

## Supporting Information.

# A Robust and Cost-Efficient Scheme for Accurate Conformational Energies of Organic Molecules

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## Tables forming the basis of Figures 3 – 7 and 9 – 10 in the main text

**Table S1. Table forming the basis of Figure 2.**

Method	PBE-D3	MNDO	AM1	RM1	PM3	PM6	PM7	QM	DFTB3	GFN-xTB	GFN2-xTB	MMFF94
Q3	0.996	0.785	0.848	0.866	0.839	0.897	0.920	0.975	0.976	0.961	0.971	0.962
Q1	0.980	-0.097	0.150	0.534	0.460	0.285	0.537	0.902	0.833	0.839	0.874	0.800
max	1.000	0.967	0.971	0.983	0.976	0.985	0.998	0.997	0.994	0.979	0.993	0.987
min	0.926	-0.920	-0.951	-0.786	-0.887	-0.314	-0.294	0.264	0.185	-0.153	0.340	0.586
median	0.990	0.523	0.669	0.709	0.726	0.614	0.760	0.949	0.922	0.916	0.927	0.918
Worst compound	18crown6	ProstaglandinE1	Liu_catal	Glufosinate	Glufosinate	Cryptand222	Glufosinate	UH232	18crown6	XylitolPentanit	18crown6	Cimetidine
Best compound	Isradipine	Bispidine	Triacetin	Knoevenagel_cond	Indometacin	Indometacin	Indometacin	Isradipine	Indometacin	Isradipine	Cembrene	XylitolPentanit
median Opt		0.655	0.598	0.753	0.514	0.631	0.599	0.864	0.895	0.853	0.879	0.878
<b>Pearson correlation coefficient (<math>\rho</math>)<sup>a</sup></b>												
18crown6	0.93	0.48	0.41	0.47	0.17	0.14	0.15	0.84	0.19	0.26	0.34	0.67
Artemizinin	1.00	0.73	0.82	0.80	0.84	0.85	0.90	0.96	0.95	0.96	0.97	0.90
Atropine	1.00	-0.19	0.40	0.71	0.74	0.56	0.83	0.97	0.99	0.87	0.97	0.90
Bispidine	1.00	0.97	0.75	0.86	0.79	0.57	0.62	0.99	0.98	0.94	0.96	0.96
Cembrene	1.00	0.95	0.96	0.98	0.96	0.96	0.94	0.98	0.98	0.98	0.99	0.98
Cimetidine	1.00	-0.90	0.48	0.43	-0.15	0.57	0.64	0.97	0.97	0.98	0.93	0.59
Cryptand222	0.97	0.76	0.15	0.22	0.21	-0.31	-0.26	0.36	0.58	0.57	0.66	0.74
Dapiprazole	0.99	0.64	0.56	0.29	0.51	-0.25	0.09	0.87	0.94	0.76	0.86	0.61
Eperisone	0.99	0.71	0.85	0.74	0.83	0.63	0.69	0.97	0.92	0.90	0.91	0.82
Fotemustin	0.99	0.28	-0.04	0.47	0.53	0.48	0.68	0.93	0.90	0.90	0.89	0.93
GABA	0.96	-0.36	0.08	0.87	0.46	0.77	0.88	0.98	0.82	0.96	0.95	0.97
Gemcitabine	0.98	0.10	0.67	0.54	0.64	0.59	0.70	0.84	0.92	0.94	0.86	0.92
Glufosinate	0.98	-0.83	-0.78	-0.79	-0.89	0.04	-0.29	0.94	0.90	0.66	0.93	0.91
Indometacin	0.97	0.52	0.92	0.94	0.98	0.98	1.00	0.96	0.99	0.96	0.98	0.97
Isradipine	1.00	0.38	0.91	0.91	0.50	0.93	1.00	1.00	0.99	0.98	0.98	0.98
Knoevenagel_cond	0.99	0.95	0.97	0.98	0.96	0.91	0.95	0.98	0.91	0.92	0.92	0.92

Liu_catal	1.00	-0.86	-0.95	0.68	-0.22	0.08	0.76	0.93	0.90	0.97	0.99	0.96
Mannosulfan	0.98	0.86	0.39	0.70	0.08	0.61	0.75	0.90	0.88	0.94	0.86	0.93
N_iprit	0.97	0.94	0.92	0.77	0.90	0.70	0.08	0.99	0.61	0.94	0.92	0.84
Oseltamivir	1.00	0.95	0.69	0.89	0.84	0.35	0.76	0.98	0.98	0.84	0.89	0.96
Pantoprazole	0.94	0.72	0.68	0.86	0.76	0.89	0.87	0.90	0.95	0.96	0.97	0.72
Propranolol	0.99	-0.10	-0.04	0.77	0.49	0.61	0.89	0.92	0.97	0.93	0.95	0.96
ProstaglandinE1	1.00	-0.92	-0.84	-0.07	-0.48	0.97	0.97	0.97	0.98	0.98	0.98	0.88
Pyridoxalphosphate	0.98	-0.72	-0.34	0.64	0.28	0.82	0.76	0.93	0.95	0.84	0.92	0.77
Quinuclidinyl	0.99	0.42	0.59	0.69	0.88	0.51	0.83	0.96	0.99	0.84	0.97	0.95
RAMBO	1.00	0.82	-0.12	-0.42	0.76	0.05	0.73	0.94	0.30	0.65	0.76	0.93
RH34	0.98	0.30	0.82	0.87	0.73	0.73	0.83	0.95	0.92	0.96	0.93	0.94
Rolipram	0.96	0.34	0.85	0.66	0.56	-0.17	0.22	0.55	0.74	0.52	0.87	0.77
Rosmaridiphenol	1.00	0.67	0.84	0.80	0.83	0.95	0.96	0.95	0.98	0.95	0.98	0.92
Takemoto	0.99	-0.32	0.79	0.94	0.95	0.93	0.92	0.99	0.90	0.98	0.97	0.92
Triacetin	1.00	0.79	0.97	0.96	0.92	0.92	0.97	0.99	0.97	0.93	0.97	0.98
Tsogoeva	0.99	-0.69	-0.33	0.06	-0.15	0.29	0.54	0.72	0.73	0.84	0.68	0.95
UH232	0.99	0.83	0.94	0.98	0.94	0.92	0.93	0.26	0.98	0.89	0.96	0.98
VX	0.98	0.79	0.85	0.68	0.52	0.88	0.43	0.95	0.83	0.62	0.66	0.65
WAY100635	1.00	0.39	0.59	0.53	0.63	-0.07	0.31	0.84	0.92	0.85	0.88	0.80
XylitolPentanit	0.99	0.72	0.38	0.55	0.78	-0.09	0.37	0.73	0.69	-0.15	0.86	0.99
shiepox_depox	0.99	0.56	0.80	0.83	0.74	0.90	0.93	0.96	0.83	0.87	0.90	0.78

<sup>a</sup> Correlation coefficient is calculated with respect to DLPNO-CCSD(T)/cc-pVTZ conformational energies

**Table S2. Table forming the basis of Figure 3.**

Method	PBE-D3	MNDO	AM1	RM1	PM3	PM6	PM7	QM	DFTB3	GFN-xTB	GFN2-xTB	MMFF94
Q3	0.96	5.38	4.21	3.08	3.86	3.90	3.10	2.35	2.41	2.41	2.04	3.43
Q1	0.45	2.21	1.90	1.60	1.83	1.79	1.53	0.78	1.20	1.29	0.94	1.98
max	1.60	33.44	10.18	7.31	8.16	9.58	6.05	6.53	4.97	4.46	4.22	6.15
min	0.20	1.19	0.81	0.63	0.59	0.70	0.70	0.24	0.50	0.57	0.67	0.61
median	0.61	3.51	2.90	2.46	2.96	2.66	2.25	1.35	1.71	1.82	1.52	2.69
Worst compound	GABA	Prostagla ndinE1	Glufosi- nate	Glufosi- nate	Glufosi- nate	18crown 6	18crown 6	Cryptand 222	RAMBO	18crown6	RAMBO	Oseltamivi r
Best compound	Rosmari diphenol	Cembren e	Indomet acin	Indomet acin	UH232	Rosmari diphenol	Rosmari diphenol	Isradipin e	Indomet acin	RH34	GABA	Liu_catal
median Opt		2.78	2.99	2.79	3.30	2.93	2.44	2.20	2.15	1.86	1.62	2.36
<b>Mean absolute error in conformational energy (MAE)<sup>a</sup></b>												
18crown6	0.7	7.0	3.0	2.6	3.6	9.6	6.1	4.2	2.6	4.5	3.3	1.5
Artemizinin	0.5	2.2	2.5	2.0	4.4	2.7	2.4	1.0	2.9	2.2	2.0	3.1
Atropine	1.0	5.4	4.8	3.8	4.9	4.9	2.7	2.1	2.0	1.9	2.0	1.6
Bispidin	0.4	1.6	3.2	1.7	2.3	4.1	3.5	2.7	1.7	1.7	1.2	5.2
Cembrene	0.5	1.2	2.6	1.6	3.1	3.4	3.1	1.0	2.4	2.0	1.5	2.6
Cimetidine	0.4	6.0	3.4	3.3	4.6	3.2	4.1	1.5	1.4	1.3	1.7	3.0
Cryptand222	0.9	6.7	3.1	3.0	2.7	6.0	5.4	6.5	2.3	2.4	2.4	3.6
Dapiprazole	0.5	2.9	4.2	4.2	3.5	4.3	3.8	1.4	3.6	3.7	3.6	3.1
Eperisone	0.4	1.8	1.3	1.5	1.7	1.8	1.8	0.7	1.2	1.4	0.9	2.9
Fotemustin	0.5	4.8	2.9	2.5	2.6	2.8	2.8	2.5	1.6	1.9	1.4	3.6
GABA	1.6	4.9	3.6	1.2	2.1	2.0	1.6	0.7	1.1	1.3	0.7	2.3
Gemcitabine	0.6	3.4	2.4	3.2	3.1	2.6	1.9	2.4	1.0	0.9	1.2	0.9
Glufosinate	1.0	14.8	10.2	7.3	8.2	3.3	4.0	4.1	1.2	2.7	1.9	1.4
Indometacin	0.7	1.9	0.8	0.6	1.2	0.8	1.0	0.6	0.5	1.8	0.7	2.0
Isradipine	0.5	5.1	2.8	2.7	5.0	2.5	1.2	0.2	0.5	1.6	1.3	2.0
Knoevenagel_cond	1.1	3.6	1.9	0.9	1.8	2.9	2.0	0.5	3.3	2.5	2.5	2.7
Liu_catal	0.5	8.7	4.7	3.3	3.9	4.1	2.0	0.9	3.7	2.1	2.4	0.6
Mannosulfan	1.2	5.4	4.2	3.0	4.9	3.7	2.5	5.0	4.2	2.4	2.0	3.2
N_iprit	1.4	1.3	2.1	1.5	0.7	1.4	2.0	0.4	1.7	1.2	1.5	4.0
Oseltamivir	1.0	1.6	3.9	2.2	5.0	4.3	4.1	0.8	3.1	3.5	3.7	6.1
Pantoprazole	0.9	6.2	2.4	1.7	1.7	1.4	2.4	2.6	0.9	1.7	1.0	3.6
Propranolol	0.3	3.4	3.3	2.1	2.5	2.0	1.0	0.8	1.3	0.7	0.9	3.1

ProstaglandinE1	1.0	33.4	9.6	5.6	6.6	1.1	3.0	3.9	1.6	1.0	0.8	3.3
Pyridoxalphosphate	1.2	8.1	4.4	2.4	3.8	2.1	3.1	1.9	2.7	1.8	2.3	2.5
Quinuclidinyl	0.5	5.0	3.2	2.7	3.0	3.1	2.3	1.0	1.3	1.9	0.9	2.4
RAMBO	0.6	1.3	4.4	4.7	3.8	4.5	3.2	0.7	5.0	4.3	4.2	5.5
RH34	0.4	2.4	1.7	1.4	1.5	1.5	1.2	1.3	1.0	0.6	0.8	1.0
Rolipram	1.1	3.1	1.4	1.5	1.7	1.9	1.5	1.3	1.9	1.4	0.8	2.0
Rosmaridiphenol	0.2	4.1	1.2	2.5	1.8	0.7	0.7	1.5	0.8	0.9	0.7	4.5
Takemoto	0.8	4.6	2.5	2.7	2.2	2.5	1.5	0.5	1.5	2.1	1.6	1.4
Triacetin	0.5	2.6	1.4	1.6	3.0	2.1	1.1	1.7	2.4	1.6	1.4	2.2
Tsogoeva	0.6	10.0	7.5	2.8	3.3	3.1	2.2	2.5	1.7	1.7	1.8	3.2
UH232	0.8	1.5	0.9	2.0	0.6	1.7	1.4	2.1	1.0	1.1	1.1	4.5
VX	0.4	3.0	0.9	1.6	1.1	0.8	1.6	0.8	1.3	1.3	0.9	1.5
WAY100635	0.3	3.5	4.4	4.1	4.1	4.6	3.9	1.8	1.8	3.1	2.4	2.4
XylitolPentanit	0.7	3.0	2.9	3.1	1.8	3.9	2.6	1.8	1.9	3.7	1.5	2.2
shiepox_depox	0.3	1.9	1.7	1.5	2.0	0.9	0.9	0.8	1.7	1.0	1.2	3.4

<sup>a</sup> Mean absolute error is calculated with respect to DLPNO-CCSD(T)/cc-pVTZ conformational energies

**Table S3. Table forming the basis of Figure 4.**

Symmetry	CCSD(T)	PBE-D3	MNDO	AM1	RM1	PM3	PM6	PM7	QM	DFTB3	GFN-xTB	GFN2-xTB	MMFF94	
S <sub>6</sub>	0.0	0.0	10.5	0.0	0.0	2.0	2.1	1.8	0.0	2.8	0.0	0.0	0.6	
C <sub>i</sub>	0.2	0.9	3.9	5.1	3.3	6.2	13.2	8.4	5.5	1.1	5.4	4.5	0.6	
D <sub>3d</sub>	4.3	2.7	0.0	13.2	12.1	11.1	34.4	23.2	6.8	9.2	21.4	16.4	8.8	
C <sub>2</sub>	2.0	2.6	6.2	4.0	1.7	3.9	9.6	6.3	7.0	2.6	5.3	4.5	1.8	
C <sub>s</sub>	2.4	2.7	8.1	7.4	7.5	7.2	14.3	10.4	7.4	3.6	6.9	5.7	3.7	
C <sub>3</sub>	3.0	3.9	6.2	3.0	2.6	5.2	6.9	5.1	8.8	0.0	2.4	2.9	0.0	
D <sub>2</sub>	5.0	5.0	18.4	2.4	1.4	0.0	0.0	0.0	8.7	2.9	1.2	1.1	4.7	
C <sub>2h</sub>	6.4	8.1	17.0	6.8	7.0	6.6	9.4	6.5	12.5	1.3	5.1	6.0	4.5	
<b>Energy difference <math>\Delta\Delta E = (E_{D_{3d}}^{method} - E_{D_{2/S_6}}^{method}) - (E_{D_{3d}}^{DLPNO-CCSD(T)} - E_{D_{2/S_6}}^{DLPNO-CCSD(T)})</math> (kcal/mol)</b>														
		PBE-D3	MNDO	AM1	RM1	PM3	PM6	PM7	QM	DFTB3	GFN-xTB	GFN2-xTB	MMFF94	MM3
$\Delta\Delta E(D_{3d}-D_2)$		-1.6	-17.7	11.5	11.4	11.8	35.1	23.9	-1.2	7.01	20.8	16.1	6.4	-12.7
$\Delta\Delta E(D_{3d}-S_6)$		-1.6	-14.8	8.9	7.8	4.8	28	17.1	2.5	2.13	17.1	12.1	5.1	-12.2

**Table S4. Table forming the basis of Figure 5.**

Method	MNDO	AM1	RM1	PM3	PM6	PM7	QM	DFTB3	GFN-xTB	GFN2-xTB	MMFF94
Q3	24	18	13	18	22	21	11	8	13	9	9
Q1	16	7	7	8	7	10	3	4	5	3	5
max	52	32	42	37	35	45	27	23	26	18	19
min	9	4	2	5	3	3	2	2	2	1	3
median	20	13	10	12	13	12	7	6	7	5	7
Worst compound	Pantoprazole	Pantoprazole	Pyridoxal phosphate	Pyridoxal phosphate	Gemcitabine	Pyridoxal phosphate	RH34	Gemcitabine	VX	Fotemustin	Pyridoxal phosphate
Best compound	Liu_catal	UH232	Bispidin	Knoevenagel_cond	WAY100635	Bispidin	Bispidin	Bispidin	Atropine	Atropine	18crown6
<b>Mean absolute deviation in rotatable bond dihedral (MAE)<sup>a</sup></b>											
18crown6	10	9	10	12	26	17	6	5	18	5	3
Artemizinin	14	7	8	10	7	10	7	5	5	5	7
Atropine	15	9	4	8	5	3	3	2	2	1	4
Bispidin	16	15	2	7	4	3	2	2	2	1	4
Cembrene	21	9	12	9	11	11	2	2	5	5	3
Cimetidine	24	13	9	16	12	14	6	8	6	6	12
Cryptand222	17	10	10	16	29	29	13	5	10	9	4
Dapiprazole	17	7	6	6	8	13	5	2	3	2	6
Eperisone	23	18	11	12	21	21	8	5	5	5	6
Fotemustin	29	19	23	22	22	23	11	18	22	18	9
GABA	13	8	7	6	7	10	3	4	2	3	5
Gemcitabine	23	16	16	21	35	23	9	23	18	10	9
Glufosinate	16	23	16	20	11	10	8	7	6	5	7
Indometacin	35	25	18	22	28	26	8	12	15	8	6
Isradipine	30	6	11	24	5	12	12	3	3	3	15
Knoevenagel_cond	16	5	6	5	11	5	6	4	10	3	5
Liu_catal	9	12	6	7	5	10	6	6	3	2	5
Mannosulfan	28	16	19	14	20	18	17	8	14	14	14
N_iprit	17	6	10	8	13	13	3	10	6	8	9
Oseltamivir	23	5	7	10	11	8	2	3	5	2	5
Pantoprazole	52	32	33	35	31	24	12	16	12	14	18
Propranolol	28	7	10	12	16	17	3	6	7	3	8



ProstaglandinE1	15	7	6	8	7	6	3	4	4	2	5
Pyridoxalphosphate	46	24	42	37	32	45	15	10	23	14	19
Quinuclidinyl	16	17	7	11	15	12	8	6	6	3	13
RAMBO	18	14	7	15	12	8	3	7	5	5	5
RH34	24	18	12	10	18	19	27	8	11	10	14
Rolipram	33	18	13	20	18	18	12	11	13	7	15
Rosmaridiphenol	9	7	8	7	7	6	7	5	6	4	10
Takemoto	20	12	9	11	12	10	10	7	9	8	6
Triacetin	14	14	10	8	24	22	12	5	16	12	8
Tsogoeva	26	16	12	18	14	12	9	12	11	8	8
UH232	22	4	7	6	6	7	3	4	8	6	3
VX	23	22	25	14	26	21	6	7	26	13	9
WAY100635	19	5	9	8	3	6	2	3	3	2	7
XylitolPentanit	13	13	8	15	19	12	25	7	10	17	7
shiepox_depox	22	23	19	24	23	26	5	8	15	9	8

<sup>a</sup> Mean absolute error is calculated with respect to corresponding PBE-D3/def2-tzvp dihedrals

## Tabulated relative conformational energies (kcal/mol)

**Table S5. Tabulated relative conformational energies (kcal/mol) obtained with SP DLPNO-CCSD(T)/cc-pVTZ, PBE-D3/def2-tzvp, AM1, MNDO, PM3, PM6, PM7, RM1, QM, MMFF94, DFTB3, GFN-xTB, GFN2-xTB.**

	No	DLPNO/TZ	PBE-D3	AM1	MNDO	PM3	PM6	PM7	RM1	QM	MMFF94	DFTB3	GFN-xTB	GFN2-xTB
18crown6	1	0.0	0.0	0.0	10.5	2.0	2.1	1.8	0.0	0.0	0.6	2.8	0.0	0.0
18crown6	2	0.2	0.9	5.1	3.9	6.2	13.2	8.4	3.3	5.5	0.6	1.1	5.4	4.5
18crown6	3	4.3	2.7	13.2	0.0	11.1	34.4	23.2	12.1	6.8	8.8	9.2	21.4	16.4
18crown6	4	2.0	2.6	4.0	6.2	3.9	9.6	6.3	1.7	7.0	1.8	2.6	5.3	4.5
18crown6	5	2.4	2.7	7.4	8.1	7.2	14.3	10.4	7.5	7.4	3.7	3.6	6.9	5.7
18crown6	6	3.0	3.9	3.0	6.2	5.2	6.9	5.1	2.6	8.8	0.0	0.0	2.4	2.9
18crown6	7	5.0	5.0	2.4	18.4	0.0	0.0	0.0	1.4	8.7	4.7	2.9	1.2	1.1
18crown6	8	6.4	8.1	6.8	17.0	6.6	9.4	6.5	7.0	12.5	4.5	1.3	5.1	6.0
Artemizinin	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Artemizinin	2	7.4	7.0	4.3	6.3	0.8	5.1	5.3	5.0	9.1	12.9	4.9	6.1	5.5
Artemizinin	3	8.2	7.3	10.1	8.2	4.6	8.8	8.5	11.5	9.0	9.5	6.8	7.0	7.7
Artemizinin	4	7.7	7.2	5.8	5.2	4.0	4.1	4.5	5.7	8.0	9.7	4.6	4.4	5.4
Artemizinin	5	10.0	9.0	7.0	5.5	5.0	6.9	6.9	9.0	10.2	11.7	5.7	7.5	7.3
Artemizinin	6	10.3	9.6	7.9	8.1	6.6	6.6	6.2	9.5	9.9	13.3	5.8	7.2	7.1
Artemizinin	7	11.3	11.0	8.4	11.4	5.2	8.6	8.9	10.0	14.8	20.3	8.2	8.9	8.9
Artemizinin	8	11.2	11.0	6.3	3.9	5.0	5.9	7.3	6.2	11.9	13.5	7.3	7.4	8.3
Atropine	1	0.0	0.0	0.0	9.6	0.4	0.1	0.0	1.3	0.0	0.0	0.0	0.0	0.0
Atropine	2	1.5	0.7	1.5	9.6	1.2	0.6	1.0	2.4	0.5	2.3	0.4	2.2	0.6
Atropine	3	2.9	2.3	1.2	3.3	0.0	0.0	1.4	0.0	4.3	0.5	2.5	4.3	2.1
Atropine	4	8.1	7.2	0.2	0.0	0.4	1.6	4.9	2.0	13.2	6.2	6.3	7.1	5.7
Atropine	5	9.6	8.5	1.0	5.2	2.2	2.2	5.5	4.7	12.7	9.4	6.8	6.9	6.4
Atropine	6	11.6	10.5	2.0	8.3	3.5	1.8	5.3	5.3	13.1	10.3	7.5	6.6	7.1
Atropine	7	11.6	9.9	2.9	5.5	3.5	2.4	6.5	4.7	15.3	9.3	8.1	9.1	8.3
Atropine	8	8.7	6.7	6.6	11.5	4.4	6.5	9.7	7.9	9.4	12.6	6.2	10.3	7.7
Bispidin	1	0.0	0.0	2.1	0.0	2.1	3.9	2.6	0.0	0.0	0.0	0.0	0.0	0.0
Bispidin	2	3.4	3.0	0.0	0.1	0.0	0.0	0.0	0.3	4.6	4.0	1.6	1.4	1.6
Bispidin	3	4.1	3.9	7.0	3.8	6.2	8.4	7.6	4.9	6.8	11.2	3.9	3.9	4.4
Bispidin	4	4.3	4.4	11.1	3.6	8.5	12.8	11.4	7.5	6.7	10.1	4.2	5.1	5.4
Bispidin	5	5.2	4.7	7.0	4.3	5.4	7.4	6.4	5.5	8.5	10.3	3.7	4.1	4.6

Bispidin	6	11.3	10.6	8.0	8.6	7.4	6.1	5.8	7.3	15.7	19.1	7.8	6.9	8.6
Bispidin	7	11.5	10.9	12.2	8.9	10.1	11.2	10.3	10.2	15.4	18.8	8.1	8.3	9.6
Bispidin	8	11.9	11.5	16.6	9.5	13.2	16.5	15.0	13.1	15.3	19.4	8.6	9.9	11.0
Cembrene	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Cembrene	2	3.0	2.8	1.5	2.2	2.0	1.6	2.0	1.9	3.6	3.3	2.6	2.6	2.7
Cembrene	3	3.8	3.2	1.7	2.5	0.8	1.4	0.9	2.7	1.8	6.1	2.4	1.6	2.2
Cembrene	4	6.0	5.6	2.1	4.1	2.3	1.7	3.0	3.0	5.1	6.7	3.8	4.8	4.7
Cembrene	5	6.0	5.8	3.4	8.2	2.0	1.9	1.2	4.5	4.6	9.1	3.2	3.8	4.2
Cembrene	6	9.5	8.8	5.0	7.8	6.0	5.0	6.3	6.7	8.4	12.1	5.6	6.7	7.5
Cembrene	7	10.4	9.7	7.1	11.0	6.4	5.7	6.2	9.3	9.7	16.7	7.2	7.9	8.2
Cembrene	8	12.5	11.1	9.4	13.5	7.3	6.4	6.9	10.3	11.4	17.9	7.1	8.1	9.3
Cimetidine	1	0.0	0.0	0.0	11.8	5.3	1.0	4.9	1.6	0.0	3.4	0.0	0.0	0.0
Cimetidine	2	2.2	2.4	0.9	11.4	5.7	1.8	2.7	2.1	3.1	3.0	0.8	1.9	1.6
Cimetidine	3	5.2	5.4	1.9	6.6	3.1	0.0	0.0	2.1	6.0	0.0	5.2	3.3	3.0
Cimetidine	4	8.1	7.6	0.6	5.6	0.0	0.3	3.2	0.0	8.6	4.0	6.8	5.0	4.0
Cimetidine	5	5.6	6.0	4.9	11.3	9.2	6.4	11.8	7.2	6.8	7.9	4.0	4.8	4.8
Cimetidine	6	6.4	7.2	1.7	6.5	4.4	0.5	4.4	1.6	9.2	7.0	5.0	5.1	3.5
Cimetidine	7	6.0	6.3	5.7	10.0	10.3	6.8	10.0	6.9	5.7	12.0	3.3	4.6	3.4
Cimetidine	8	13.4	14.1	4.2	0.0	5.2	9.4	18.1	7.2	18.9	11.8	11.1	11.8	13.0
Cryptand222	1	0.0	0.0	3.4	7.5	6.4	14.1	12.0	9.5	0.0	7.5	5.2	4.0	1.6
Cryptand222	2	1.0	1.5	3.7	0.0	2.0	9.8	8.2	2.9	9.8	0.1	1.3	2.8	1.6
Cryptand222	3	0.4	1.6	2.9	6.5	1.4	7.3	6.1	1.5	9.5	0.0	0.0	1.6	0.2
Cryptand222	4	2.5	3.0	5.3	2.6	3.1	11.7	10.0	5.6	11.1	2.7	1.8	3.7	2.5
Cryptand222	5	3.2	4.7	0.2	15.3	1.9	0.0	0.0	1.4	8.3	1.9	0.3	0.0	0.0
Cryptand222	6	5.9	6.9	0.0	13.5	0.0	0.5	0.1	0.0	16.1	1.8	0.7	2.7	1.2
Cryptand222	7	4.9	6.2	5.2	15.9	2.4	4.7	3.8	5.4	13.0	6.7	1.8	1.1	1.2
Cryptand222	8	9.9	8.9	5.5	18.2	6.7	9.6	8.9	9.6	7.6	22.6	10.2	8.9	5.2
Dapiprazole	1	0.0	0.0	0.8	2.2	1.1	6.3	4.1	2.3	0.0	1.0	0.0	0.3	0.1
Dapiprazole	2	2.3	2.1	0.9	0.0	1.9	6.5	6.3	2.1	2.9	0.0	1.5	2.4	1.4
Dapiprazole	3	5.3	4.4	0.7	3.5	2.7	4.6	5.4	1.5	5.3	2.6	1.8	2.1	1.7
Dapiprazole	4	5.2	5.0	0.0	2.0	0.0	2.0	2.9	0.0	4.7	2.4	2.5	1.8	2.1
Dapiprazole	5	4.8	4.4	2.9	9.1	3.8	2.3	0.0	2.9	0.3	10.2	1.5	0.0	0.0
Dapiprazole	6	8.8	7.6	0.3	5.8	0.3	0.0	1.4	0.5	7.2	4.8	3.1	2.2	2.4
Dapiprazole	7	10.5	9.6	3.8	6.0	5.3	6.6	7.5	3.5	10.9	5.7	3.4	5.3	5.2
Dapiprazole	8	9.5	9.6	4.9	11.2	5.4	5.0	4.5	4.8	6.3	11.2	4.2	3.8	4.8
Eperisone	1	0.0	0.1	0.7	0.0	0.3	2.3	2.4	1.2	0.5	0.0	1.2	0.7	1.4

Eperisone	2	0.9	0.0	0.0	1.7	0.0	0.0	0.0	0.0	0.0	6.4	0.0	0.0	0.0
Eperisone	3	1.3	0.9	1.6	4.4	1.3	2.3	2.6	2.0	1.2	5.3	1.2	2.1	2.2
Eperisone	4	3.5	3.1	4.4	2.1	2.4	4.1	5.1	6.1	2.9	6.4	3.1	3.9	4.1
Eperisone	5	5.0	4.0	2.4	3.1	1.6	1.3	2.3	3.8	3.8	9.5	2.4	3.3	2.9
Eperisone	6	5.9	6.0	4.9	3.7	4.6	6.9	6.8	5.1	6.9	8.4	4.9	4.4	6.0
Eperisone	7	6.0	6.1	5.1	3.7	4.3	6.9	6.8	5.1	6.8	8.7	5.2	4.3	6.2
Eperisone	8	7.8	7.8	4.4	4.8	3.0	3.9	4.4	4.0	8.0	9.0	5.1	4.3	6.5
Fotemustin	1	0.0	0.0	0.0	0.0	0.0	2.5	2.8	0.0	0.0	0.0	2.8	1.0	2.0
Fotemustin	2	0.2	1.1	14.6	6.4	5.6	4.4	3.2	6.7	0.1	1.6	0.0	0.0	0.0
Fotemustin	3	3.8	2.8	4.8	13.2	2.6	0.0	0.0	4.9	0.5	4.8	1.7	1.6	2.2
Fotemustin	4	4.4	4.9	4.1	11.2	0.9	6.8	2.5	1.1	1.4	4.1	4.2	2.7	4.4
Fotemustin	5	5.2	4.9	4.6	11.7	5.9	3.9	2.8	7.5	2.0	13.5	2.8	2.5	3.6
Fotemustin	6	7.5	7.3	7.3	10.4	5.1	6.7	5.0	7.1	5.0	10.3	5.6	3.6	6.2
Fotemustin	7	9.0	8.5	4.7	5.7	5.7	3.4	5.2	5.7	4.4	16.9	7.1	6.1	6.5
Fotemustin	8	9.7	10.2	7.4	6.6	5.5	8.1	7.8	6.4	6.5	17.1	8.4	9.0	11.7
GABA	1	0.3	0.0	11.2	14.4	4.0	4.0	2.5	2.3	0.2	0.0	3.4	0.8	0.8
GABA	2	0.0	0.6	10.4	15.9	5.1	4.7	2.9	3.2	0.0	0.6	2.2	1.8	1.3
GABA	3	0.0	2.3	1.9	0.6	1.7	1.5	1.0	0.6	1.3	2.1	0.0	0.0	0.0
GABA	4	1.4	3.4	0.0	0.0	0.0	0.0	0.0	0.0	2.9	4.7	0.6	0.6	0.1
GABA	5	2.3	4.4	1.8	2.0	3.2	2.5	1.6	2.8	2.3	3.9	1.3	1.9	1.5
GABA	6	5.1	7.0	6.8	2.3	4.3	7.2	7.1	5.3	6.2	8.8	5.2	7.7	5.6
GABA	7	5.3	7.3	7.0	5.7	6.0	7.2	6.8	6.5	5.6	9.0	4.9	7.8	6.0
GABA	8	6.9	8.5	7.3	3.5	4.6	7.7	7.7	7.0	7.8	10.3	5.8	9.1	6.6
Gemcitabine	1	0.0	0.0	3.2	6.4	0.1	0.0	0.0	1.1	0.0	0.0	0.0	0.0	0.0
Gemcitabine	2	2.2	2.4	0.0	2.1	0.0	0.3	1.9	0.0	5.8	3.9	3.1	2.4	2.1
Gemcitabine	3	3.1	4.1	2.6	0.0	1.6	2.8	4.1	1.9	6.3	2.5	3.4	3.5	4.6
Gemcitabine	4	5.2	4.9	2.3	3.0	2.1	1.7	3.3	2.1	8.4	6.1	3.9	3.6	3.2
Gemcitabine	5	4.7	5.8	4.1	7.4	3.6	4.4	5.4	3.5	7.4	3.6	6.4	5.8	7.2
Gemcitabine	6	6.9	6.8	4.0	9.7	0.8	3.7	4.4	2.2	5.2	7.8	7.3	6.1	5.5
Gemcitabine	7	7.0	7.7	3.4	8.3	2.5	0.8	2.2	1.8	8.9	6.6	8.5	6.3	7.6
Gemcitabine	8	9.3	10.6	6.4	1.0	2.9	3.7	5.5	2.4	11.8	7.4	7.8	7.3	7.8
Glufosinate	1	0.0	0.0	22.6	36.2	18.5	6.5	5.0	20.3	0.0	0.0	0.0	1.3	0.0
Glufosinate	2	1.7	2.4	16.5	23.2	14.2	3.0	4.7	9.0	5.6	3.9	4.4	0.0	1.3
Glufosinate	3	1.9	2.9	14.3	19.0	12.9	4.0	4.4	6.2	5.3	2.0	4.5	0.1	2.1
Glufosinate	4	3.1	4.4	10.0	15.4	4.6	0.0	1.2	9.1	1.2	4.4	4.0	6.6	2.4
Glufosinate	5	7.1	7.3	13.2	17.5	7.7	6.0	7.0	8.2	12.5	5.9	7.1	5.1	4.7

Glufosinate	6	8.3	8.6	11.6	13.6	3.3	7.8	4.9	4.7	15.8	7.5	8.6	5.7	5.6
Glufosinate	7	7.6	9.9	0.0	0.0	0.1	2.3	0.0	0.0	13.1	10.7	6.7	3.4	3.4
Glufosinate	8	8.7	10.7	1.0	0.4	0.0	1.9	0.5	0.4	14.2	11.6	6.9	4.5	4.1
Indometacin	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Indometacin	2	1.4	1.3	0.7	2.0	0.6	1.1	1.7	1.6	0.7	3.2	1.0	1.8	1.1
Indometacin	3	3.4	1.7	5.4	7.6	3.0	4.6	4.1	5.3	4.5	4.9	2.6	6.0	4.6
Indometacin	4	3.6	2.2	6.0	8.5	3.2	4.8	4.3	5.7	4.6	7.0	2.8	7.2	5.1
Indometacin	5	5.2	4.8	6.2	3.6	3.8	6.5	6.7	5.7	4.8	7.4	5.1	7.4	6.3
Indometacin	6	6.0	5.4	6.4	6.1	4.4	6.9	7.7	6.2	5.2	8.7	5.2	8.0	6.2
Indometacin	7	6.5	6.0	6.5	4.1	4.3	7.2	7.8	6.4	6.0	8.6	6.0	8.4	7.4
Indometacin	8	7.3	6.6	7.3	6.0	4.9	7.8	8.8	7.3	6.7	9.9	6.7	9.3	7.7
Isradipine	1	0.0	0.0	0.8	2.0	0.9	0.8	0.0	1.0	0.0	0.0	0.0	0.0	0.0
Isradipine	2	0.1	0.2	1.0	2.5	1.3	1.1	0.7	1.3	0.3	0.6	0.0	0.2	0.2
Isradipine	3	3.3	3.1	0.0	0.0	0.0	0.0	2.1	0.0	2.8	6.3	3.0	1.4	0.9
Isradipine	4	4.6	4.4	1.0	0.0	0.8	1.4	4.0	1.4	4.6	6.9	5.5	3.4	2.5
Isradipine	5	4.6	4.4	1.0	0.0	0.8	1.4	4.0	1.4	4.6	6.9	5.5	3.4	2.5
Isradipine	6	10.3	9.2	7.6	3.8	2.5	8.1	8.3	8.3	10.4	13.4	9.8	12.8	9.1
Isradipine	7	9.9	8.9	6.0	2.1	1.2	7.1	8.3	6.6	9.7	13.4	9.6	12.7	8.1
Isradipine	8	12.0	11.0	8.3	2.5	1.3	8.9	9.4	8.0	12.9	13.0	10.9	15.1	11.6
Knoevenagel_cond	1	0.0	0.0	0.0	0.0	0.0	0.8	0.0	0.0	0.0	0.0	0.5	1.1	0.8
Knoevenagel_cond	2	3.4	1.7	1.3	8.9	3.2	0.0	1.2	4.8	4.3	1.7	0.3	0.0	0.0
Knoevenagel_cond	3	4.2	3.1	1.7	6.8	1.8	1.8	1.5	3.4	3.4	2.5	0.0	1.0	0.4
Knoevenagel_cond	4	5.8	4.9	2.6	6.9	3.1	1.3	1.9	5.5	6.7	2.8	2.7	2.6	3.3
Knoevenagel_cond	5	4.9	4.5	2.4	7.0	2.5	1.5	2.7	5.3	6.1	0.3	3.1	2.8	3.3
Knoevenagel_cond	6	7.5	5.9	5.3	9.8	6.1	4.3	5.1	8.3	7.3	8.2	3.5	4.6	3.7
Knoevenagel_cond	7	8.9	7.7	7.7	14.9	6.2	6.2	8.2	9.6	9.0	14.1	5.2	7.6	7.2
Knoevenagel_cond	8	12.6	10.8	11.1	22.1	9.7	9.8	10.9	15.5	12.6	17.3	6.8	9.6	10.4
Liu_catal	1	0.0	0.0	6.6	14.0	2.2	0.4	0.0	1.4	0.0	0.0	0.0	0.0	0.0
Liu_catal	2	2.4	2.6	5.8	9.4	0.9	0.6	3.3	0.0	3.0	1.4	0.8	0.9	1.2
Liu_catal	3	2.6	2.9	6.9	11.3	1.9	1.3	4.1	0.9	3.2	3.2	0.6	0.9	1.5
Liu_catal	4	3.9	3.8	4.9	15.1	0.0	0.0	2.1	0.2	5.2	5.2	0.3	1.2	1.6
Liu_catal	5	3.9	3.9	4.9	13.6	0.1	0.0	2.4	0.2	5.2	5.4	0.5	1.4	1.7
Liu_catal	6	5.9	6.9	1.4	3.9	2.2	0.1	1.8	3.7	4.1	6.0	0.8	3.8	2.2
Liu_catal	7	8.4	9.6	0.2	0.4	1.1	0.6	5.8	2.8	7.4	8.3	1.6	5.5	4.0
Liu_catal	8	8.7	10.1	0.0	0.0	0.9	0.9	5.5	3.0	7.8	8.3	2.0	5.6	4.2
Mannosulfan	1	0.0	0.0	0.0	0.0	5.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0

Mannosulfan	2	2.0	2.0	1.8	4.5	6.2	1.8	2.3	4.0	3.7	0.7	1.9	1.4	4.4
Mannosulfan	3	3.5	2.7	1.0	6.2	0.0	1.1	3.3	1.5	11.7	4.7	1.3	3.1	6.0
Mannosulfan	4	7.8	4.8	6.5	20.1	3.5	6.4	9.6	5.8	13.7	13.1	1.9	4.0	4.8
Mannosulfan	5	2.6	1.8	5.1	12.0	5.6	5.7	8.0	6.0	2.7	8.3	0.9	1.5	3.9
Mannosulfan	6	9.7	9.3	0.9	13.7	7.7	0.9	5.4	9.0	19.2	8.7	2.7	4.2	9.1
Mannosulfan	7	11.2	9.3	0.0	12.9	4.8	4.4	7.4	4.6	22.3	15.3	5.1	8.9	10.7
Mannosulfan	8	14.7	11.8	7.5	25.1	4.6	8.0	10.2	7.5	18.1	22.0	4.0	9.1	8.8
N_iprit	1	0.0	0.0	0.3	0.0	0.0	0.2	3.0	0.1	0.0	0.0	2.1	0.0	0.8
N_iprit	2	1.1	0.1	0.0	1.9	1.3	1.5	3.0	0.5	1.1	3.6	0.0	0.5	0.0
N_iprit	3	1.6	0.7	0.1	1.2	1.2	0.0	1.9	0.0	1.6	5.0	3.1	0.9	0.9
N_iprit	4	2.2	0.4	0.2	3.3	2.0	1.4	0.0	0.4	1.3	9.0	5.2	0.7	0.5
N_iprit	5	2.2	0.9	0.4	4.2	2.1	4.0	3.4	2.5	2.1	7.0	0.4	1.0	1.3
N_iprit	6	3.4	1.9	0.4	5.3	1.8	0.0	2.5	0.5	3.0	5.7	2.5	1.1	1.2
N_iprit	7	3.3	1.1	1.0	6.3	5.1	4.0	1.9	2.8	3.0	10.6	2.5	2.9	1.4
N_iprit	8	7.7	5.3	3.0	9.1	6.6	5.4	3.0	3.6	6.1	12.7	5.5	4.9	4.7
Oseltamivir	1	0.0	0.0	0.0	0.0	0.0	3.2	0.1	0.0	0.0	0.0	0.0	0.5	0.0
Oseltamivir	2	2.6	1.8	1.4	1.4	0.2	4.2	1.7	2.5	3.1	8.0	1.8	2.4	1.0
Oseltamivir	3	4.2	3.7	0.1	7.8	0.0	0.0	0.0	2.3	2.2	8.6	1.4	0.0	0.2
Oseltamivir	4	6.4	5.2	4.8	6.0	2.7	7.9	5.5	6.7	7.3	10.1	4.1	5.5	5.1
Oseltamivir	5	8.1	6.7	0.8	9.1	1.2	3.3	1.9	3.8	7.0	17.6	4.5	3.6	2.4
Oseltamivir	6	8.5	7.6	4.8	9.9	3.1	3.6	3.2	5.1	8.2	13.6	5.0	3.1	4.1
Oseltamivir	7	11.6	10.4	2.1	11.9	2.2	4.2	4.4	6.3	11.9	19.6	5.7	6.0	5.4
Oseltamivir	8	12.8	10.6	9.1	18.0	4.8	6.0	5.1	10.3	14.1	25.9	6.6	6.4	6.3
Pantoprazole	1	0.0	0.0	0.0	0.0	0.0	0.0	4.7	0.0	4.9	0.0	0.0	2.5	0.0
Pantoprazole	2	4.7	1.8	7.8	13.6	2.6	6.5	6.7	1.6	8.9	5.5	4.6	7.4	2.6
Pantoprazole	3	1.1	1.3	6.4	5.4	4.1	4.5	5.3	3.8	1.7	5.3	0.9	3.3	0.3
Pantoprazole	4	4.5	5.2	3.0	3.5	3.1	2.8	7.9	2.7	8.9	1.7	5.2	6.5	4.8
Pantoprazole	5	0.6	2.4	7.5	14.3	6.7	1.1	0.0	3.6	0.0	9.6	2.2	0.0	1.1
Pantoprazole	6	8.2	7.3	8.6	15.7	8.8	7.2	10.3	9.8	11.1	11.6	5.2	9.2	5.8
Pantoprazole	7	7.2	7.0	8.3	13.1	7.2	6.0	6.5	6.4	8.7	12.7	6.9	9.1	6.3
Pantoprazole	8	11.1	11.0	10.3	19.6	10.6	9.2	12.6	12.0	13.0	14.0	9.6	12.0	10.1
Propranolol	1	0.5	0.1	4.9	3.4	0.6	1.8	2.3	0.6	1.0	0.0	0.0	0.9	0.0
Propranolol	2	0.0	0.0	4.1	8.2	1.0	0.0	0.0	1.2	0.0	3.7	0.4	0.0	0.6
Propranolol	3	1.9	2.2	0.0	1.7	0.7	0.8	2.8	0.0	0.6	2.7	0.8	2.3	1.7
Propranolol	4	3.0	3.2	1.1	0.8	0.5	1.5	3.9	1.1	3.0	5.2	1.5	3.2	2.0
Propranolol	5	1.3	0.5	6.2	3.9	1.9	5.1	4.8	2.2	0.5	2.8	1.9	3.7	2.3

Propranolol	6	5.9	5.6	2.4	0.0	0.0	3.5	5.8	1.5	5.8	9.9	4.3	6.1	4.4
Propranolol	7	6.8	6.6	4.9	2.0	1.8	4.7	6.8	3.8	7.5	12.2	4.9	6.8	6.7
Propranolol	8	7.6	7.3	3.9	8.3	4.1	4.0	6.5	4.4	4.8	14.5	5.2	5.9	5.5
ProstaglandinE1	1	0.0	0.1	9.5	48.8	4.4	0.0	1.5	0.8	0.0	0.0	1.3	0.6	0.2
ProstaglandinE1	2	0.1	0.0	11.5	54.4	5.9	0.7	1.5	2.5	0.6	1.3	0.5	0.0	0.0
ProstaglandinE1	3	0.9	0.5	12.6	58.4	4.3	0.9	0.0	3.8	1.3	2.0	0.0	0.0	0.6
ProstaglandinE1	4	1.1	1.2	12.6	58.6	4.4	1.6	0.1	3.3	1.9	3.6	0.8	1.1	1.4
ProstaglandinE1	5	9.7	11.4	8.8	34.1	4.8	8.6	11.0	5.5	17.5	7.2	10.9	10.2	11.3
ProstaglandinE1	6	12.8	14.5	5.0	18.7	5.5	11.1	18.1	4.0	15.4	5.4	14.4	12.9	12.0
ProstaglandinE1	7	9.8	11.5	0.0	5.3	0.0	13.1	17.9	1.1	19.9	5.9	14.9	14.4	12.8
ProstaglandinE1	8	14.6	16.5	0.6	0.0	1.1	13.4	18.8	0.0	23.7	6.5	16.5	15.5	14.4
Pyridoxalphosphate	1	0.0	0.0	5.4	13.6	2.9	2.9	6.9	3.2	0.0	0.8	0.0	2.6	0.7
Pyridoxalphosphate	2	2.0	2.0	3.6	13.4	0.0	0.2	1.1	0.0	1.5	2.2	0.2	1.3	0.8
Pyridoxalphosphate	3	3.1	2.7	6.4	15.2	2.7	0.0	0.0	2.3	1.0	2.8	0.2	0.0	0.0
Pyridoxalphosphate	4	4.9	7.4	0.0	10.0	1.0	5.2	9.6	5.1	1.7	0.0	2.4	5.2	3.5
Pyridoxalphosphate	5	5.8	7.6	7.1	12.8	8.0	3.6	7.3	11.6	1.9	0.8	1.8	5.0	2.2
Pyridoxalphosphate	6	8.4	9.5	4.1	12.5	2.1	7.5	10.9	6.3	6.0	7.7	5.3	6.1	5.4
Pyridoxalphosphate	7	9.3	10.8	1.4	8.1	1.7	5.9	12.4	5.7	7.2	6.6	5.1	6.6	5.9
Pyridoxalphosphate	8	10.2	12.8	3.5	0.0	5.5	7.9	12.3	8.7	9.6	15.5	7.3	12.3	8.6
Quinuclidinyl	1	0.0	0.0	4.3	6.3	0.7	2.4	0.5	1.8	0.0	1.1	0.0	0.0	0.0
Quinuclidinyl	2	0.4	0.3	4.7	5.5	0.7	3.5	1.4	1.8	1.2	0.0	0.2	0.3	0.4
Quinuclidinyl	3	2.8	3.4	0.0	1.7	0.0	0.0	0.0	0.6	2.1	2.2	2.5	1.7	2.9
Quinuclidinyl	4	4.4	4.7	4.1	0.0	0.7	3.3	2.2	0.0	5.3	7.5	3.7	4.9	4.8
Quinuclidinyl	5	6.8	6.0	8.8	12.2	5.1	4.4	5.1	7.3	5.2	9.1	4.8	2.8	4.3
Quinuclidinyl	6	8.8	8.8	4.5	8.2	4.5	2.1	4.7	6.0	6.6	9.8	7.1	4.5	7.1
Quinuclidinyl	7	8.6	7.0	13.4	19.7	4.6	9.4	9.9	10.8	9.6	15.1	6.0	9.9	8.1
Quinuclidinyl	8	11.0	10.5	8.1	5.2	4.8	5.2	6.6	4.9	10.6	14.9	8.0	7.4	8.8
RAMBO	1	0.0	0.0	3.8	0.0	0.0	1.5	0.0	3.6	0.0	0.0	0.9	0.0	0.0
RAMBO	2	5.0	4.5	0.0	4.6	2.2	0.0	0.4	1.0	2.4	10.0	0.0	0.4	0.4
RAMBO	3	6.2	5.2	1.9	3.8	2.4	4.0	4.5	2.2	5.6	13.8	0.9	2.9	2.6
RAMBO	4	5.5	5.0	0.1	6.3	1.6	0.5	1.7	0.0	4.9	14.1	0.8	1.3	0.9
RAMBO	5	6.0	5.4	3.0	4.1	2.4	2.5	3.6	1.7	6.4	10.8	0.2	1.3	1.4
RAMBO	6	6.6	5.6	2.4	8.3	2.3	0.6	2.7	1.9	5.8	14.1	1.3	1.2	1.6
RAMBO	7	7.2	6.6	2.4	5.7	1.2	0.7	2.5	1.1	7.2	12.0	0.9	1.1	1.3
RAMBO	8	8.1	7.4	3.7	6.3	2.2	1.7	3.7	2.7	7.7	14.1	1.6	1.9	2.7
RH34	1	0.0	0.0	1.4	3.7	2.2	3.8	2.8	1.9	1.0	0.0	0.0	0.9	0.6

RH34	2	0.4	0.4	0.0	2.3	0.0	0.0	0.0	0.0	0.0	0.4	0.7	0.0	0.0
RH34	3	2.7	2.2	0.7	1.6	0.0	0.5	0.6	1.0	3.7	0.9	2.1	1.2	1.1
RH34	4	1.2	1.5	5.5	6.5	3.3	2.3	1.4	4.4	1.8	3.0	1.5	0.3	1.3
RH34	5	4.0	4.5	4.2	5.3	4.1	4.9	2.6	5.2	2.8	4.0	6.8	4.1	5.7
RH34	6	3.8	4.7	5.7	0.0	2.3	5.9	5.3	3.7	4.6	4.1	7.0	3.9	5.4
RH34	7	6.4	5.9	8.0	4.8	5.1	6.3	7.0	7.9	9.0	7.9	7.0	6.3	6.3
RH34	8	7.0	6.5	8.4	6.7	5.1	6.0	6.5	8.4	10.1	9.3	7.4	6.5	6.9
Rolipram	1	0.0	0.0	0.5	2.6	0.0	2.2	1.3	0.7	1.6	1.6	0.0	0.5	0.0
Rolipram	2	1.2	1.1	1.0	0.0	0.0	2.5	1.9	1.1	2.6	0.9	0.2	0.9	0.7
Rolipram	3	0.4	1.6	0.0	3.5	0.7	1.1	0.6	0.0	0.2	1.1	3.3	0.0	1.5
Rolipram	4	0.4	1.8	0.2	3.2	1.0	1.4	0.9	0.1	0.0	1.3	3.7	0.3	1.8
Rolipram	5	3.2	5.4	0.4	0.1	1.3	1.8	2.5	0.9	0.9	0.0	7.3	2.6	4.3
Rolipram	6	4.4	5.5	1.9	0.6	0.4	3.1	3.1	0.6	3.8	6.0	5.6	2.5	3.7
Rolipram	7	4.1	5.0	2.0	8.9	1.3	0.9	1.0	2.5	3.7	7.9	4.2	0.8	2.9
Rolipram	8	4.4	6.2	2.1	7.5	1.6	0.0	0.0	1.8	1.0	8.2	6.7	0.3	4.0
Rosmaridiphenol	1	0.0	0.0	0.0	4.0	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Rosmaridiphenol	2	2.1	2.2	0.7	0.0	0.0	2.3	2.1	0.1	3.5	1.1	2.9	3.2	3.0
Rosmaridiphenol	3	1.4	1.7	5.1	10.6	5.9	4.0	2.7	2.9	1.6	5.5	1.5	1.4	2.2
Rosmaridiphenol	4	4.7	4.8	2.9	5.2	2.8	4.1	4.4	2.2	6.9	5.2	4.0	4.8	4.9
Rosmaridiphenol	5	6.0	6.0	5.0	9.3	6.7	6.6	5.8	2.8	8.9	14.1	5.2	4.8	5.2
Rosmaridiphenol	6	7.0	7.4	6.6	13.5	9.4	8.1	7.1	3.4	10.2	16.4	6.4	6.2	6.6
Rosmaridiphenol	7	7.1	7.1	8.2	11.3	8.4	7.4	5.3	5.0	6.6	12.8	5.6	4.9	6.1
Rosmaridiphenol	8	8.7	9.4	8.2	11.8	9.6	8.5	6.8	3.9	10.0	15.6	7.1	6.7	7.3
Takemoto	1	1.7	0.3	5.0	6.3	0.0	0.8	0.0	0.0	0.0	1.1	4.0	0.8	0.4
Takemoto	2	0.0	0.0	0.0	5.2	1.3	0.0	1.6	0.3	0.3	0.0	0.0	0.0	0.0
Takemoto	3	3.8	2.4	3.9	2.4	2.1	1.2	2.7	3.0	3.7	2.1	3.2	2.1	2.5
Takemoto	4	4.4	3.5	2.6	3.4	4.3	1.4	1.2	0.9	4.3	3.9	3.9	3.1	2.7
Takemoto	5	7.6	6.4	3.5	0.0	5.3	4.9	7.7	4.1	7.5	6.9	5.1	5.0	4.9
Takemoto	6	6.9	7.2	4.9	7.4	4.3	6.1	7.6	5.0	7.1	4.6	6.4	3.7	6.2
Takemoto	7	11.1	10.1	6.5	0.3	6.9	5.9	10.0	6.3	10.0	6.7	6.7	6.5	7.5
Takemoto	8	11.4	11.4	7.2	5.7	8.0	6.6	8.8	6.5	10.7	12.4	10.4	8.7	9.6
Triacetin	1	0.0	0.0	0.0	1.5	0.1	0.3	0.0	0.5	0.0	0.0	0.3	0.2	0.0
Triacetin	2	0.8	1.0	0.1	0.0	0.0	0.3	0.2	0.1	2.3	1.4	0.0	0.6	0.3
Triacetin	3	2.5	2.5	2.6	2.3	2.0	1.8	2.0	2.5	3.8	4.0	0.2	1.3	0.7
Triacetin	4	3.6	3.7	0.8	1.8	0.4	0.0	1.1	0.0	5.3	1.9	1.0	0.0	1.5
Triacetin	5	4.5	4.4	3.1	6.1	3.3	0.5	1.7	2.5	4.9	5.7	0.9	0.4	1.2



Triacetin	6	8.7	7.9	6.4	1.6	2.7	7.6	8.8	6.5	12.7	11.7	6.3	9.5	8.8
Triacetin	7	9.0	8.1	8.4	6.3	4.5	7.2	9.0	7.9	10.3	13.8	6.6	9.1	8.4
Triacetin	8	14.2	12.7	10.7	9.1	6.7	9.1	11.9	11.8	17.3	18.8	9.8	11.4	11.9
Tsogoeva	1	0.0	0.0	9.8	21.8	6.3	1.1	0.0	3.7	0.0	0.0	0.0	0.0	0.0
Tsogoeva	2	2.5	2.3	13.0	16.3	5.1	5.3	4.5	4.7	4.5	4.1	2.7	5.4	2.7
Tsogoeva	3	1.4	1.9	11.2	16.1	5.1	3.3	1.9	2.2	3.8	2.5	2.8	3.7	2.7
Tsogoeva	4	2.5	3.4	12.8	11.6	3.4	5.3	4.1	1.6	7.8	4.8	5.3	7.0	4.9
Tsogoeva	5	4.8	6.3	0.0	0.0	0.0	0.0	0.8	0.0	7.4	11.6	1.8	4.9	2.3
Tsogoeva	6	6.5	6.4	15.6	12.8	4.6	8.9	7.6	5.2	11.3	10.2	6.1	10.4	6.5
Tsogoeva	7	7.2	7.6	8.2	11.0	7.2	4.3	4.7	7.3	7.6	9.2	4.3	7.4	4.3
Tsogoeva	8	9.6	11.0	4.8	4.0	3.6	3.7	3.9	0.9	7.4	17.9	6.3	9.7	4.7
UH232	1	0.0	0.0	1.1	1.3	0.0	2.0	1.8	0.0	1.6	0.0	0.0	0.7	0.4
UH232	2	1.1	0.8	0.0	0.0	0.0	0.0	0.0	0.2	2.4	2.5	0.5	1.2	0.0
UH232	3	1.9	1.1	2.8	0.5	1.8	3.8	3.2	3.6	0.0	6.0	0.7	0.0	0.8
UH232	4	2.4	2.1	3.2	3.1	2.1	3.6	3.1	3.7	2.8	5.1	2.1	1.9	2.0
UH232	5	3.8	2.8	4.6	1.8	3.4	5.5	5.3	6.2	0.9	8.5	2.4	2.1	2.3
UH232	6	4.4	3.1	5.7	7.7	5.6	6.0	4.9	6.6	2.6	12.9	2.9	3.9	3.2
UH232	7	5.1	3.9	6.3	6.7	4.8	8.1	7.8	8.9	1.7	11.9	3.5	3.2	3.9
UH232	8	6.2	4.8	6.2	6.5	4.9	7.6	8.1	9.6	2.7	14.0	4.5	4.4	4.6
VX	1	0.0	0.0	0.5	0.0	0.5	0.0	1.2	0.3	0.0	0.0	0.0	0.0	0.3
VX	2	1.9	1.2	0.0	0.4	1.2	0.7	1.9	0.0	1.0	1.7	1.0	1.0	0.0
VX	3	1.7	1.6	2.5	3.8	2.6	2.1	3.1	3.0	1.1	3.2	1.8	2.6	2.6
VX	4	2.6	2.1	3.4	4.6	4.2	1.7	3.6	6.0	2.7	5.6	1.0	1.7	2.8
VX	5	2.4	2.7	3.2	9.4	2.5	3.1	4.3	0.7	1.4	1.1	1.9	2.3	1.8
VX	6	4.0	3.3	3.2	5.1	0.0	2.8	0.0	4.4	3.0	4.8	1.9	1.0	4.2
VX	7	4.3	3.8	5.5	7.0	4.2	2.9	2.6	7.2	3.5	7.5	1.8	1.9	4.0
VX	8	5.3	5.1	5.4	12.8	4.5	4.9	6.8	4.3	3.5	3.2	2.4	3.2	2.2
WAY100635	1	0.0	0.0	1.6	5.9	1.2	6.1	7.1	2.3	0.0	2.4	0.0	0.6	0.0
WAY100635	2	3.6	4.1	0.0	0.0	0.0	4.1	9.0	0.8	6.3	0.0	3.0	2.8	3.5
WAY100635	3	5.7	5.7	0.1	1.5	0.0	2.8	7.7	0.0	6.9	2.3	2.9	2.8	3.0
WAY100635	4	6.7	7.1	1.0	6.7	0.5	1.7	7.3	1.0	8.1	2.9	6.6	3.6	4.0
WAY100635	5	8.4	8.1	5.7	7.5	5.7	9.2	13.8	7.8	10.4	8.0	5.1	6.4	7.4
WAY100635	6	5.2	5.6	3.9	17.7	4.9	0.0	0.0	3.0	0.9	7.0	2.7	0.0	0.5
WAY100635	7	10.6	11.4	4.0	9.9	4.9	3.9	9.0	4.9	9.6	9.0	9.5	6.5	8.1
WAY100635	8	12.4	12.4	4.2	12.1	4.9	2.9	8.7	4.4	10.8	9.9	8.2	6.4	7.2
XylitolPentanit	1	0.0	0.0	0.1	3.8	0.6	3.1	0.0	0.5	0.0	0.4	1.8	2.0	1.1

XylitolPentanit	2	1.0	1.6	2.0	4.0	2.8	4.8	1.4	1.3	1.9	0.0	0.0	1.3	0.0
XylitolPentanit	3	2.9	3.3	0.6	0.0	0.0	6.1	1.6	0.2	6.7	4.1	3.0	6.2	2.6
XylitolPentanit	4	4.0	4.5	1.1	6.1	2.1	3.0	1.7	0.3	3.7	5.7	2.6	0.0	2.0
XylitolPentanit	5	3.8	5.2	0.0	1.9	2.3	4.3	1.8	0.0	2.3	5.3	1.7	0.7	1.8
XylitolPentanit	6	5.5	6.7	5.2	3.6	8.5	12.2	7.0	3.9	0.6	7.8	6.1	6.8	7.1
XylitolPentanit	7	7.3	8.1	0.0	13.3	5.5	0.0	0.0	0.9	6.6	12.6	3.5	0.1	4.5
XylitolPentanit	8	8.9	9.9	3.4	11.2	7.8	3.3	3.3	3.3	10.8	13.4	4.2	0.5	7.5
shiepox_depox	1	0.0	0.0	0.8	5.0	1.1	0.0	0.0	1.0	0.4	1.4	0.0	0.0	0.0
shiepox_depox	2	0.5	0.3	1.8	3.3	1.3	1.4	0.7	1.5	0.1	0.0	1.1	1.3	1.3
shiepox_depox	3	1.3	2.0	0.1	0.5	0.5	1.7	0.7	0.0	0.9	7.6	0.5	0.6	0.2
shiepox_depox	4	1.7	2.3	0.0	0.0	0.0	1.4	0.4	0.1	0.0	6.2	0.9	1.2	0.9
shiepox_depox	5	3.7	3.8	1.4	5.5	2.4	1.7	2.0	1.8	5.0	9.8	0.6	1.2	1.2
shiepox_depox	6	4.8	4.6	3.6	5.0	2.0	4.8	4.8	4.0	4.7	6.0	3.1	4.4	3.5
shiepox_depox	7	5.1	5.0	2.1	3.4	2.2	3.0	2.8	2.3	5.0	8.7	1.7	2.7	2.8
shiepox_depox	8	6.6	6.7	4.9	7.6	2.3	5.1	5.8	5.2	8.2	10.5	2.9	7.0	5.7

**Table S6. Tabulated relative conformational energies (kcal/mol) obtained with SP DLPNO-CCSD(T)/cc-pVTZ and PBE-D3/def2-tzvp on PBE-D3/def2-tzvp geometries and OPT AM1, MNDO, PM3, PM6, PM7, RM1, QM, MMFF94, DFTB3, GFN-xTB, GFN2-xTB starting from PBE-D3/def2-tzvp geometries.**

	No	DLPNO/TZ	PBE-D3	AM1	MNDO	PM3	PM6	PM7	RM1	QM1	MMFF94	DFTB3	GFN-xTB	GFN2-xTB
18crown6	1	0.0	0.0	0.0	1.5	0.4	0.0	0.0	0.0	0.0	0.0	0.2	0.0	0.0
18crown6	2	0.2	0.9	7.2	0.0	7.9	4.6	6.4	8.9	7.6	1.4	1.0	7.2	6.3
18crown6	3	4.3	2.7	11.4	1.4	12.8	14.7	16.1	18.0	9.6	10.3	9.2	8.0	16.6
18crown6	4	2.0	2.6	5.3	2.7	4.4	1.9	6.9	6.7	6.9	2.7	2.7	4.3	5.9
18crown6	5	2.4	2.7	6.5	2.3	7.0	6.3	7.7	10.3	8.5	4.8	3.0	7.2	5.8
18crown6	6	3.0	3.9	5.4	1.8	0.4	6.7	7.7	3.7	11.2	1.4	0.0	4.8	4.6
18crown6	7	5.0	5.0	3.5	7.9	0.0	1.5	3.6	4.4	8.9	6.6	2.3	2.9	1.7
18crown6	8	6.4	8.1	9.1	5.0	7.8	0.5	11.4	11.2	14.5	5.4	2.1	8.0	8.5
Artemizinin	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Artemizinin	2	7.4	7.0	4.8	0.0	0.9	5.5	5.5	3.6	10.6	10.7	5.0	5.7	5.4
Artemizinin	3	8.2	7.3	4.8	0.0	0.9	5.5	5.5	3.6	9.2	9.1	5.0	6.5	7.2
Artemizinin	4	7.7	7.2	4.6	3.4	3.6	3.7	4.3	4.6	7.9	9.4	4.7	4.3	5.2
Artemizinin	5	10.0	9.0	5.7	5.9	4.6	6.4	6.4	8.0	8.5	11.6	5.8	7.5	7.0
Artemizinin	6	10.3	9.6	5.7	5.9	6.5	6.3	6.0	8.4	8.5	13.0	5.7	7.1	6.8
Artemizinin	7	11.3	11.0	8.1	6.7	5.6	9.0	9.5	9.4	16.5	17.2	8.3	8.7	8.8
Artemizinin	8	11.2	11.0	6.9	3.4	3.6	6.9	4.3	6.2	13.3	12.4	7.6	4.3	5.2
Atropine	1	0.0	0.0	0.1	1.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Atropine	2	1.5	0.7	1.2	2.8	1.2	0.8	1.1	1.5	0.8	2.3	0.3	2.3	0.8
Atropine	3	2.9	2.3	1.1	2.5	0.3	1.3	2.6	0.8	4.9	0.5	2.9	4.6	2.4
Atropine	4	8.1	7.2	0.1	0.8	0.4	2.2	5.9	2.4	15.5	5.1	7.3	7.4	6.2
Atropine	5	9.6	8.5	0.0	0.0	0.3	2.3	5.6	3.3	14.3	8.1	7.6	6.9	6.7
Atropine	6	11.6	10.5	2.8	3.3	1.2	2.3	5.3	3.9	14.1	8.1	8.2	6.4	7.0
Atropine	7	11.6	9.9	4.3	6.3	2.6	3.3	7.2	5.0	16.9	8.7	8.8	9.3	8.6
Atropine	8	8.7	6.7	5.3	7.8	0.3	7.2	9.9	6.3	9.4	12.1	6.5	10.3	7.8
Bispidin	1	0.0	0.0	1.0	0.0	0.8	2.9	2.0	0.0	0.0	0.0	0.0	0.0	0.0
Bispidin	2	3.4	3.0	0.0	2.1	0.0	0.0	0.0	0.8	4.8	4.9	1.5	1.3	1.5
Bispidin	3	4.1	3.9	5.9	3.8	4.7	7.1	6.9	5.0	7.5	10.3	3.9	3.8	4.7
Bispidin	4	4.3	4.4	1.0	2.7	6.2	10.9	10.7	7.3	7.4	8.1	4.2	5.1	5.9
Bispidin	5	5.2	4.7	5.0	3.9	3.9	6.0	5.9	5.2	9.1	9.9	3.7	4.2	4.9
Bispidin	6	11.3	10.6	7.1	7.5	5.5	5.4	5.6	7.5	16.4	19.3	7.8	6.8	8.7

Bispidin	7	11.5	10.9	2.2	2.7	7.4	9.7	9.7	9.8	16.0	17.9	8.1	8.2	9.7
Bispidin	8	11.9	11.5	12.1	6.8	9.7	14.2	14.1	12.2	15.9	17.1	8.5	9.7	11.2
Cembrene	1	0.0	0.0	0.0	0.0	0.6	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0
Cembrene	2	3.0	2.8	1.1	0.5	2.1	1.7	2.1	1.5	3.5	3.0	2.8	2.5	3.0
Cembrene	3	3.8	3.2	1.2	2.9	1.2	1.1	0.0	2.2	1.7	6.0	2.4	1.8	2.1
Cembrene	4	6.0	5.6	1.2	0.0	1.2	1.3	2.9	1.7	5.0	6.9	3.8	5.1	4.9
Cembrene	5	6.0	5.8	2.7	7.8	0.0	1.3	0.7	3.3	4.3	8.2	2.7	3.8	4.1
Cembrene	6	9.5	8.8	4.4	3.8	2.8	4.6	5.5	4.8	8.2	11.2	5.7	6.4	7.4
Cembrene	7	10.4	9.7	5.9	9.3	3.2	5.0	5.4	9.0	9.4	14.8	6.9	7.9	8.4
Cembrene	8	12.5	11.1	7.7	6.9	4.6	6.0	6.1	8.4	11.1	15.9	7.1	8.3	9.3
Cimetidine	1	0.0	0.0	0.0	0.6	0.0	0.0	0.0	0.0	0.0	4.3	0.0	0.0	0.0
Cimetidine	2	2.2	2.4	0.9	0.0	1.9	1.8	1.6	1.1	3.6	3.1	0.8	2.0	2.1
Cimetidine	3	5.2	5.4	2.1	1.5	3.0	1.3	0.0	1.0	5.9	0.0	5.3	3.7	4.0
Cimetidine	4	8.1	7.6	3.7	2.3	2.2	2.2	2.6	0.9	8.9	4.8	7.0	5.5	5.0
Cimetidine	5	5.6	6.0	3.2	0.8	1.4	5.1	5.7	4.6	6.4	9.6	3.9	5.3	5.2
Cimetidine	6	6.4	7.2	3.3	1.1	2.3	1.8	4.0	2.8	9.8	6.9	4.2	4.4	2.9
Cimetidine	7	6.0	6.3	2.2	1.2	1.7	3.7	6.7	1.6	4.9	6.5	3.0	4.3	3.8
Cimetidine	8	13.4	14.1	5.8	1.8	3.9	8.4	2.7	7.2	19.9	15.1	11.9	11.5	13.9
Cryptand222	1	0.0	0.0	0.9	0.8	10.8	7.3	3.8	4.7	0.0	6.7	3.2	1.1	0.0
Cryptand222	2	1.0	1.5	3.9	0.0	0.0	1.6	2.9	2.7	18.9	0.2	1.3	2.0	3.8
Cryptand222	3	0.4	1.6	3.9	1.4	9.4	4.2	4.4	1.8	18.4	0.0	0.4	1.8	2.7
Cryptand222	4	2.5	3.0	2.8	0.7	7.4	2.9	0.9	0.5	7.7	1.5	1.8	1.5	2.4
Cryptand222	5	3.2	4.7	0.0	4.1	4.1	0.0	2.2	1.3	15.4	1.1	0.0	0.6	0.7
Cryptand222	6	5.9	6.9	0.3	3.6	5.3	0.5	1.1	0.0	25.4	1.3	0.9	3.0	2.9
Cryptand222	7	4.9	6.2	6.0	5.1	4.8	2.5	0.0	1.6	15.5	4.9	1.7	0.0	1.5
Cryptand222	8	9.9	8.9	4.1	5.2	5.7	3.2	4.6	5.6	6.5	18.8	8.4	7.5	5.5
Dapiprazole	1	0.0	0.0	1.5	0.1	1.6	4.8	3.5	1.0	0.0	1.4	0.0	0.0	0.0
Dapiprazole	2	2.3	2.1	1.4	0.0	2.1	5.6	3.7	1.0	6.0	0.0	1.7	2.6	1.6
Dapiprazole	3	5.3	4.4	0.6	2.4	1.1	3.3	6.1	0.9	8.4	2.9	2.2	2.5	2.1
Dapiprazole	4	5.2	5.0	0.5	1.9	1.0	2.6	2.3	0.0	7.7	2.5	2.7	2.3	2.1
Dapiprazole	5	4.8	4.4	1.2	4.6	1.6	1.9	0.9	2.8	1.2	9.4	0.9	0.0	0.1
Dapiprazole	6	8.8	7.6	0.0	4.1	0.0	0.0	0.0	0.6	10.2	4.8	3.4	2.8	2.4
Dapiprazole	7	10.5	9.6	2.4	3.7	4.7	5.4	4.9	3.5	13.7	5.8	3.7	6.0	5.3
Dapiprazole	8	9.5	9.6	2.7	3.4	4.3	3.5	5.2	3.9	8.3	9.8	4.7	3.8	4.8
Eperisone	1	0.0	0.1	1.2	0.0	0.9	2.0	2.8	1.4	4.3	0.0	1.5	0.6	1.9
Eperisone	2	0.9	0.0	0.0	1.7	0.0	0.0	0.0	0.8	2.9	5.7	0.0	0.0	0.0

Eperisone	3	1.3	0.9	0.8	0.0	1.9	0.0	0.0	0.0	0.0	4.6	1.2	0.8	2.6
Eperisone	4	3.5	3.1	3.3	2.2	1.1	3.0	3.9	2.7	6.5	3.6	3.1	3.3	3.9
Eperisone	5	5.0	4.0	1.9	3.6	0.2	0.4	2.1	2.5	4.5	8.9	2.7	3.0	3.0
Eperisone	6	5.9	6.0	3.2	2.1	3.6	1.9	3.5	5.2	10.7	7.0	5.1	4.5	6.5
Eperisone	7	6.0	6.1	2.9	2.0	4.5	1.8	2.2	5.5	10.5	7.4	5.5	4.3	6.7
Eperisone	8	7.8	7.8	3.2	2.1	3.6	1.9	3.2	4.3	11.8	8.2	5.5	4.3	6.8
Fotemustin	1	0.0	0.0	0.6	0.0	0.0	0.0	5.8	0.0	5.9	0.0	6.3	2.4	3.6
Fotemustin	2	0.2	1.1	6.3	2.5	6.2	0.2	6.6	4.3	4.3	1.0	5.6	3.2	0.0
Fotemustin	3	3.8	2.8	0.0	2.8	2.9	0.6	3.1	2.1	5.4	3.5	3.9	4.2	3.8
Fotemustin	4	4.4	4.9	5.2	2.2	2.9	7.0	0.0	0.1	6.1	4.5	0.0	0.0	1.3
Fotemustin	5	5.2	4.9	3.6	3.3	2.5	2.4	1.8	4.9	0.0	10.7	6.7	6.2	4.6
Fotemustin	6	7.5	7.3	5.9	2.6	2.0	1.2	8.7	4.2	9.6	10.4	10.2	7.1	4.4
Fotemustin	7	9.0	8.5	4.1	2.3	2.7	1.9	2.9	4.9	8.5	14.3	9.5	8.2	5.8
Fotemustin	8	9.7	10.2	4.2	3.0	2.1	3.2	10.0	4.4	6.0	15.8	13.7	10.6	11.8
GABA	1	0.3	0.0	5.7	3.7	1.2	4.1	2.4	2.0	0.1	0.0	2.5	0.4	0.8
GABA	2	0.0	0.6	5.7	3.4	3.3	4.4	3.1	2.4	0.0	1.3	1.8	1.5	1.4
GABA	3	0.0	2.3	1.6	0.4	1.6	1.3	1.1	1.1	2.1	2.6	0.0	0.0	0.0
GABA	4	1.4	3.4	0.0	0.7	0.0	0.0	0.0	0.4	3.9	5.6	0.5	0.4	0.1
GABA	5	2.3	4.4	1.4	0.0	2.4	2.0	0.2	0.0	2.9	4.4	1.3	2.0	1.4
GABA	6	5.1	7.0	6.6	4.4	3.3	7.7	7.9	5.6	6.7	8.6	5.6	7.5	5.7
GABA	7	5.3	7.3	6.9	4.7	3.5	7.9	7.7	6.4	5.8	9.3	5.0	7.8	6.0
GABA	8	6.9	8.5	7.0	4.3	3.2	8.2	2.4	6.7	8.2	10.1	5.8	8.9	6.6
Gemcitabine	1	0.0	0.0	2.3	0.0	0.2	0.0	0.0	1.4	0.0	0.6	0.0	0.0	0.0
Gemcitabine	2	2.2	2.4	0.7	0.8	0.0	1.4	2.8	1.2	4.7	0.7	3.4	1.4	2.4
Gemcitabine	3	3.1	4.1	2.1	1.0	0.7	1.4	3.9	1.2	6.3	0.0	3.4	2.2	4.2
Gemcitabine	4	5.2	4.9	2.9	3.2	0.5	4.0	5.6	3.5	7.1	0.9	4.2	2.9	2.4
Gemcitabine	5	4.7	5.8	4.9	2.6	2.3	1.4	5.0	5.0	7.4	1.9	6.9	7.3	7.9
Gemcitabine	6	6.9	6.8	0.0	0.4	0.5	0.3	1.4	0.0	3.8	8.5	0.0	1.5	5.7
Gemcitabine	7	7.0	7.7	4.0	2.3	0.9	0.7	5.7	3.9	8.6	5.2	6.9	7.3	8.1
Gemcitabine	8	9.3	10.6	4.7	0.7	3.4	0.9	0.0	2.8	11.9	6.2	3.3	8.4	3.1
Glufosinate	1	0.0	0.0	8.1	7.7	6.8	3.2	1.5	4.5	0.0	0.0	0.0	2.5	0.0
Glufosinate	2	1.7	2.4	0.0	1.9	4.4	0.0	0.4	3.8	6.8	5.1	4.8	0.0	2.3
Glufosinate	3	1.9	2.9	2.4	0.9	8.9	0.5	1.8	3.7	5.5	2.7	4.0	1.6	3.1
Glufosinate	4	3.1	4.4	5.7	3.8	2.8	1.2	0.9	7.3	2.0	5.3	5.6	7.4	3.7
Glufosinate	5	7.1	7.3	4.4	4.0	1.5	4.2	4.2	6.2	5.5	5.9	8.0	6.7	6.0
Glufosinate	6	8.3	8.6	6.0	4.8	1.5	6.3	0.3	4.1	17.5	7.8	9.4	7.7	7.0

Glufosinate	7	7.6	9.9	0.6	0.0	0.0	3.3	0.0	0.0	15.4	11.6	8.0	5.3	4.9
Glufosinate	8	8.7	10.7	1.2	0.8	0.2	2.8	0.0	0.5	16.4	12.7	8.1	5.9	5.6
Indometacin	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Indometacin	2	1.4	1.3	1.4	0.4	1.1	1.5	2.3	1.8	2.1	3.2	1.5	2.4	1.7
Indometacin	3	3.4	1.7	4.8	5.0	1.0	5.0	4.6	3.8	5.0	3.9	3.1	6.8	5.2
Indometacin	4	3.6	2.2	5.2	4.9	1.2	5.0	5.0	4.1	4.0	6.2	3.5	7.7	5.1
Indometacin	5	5.2	4.8	4.4	4.3	0.7	4.5	3.8	6.0	5.9	7.1	6.0	6.2	7.0
Indometacin	6	6.0	5.4	5.2	3.9	1.0	5.2	5.3	4.4	6.3	8.2	6.1	7.7	6.8
Indometacin	7	6.5	6.0	4.8	4.2	0.8	5.0	4.6	3.8	7.0	8.3	3.1	6.7	8.0
Indometacin	8	7.3	6.6	5.3	4.4	1.2	5.2	5.5	4.3	7.6	9.9	3.8	7.8	4.8
Isradipine	1	0.0	0.0	0.7	0.6	0.3	1.8	0.0	2.1	0.0	0.0	0.0	0.0	0.0
Isradipine	2	0.1	0.2	0.8	1.3	0.6	2.0	0.0	2.5	0.0	0.6	0.2	0.3	0.2
Isradipine	3	3.3	3.1	0.0	2.0	1.4	0.0	0.7	0.0	4.1	5.3	3.3	1.3	0.7
Isradipine	4	4.6	4.4	0.8	2.2	1.3	1.2	2.1	0.7	6.4	7.3	6.1	3.4	2.5
Isradipine	5	4.6	4.4	0.8	2.2	1.3	1.2	2.1	0.7	6.4	7.6	6.1	3.4	2.5
Isradipine	6	10.3	9.2	6.7	5.2	0.0	7.9	7.4	6.9	0.0	12.0	9.5	11.9	8.5
Isradipine	7	9.9	8.9	5.5	0.0	1.6	7.3	6.5	6.1	10.4	13.6	9.8	12.3	8.0
Isradipine	8	12.0	11.0	7.8	8.8	2.0	9.7	8.6	9.2	13.9	15.9	11.1	14.7	11.2
Knoevenagel_cond	1	0.0	0.0	0.0	0.0	0.0	0.1	0.6	0.0	0.0	0.0	0.5	0.0	1.2
Knoevenagel_cond	2	3.4	1.7	0.8	5.8	1.6	0.0	0.0	1.6	4.0	2.6	0.0	0.8	0.0
Knoevenagel_cond	3	4.2	3.1	1.3	3.5	2.0	1.8	1.2	2.2	3.0	3.0	0.0	1.2	0.4
Knoevenagel_cond	4	5.8	4.9	2.1	5.5	3.2	1.0	1.5	3.9	7.2	3.8	2.6	3.6	3.3
Knoevenagel_cond	5	4.9	4.5	1.2	3.7	2.0	0.0	1.9	2.5	4.8	1.2	2.4	2.5	3.4
Knoevenagel_cond	6	7.5	5.9	3.5	6.9	5.8	3.1	2.9	4.7	6.2	7.9	1.4	3.2	3.5
Knoevenagel_cond	7	8.9	7.7	6.3	1.6	1.1	6.4	7.7	7.4	7.9	13.5	5.3	8.5	7.4
Knoevenagel_cond	8	12.6	10.8	8.2	2.2	5.5	7.8	9.1	12.8	10.7	17.5	6.3	9.7	10.3
Liu_catal	1	0.0	0.0	0.3	0.2	1.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Liu_catal	2	2.4	2.6	1.3	0.0	1.9	1.0	3.5	0.4	3.1	1.3	1.0	0.6	1.3
Liu_catal	3	2.6	2.9	2.6	1.2	2.9	2.0	4.5	1.5	3.4	3.2	0.9	0.7	1.6
Liu_catal	4	3.9	3.8	0.3	3.7	1.1	1.0	2.5	0.6	5.6	4.6	0.6	1.1	1.8
Liu_catal	5	3.9	3.9	0.0	2.5	0.0	0.9	2.7	0.8	5.6	4.8	0.8	1.3	1.9
Liu_catal	6	5.9	6.9	0.1	1.9	3.6	2.6	3.5	3.9	4.6	7.0	2.9	4.0	2.8
Liu_catal	7	8.4	9.6	0.1	1.1	2.1	3.1	5.7	4.0	8.1	8.7	3.4	5.5	4.6
Liu_catal	8	8.7	10.1	0.1	1.4	2.3	3.5	5.7	4.1	7.9	9.3	3.6	5.7	4.9
Mannosulfan	1	0.0	0.0	1.7	2.9	2.0	0.5	1.6	0.0	8.8	0.0	0.7	0.3	0.0
Mannosulfan	2	2.0	2.0	2.3	2.2	2.3	3.6	0.9	2.5	13.6	2.0	2.6	1.9	6.3

Mannosulfan	3	3.5	2.7	2.3	5.6	0.0	1.1	2.2	3.9	0.0	5.3	0.0	0.4	3.3
Mannosulfan	4	7.8	4.8	3.1	11.2	3.2	7.8	9.1	2.7	12.7	7.8	3.5	1.9	5.6
Mannosulfan	5	2.6	1.8	6.1	8.7	3.8	8.8	6.0	6.7	6.7	9.0	1.4	0.0	2.3
Mannosulfan	6	9.7	9.3	3.6	3.8	1.1	1.7	3.2	5.1	22.0	8.2	4.0	1.0	9.3
Mannosulfan	7	11.2	9.3	2.6	0.0	3.3	0.4	0.0	7.3	31.3	13.2	6.1	8.3	12.2
Mannosulfan	8	14.7	11.8	0.0	10.9	4.9	0.0	7.7	10.3	27.2	19.3	4.0	6.7	9.9
N_iprit	1	0.0	0.0	0.9	0.0	0.7	2.0	6.7	2.1	0.0	0.0	6.2	0.8	1.3
N_iprit	2	1.1	0.1	0.2	1.3	0.7	3.1	5.9	3.3	0.9	3.1	1.6	0.8	0.3
N_iprit	3	1.6	0.7	0.0	1.4	0.2	0.7	4.4	1.4	1.3	3.4	6.8	0.8	1.1
N_iprit	4	2.2	0.4	0.0	2.4	0.0	0.0	0.6	0.0	0.6	9.0	5.0	0.0	0.0
N_iprit	5	2.2	0.9	0.9	2.6	1.3	2.4	4.5	3.0	1.8	8.2	0.0	1.0	1.4
N_iprit	6	3.4	1.9	0.0	1.4	0.1	0.9	0.0	2.1	2.6	7.1	6.3	1.4	1.2
N_iprit	7	3.3	1.1	0.1	2.7	0.6	2.4	2.2	2.7	2.1	11.1	4.4	1.6	0.9
N_iprit	8	7.7	5.3	2.0	4.1	2.8	3.4	2.6	3.0	5.2	13.1	9.2	3.8	1.4
Oseltamivir	1	0.0	0.0	0.0	0.3	4.3	2.6	0.5	0.3	0.0	0.0	0.0	1.5	0.3
Oseltamivir	2	2.6	1.8	1.3	1.9	0.0	2.8	0.7	0.8	4.0	7.7	1.9	3.0	1.6
Oseltamivir	3	4.2	3.7	0.1	4.5	1.8	0.0	0.0	0.0	1.5	8.2	0.9	0.0	0.0
Oseltamivir	4	6.4	5.2	4.5	5.5	4.3	3.3	4.9	4.9	7.6	9.8	3.9	6.3	5.6
Oseltamivir	5	8.1	6.7	0.9	5.7	1.0	1.5	1.3	0.8	6.8	17.2	3.9	4.0	1.9
Oseltamivir	6	8.5	7.6	3.9	4.7	3.1	2.3	2.2	2.0	7.7	13.2	4.6	3.3	4.3
Oseltamivir	7	11.6	10.4	2.1	4.3	0.0	3.9	2.2	4.0	12.6	18.3	5.5	6.7	5.9
Oseltamivir	8	12.8	10.6	7.7	0.0	4.7	4.8	3.9	7.0	14.6	23.5	5.9	7.0	6.4
Pantoprazole	1	0.0	0.0	0.2	0.1	0.0	0.0	4.6	0.0	9.4	0.0	4.7	3.2	0.7
Pantoprazole	2	4.7	1.8	4.5	0.7	3.7	3.7	5.8	3.6	13.0	5.1	6.6	5.5	1.8
Pantoprazole	3	1.1	1.3	0.0	0.0	0.1	3.1	3.8	3.0	0.3	4.6	0.0	2.4	0.0
Pantoprazole	4	4.5	5.2	0.8	0.0	0.4	0.6	6.7	1.1	2.8	1.2	9.3	7.5	1.6
Pantoprazole	5	0.6	2.4	2.2	0.1	2.1	0.1	0.0	1.4	0.0	6.3	6.6	0.0	1.5
Pantoprazole	6	8.2	7.3	1.9	0.4	0.7	4.2	0.5	0.1	14.8	12.1	8.6	8.1	5.4
Pantoprazole	7	7.2	7.0	5.6	0.5	0.9	0.4	5.5	1.1	4.5	8.2	8.9	5.6	2.7
Pantoprazole	8	11.1	11.0	1.9	0.4	0.8	0.4	0.5	1.1	16.0	12.9	14.2	11.8	10.2
Propranolol	1	0.5	0.1	1.4	0.0	1.4	0.7	2.9	0.2	1.6	0.0	0.0	0.9	0.0
Propranolol	2	0.0	0.0	1.3	1.1	0.0	0.0	0.0	0.0	0.0	3.4	0.3	0.0	0.4
Propranolol	3	1.9	2.2	0.0	0.8	2.1	1.6	4.3	0.6	1.2	2.8	1.2	2.6	1.7
Propranolol	4	3.0	3.2	1.6	0.2	1.7	1.0	3.6	1.6	3.8	5.0	1.5	3.4	2.0
Propranolol	5	1.3	0.5	2.7	0.1	2.7	0.7	2.9	1.4	1.0	2.6	1.9	3.3	2.2
Propranolol	6	5.9	5.6	2.4	2.3	1.5	1.0	3.6	1.7	6.6	5.0	1.5	3.4	4.4

Propranolol	7	6.8	6.6	4.9	1.5	3.8	2.0	5.0	2.3	8.3	11.9	5.1	3.5	6.5
Propranolol	8	7.6	7.3	3.2	2.8	4.3	2.9	6.7	5.6	4.4	14.3	5.0	5.8	5.3
ProstaglandinE1	1	0.0	0.1	0.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	1.4	0.6	0.2
ProstaglandinE1	2	0.1	0.0	1.1	1.7	0.6	0.5	0.1	1.5	0.3	1.2	0.3	0.0	0.0
ProstaglandinE1	3	0.9	0.5	1.6	6.1	2.7	0.8	0.8	2.0	1.3	2.2	0.0	0.6	0.5
ProstaglandinE1	4	1.1	1.2	2.3	5.6	1.6	1.9	1.4	2.5	2.1	3.6	1.2	1.6	1.6
ProstaglandinE1	5	9.7	11.4	6.3	3.9	3.2	8.0	13.4	3.1	19.1	7.5	12.2	11.2	11.5
ProstaglandinE1	6	12.8	14.5	3.7	0.0	7.4	9.3	16.4	2.6	17.2	6.0	15.6	13.2	12.4
ProstaglandinE1	7	9.8	11.5	4.7	0.6	1.8	11.2	15.0	2.1	23.5	8.5	14.7	15.1	13.8
ProstaglandinE1	8	14.6	16.5	3.8	0.3	2.9	12.7	17.9	1.3	26.9	9.4	17.7	16.6	15.2
Pyridoxalphosphate	1	0.0	0.0	0.0	0.0	0.0	5.1	0.0	0.3	1.2	2.1	0.4	2.3	1.5
Pyridoxalphosphate	2	2.0	2.0	1.2	2.0	1.2	6.8	0.5	0.0	0.0	1.9	0.0	0.0	0.0
Pyridoxalphosphate	3	3.1	2.7	2.5	1.1	1.1	7.2	1.8	1.2	1.2	3.8	0.9	0.3	0.9
Pyridoxalphosphate	4	4.9	7.4	3.2	0.5	6.0	10.8	4.3	1.8	3.2	1.1	3.3	2.1	4.9
Pyridoxalphosphate	5	5.8	7.6	4.2	2.2	4.2	0.0	9.3	1.9	2.6	1.1	0.7	4.0	1.6
Pyridoxalphosphate	6	8.4	9.5	5.7	1.1	7.4	16.6	8.1	3.4	7.0	0.0	5.6	5.2	5.7
Pyridoxalphosphate	7	9.3	10.8	4.0	0.1	6.8	15.1	0.6	2.1	8.7	1.2	5.8	7.0	6.4
Pyridoxalphosphate	8	10.2	12.8	7.2	1.1	3.6	11.9	8.0	1.8	10.6	16.6	7.9	9.4	9.9
Quinuclidinyl	1	0.0	0.0	1.1	0.6	0.0	0.1	0.0	0.2	0.0	0.9	0.0	0.0	0.0
Quinuclidinyl	2	0.4	0.3	1.2	0.8	0.8	0.8	0.9	0.6	0.8	0.0	0.4	0.5	0.5
Quinuclidinyl	3	2.8	3.4	0.0	0.0	0.2	0.0	1.7	0.0	2.1	2.3	2.0	2.0	3.1
Quinuclidinyl	4	4.4	4.7	3.8	1.6	1.4	3.7	4.0	0.9	5.2	4.2	3.5	5.2	4.5
Quinuclidinyl	5	6.8	6.0	5.4	8.5	5.0	2.5	4.3	4.9	5.1	8.5	4.5	3.3	4.2
Quinuclidinyl	6	8.8	8.8	4.2	6.0	4.6	1.9	5.4	4.0	6.2	9.5	6.1	4.5	7.0
Quinuclidinyl	7	8.6	7.0	10.2	0.6	1.7	8.4	10.1	8.2	9.3	14.1	6.3	10.4	8.1
Quinuclidinyl	8	11.0	10.5	8.1	8.0	5.6	6.0	8.0	5.1	10.4	11.0	7.6	7.5	8.4
RAMBO	1	0.0	0.0	2.6	0.0	0.0	0.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0
RAMBO	2	5.0	4.5	1.3	8.0	5.1	2.4	2.2	1.0	3.9	8.9	2.0	0.0	3.6
RAMBO	3	6.2	5.2	0.0	8.1	1.9	0.0	5.0	2.3	8.0	12.9	3.1	2.4	5.9
RAMBO	4	5.5	5.0	0.6	8.2	1.3	2.0	2.6	0.3	7.0	12.8	2.9	0.7	4.1
RAMBO	5	6.0	5.4	4.1	8.8	5.0	4.7	5.5	2.0	9.0	9.6	2.4	1.1	4.8
RAMBO	6	6.6	5.6	3.2	11.0	3.8	2.7	4.0	1.6	8.0	12.7	3.2	0.5	4.7
RAMBO	7	7.2	6.6	3.4	9.4	3.3	3.3	4.5	1.1	9.7	10.9	3.1	0.8	4.6
RAMBO	8	8.1	7.4	4.2	8.8	3.5	3.7	5.1	2.5	10.2	9.6	3.6	1.2	5.5
RH34	1	0.0	0.0	1.6	0.0	0.5	5.0	6.0	0.8	0.7	0.0	0.0	0.7	0.8
RH34	2	0.4	0.4	0.0	1.4	0.0	0.0	0.0	0.0	11.8	0.4	0.7	0.0	0.0



RH34	3	2.7	2.2	2.1	1.8	0.3	0.3	4.5	1.1	16.1	1.5	2.0	1.0	1.3
RH34	4	1.2	1.5	6.3	1.8	3.7	4.7	5.4	3.7	12.9	2.2	1.3	0.3	1.2
RH34	5	4.0	4.5	5.0	0.5	3.5	6.8	6.2	3.6	11.3	3.8	2.6	3.8	5.4
RH34	6	3.8	4.7	6.8	1.2	2.9	8.1	2.9	3.5	0.0	3.8	2.1	3.9	5.5
RH34	7	6.4	5.9	2.1	1.4	2.2	6.9	8.3	5.5	16.2	5.3	6.9	4.6	5.3
RH34	8	7.0	6.5	2.0	1.4	2.3	3.6	8.3	0.7	14.6	2.9	7.2	4.6	5.7
Rolipram	1	0.0	0.0	0.5	0.0	0.7	2.4	3.0	1.0	0.4	1.7	0.0	2.3	0.0
Rolipram	2	1.2	1.1	0.6	0.0	0.6	2.4	2.9	0.9	1.6	1.6	0.0	2.1	0.0
Rolipram	3	0.4	1.6	0.0	0.1	0.4	0.5	1.3	0.2	0.8	0.0	1.4	0.0	0.3
Rolipram	4	0.4	1.8	0.2	0.0	0.7	0.5	1.3	0.2	0.6	0.3	1.4	0.0	0.3
Rolipram	5	3.2	5.4	0.0	0.0	1.1	2.1	3.2	0.0	0.0	0.1	6.9	4.0	4.0
Rolipram	6	4.4	5.5	1.5	0.0	0.1	2.9	3.0	0.7	4.9	7.1	5.2	3.8	3.5
Rolipram	7	4.1	5.0	1.2	0.0	0.1	0.0	0.0	1.0	4.5	7.2	1.8	0.4	1.8
Rolipram	8	4.4	6.2	1.3	0.0	0.0	0.0	0.0	0.2	0.8	6.9	1.8	0.4	2.7
Rosmaridiphenol	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Rosmaridiphenol	2	2.1	2.2	0.5	0.1	0.1	1.9	1.8	0.6	3.6	1.6	3.4	3.1	2.5
Rosmaridiphenol	3	1.4	1.7	3.8	2.8	3.5	3.5	2.8	2.0	0.0	4.3	1.2	0.9	1.7
Rosmaridiphenol	4	4.7	4.8	3.1	3.5	3.6	4.6	5.2	3.7	7.2	5.1	4.4	5.0	5.2
Rosmaridiphenol	5	6.0	6.0	5.2	5.0	7.4	6.9	6.5	4.2	8.9	11.7	5.5	4.9	5.4
Rosmaridiphenol	6	7.0	7.4	6.1	7.1	8.6	7.8	6.8	4.7	10.3	15.2	6.8	5.9	6.1
Rosmaridiphenol	7	7.1	7.1	5.2	4.3	6.4	6.2	4.8	4.7	3.6	8.8	4.7	4.3	5.9
Rosmaridiphenol	8	8.7	9.4	7.7	7.0	8.1	8.6	6.3	4.8	5.4	15.4	7.5	6.7	7.1
Takemoto	1	1.7	0.3	1.0	1.6	0.4	0.0	0.0	0.0	0.0	2.7	2.3	1.2	0.4
Takemoto	2	0.0	0.0	0.0	3.3	0.3	1.7	3.4	1.0	0.1	0.0	0.0	0.0	0.0
Takemoto	3	3.8	2.4	1.9	3.3	0.0	1.4	3.6	2.5	3.4	4.0	3.3	2.8	2.8
Takemoto	4	4.4	3.5	2.4	4.0	2.5	2.6	2.4	2.4	4.6	5.3	3.7	3.1	3.0
Takemoto	5	7.6	6.4	0.9	0.0	2.6	1.5	4.9	4.7	5.6	6.8	2.8	2.1	1.8
Takemoto	6	6.9	7.2	7.6	6.6	5.4	4.5	7.9	7.2	5.8	5.8	6.0	4.6	6.5
Takemoto	7	11.1	10.1	2.6	2.3	6.3	2.5	5.9	3.5	7.8	7.6	6.8	2.9	4.0
Takemoto	8	11.4	11.4	3.8	2.5	8.9	3.4	3.8	4.7	4.8	12.8	4.1	3.7	3.6
Triacetin	1	0.0	0.0	0.2	0.0	0.0	1.6	0.5	0.0	0.0	0.0	0.1	1.7	1.4
Triacetin	2	0.8	1.0	0.1	0.7	0.5	0.5	0.3	1.0	3.4	1.3	0.0	1.9	0.0
Triacetin	3	2.5	2.5	0.0	1.9	2.4	0.2	0.3	3.6	4.7	3.9	0.4	2.5	2.1
Triacetin	4	3.6	3.7	0.0	2.8	1.1	0.0	0.0	0.8	5.9	2.0	1.2	0.0	0.3
Triacetin	5	4.5	4.4	0.0	4.2	0.9	0.3	0.5	2.7	5.3	5.1	1.2	3.2	2.5
Triacetin	6	8.7	7.9	5.7	5.0	1.1	4.9	5.5	7.7	13.1	11.9	7.1	7.3	8.2

Triacetin	7	9.0	8.1	7.2	6.3	2.0	7.7	10.2	7.6	10.7	13.2	6.9	11.9	9.8
Triacetin	8	14.2	12.7	8.7	0.3	4.8	6.9	8.9	3.0	0.8	2.6	6.6	10.1	8.4
Tsogoeva	1	0.0	0.0	4.5	3.5	3.9	0.2	0.0	0.9	0.0	0.0	0.0	0.0	0.0
Tsogoeva	2	2.5	2.3	4.9	2.8	2.9	1.4	1.9	2.1	5.5	3.1	2.3	3.0	2.5
Tsogoeva	3	1.4	1.9	5.2	3.0	2.9	1.4	2.1	1.9	4.6	2.2	2.7	3.2	1.8
Tsogoeva	4	2.5	3.4	5.6	2.1	1.9	2.6	3.8	3.3	9.0	5.2	1.1	6.1	3.6
Tsogoeva	5	4.8	6.3	0.0	0.0	0.0	0.0	1.2	0.0	9.6	12.1	2.0	4.5	1.9
Tsogoeva	6	6.5	6.4	6.8	3.9	2.2	3.4	5.0	4.6	9.4	9.9	5.2	7.5	4.5
Tsogoeva	7	7.2	7.6	5.6	5.3	4.0	4.3	5.5	4.1	9.2	7.5	4.8	7.0	4.4
Tsogoeva	8	9.6	11.0	1.2	3.1	1.0	2.7	4.0	1.9	9.0	12.7	4.4	8.0	4.1
UH232	1	0.0	0.0	1.3	2.1	0.0	2.1	2.0	0.4	2.0	0.0	0.0	0.4	0.4
UH232	2	1.1	0.8	0.0	0.0	1.0	0.0	0.0	0.0	3.0	2.4	0.3	1.3	0.3
UH232	3	1.9	1.1	2.6	5.0	3.3	3.8	2.6	3.2	0.0	5.4	0.6	0.0	0.9
UH232	4	2.4	2.1	2.8	2.8	0.8	3.3	2.4	0.9	3.4	4.9	2.1	1.5	2.1
UH232	5	3.8	2.8	3.7	5.5	3.9	4.7	3.8	4.8	0.8	7.6	2.5	2.0	2.3
UH232	6	4.4	3.1	3.0	5.3	4.0	4.0	1.6	4.5	1.9	9.6	0.4	0.2	0.0
UH232	7	5.1	3.9	4.5	5.4	4.7	5.6	5.1	6.5	1.4	10.5	3.4	2.8	3.5
UH232	8	6.2	4.8	4.8	3.9	4.7	5.1	5.4	7.9	2.4	12.9	4.6	2.3	3.9
VX	1	0.0	0.0	0.6	0.0	2.4	0.0	1.4	0.0	0.2	0.0	0.2	0.0	0.2
VX	2	1.9	1.2	0.0	0.3	0.0	0.1	1.7	1.3	0.0	2.9	0.9	1.0	0.0
VX	3	1.7	1.6	1.6	2.0	3.6	1.7	3.2	1.1	0.9	1.4	0.6	0.4	0.6
VX	4	2.6	2.1	5.1	2.9	3.5	1.7	0.7	1.7	2.6	3.7	0.0	1.3	2.6
VX	5	2.4	2.7	4.8	0.0	5.5	4.3	4.7	2.0	1.4	0.7	1.4	1.7	2.1
VX	6	4.0	3.3	4.0	3.3	4.1	2.6	0.0	0.9	2.7	6.0	0.9	1.2	2.8
VX	7	4.3	3.8	1.4	6.4	2.4	4.3	0.2	1.5	3.7	6.8	1.3	2.4	3.6
VX	8	5.3	5.1	5.5	3.5	5.1	4.6	6.3	5.1	3.4	3.1	2.1	4.2	2.6
WAY100635	1	0.0	0.0	1.1	0.9	1.4	5.0	7.2	2.0	0.0	3.0	0.0	1.0	0.0
WAY100635	2	3.6	4.1	0.3	0.0	0.7	3.3	9.3	0.0	7.8	0.0	1.9	3.0	3.3
WAY100635	3	5.7	5.7	0.0	1.0	0.8	2.7	8.4	0.6	8.6	1.8	1.7	3.2	2.9
WAY100635	4	6.7	7.1	0.1	2.0	0.0	0.8	6.7	1.1	9.9	2.0	6.8	4.3	4.4
WAY100635	5	8.4	8.1	5.3	3.3	4.7	7.2	12.6	5.8	11.4	8.7	4.7	6.7	6.9
WAY100635	6	5.2	5.6	0.7	6.9	1.6	0.0	0.0	0.0	0.5	4.2	2.0	0.0	0.1
WAY100635	7	10.6	11.4	2.3	5.4	3.3	2.6	6.2	4.1	10.6	8.2	8.3	6.7	7.7
WAY100635	8	12.4	12.4	2.4	6.6	1.8	2.5	8.3	4.3	11.5	9.4	7.7	6.5	6.6
XylitolPentanit	1	0.0	0.0	2.5	0.5	1.4	3.1	0.1	1.5	1.6	0.0	1.2	1.5	0.0
XylitolPentanit	2	1.0	1.6	5.1	0.0	2.6	3.1	0.0	4.4	2.4	0.2	0.0	1.3	1.3

XylitolPentanit	3	2.9	3.3	2.1	0.2	0.0	0.1	1.0	3.7	8.9	4.6	1.3	2.1	3.1
XylitolPentanit	4	4.0	4.5	2.0	3.5	3.0	2.9	1.3	0.0	13.2	5.5	0.8	0.0	3.2
XylitolPentanit	5	3.8	5.2	2.1	2.8	5.1	6.0	2.5	2.7	0.0	4.3	0.3	0.6	3.9
XylitolPentanit	6	5.5	6.7	6.1	1.5	3.0	7.6	4.3	3.2	3.8	5.3	4.4	2.6	5.8
XylitolPentanit	7	7.3	8.1	1.1	12.7	1.3	0.0	0.2	1.7	2.9	11.0	2.6	0.5	7.8
XylitolPentanit	8	8.9	9.9	0.0	5.0	1.8	1.9	2.4	5.1	14.0	12.8	2.2	0.6	6.2
shiepox_depox	1	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.2	1.2	1.8	0.0	0.0	0.0
shiepox_depox	2	0.5	0.3	0.0	0.0	0.1	0.0	0.0	0.4	0.9	0.0	1.0	0.0	1.0
shiepox_depox	3	1.3	2.0	0.0	0.0	0.1	0.0	0.0	0.0	0.3	6.5	0.4	0.9	0.0
shiepox_depox	4	1.7	2.3	0.0	0.0	0.1	0.0	0.0	0.3	0.0	6.0	0.9	0.9	0.6
shiepox_depox	5	3.7	3.8	1.2	2.7	0.4	1.4	2.0	0.9	7.0	8.1	0.7	0.0	1.1
shiepox_depox	6	4.8	4.6	1.2	2.7	0.3	1.4	2.0	0.9	6.4	5.8	2.2	0.0	0.6
shiepox_depox	7	5.1	5.0	1.2	2.7	0.3	1.4	2.0	0.9	6.9	7.0	2.0	0.0	1.1
shiepox_depox	8	6.6	6.7	1.2	2.7	0.0	6.8	2.0	0.9	8.9	8.7	3.5	7.3	5.6