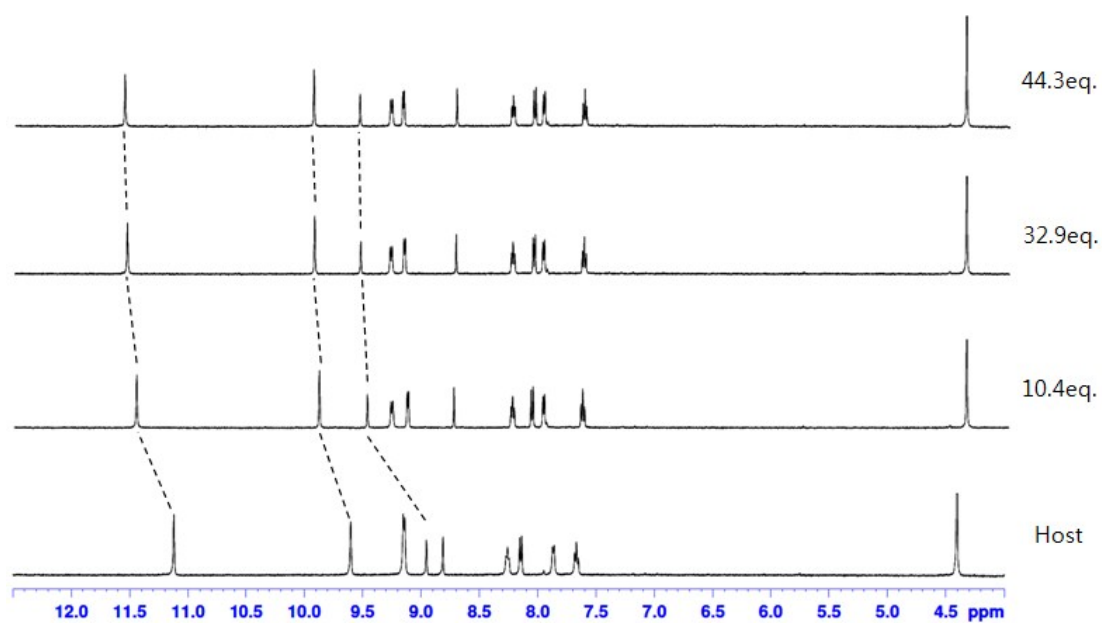


Fig S15. ¹H NMR spectra of 2 mM of receptor **1** containing increasing amounts of tetrabutylammonium chloride (0 –44 equiv.) in DMSO-d₆

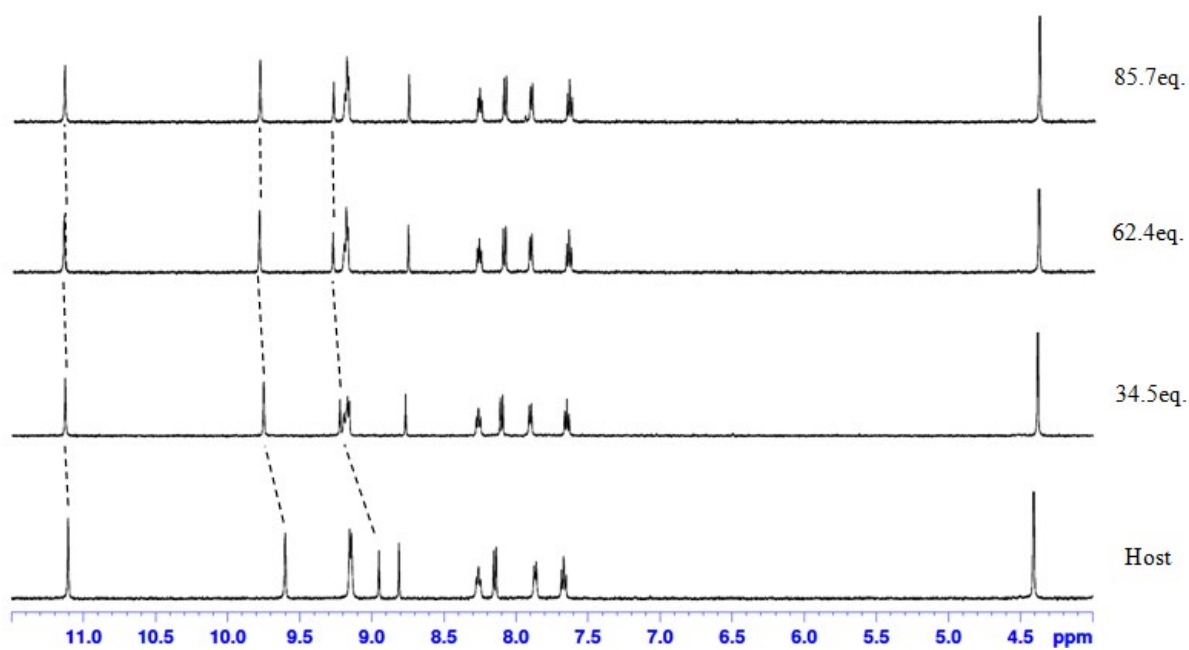


Fig S16. ¹H NMR spectra of 2 mM of receptor **1** containing increasing amounts of tetrabutylammonium bromide (0 –86 equiv.) in DMSO-d₆

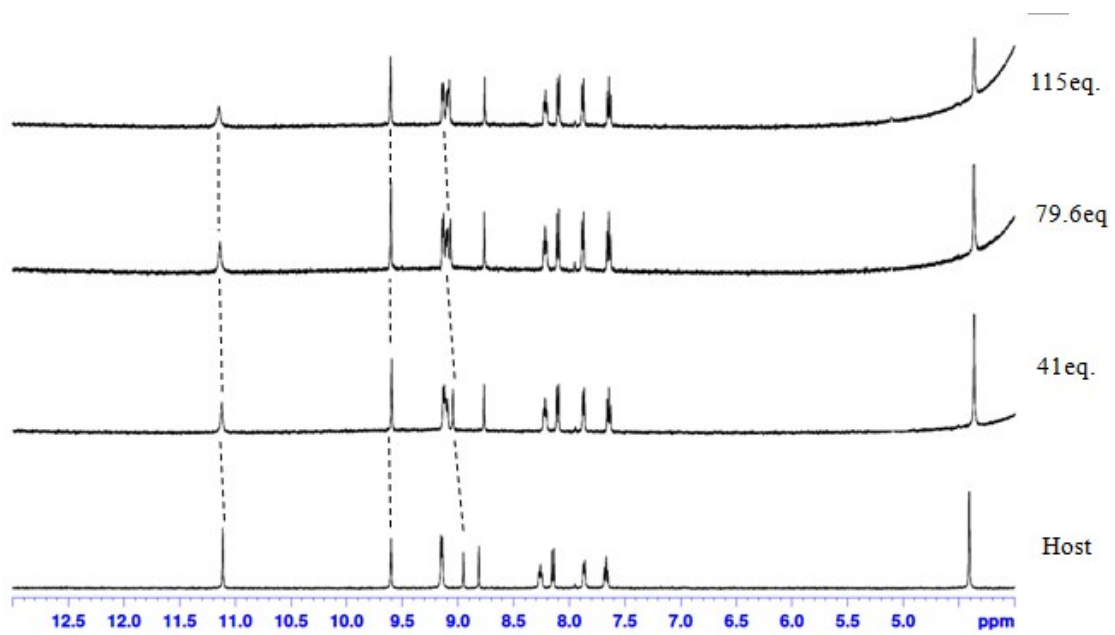


Fig S17. ¹H NMR spectra of 2 mM of receptor **1** containing increasing amounts of tetrabutylammonium hydrogen sulfate (0–115 equiv.) in DMSO-d₆

Table 1. PDB file of receptor 1 in complex with H₂PO₄⁻ anion.

TITLE	COMPLEX1 WITH H2PO4-					
HETATM	1	C	0	-3.936	4.000	-0.341
HETATM	2	C	0	-4.720	2.893	-0.182
HETATM	3	C	0	-4.134	1.592	-0.128
HETATM	4	C	0	-2.701	1.453	-0.240
HETATM	5	C	0	-1.914	2.652	-0.371
HETATM	6	C	0	-2.524	3.880	-0.428
HETATM	7	C	0	-4.918	0.443	0.004
HETATM	8	C	0	-2.141	0.169	-0.242
HETATM	9	C	0	-2.931	-0.981	-0.119
HETATM	10	C	0	-4.362	-0.839	0.010
HETATM	11	C	0	-5.178	-2.007	0.108
HETATM	12	H	0	-6.251	-1.885	0.203
HETATM	13	C	0	-4.615	-3.251	0.073
HETATM	14	C	0	-3.208	-3.404	-0.047
HETATM	15	C	0	-2.385	-2.312	-0.138
HETATM	16	H	0	-5.995	0.549	0.086
HETATM	17	H	0	-4.382	4.986	-0.387
HETATM	18	H	0	-5.798	2.982	-0.101
HETATM	19	H	0	-1.919	4.770	-0.532
HETATM	20	H	0	-1.072	0.061	-0.356
HETATM	21	H	0	-5.235	-4.137	0.146
HETATM	22	H	0	-2.779	-4.397	-0.064
HETATM	23	N	0	-0.496	2.541	-0.402
HETATM	24	C	0	0.285	3.178	-1.313
HETATM	25	H	0	-0.080	1.975	0.351
HETATM	26	O	0	-0.138	3.830	-2.260
HETATM	27	C	0	1.786	3.059	-1.135
HETATM	28	C	0	2.593	3.419	-2.217
HETATM	29	C	0	2.401	2.674	0.047
HETATM	30	H	0	2.119	3.739	-3.135
HETATM	31	H	0	1.853	2.365	0.932
HETATM	32	C	0	4.536	2.975	-0.901
HETATM	33	H	0	5.603	2.922	-0.738
HETATM	34	C	0	3.977	3.367	-2.100

HETATM	35	N	0	-0.972	-2.476	-0.236
HETATM	36	C	0	-0.389	-3.250	-1.187
HETATM	37	H	0	-0.417	-1.999	0.488
HETATM	38	O	0	-0.998	-3.832	-2.076
HETATM	39	C	0	1.122	-3.383	-1.145
HETATM	40	C	0	1.739	-4.004	-2.233
HETATM	41	C	0	1.918	-2.953	-0.092
HETATM	42	H	0	1.120	-4.350	-3.050
HETATM	43	H	0	1.526	-2.458	0.793
HETATM	44	C	0	3.863	-3.719	-1.181
HETATM	45	H	0	4.938	-3.817	-1.125
HETATM	46	C	0	3.119	-4.168	-2.251
HETATM	47	H	0	3.620	-4.640	-3.085
HETATM	48	H	0	4.624	3.630	-2.925
HETATM	49	C	0	4.368	2.267	1.440
HETATM	50	H	0	3.706	1.582	1.965
HETATM	51	H	0	5.322	1.786	1.243
HETATM	52	H	0	4.520	3.173	2.027
HETATM	53	C	0	4.074	-2.687	1.028
HETATM	54	H	0	5.023	-2.302	0.662
HETATM	55	H	0	3.537	-1.906	1.557
HETATM	56	H	0	4.243	-3.540	1.684
HETATM	57	N	0	3.256	-3.125	-0.130
HETATM	58	N	0	3.747	2.640	0.143
HETATM	59	P	0	0.725	-0.037	2.603
HETATM	60	O	0	0.504	1.251	1.856
HETATM	61	O	0	0.389	-1.358	1.949
HETATM	62	O	0	2.302	0.005	3.044
HETATM	63	H	0	2.539	-0.685	3.676
HETATM	64	O	0	-0.013	0.027	4.059
HETATM	65	H	0	-0.686	-0.660	4.135

END

CONNECT	1	2	6	17
CONNECT	2	1	3	18
CONNECT	3	2	4	7
CONNECT	4	3	5	8
CONNECT	5	4	6	23
CONNECT	6	1	5	19

CONECT	7	3	10	16
CONECT	8	4	9	20
CONECT	9	8	10	15
CONECT	10	7	9	11
CONECT	11	10	12	13
CONECT	12	11		
CONECT	13	11	14	21
CONECT	14	13	15	22
CONECT	15	14	9	35
CONECT	16	7		
CONECT	17	1		
CONECT	18	2		
CONECT	19	6		
CONECT	20	8		
CONECT	21	13		
CONECT	22	14		
CONECT	23	5	24	25
CONECT	24	23	26	27
CONECT	25	23		
CONECT	26	24		
CONECT	27	24	28	29
CONECT	28	27	30	34
CONECT	29	27	31	58
CONECT	30	28		
CONECT	31	29		
CONECT	32	33	34	58
CONECT	33	32		
CONECT	34	28	32	48
CONECT	35	15	36	37
CONECT	36	35	38	39
CONECT	37	35		
CONECT	38	36		
CONECT	39	36	40	41
CONECT	40	39	42	46
CONECT	41	39	43	57
CONECT	42	40		
CONECT	43	41		
CONECT	44	45	46	57

CONECT	45	44			
CONECT	46	40	44	47	
CONECT	47	46			
CONECT	48	34			
CONECT	49	50	51	52	58
CONECT	50	49			
CONECT	51	49			
CONECT	52	49			
CONECT	53	54	55	56	57
CONECT	54	53			
CONECT	55	53			
CONECT	56	53			
CONECT	57	41	44	53	
CONECT	58	29	32	49	
CONECT	59	60	61	62	64
CONECT	60	59			
CONECT	61	59			
CONECT	62	59	63		
CONECT	63	62			
CONECT	64	59	65		
CONECT	65	64			

Table 2. PDB file of receptor 1 in complex with Cl⁻ anion.

TITLE	COMPLEX1 WITH CL-					
HETATM	1	C	0	-0.183	4.204	3.648
HETATM	2	C	0	-0.256	4.889	2.468
HETATM	3	C	0	-0.205	4.191	1.224
HETATM	4	C	0	-0.062	2.755	1.224
HETATM	5	C	0	-0.026	2.074	2.493
HETATM	6	C	0	-0.078	2.788	3.664
HETATM	7	C	0	-0.266	4.864	0.000
HETATM	8	C	0	0.035	2.084	0.000
HETATM	9	C	0	-0.062	2.755	-1.224
HETATM	10	C	0	-0.205	4.191	-1.224
HETATM	11	C	0	-0.256	4.889	-2.468
HETATM	12	H	0	-0.352	5.969	-2.457
HETATM	13	C	0	-0.183	4.204	-3.648
HETATM	14	C	0	-0.078	2.788	-3.664
HETATM	15	C	0	-0.026	2.074	-2.493
HETATM	16	H	0	-0.359	5.945	0.000
HETATM	17	H	0	-0.221	4.734	4.592
HETATM	18	H	0	-0.352	5.969	2.457
HETATM	19	H	0	-0.048	2.268	4.610
HETATM	20	H	0	0.195	1.017	0.000
HETATM	21	H	0	-0.221	4.734	-4.592
HETATM	22	H	0	-0.048	2.268	-4.610
HETATM	23	N	0	0.031	0.655	2.502
HETATM	24	C	0	0.793	-0.083	3.359
HETATM	25	H	0	-0.480	0.166	1.765
HETATM	26	O	0	1.574	0.375	4.179
HETATM	27	C	0	0.639	-1.581	3.248
HETATM	28	C	0	1.730	-2.390	3.575
HETATM	29	C	0	-0.558	-2.190	2.908
HETATM	30	H	0	2.659	-1.924	3.875
HETATM	31	H	0	-1.455	-1.642	2.666
HETATM	32	C	0	0.395	-4.326	3.148
HETATM	33	H	0	0.233	-5.392	3.082
HETATM	34	C	0	1.607	-3.772	3.511

HETATM	35	N	0	0.031	0.655	-2.502
HETATM	36	C	0	0.793	-0.083	-3.359
HETATM	37	H	0	-0.480	0.166	-1.765
HETATM	38	O	0	1.574	0.375	-4.179
HETATM	39	C	0	0.639	-1.581	-3.248
HETATM	40	C	0	1.730	-2.390	-3.575
HETATM	41	C	0	-0.558	-2.190	-2.908
HETATM	42	H	0	2.659	-1.924	-3.875
HETATM	43	H	0	-1.455	-1.642	-2.666
HETATM	44	C	0	0.395	-4.326	-3.148
HETATM	45	H	0	0.233	-5.392	-3.082
HETATM	46	C	0	1.607	-3.772	-3.511
HETATM	47	Cl	0	-1.684	-1.021	0.000
HETATM	48	H	0	2.438	-4.424	-3.741
HETATM	49	H	0	2.438	-4.424	3.741
HETATM	50	C	0	-1.969	-4.131	2.499
HETATM	51	H	0	-2.343	-3.616	1.615
HETATM	52	H	0	-1.829	-5.187	2.288
HETATM	53	H	0	-2.659	-4.005	3.332
HETATM	54	C	0	-1.969	-4.131	-2.499
HETATM	55	H	0	-1.829	-5.187	-2.288
HETATM	56	H	0	-2.343	-3.616	-1.615
HETATM	57	H	0	-2.659	-4.005	-3.332
HETATM	58	N	0	-0.661	-3.536	-2.863
HETATM	59	N	0	-0.661	-3.536	2.863

END

CONECT	1	2	6	17
CONECT	2	1	3	18
CONECT	3	2	4	7
CONECT	4	3	5	8
CONECT	5	4	6	23
CONECT	6	5	1	19
CONECT	7	3	10	16
CONECT	8	4	9	20
CONECT	9	8	10	15
CONECT	10	9	7	11
CONECT	11	10	12	13
CONECT	12	11		

CONECT	13	11	14	21	
CONECT	14	13	15	22	
CONECT	15	14	9	35	
CONECT	16	7			
CONECT	17	1			
CONECT	18	2			
CONECT	19	6			
CONECT	20	8			
CONECT	21	13			
CONECT	22	14			
CONECT	23	5	24	25	
CONECT	24	23	26	27	
CONECT	25	23			
CONECT	26	24			
CONECT	27	24	28	29	
CONECT	28	27	30	34	
CONECT	29	27	31	59	
CONECT	30	28			
CONECT	31	29			
CONECT	32	33	34	59	
CONECT	33	32			
CONECT	34	32	28	49	
CONECT	35	15	36	37	
CONECT	36	35	38	39	
CONECT	37	35			
CONECT	38	36			
CONECT	39	36	40	41	
CONECT	40	39	42	46	
CONECT	41	39	43	58	
CONECT	42	40			
CONECT	43	41			
CONECT	44	45	46	58	
CONECT	45	44			
CONECT	46	44	40	48	
CONECT	48	46			
CONECT	49	34			
CONECT	50	51	52	53	59
CONECT	51	50			

CONECT	52	50			
CONECT	53	50			
CONECT	54	55	56	57	58
CONECT	55	54			
CONECT	56	54			
CONECT	57	54			
CONECT	58	54	41	44	
CONECT	59	50	29	32	