

Supporting Information

Fungal bis-Naphthopyrones as Inhibitors of Botulinum Neurotoxin Serotype A

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In Silico Screen

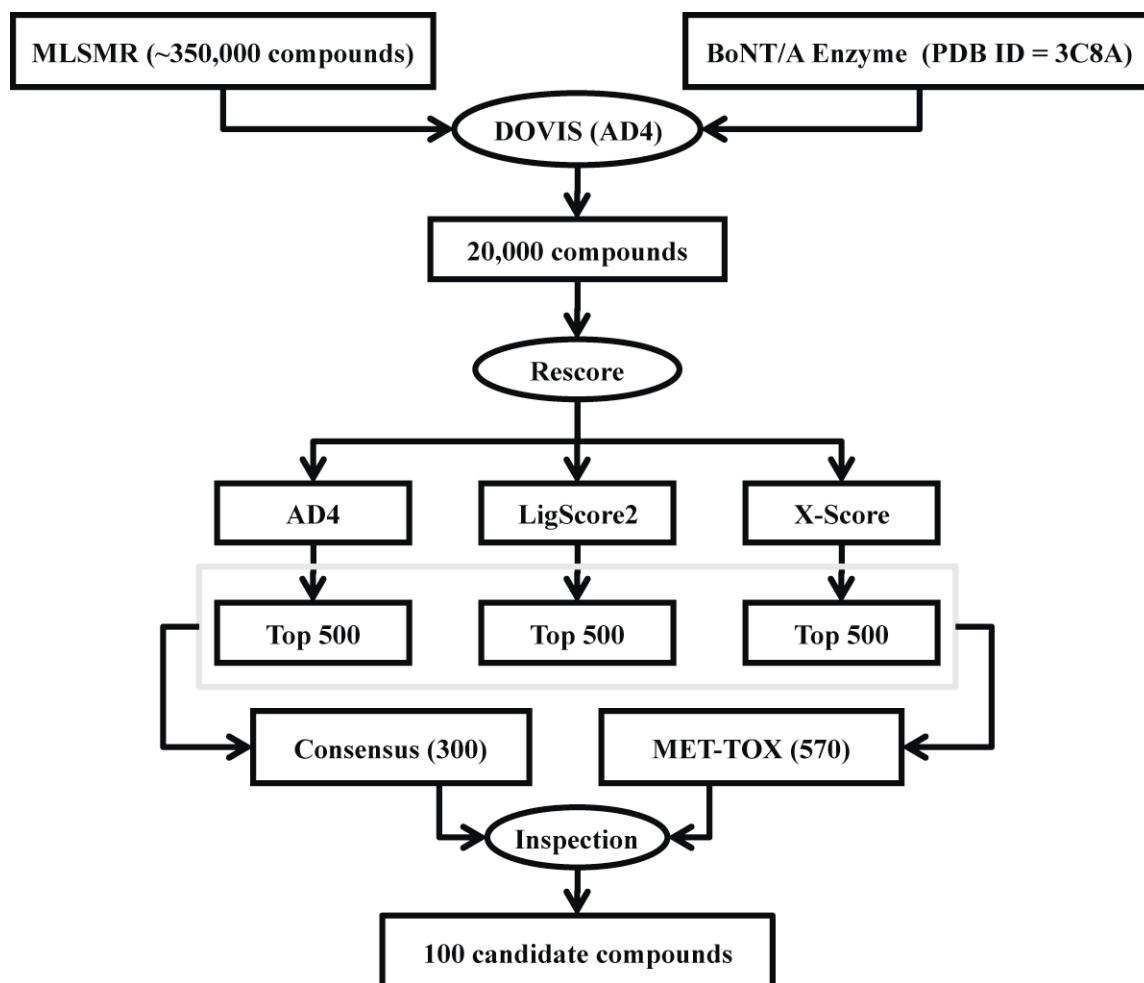


Figure 1S. Virtual screening protocol and flow diagram.

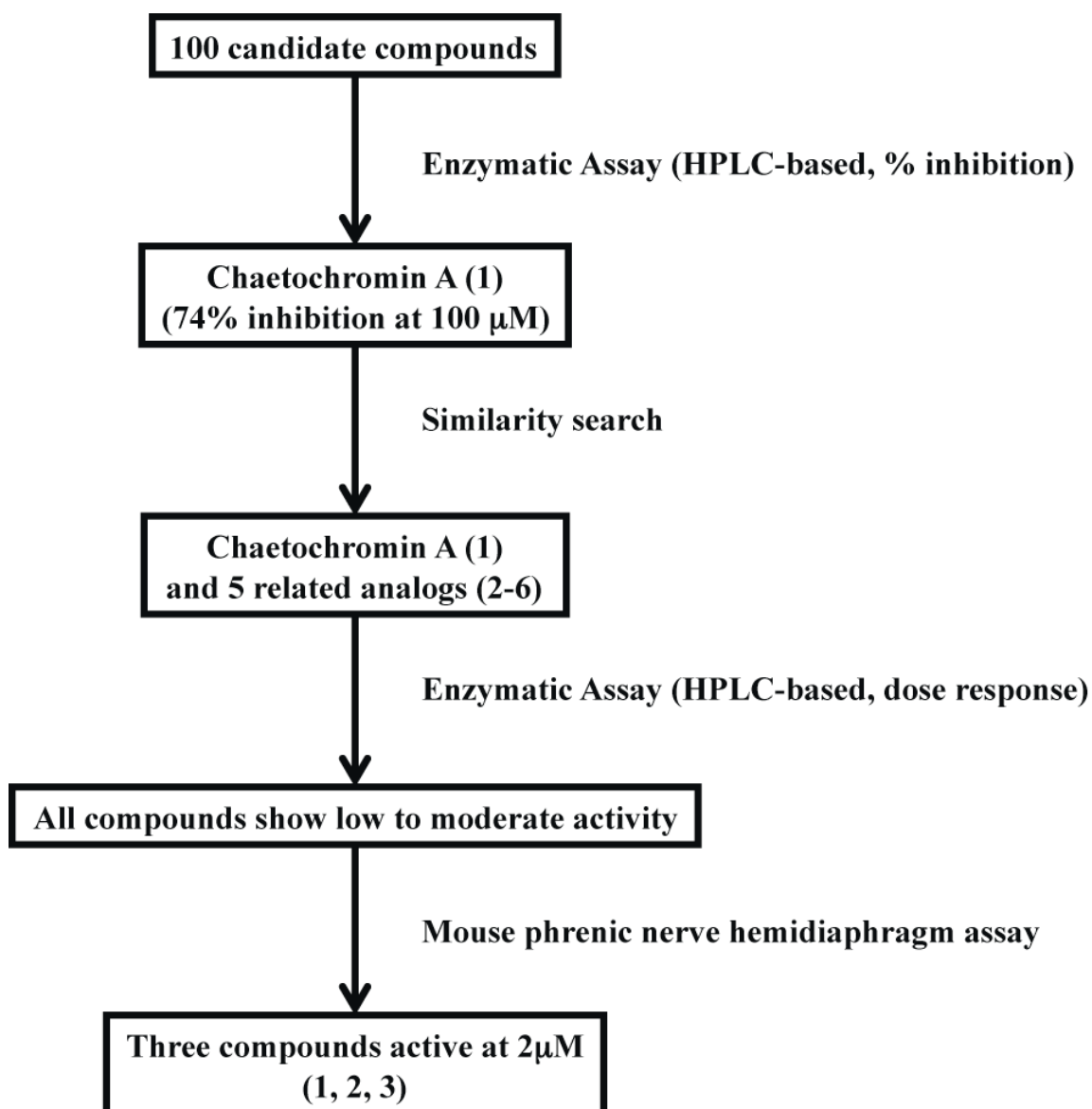


Figure 2S. Bioassay evaluation of candidate compounds.

Bioassays

Materials. BoNT/A (Hall strain) was obtained from Metabionics (Madison, WI). The specific toxicity of the toxin was 2.4×10^8 mouse i.p. LD₅₀/mg of protein, as determined by a toxin titration procedure described previously.¹ The substrate for the HPLC protease assay was a peptide containing residues 187-203 of SNAP-25.² An Alliance HPLC System (2695 XE Separation Module and 2996 Photodiode Array Detector) and Empower/Millennium software were purchased from Waters (Milford, MA). HPLC columns (Hi-Pore C18, 0.45 x 25 cm) were obtained from Bio-Rad Laboratories (Hercules, CA).

Botulinum neurotoxin (BoNT) inhibition assays. HPLC assays of the protease activity of BoNT/A were carried out as previously described.³ IC₅₀ (50% inhibitory concentration) values against BoNT/A LC were calculated from nine concentrations of compound by a log-probit analysis program using the statistical software GraphPad Prism 4 (GraphPad Software, La Jolla, CA).

The mouse phrenic nerve hemidiaphragm assay (MPNHDA) was conducted as previously described.³ BoNT/A-induced paralysis was measured as a 50% loss of twitch tension evoked by nerve stimulation. Estimates of statistical significance were based on unpaired t-test (two-tailed). *P* values of <0.05 were considered significant. Statistical analysis was performed using SigmaPlot 10 (Systat Software, San Jose, CA).

Procedures used to obtain mouse tissues were conducted in compliance with the Animal Welfare Act and other federal statutes and regulations relating to animals and experiments involving animals, and adhered to principles stated in the *Guide for the Care and Use of Laboratory Animals*, National Research Council, 1996. The facility where this research was conducted is fully accredited by the Association for Assessment and Accreditation of Laboratory Animal Care International.

Isolation of Fungal Natural Products

Chaetochromin A. A culture of *Chaetomium arcuatum* (syn. *Chaetomium virescens*) (NRRL 25243 = IMI 86456), originally isolated from soil collected in Lucknow, India) was subjected to solid-substrate fermentation on rice (8 x 50 g rice in 500-mL Erlenmeyer flasks). After 45 days, the cultures were extracted with 2 L of EtOAc, and the solvent was evaporated to afford approximately 3 g of an extract that was then partitioned between acetonitrile and hexane. A 100-mg portion of the acetonitrile-soluble fraction (2.2 g) was placed onto a silica gel column and eluted sequentially with hexane-acetone-MeOH mixtures of increasing polarity. A fraction eluting with 1:1 hexane-acetone (45 mg) was identified as chaetochromin A (**1**) by comparison of its ¹H NMR, ¹³C NMR, CD, and MS data with literature values.⁴

Talaroderxines A and B. The culture of *Delitschia* sp. was a subculture of an isolate originally obtained from a sample of kangaroo dung collected by J. A Scott from scrubland in Australia at a site 20 km west of Balranald in March 1994. This isolate was assigned the accession number JS

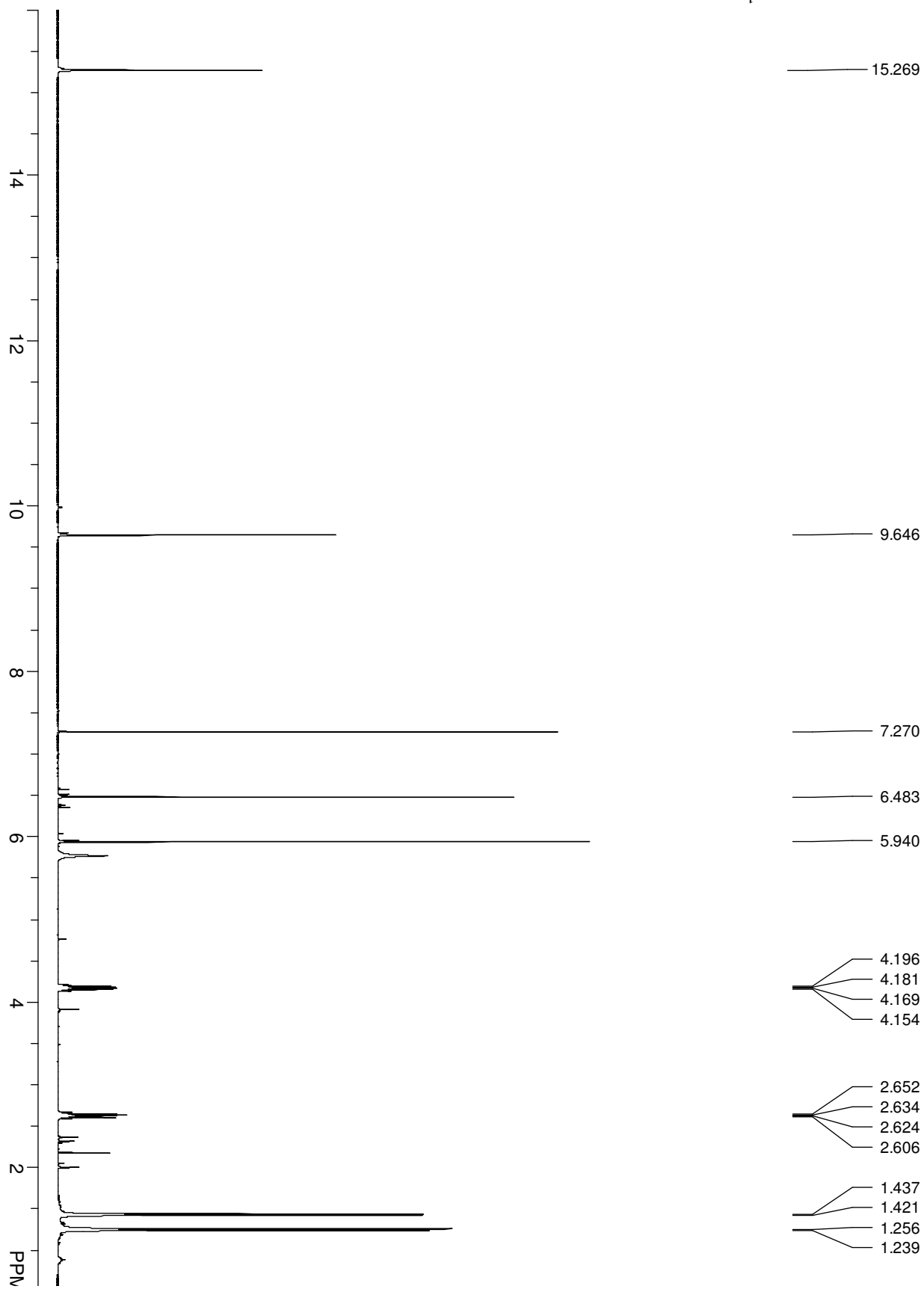
300 in the D. Malloch culture collection at the University of Toronto. Six 2-L Erlenmeyer flasks containing PDB (400 mL each) were each inoculated with one-cm² agar plugs of *Delitschia* sp. from stock cultures maintained on PDA. The flasks were incubated at rt on rotary shakers at 150 rpm for 20 days. The culture broth was filtered to separate the mycelia from the broth. The broth (2.4 L) was extracted with EtOAc (3 x 1 L), and the organic layer was dried with MgSO₄ and evaporated to yield 1.45 g of an antibacterial extract. The EtOAc extract was subjected to VLC on silica gel using a step gradient from 100% hexane to 100% CHCl₃ and finally to 50% MeOH in CHCl₃. The fraction eluting at 100% CHCl₃ (366 mg) was further purified by preparative reversed phase HPLC (C₁₈, 8μ particles, 2.14 x 25 cm, 10 mL/min, 20-100% MeOH in H₂O over 30 min) to afford talaroderxine A (**2**; t_R 31.8 min; 12.6 mg) and talaroderxine B (**3**; t_R 29.8 min; 18.2 mg). Talaroderxines A and B were identified by analysis of ¹H NMR, ¹³C NMR, MS, CD, and [α]_D data, and by comparison to literature values.⁵

Secalonic acid A. A culture of *Setophoma terrestris* (NRRL 25008), obtained as a colonist of an *Aspergillus flavus* sclerotium that was buried in field soil near Kilbourne, Illinois, was subjected to solid-substrate fermentation on rice (2x 200 g rice in 2.8-L Fernbach flasks). After 45 days, the cultures were extracted with 2 L of EtOAc. After evaporation of the solvent, the resulting extract (2.27 of 2.41 g) was subjected to VLC on silica gel, eluting sequentially with hexane-CH₂Cl₂-MeOH mixtures of increasing polarity. The fraction eluting with 15% MeOH in CH₂Cl₂ (47 mg) was subjected to reversed phase (C₁₈) HPLC using a linear gradient of 20-90% acetonitrile in 0.1% aqueous formic acid. A peak eluting at 26 min was identified as secalonic acid A (**4**; 5.0 mg) by comparison of its ¹H NMR, ¹³C NMR, and MS data with literature values.⁶

Cephalochromin. A culture of *Cosmospora* sp. (syn. *Acremonium butyri*; NRRL 28291), obtained as a colonist of the stromata of *Hypoxylon* sp. found on a fallen hardwood log, Picture Ridge Drive, Peoria, IL collected by H.D. Thiers in May 1996) was subjected to solid-substrate fermentation on rice (50 g rice in a 500-mL Erlenmeyer flask). After 45 days, the culture was extracted with 0.5 L of EtOAc. A portion of the resulting extract (approximately 50 mg of 1.57 g total) was placed directly onto a column of Sephadex LH-20 and eluted sequentially with 4:1 CH₂Cl₂-hexane (with which the column had been packed), 3:2 CH₂Cl₂-acetone, and 1:4 CH₂Cl₂-acetone; five fractions were collected. The second fraction (41 mg) was essentially a pure compound, as judged by ¹H and ¹³C NMR analysis. It had an orange color and accounted for most of the mass of the extract. This compound was identified as cephalochromin (**5**) by comparison of its NMR, MS, and optical rotation data with literature values.^{6,7}

Skyrin. A culture of *Geomyces pannorum* (NRRL 22978), obtained as a colonist of an *Aspergillus flavus* sclerotium buried in sandy field soil near Kilbourne, Illinois, was subjected to solid-substrate fermentation on rice (2 x 50 g rice in 500-mL Erlenmeyer flasks). After 45 days, the cultures were extracted with 0.5 L of EtOAc. After evaporation of the solvent, the resulting extract (409 mg) was placed directly onto a column of Sephadex LH-20 and eluted sequentially with 4:1 CH₂Cl₂-hexane (with which the column had been packed), 4:1 CH₂Cl₂-acetone, 3:2 CH₂Cl₂-acetone, and 1:4 CH₂Cl₂-acetone; 14 fractions were collected. The eighth fraction (82 mg), eluted with 3:2 CH₂Cl₂-acetone, was essentially a pure compound, as judged by ¹H and ¹³C NMR analysis. This metabolite was identified as *S*-(+)-skyrin (**6**) by comparison of its NMR, MS, and [α]_D data with literature values.⁸

Figure 3S. ^1H NMR spectrum of chaetochromin A (1) at 400 MHz in CDCl_3



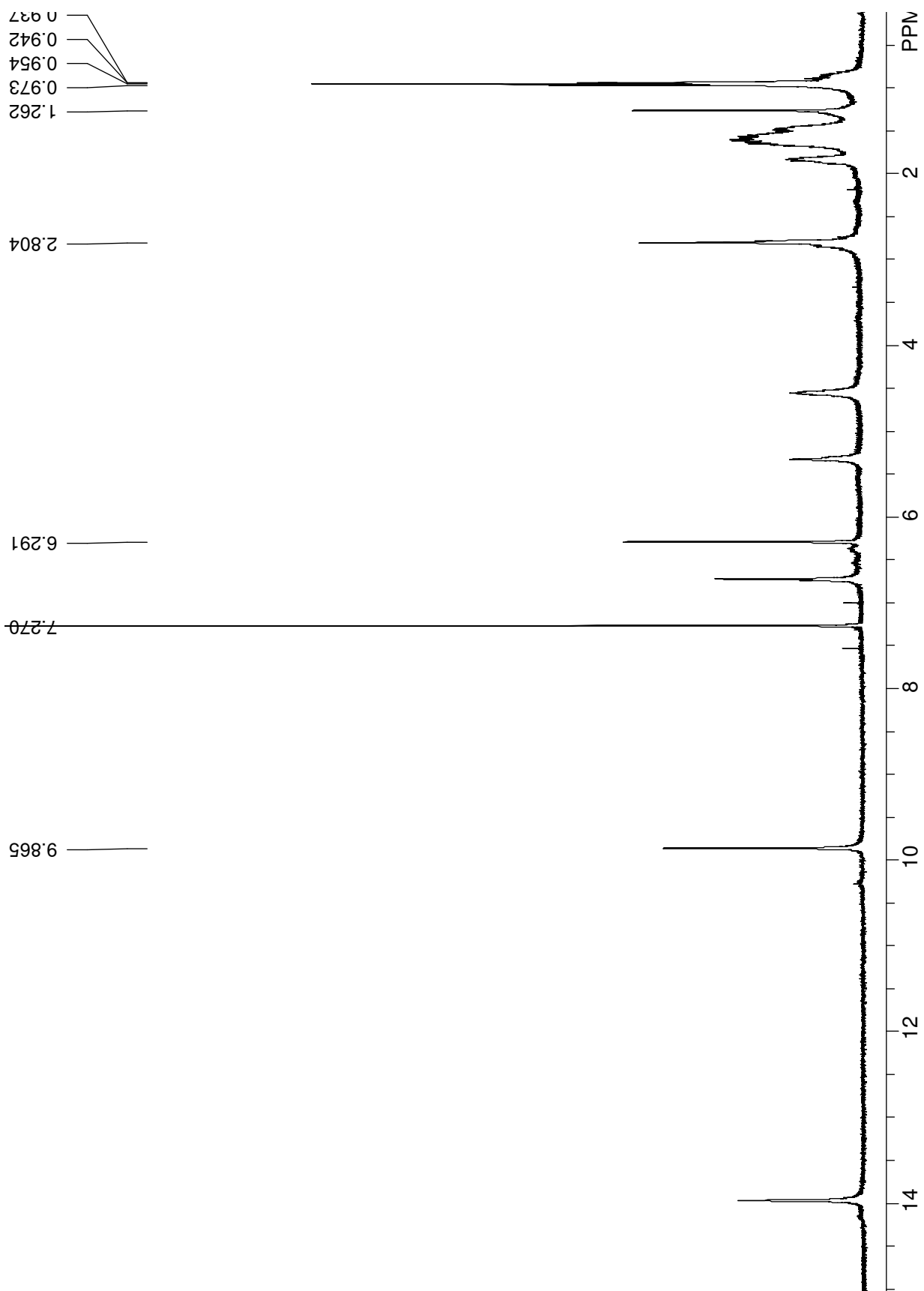
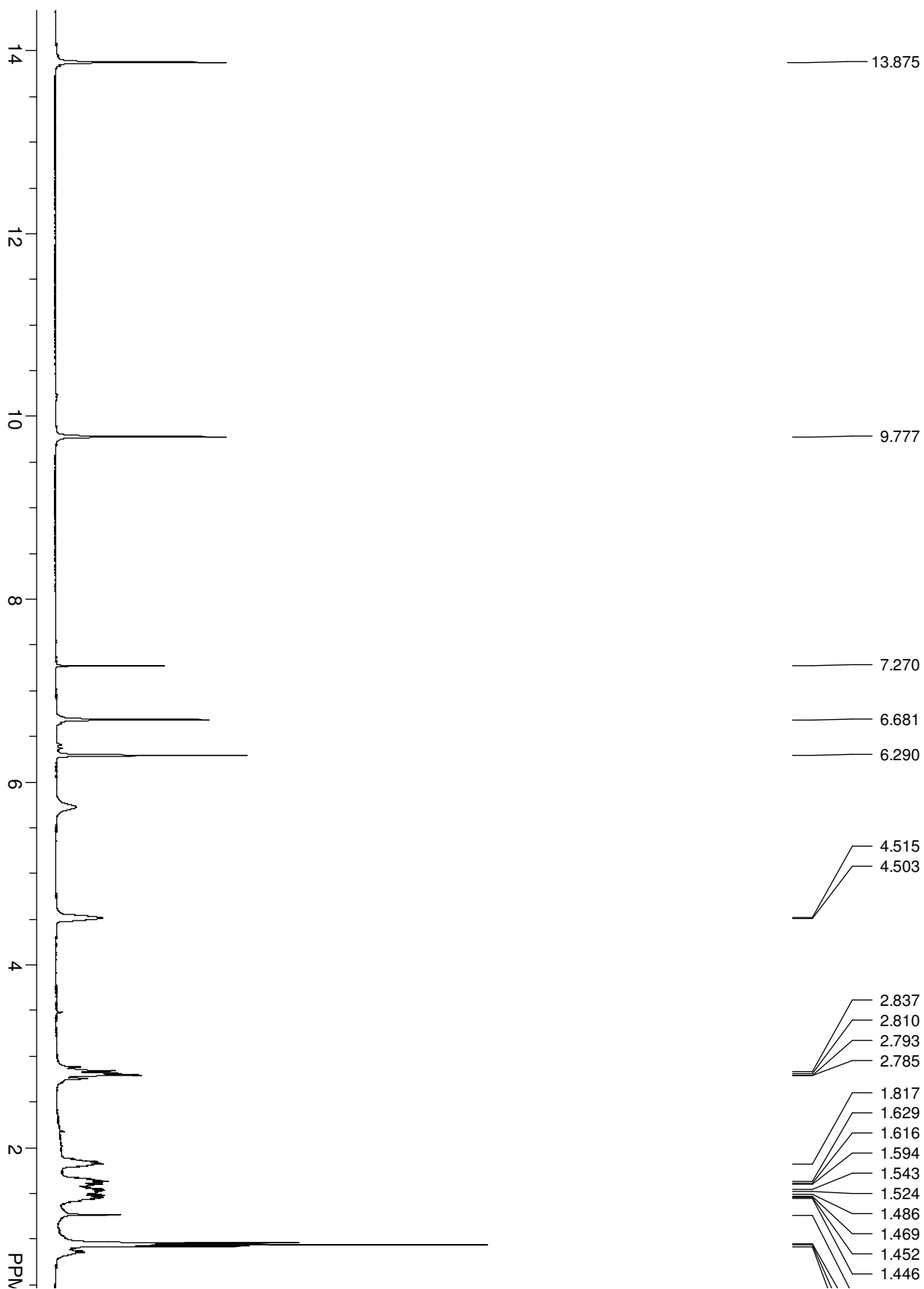


Figure 4S. ^1H NMR spectrum of talaroderxine A (**2**) at 400 MHz in CDCl_3

Figure S5. ^1H NMR spectrum of talaroderxine B (**3**) at 400 MHz in CDCl_3



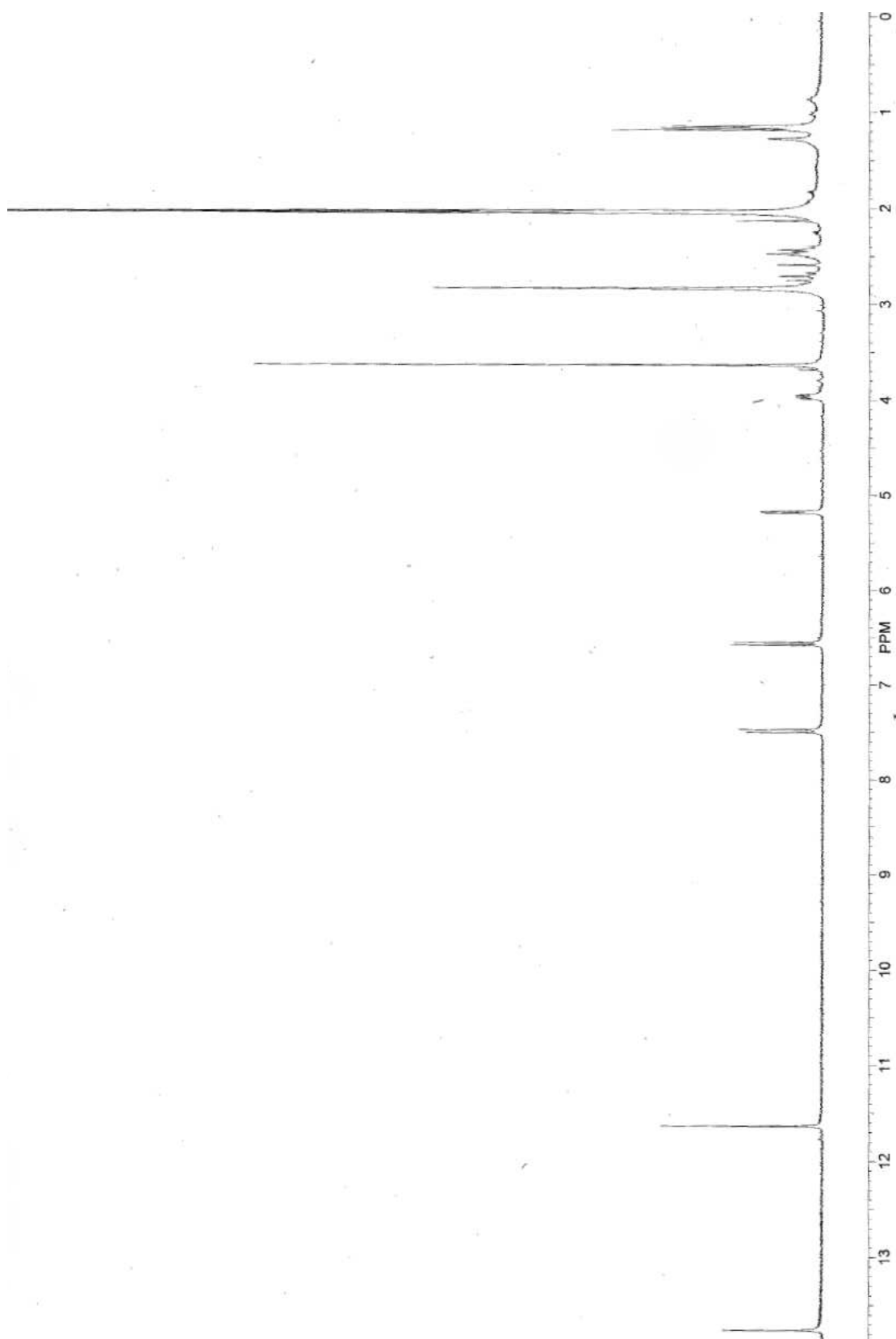
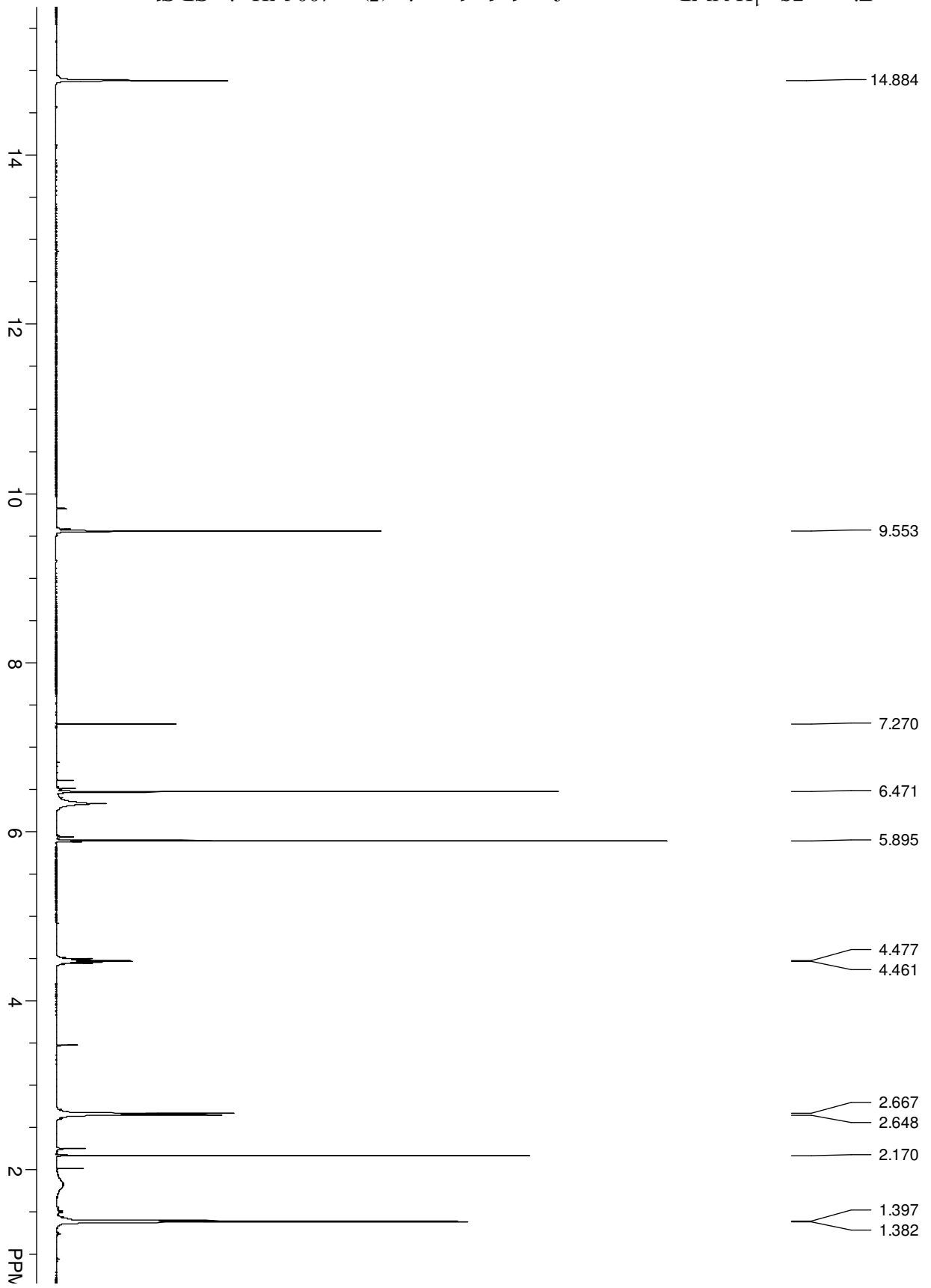


Figure 6S. ^1H NMR Spectrum of Secalonic Acid (4) at 400 MHz in CDCl_3

Figure 7S. ^1H NMR spectrum of cephalochromin (**5**) at 400 MHz in CDCl_3



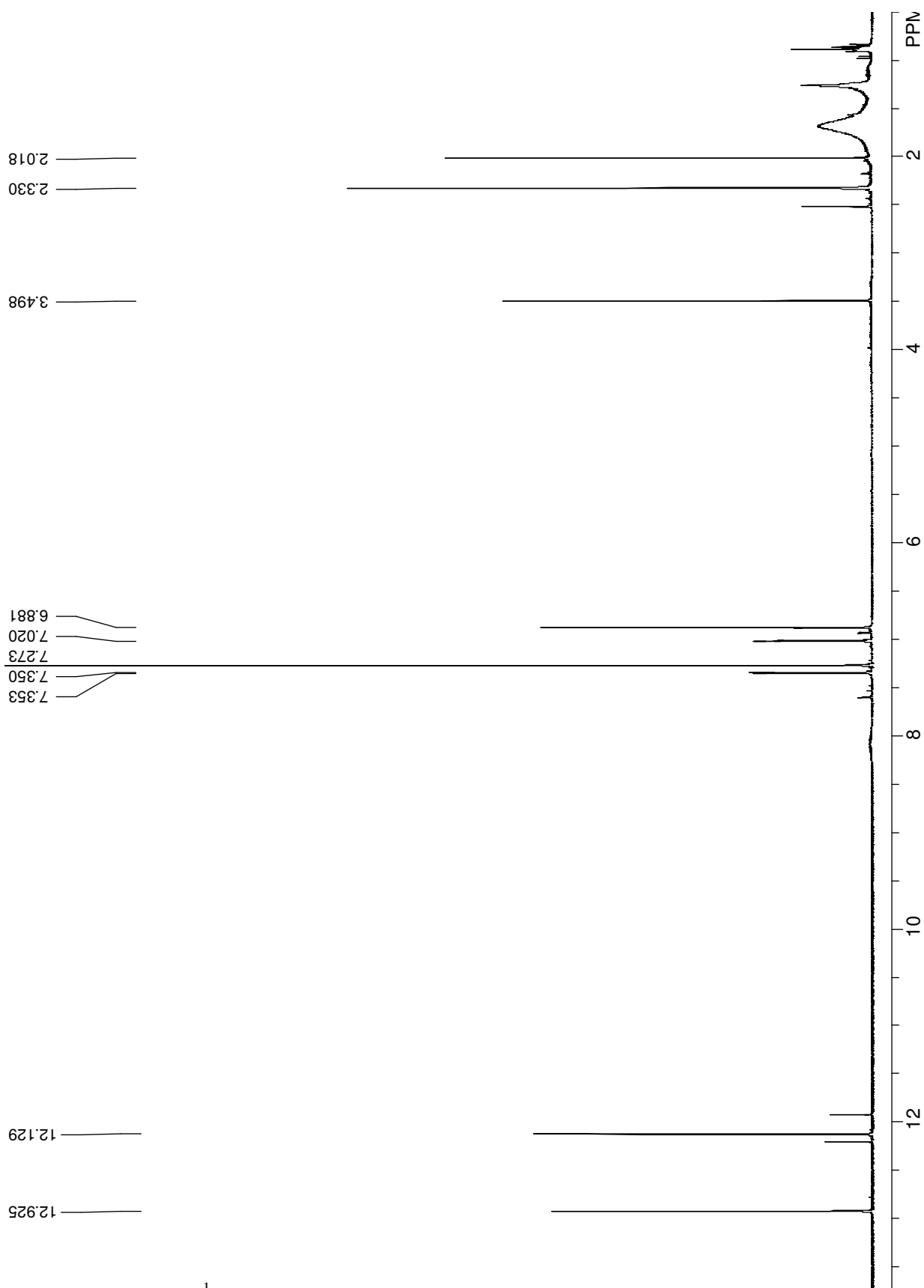


Figure 8S. ^1H NMR spectrum of skyrin (6) at 400 MHz in CDCl_3

The details of the BoNT/A structure setup were the same as in our previous work.⁹ Small molecule parameters were derived using GAFF forcefield.¹⁰ Charges were derived using AM1-BCC approach^{11,12} as implemented in our in-house software described previously.^{9,13,14} The parameters of all ligands are provided at the end of this document in mol2 format. For this work, we designed and parameterized a special reference molecule shown in Scheme 1. This molecule contains the 1,1'-binaphthyl core fused to cyclohexenone instead of pyrone.

All compounds, including reference compound, were docked into the BoNT/A structure using Autodock Vina.¹⁵ Nine poses were saved for each compound and visually inspected to find common binding poses. Skyrin was the only compound that differed substantially in its binding poses from all the other compounds. It is worth noting that the docking software breaks the axial symmetry of the compounds as it is not cognizant of the higher order chirality. Therefore, docking poses contain both stereoisomers of the compounds with respect to the chiral axis. Skyrin differs from the rest of the compounds in the pattern of carbonyl and hydroxyl groups around the ring. To get the complex of skyrin in the selected consensus pose corresponding to A stereoisomer, its coordinates were rebuilt from the corresponding complex of BoNT/A with the reference compound.

For accelerated sampling of the multiple hydroxyl groups of the compounds studied in this work, we derive the unphysical reference state for Thermodynamic Integration by collapsing all of hydrogen atoms of the reference compound (Scheme 1) onto the corresponding oxygen and carbon atoms. Note that the reference structure lacks asymmetric carbon atoms. The lack of rotatable OH groups in the unphysical reference structure helps boost rotations of the hydroxyl groups in all of the real compounds studied here. This is achieved with the help of recently developed Single Reference Thermodynamic Integration (SRTI) method augmented with Hamiltonian Replica Exchange (HREX).^{13,14}

For the HREX-SRTI simulations reported in this work, we used soft-core potentials with $p = 1$ (refer to GROMACS manual for notation used in soft-core potentials) and optimized parameters.⁹ Specifically, we used previously optimized¹⁴ values of the $\alpha = 0.4$. Furthermore, due to the presence of large number of hydroxyl groups in the compounds the H atoms of which are lacking the vdW parameters, we optimized the value of σ from its default value of 0.3 to 0.25 that gives better results. This is first time we attempted optimization of the σ parameter of the soft core potential within HREX-SRTI.

HREX-SRTI simulations involved running two independent simulations in water and solvated protein environment for each ligand. Hence the standard deviations provided were derived from two independent simulations in each leg of cycle and relative to the chaetochromin A. We employed 24 windows in both protein and water simulations. The production runs were 1 ns long in each window with a total of 1000 exchange steps. Each window was running on 8 cpu cores for approximately 48 hours. All the simulations were performed on DoD's Mana MHPCC supercomputer. Coordinates were saved every 1 ps. Unless otherwise stated, the details of MD simulations were the same as in our previous work.¹⁴ The HREX-SRTI trajectories for the physical state of the complexes were merged using our in-house tools prior to analysis. GROMOS-style clustering was done using *g_cluster* program of GROMACS with a RMSD cutoff of 0.05 nm.^{16,17} During clustering, the complex frames were fitted using Zn^{2+} , heavy atoms of the bound ligands and $C\alpha$ atoms of the rigid structural elements of the protein (residues 164-167; 184-187; 192-196; 217-233 and 260-266). Centroids of the clusters that had at least 10% of the total population were retained. The top clusters were used to generate the figures in the main text. Ligand-protein interactions were analyzed in 2D projections using MOE software.¹⁸

Figure 9S. The compound used to generate parameters of the single reference for HREX-SRTI simulations.

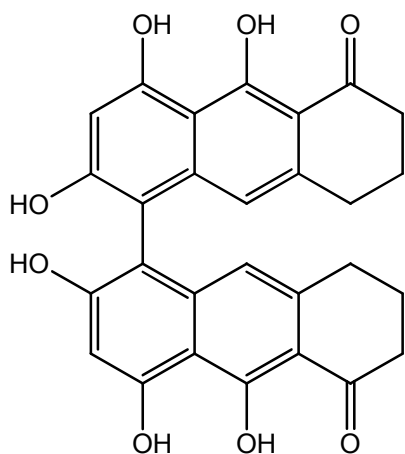
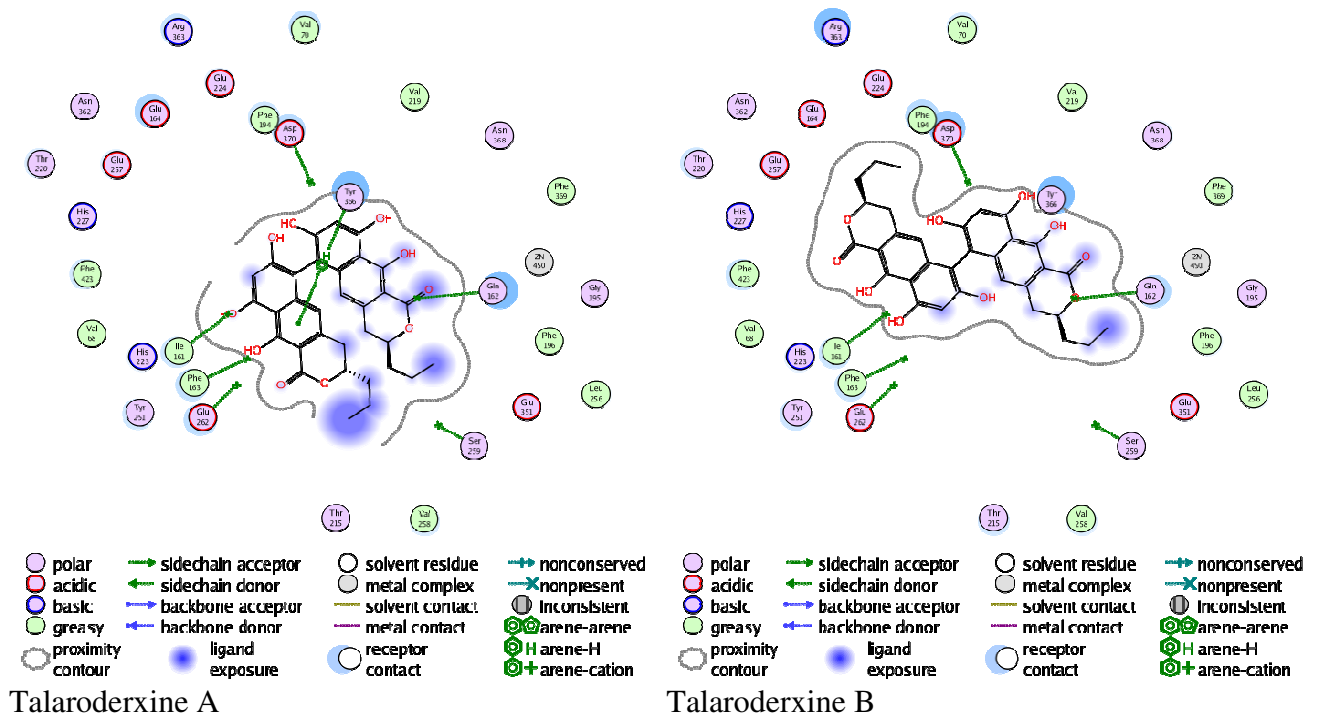
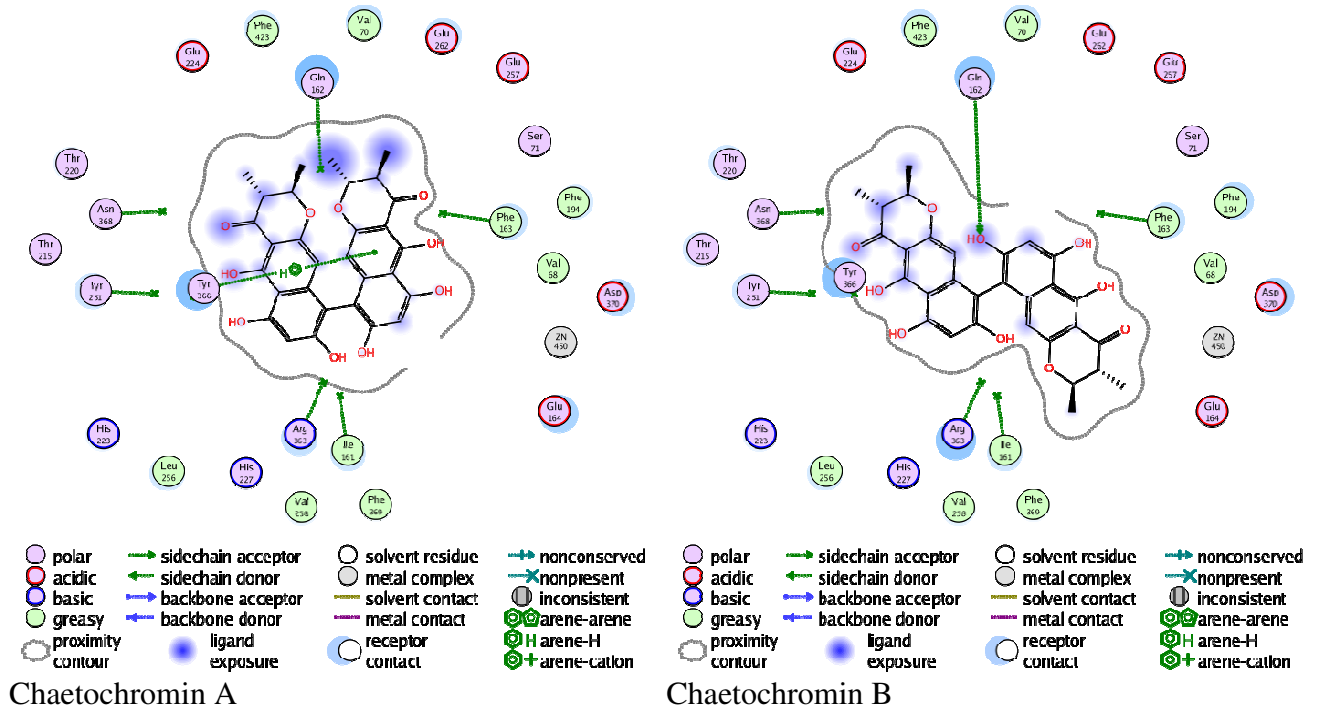


Figure 10S. 2-D depiction of the ligand-protein interactions of the A and B stereoisomers of selected compounds in corresponding consensus bound states with BoNT/A from Autodock Vina.



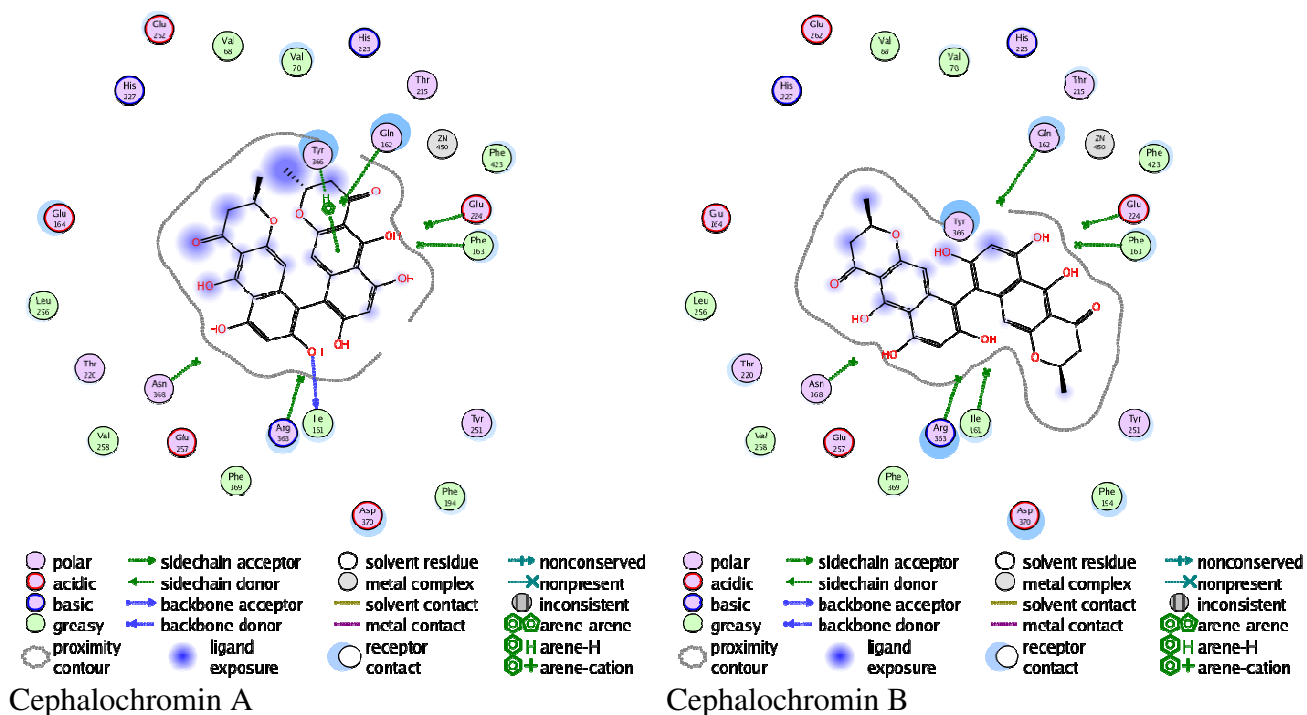
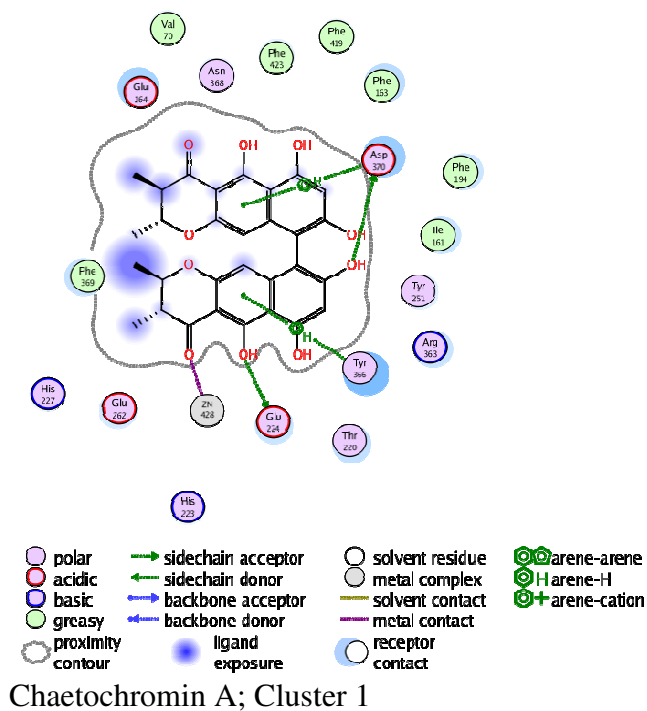
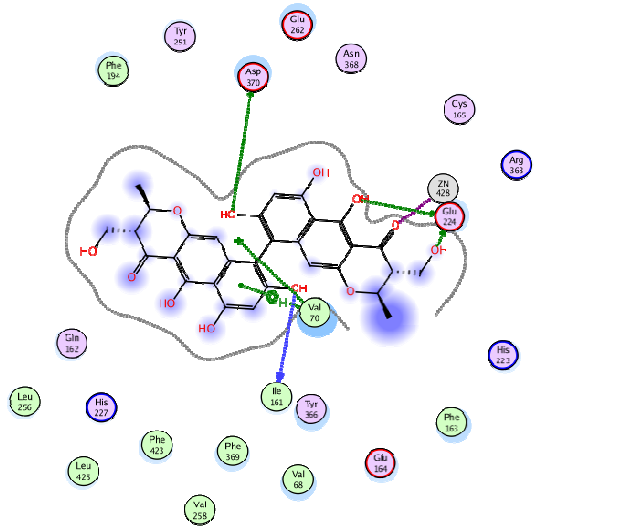


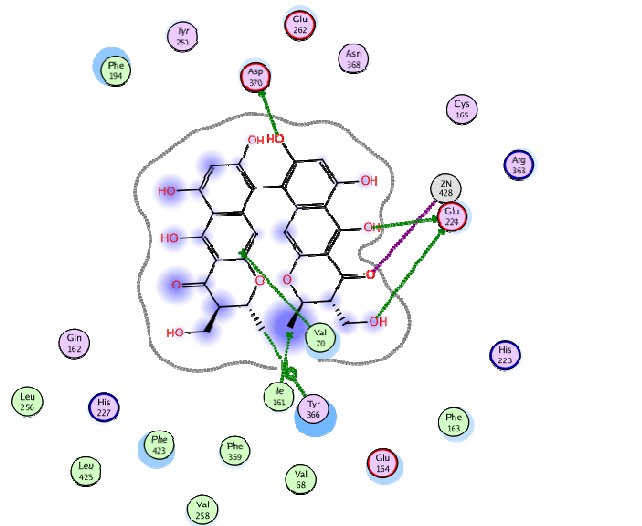
Figure 2S. 2-D depiction of the ligand-protein interactions of the A stereoisomers of studied compounds with BoNT/A in the most populated clusters from HREX-SRTI simulations.





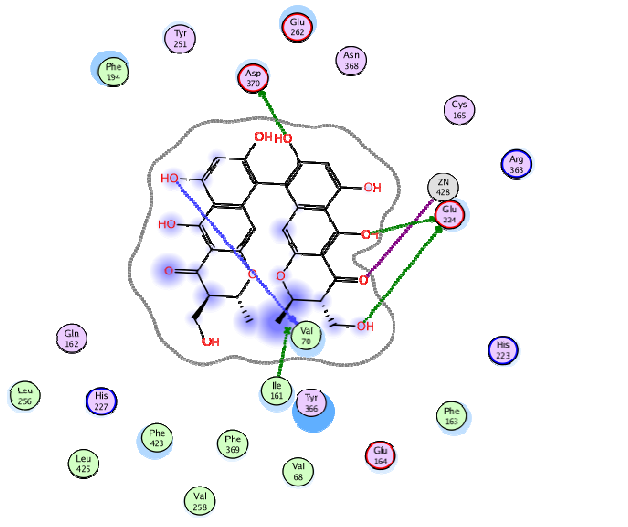
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- basic
- greasy
- proximity contour
- sidechain acceptor
- sidechain donor
- backbone acceptor
- backbone donor
- ligand exposure
- solvent residue
- metal complex
- solvent contact
- metal contact
- receptor contact
- nonconserved
- X nonpresent
- inconsistent
- arene-arene
- H arene-H
- + arene-cation

Hypothetical 7H A; Cluster 1



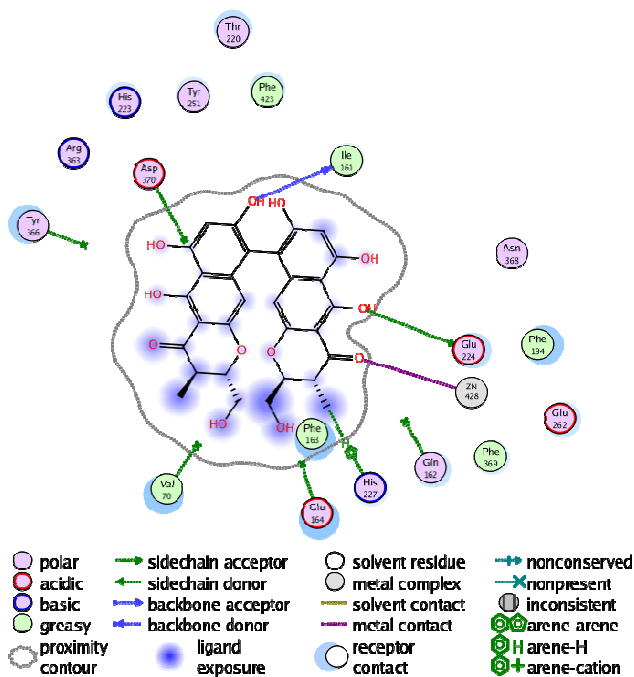
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- basic
- greasy
- proximity contour
- sidechain acceptor
- sidechain donor
- backbone acceptor
- backbone donor
- ligand exposure
- solvent residue
- metal complex
- solvent contact
- metal contact
- receptor contact
- nonconserved
- X nonpresent
- inconsistent
- arene-arene
- H arene-H
- + arene-cation

Hypothetical 7H A; Cluster 2

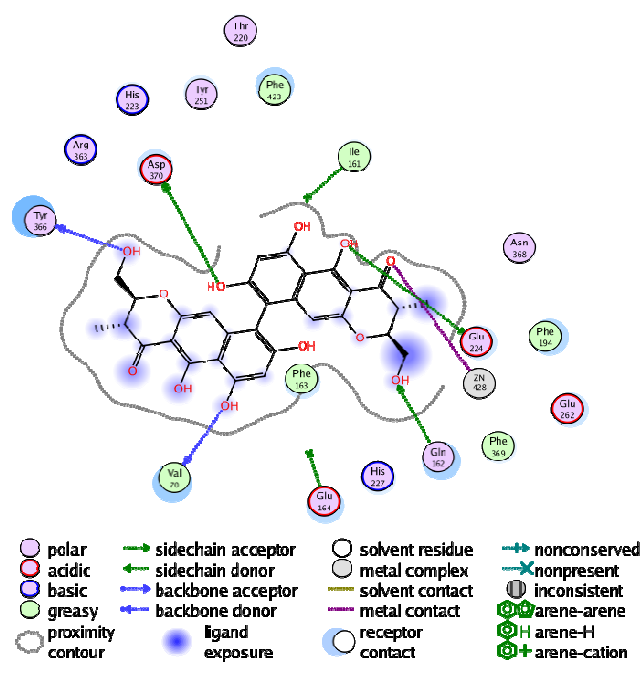


- polar
- acidic
- basic
- greasy
- proximity contour
- sidechain acceptor
- sidechain donor
- backbone acceptor
- backbone donor
- ligand exposure
- solvent residue
- metal complex
- solvent contact
- metal contact
- receptor contact
- nonconserved
- X nonpresent
- inconsistent
- arene-arene
- H arene-H
- + arene-cation

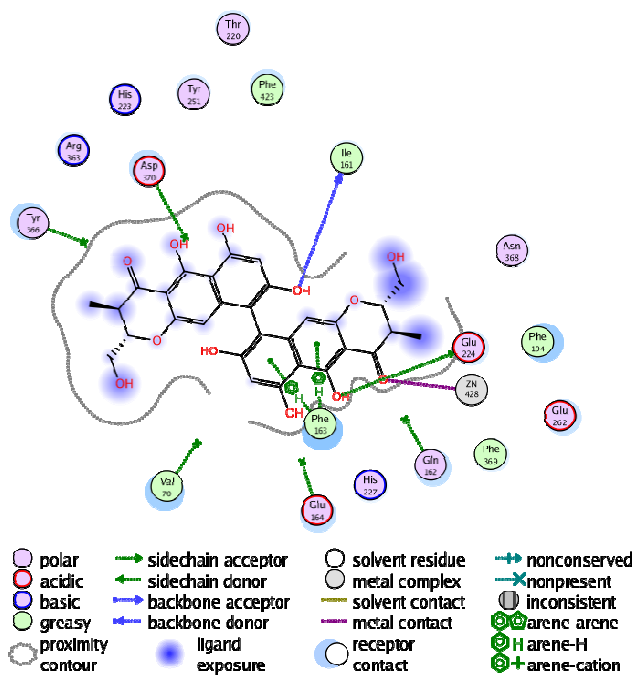
Hypothetical 7H A; Cluster 3



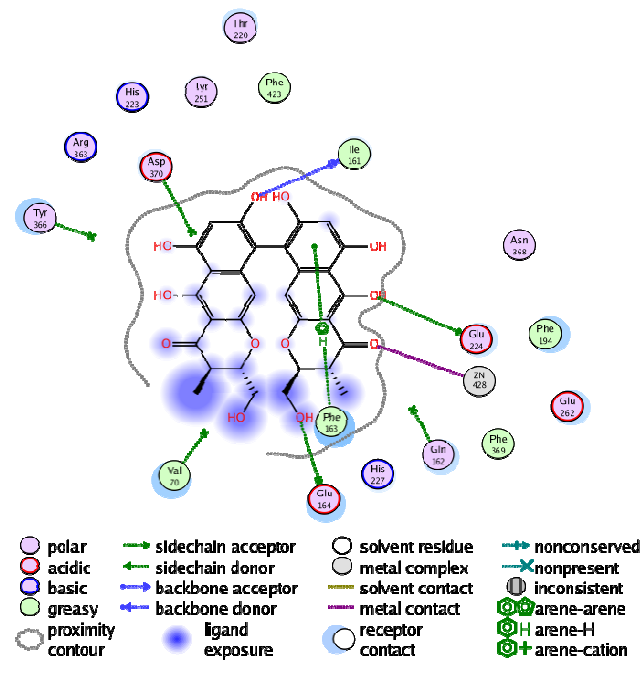
Hypothetical 8H A; Cluster 1



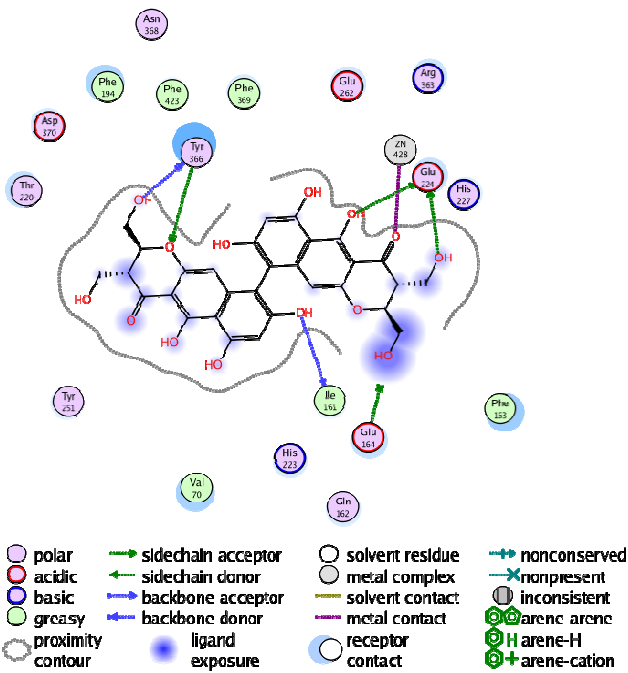
Hypothetical 8H A; Cluster 2



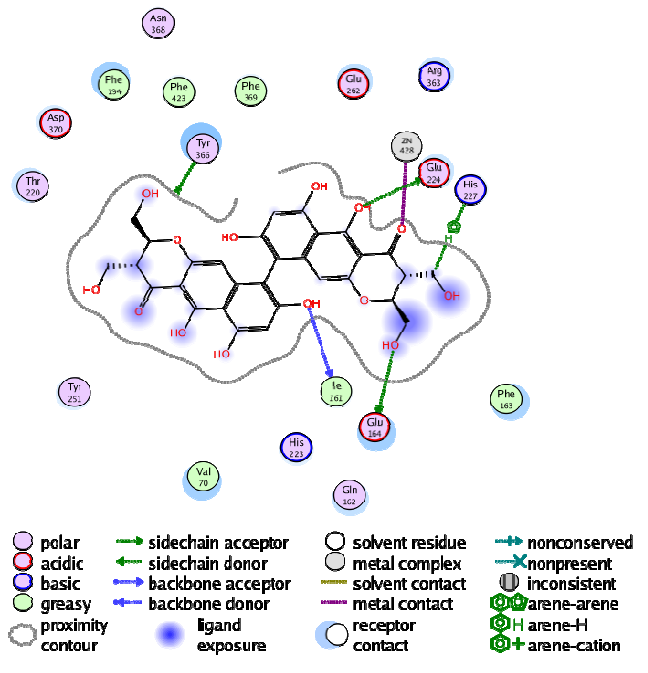
Hypothetical 8H A; Cluster 3



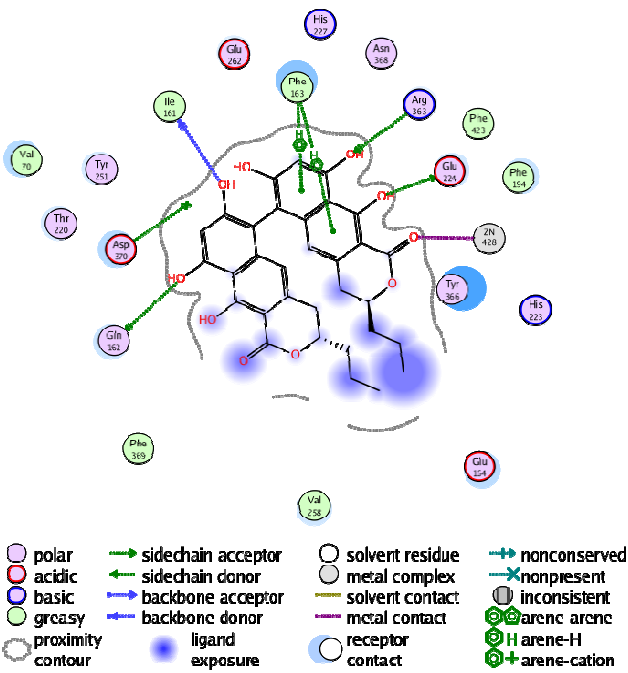
Hypothetical 8H A; Cluster 4



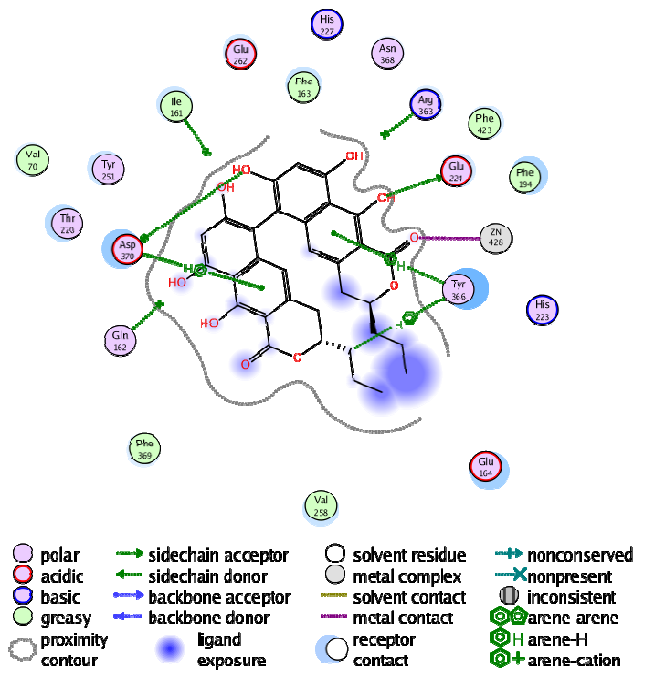
Hypothetical 9H A; Cluster 1



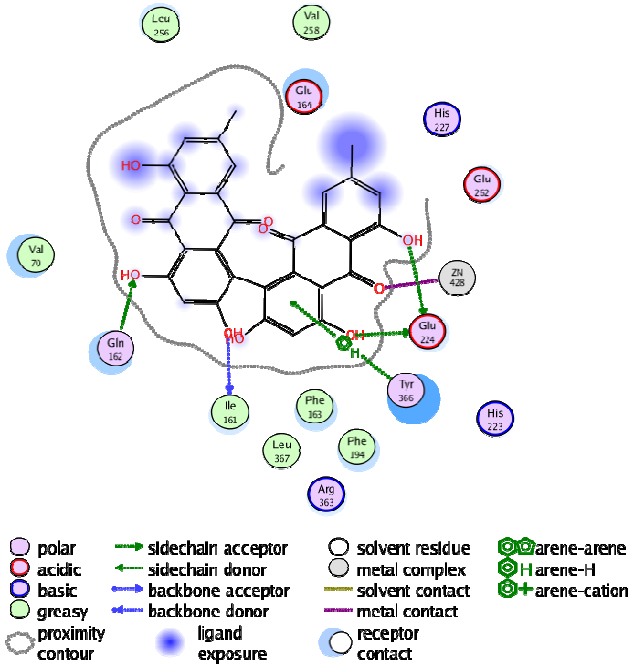
Hypothetical 9H A; Cluster 2



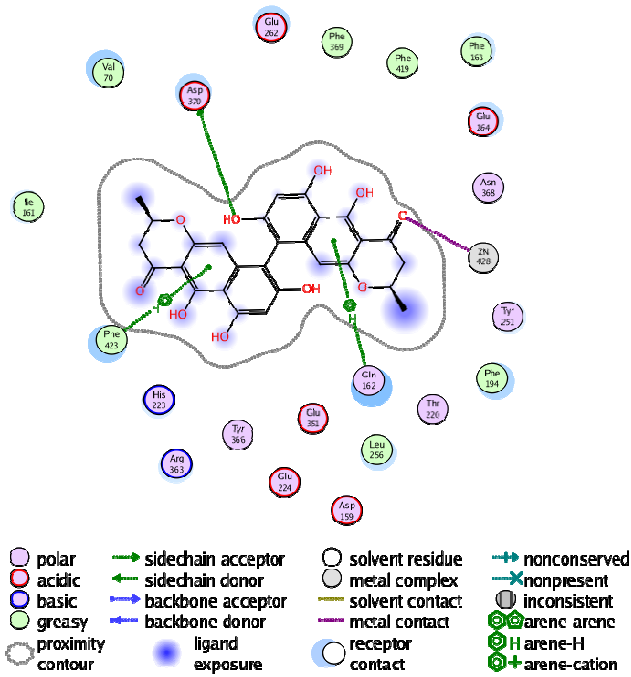
Talaroderxine A; Cluster 1



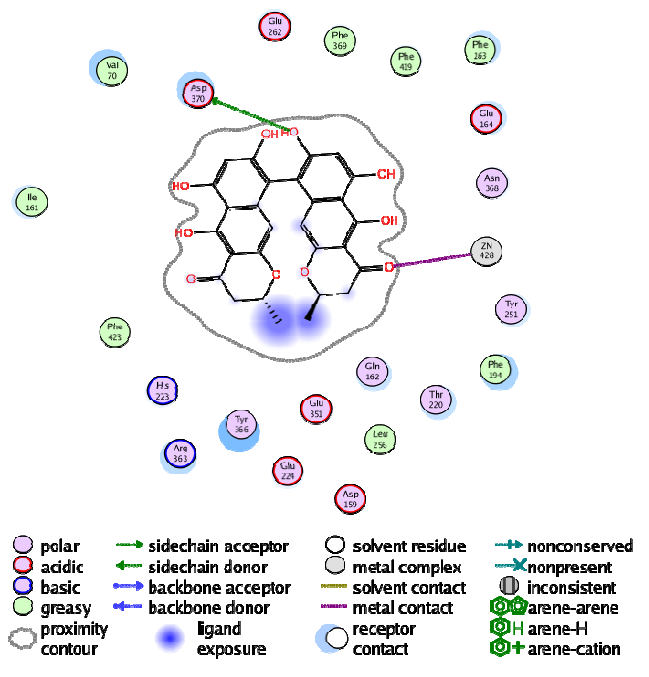
Talaroderxine A; Cluster 2



Skyrin; Cluster 1



Cephalochromin A; Cluster 1



Cephalochromin; Cluster 2

Figure 11S. The structure of the complex of BoNT/A with skyrin from HREX-SRTI simulations.

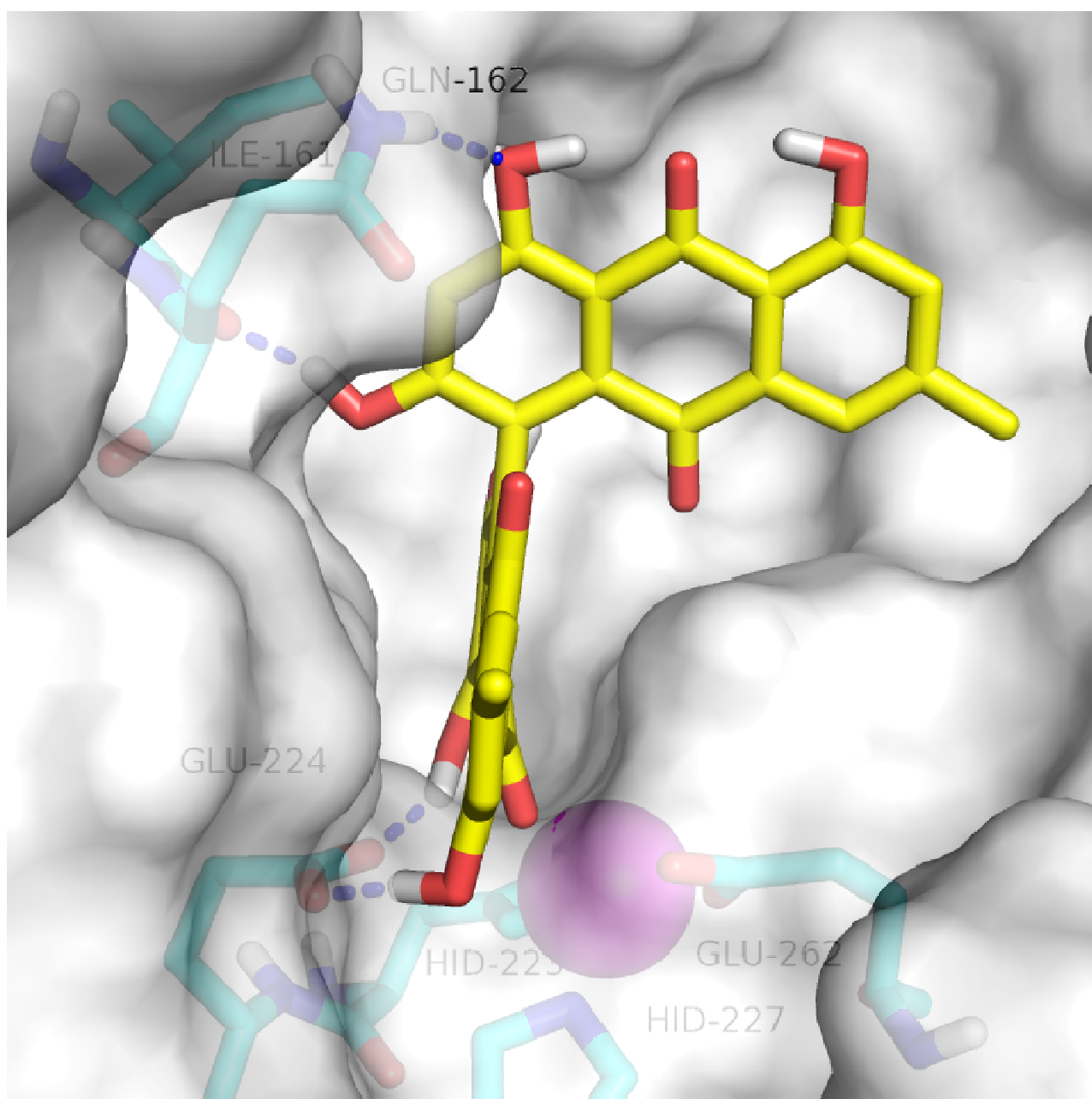


Table S1. Coordinates and Partial Charges for the Ligands *SRTI Reference Molecule*

lig_00000001.mol2

@<TRIPOS>MOLECULE

UNK

58 63 1 0 0

SMALL

No Charge or Current Charge

@<TRIPOS>ATOM

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2	C2	1.8410	0.0800	-0.0840	ca	1 UNK	-0.194605
3	C3	-0.4810	4.1010	-0.0840	ca	1 UNK	-0.338605
4	C4	0.4820	-0.1710	-4.0980	ca	1 UNK	-0.338605
5	C5	-1.3680	1.4210	-0.0980	ca	1 UNK	0.099595
6	C6	1.3690	0.0100	-1.4240	ca	1 UNK	0.099595
7	C7	-1.9770	2.3590	0.7740	ca	1 UNK	-0.149405
8	C8	1.9780	-0.9190	-2.3060	ca	1 UNK	-0.149405
9	C9	-0.3080	1.8160	-0.9630	cp	1 UNK	-0.104705
10	C10	0.3090	0.8490	-1.8720	cp	1 UNK	-0.104705
11	C11	-3.4950	0.6000	1.6290	ca	1 UNK	-0.308005
12	C12	3.4950	-1.6640	-0.4970	ca	1 UNK	-0.308005
13	C13	-2.8630	-0.3220	0.7400	ca	1 UNK	0.022095
14	C14	2.8640	-0.7190	0.3670	ca	1 UNK	0.022095
15	C15	-1.5100	3.7080	0.7570	ca	1 UNK	0.225895
16	C16	1.5100	-0.9860	-3.6540	ca	1 UNK	0.225895
17	C17	0.1120	3.1470	-0.9450	ca	1 UNK	0.210595
18	C18	-0.1110	0.7480	-3.1990	ca	1 UNK	0.210595
19	C19	-3.0420	1.9250	1.6370	ca	1 UNK	0.275695
20	C20	3.0420	-1.7530	-1.8190	ca	1 UNK	0.275695
21	C21	-4.5810	0.1460	2.5020	c	1 UNK	0.602295
22	C22	4.5810	-2.5070	0.0100	c	1 UNK	0.602295
23	C23	-3.3140	-1.7410	0.6690	c3	1 UNK	-0.047805
24	C24	3.3140	-0.5600	1.7800	c3	1 UNK	-0.047805
25	C25	-5.0510	-1.2770	2.3750	c3	1 UNK	-0.171805
26	C26	5.0500	-2.2920	1.4230	c3	1 UNK	-0.171805
27	C27	-3.9420	-2.2130	1.9610	c3	1 UNK	-0.072805
28	C28	3.9420	-1.8200	2.3310	c3	1 UNK	-0.072805
29	O1	-5.1380	0.8930	3.3280	o	1 UNK	-0.593305
30	O2	5.1380	-3.3780	-0.6840	o	1 UNK	-0.593305
31	O3	-2.1220	4.5860	1.6110	oh	1 UNK	-0.457205
32	O4	2.1230	-1.8930	-4.4760	oh	1 UNK	-0.457205
33	O5	1.1340	3.5120	-1.7860	oh	1 UNK	-0.479505
34	O6	-1.1330	1.5650	-3.6150	oh	1 UNK	-0.479505
35	O7	-3.5350	2.9150	2.4270	oh	1 UNK	-0.477105
36	O8	3.5350	-2.6040	-2.7580	oh	1 UNK	-0.477105
37	H1	-1.3590	-0.6430	-0.7620	ha	1 UNK	0.149095
38	H2	1.3600	0.8010	0.5950	ha	1 UNK	0.149095
39	H3	-0.1280	5.1410	-0.0940	ha	1 UNK	0.138695
40	H4	0.1280	-0.2250	-5.1360	ha	1 UNK	0.138695
41	H5	-4.0600	-1.8310	-0.1680	hc	1 UNK	0.059545
42	H6	-2.4450	-2.4040	0.4120	hc	1 UNK	0.059545
43	H7	4.0620	0.2810	1.8180	hc	1 UNK	0.059545
44	H8	2.4460	-0.2610	2.4260	hc	1 UNK	0.059545
45	H9	-5.4930	-1.6060	3.3520	hc	1 UNK	0.074020
46	H10	-5.8740	-1.2920	1.6100	hc	1 UNK	0.074020
47	H11	5.4920	-3.2470	1.8120	hc	1 UNK	0.074020
48	H12	5.8740	-1.5280	1.3910	hc	1 UNK	0.074020
49	H13	-4.3520	-3.2470	1.8250	hc	1 UNK	0.051545
50	H14	-3.1640	-2.2630	2.7670	hc	1 UNK	0.051545
51	H15	4.3520	-1.6200	3.3550	hc	1 UNK	0.051545
52	H16	3.1630	-2.6210	2.4310	hc	1 UNK	0.051545
53	H17	-1.7190	5.4600	1.4870	ho	1 UNK	0.417595
54	H18	1.7190	-1.8240	-5.3560	ho	1 UNK	0.417595
55	H19	1.2920	4.4610	-1.6740	ho	1 UNK	0.419595
56	H20	-1.2910	1.3940	-4.5560	ho	1 UNK	0.419595
57	H21	-4.2550	2.5880	2.9940	ho	1 UNK	0.463495

58 H22 4.2550 -3.1500 -2.3960 ho 1 UNK 0.463495
@<TRIPOS>BOND

1	1	5	ar
2	1	13	ar
3	2	6	ar
4	2	14	ar
5	3	15	ar
6	3	17	ar
7	4	16	ar
8	4	18	ar
9	5	7	ar
10	5	9	ar
11	6	8	ar
12	6	10	ar
13	7	15	ar
14	7	19	ar
15	8	16	ar
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22	11	21	l
23	12	14	ar
24	12	20	ar
25	12	22	l
26	13	23	l
27	14	24	l
28	15	31	l
29	16	32	l
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31	18	34	l
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38	23	27	l
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56	28	51	l
57	28	52	l
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61	34	56	l
62	35	57	l
63	36	58	l

@<TRIPOS>SUBSTRUCTURE

1 UNK

1 TEMP

0 *****

0 ROOT

Chaetochromin A (1)

lig_00000002.mol2

@<TRIPOS>MOLECULE

UNK

66 71 1 0 0

SMALL

No Charge or Current Charge

@<TRIPOS>ATOM

1	C1	-1.8760	0.4500	-0.4970	ca	1	UNK	-0.224000
2	C2	1.8740	0.4500	-0.4970	ca	1	UNK	-0.224000
3	C3	-0.4110	4.4350	-0.4970	ca	1	UNK	-0.344100
4	C4	0.4100	0.0640	-4.4630	ca	1	UNK	-0.344100
5	C5	-1.3810	1.7800	-0.5130	ca	1	UNK	0.123900
6	C6	1.3790	0.3370	-1.8220	ca	1	UNK	0.123900
7	C7	-1.9720	2.7430	0.3480	ca	1	UNK	-0.168900
8	C8	1.9710	-0.6130	-2.6970	ca	1	UNK	-0.168900
9	C9	-0.3000	2.1420	-1.3680	cp	1	UNK	-0.109200
10	C10	0.2980	1.1530	-2.2660	cp	1	UNK	-0.109200
11	C11	-3.5700	1.0500	1.1860	ca	1	UNK	-0.364000
12	C12	3.5690	-1.2820	-0.9310	ca	1	UNK	-0.364000
13	C13	-2.9300	0.1070	0.3240	ca	1	UNK	0.210600
14	C14	2.9290	-0.3330	-0.0760	ca	1	UNK	0.210600
15	C15	-1.4580	4.0760	0.3350	ca	1	UNK	0.232200
16	C16	1.4580	-0.7290	-4.0250	ca	1	UNK	0.232200
17	C17	0.1590	3.4600	-1.3510	ca	1	UNK	0.217700
18	C18	-0.1610	1.0080	-3.5750	ca	1	UNK	0.217700
19	C19	-3.0650	2.3570	1.1950	ca	1	UNK	0.298600
20	C20	3.0650	-1.4170	-2.2310	ca	1	UNK	0.298600
21	C21	-4.7210	0.6340	1.9800	c	1	UNK	0.606200
22	C22	4.7220	-2.0300	-0.4400	c	1	UNK	0.606200
23	C23	-5.2220	-0.7740	1.7440	c3	1	UNK	-0.184600
24	C24	5.2230	-1.6580	0.9390	c3	1	UNK	-0.184600
25	C25	-4.0610	-1.6970	1.3850	c3	1	UNK	0.149100
26	C26	4.0610	-1.2130	1.8220	c3	1	UNK	0.149100
27	C27	-5.9840	-1.2990	2.9400	c3	1	UNK	-0.086000
28	C28	5.9860	-2.7970	1.5770	c3	1	UNK	-0.086000
29	C29	-4.5200	-3.0900	0.9970	c3	1	UNK	-0.095300
30	C30	4.5200	-0.6910	3.1710	c3	1	UNK	-0.095300
31	O1	-5.3040	1.3870	2.7800	o	1	UNK	-0.582500
32	O2	5.3050	-2.8990	-1.1120	o	1	UNK	-0.582500
33	O3	-3.3560	-1.2010	0.2300	os	1	UNK	-0.325400
34	O4	3.3550	-0.1120	1.2170	os	1	UNK	-0.325400
35	O5	-2.0510	4.9740	1.1820	oh	1	UNK	-0.460000
36	O6	2.0500	-1.6580	-4.8370	oh	1	UNK	-0.460000
37	O7	1.1930	3.7920	-2.1890	oh	1	UNK	-0.478400
38	O8	-1.1950	1.8100	-3.9870	oh	1	UNK	-0.478400
39	O9	-3.5470	3.3600	1.9720	oh	1	UNK	-0.470800
40	O10	3.5470	-2.2880	-3.1540	oh	1	UNK	-0.470800
41	H1	-1.4150	-0.3070	-1.1480	ha	1	UNK	0.164800
42	H2	1.4130	1.1710	0.1940	ha	1	UNK	0.164800
43	H3	-0.0220	5.4620	-0.5040	ha	1	UNK	0.138800
44	H4	0.0210	-0.0290	-5.4860	ha	1	UNK	0.138800
45	H5	-5.9200	-0.7140	0.8580	hc	1	UNK	0.090800
46	H6	5.9200	-0.7820	0.7920	hc	1	UNK	0.090800
47	H7	-3.3320	-1.7460	2.2420	h1	1	UNK	0.057200
48	H8	3.3330	-2.0620	1.9540	h1	1	UNK	0.057200
49	H9	-6.5200	-2.2420	2.6730	hc	1	UNK	0.048367
50	H10	-6.7320	-0.5400	3.2780	hc	1	UNK	0.048367
51	H11	-5.2900	-1.5090	3.7890	hc	1	UNK	0.048367
52	H12	6.5220	-2.4390	2.4890	hc	1	UNK	0.048367
53	H13	6.7350	-3.2060	0.8540	hc	1	UNK	0.048367
54	H14	5.2930	-3.6220	1.8680	hc	1	UNK	0.048367
55	H15	-3.6670	-3.6600	0.5560	hc	1	UNK	0.052367
56	H16	-5.3400	-3.0380	0.2410	hc	1	UNK	0.052367
57	H17	-4.8880	-3.6230	1.9050	hc	1	UNK	0.052367

58	H18	3.6660	-0.1980	3.6960	hc	1	UNK	0.052367
59	H19	5.3390	0.0570	3.0460	hc	1	UNK	0.052367
60	H20	4.8890	-1.5430	3.7890	hc	1	UNK	0.052367
61	H21	-1.6140	5.8320	1.0670	ho	1	UNK	0.418600
62	H22	1.6140	-1.6270	-5.7020	ho	1	UNK	0.418600
63	H23	1.3980	4.7300	-2.0590	ho	1	UNK	0.419900
64	H24	-1.3990	1.5890	-4.9090	ho	1	UNK	0.419900
65	H25	-4.2860	3.0490	2.5250	ho	1	UNK	0.462600
66	H26	4.2860	-2.8070	-2.7910	ho	1	UNK	0.462600

@<TRIPOS>BOND

1	1	5	ar
2	1	13	ar
3	2	6	ar
4	2	14	ar
5	3	15	ar
6	3	17	ar
7	4	16	ar
8	4	18	ar
9	5	7	ar
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21	11	19	ar
22	11	21	l
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24	12	20	ar
25	12	22	l
26	13	33	l
27	14	34	l
28	15	35	l
29	16	36	l
30	17	37	l
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36	22	24	l
37	22	32	2
38	23	25	l
39	23	27	l
40	24	26	l
41	24	28	l
42	25	29	l
43	25	33	l
44	26	30	l
45	26	34	l
46	1	41	l
47	2	42	l
48	3	43	l
49	4	44	l
50	23	45	l
51	24	46	l
52	25	47	l
53	26	48	l
54	27	49	l
55	27	50	l
56	27	51	l
57	28	52	l
58	28	53	l

59	28	54	1
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61	29	56	1
62	29	57	1
63	30	58	1
64	30	59	1
65	30	60	1
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68	37	63	1
69	38	64	1
70	39	65	1
71	40	66	1

@<TRIPOS>SUBSTRUCTURE

1 UNK 1 TEMP 0 **** * 0 ROOT

Hypothetical Compound 7H

lig_00000003.mol2

@<TRIPOS>MOLECULE

UNK

68 73 1 0 0

SMALL

No Charge or Current Charge

@<TRIPOS>ATOM

1	C1	-1.8570	0.4990	-0.5570	ca	1	UNK	-0.224209
2	C2	1.8810	0.4990	-0.5570	ca	1	UNK	-0.224209
3	C3	-0.4070	4.4900	-0.5570	ca	1	UNK	-0.343940
4	C4	0.4280	0.1330	-4.5310	ca	1	UNK	-0.343940
5	C5	-1.3670	1.8320	-0.5730	ca	1	UNK	0.125997
6	C6	1.3900	0.3920	-1.8850	ca	1	UNK	0.125997
7	C7	-1.9610	2.7930	0.2890	ca	1	UNK	-0.170414
8	C8	1.9820	-0.5560	-2.7630	ca	1	UNK	-0.170414
9	C9	-0.2880	2.1980	-1.4290	cp	1	UNK	-0.110345
10	C10	0.3120	1.2110	-2.3280	cp	1	UNK	-0.110345
11	C11	-3.5500	1.0930	1.1280	ca	1	UNK	-0.365860
12	C12	3.5720	-1.2330	-0.9950	ca	1	UNK	-0.365860
13	C13	-2.9090	0.1520	0.2640	ca	1	UNK	0.212805
14	C14	2.9340	-0.2860	-0.1350	ca	1	UNK	0.212805
15	C15	-1.4530	4.1280	0.2750	ca	1	UNK	0.233384
16	C16	1.4730	-0.6640	-4.0940	ca	1	UNK	0.233384
17	C17	0.1660	3.5170	-1.4110	ca	1	UNK	0.219882
18	C18	-0.1430	1.0730	-3.6400	ca	1	UNK	0.219882
19	C19	-3.0520	2.4030	1.1360	ca	1	UNK	0.302207
20	C20	3.0690	-1.3670	-2.2950	ca	1	UNK	0.302207
21	C21	-4.6930	0.6680	1.9230	c	1	UNK	0.610925
22	C22	4.7100	-1.9920	-0.4970	c	1	UNK	0.610925
23	C23	-5.1850	-0.7440	1.7020	c3	1	UNK	-0.230218
24	C24	5.2690	-1.5410	0.8340	c3	1	UNK	-0.230218
25	C25	-4.0290	-1.6630	1.3260	c3	1	UNK	0.154626
26	C26	4.1240	-1.1350	1.7560	c3	1	UNK	0.154626
27	C27	-4.4830	-3.0540	0.9230	c3	1	UNK	-0.100194
28	C28	4.6000	-0.5720	3.0810	c3	1	UNK	-0.100194
29	C29	-5.9200	-1.2700	2.9370	c3	1	UNK	0.148612
30	C30	6.1550	-2.6180	1.4580	c3	1	UNK	0.148612
31	O1	-5.2810	1.4130	2.7300	o	1	UNK	-0.599161
32	O2	5.2260	-2.9460	-1.1080	o	1	UNK	-0.599161
33	O3	-3.3330	-1.1570	0.1690	os	1	UNK	-0.324790
34	O4	3.3490	-0.0730	1.1610	os	1	UNK	-0.324790
35	O5	-2.0490	5.0220	1.1220	oh	1	UNK	-0.460210
36	O6	2.0690	-1.5900	-4.9070	oh	1	UNK	-0.460210
37	O7	1.2000	3.8530	-2.2500	oh	1	UNK	-0.477929
38	O8	-1.1750	1.8790	-4.0510	oh	1	UNK	-0.477929
39	O9	-3.5380	3.4020	1.9140	oh	1	UNK	-0.468310
40	O10	3.5530	-2.2400	-3.2150	oh	1	UNK	-0.468310
41	O11	-7.1830	-0.6640	3.0880	oh	1	UNK	-0.601217

42	O12	5.4580	-3.8200	1.7020	oh	1	UNK	-0.601217
43	H1	-1.3940	-0.2560	-1.2100	ha	1	UNK	0.165456
44	H2	1.4170	1.2170	0.1350	ha	1	UNK	0.165456
45	H3	-0.0220	5.5190	-0.5650	ha	1	UNK	0.139701
46	H4	0.0430	0.0460	-5.5550	ha	1	UNK	0.139701
47	H5	-5.9170	-0.6980	0.8440	hc	1	UNK	0.101850
48	H6	5.9100	-0.6360	0.6310	hc	1	UNK	0.101850
49	H7	-3.2920	-1.7180	2.1760	h1	1	UNK	0.066687
50	H8	3.4430	-2.0170	1.9220	h1	1	UNK	0.066687
51	H9	-3.6760	-3.5570	0.3350	hc	1	UNK	0.054510
52	H10	-5.4020	-3.0040	0.2930	hc	1	UNK	0.054510
53	H11	-4.6950	-3.6580	1.8370	hc	1	UNK	0.054510
54	H12	3.7620	-0.0300	3.5830	hc	1	UNK	0.054510
55	H13	5.4430	0.1430	2.9290	hc	1	UNK	0.054510
56	H14	4.9370	-1.4090	3.7370	hc	1	UNK	0.054510
57	H15	-5.2910	-1.1090	3.8520	h1	1	UNK	0.054479
58	H16	-6.1450	-2.3650	2.8160	h1	1	UNK	0.054479
59	H17	6.5180	-2.2840	2.4680	h1	1	UNK	0.054479
60	H18	7.0320	-2.8040	0.7810	h1	1	UNK	0.054479
61	H19	-1.6170	5.8830	1.0070	ho	1	UNK	0.419798
62	H20	1.6330	-1.5580	-5.7730	ho	1	UNK	0.419798
63	H21	1.3980	4.7930	-2.1230	ho	1	UNK	0.420858
64	H22	-1.3790	1.6620	-4.9730	ho	1	UNK	0.420858
65	H23	-4.2790	3.0890	2.4630	ho	1	UNK	0.461905
66	H24	4.2800	-2.7690	-2.8420	ho	1	UNK	0.461905
67	H25	-7.0370	0.2730	3.2800	ho	1	UNK	0.419618
68	H26	5.2630	-4.2150	0.8410	ho	1	UNK	0.419618

@<TRIPOS>BOND

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3	2	6	ar
4	2	14	ar
5	3	15	ar
6	3	17	ar
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8	4	18	ar
9	5	7	ar
10	5	9	ar
11	6	8	ar
12	6	10	ar
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36	22	24	l
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39	23	29	l
40	24	26	l

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66	35	61	1
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71	40	66	1
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73	42	68	1

@<TRIPOS>SUBSTRUCTURE

1 UNK 1 TEMP 0 ***** 0 ROOT

Hypothetical Compound 8H

lig_00000004.mol2

@<TRIPOS>MOLECULE

UNK

68 73 1 0 0

SMALL

No Charge or Current Charge

@<TRIPOS>ATOM

1	C1	-1.8820	0.5410	-0.5810	ca	1	UNK	-0.221852
2	C2	1.8420	0.5410	-0.5810	ca	1	UNK	-0.221852
3	C3	-0.4480	4.5380	-0.5810	ca	1	UNK	-0.343649
4	C4	0.4100	0.1970	-4.5630	ca	1	UNK	-0.343649
5	C5	-1.3970	1.8760	-0.5970	ca	1	UNK	0.122702
6	C6	1.3580	0.4430	-1.9120	ca	1	UNK	0.122702
7	C7	-1.9940	2.8330	0.2670	ca	1	UNK	-0.168049
8	C8	1.9570	-0.4990	-2.7920	ca	1	UNK	-0.168049
9	C9	-0.3210	2.2470	-1.4540	cp	1	UNK	-0.109897
10	C10	0.2820	1.2640	-2.3560	cp	1	UNK	-0.109897
11	C11	-3.5750	1.1260	1.1090	ca	1	UNK	-0.362498
12	C12	3.5380	-1.1890	-1.0180	ca	1	UNK	-0.362498
13	C13	-2.9300	0.1910	0.2430	ca	1	UNK	0.211056
14	C14	2.8910	-0.2490	-0.1600	ca	1	UNK	0.211056
15	C15	-1.4910	4.1700	0.2530	ca	1	UNK	0.232800
16	C16	1.4550	-0.6000	-4.1250	ca	1	UNK	0.232800
17	C17	0.1280	3.5680	-1.4370	ca	1	UNK	0.218351
18	C18	-0.1670	1.1310	-3.6700	ca	1	UNK	0.218351
19	C19	-3.0820	2.4380	1.1160	ca	1	UNK	0.300049
20	C20	3.0470	-1.3090	-2.3250	ca	1	UNK	0.300049
21	C21	-4.7170	0.6990	1.9100	c	1	UNK	0.606903

22	C22	4.6820	-1.9490	-0.5230	c	1	UNK	0.606903
23	C23	-5.1960	-0.7230	1.7010	c3	1	UNK	-0.180543
24	C24	5.1440	-1.6400	0.8840	c3	1	UNK	-0.180543
25	C25	-4.0310	-1.6240	1.3130	c3	1	UNK	0.106752
26	C26	3.9860	-1.1540	1.7480	c3	1	UNK	0.106752
27	C27	-5.9100	-1.2480	2.9280	c3	1	UNK	-0.088340
28	C28	5.8010	-2.8490	1.5160	c3	1	UNK	-0.088340
29	C29	-4.4930	-3.0280	0.9010	c3	1	UNK	0.135834
30	C30	4.4520	-0.6080	3.1050	c3	1	UNK	0.135834
31	O1	-5.3130	1.4530	2.6980	o	1	UNK	-0.580248
32	O2	5.2910	-2.7830	-1.2140	o	1	UNK	-0.580248
33	O3	-3.3540	-1.1190	0.1480	os	1	UNK	-0.338327
34	O4	3.3110	-0.0390	1.1370	os	1	UNK	-0.338327
35	O5	-2.0890	5.0620	1.1010	oh	1	UNK	-0.459050
36	O6	2.0550	-1.5210	-4.9410	oh	1	UNK	-0.459050
37	O7	1.1580	3.9080	-2.2760	oh	1	UNK	-0.478497
38	O8	-1.2000	1.9370	-4.0820	oh	1	UNK	-0.478497
39	O9	-3.5690	3.4360	1.8950	oh	1	UNK	-0.469651
40	O10	3.5380	-2.1680	-3.2540	oh	1	UNK	-0.469651
41	O11	-3.3930	-3.8710	0.6450	oh	1	UNK	-0.594512
42	O12	4.2630	-1.6500	4.0490	oh	1	UNK	-0.594512
43	H1	-1.4160	-0.2110	-1.2350	ha	1	UNK	0.164102
44	H2	1.3760	1.2560	0.1130	ha	1	UNK	0.164102
45	H3	-0.0680	5.5680	-0.5890	ha	1	UNK	0.139749
46	H4	0.0310	0.1150	-5.5900	ha	1	UNK	0.139749
47	H5	-5.9240	-0.6820	0.8390	hc	1	UNK	0.097103
48	H6	5.9020	-0.8090	0.7900	hc	1	UNK	0.097103
49	H7	-3.2850	-1.6930	2.1520	h1	1	UNK	0.068459
50	H8	3.2400	-1.9840	1.9030	h1	1	UNK	0.068459
51	H9	-6.4850	-2.1710	2.6770	hc	1	UNK	0.050539
52	H10	-6.6220	-0.4770	3.3130	hc	1	UNK	0.050539
53	H11	-5.1790	-1.4890	3.7370	hc	1	UNK	0.050539
54	H12	6.1110	-2.6130	2.5630	hc	1	UNK	0.050539
55	H13	6.6960	-3.1500	0.9190	hc	1	UNK	0.050539
56	H14	5.0900	-3.7100	1.5430	hc	1	UNK	0.050539
57	H15	-5.1570	-2.9620	-0.0010	h1	1	UNK	0.059805
58	H16	-5.0400	-3.5110	1.7530	h1	1	UNK	0.059805
59	H17	3.8420	0.2950	3.3770	h1	1	UNK	0.059805
60	H18	5.5380	-0.3270	3.0710	h1	1	UNK	0.059805
61	H19	-1.6620	5.9250	0.9840	ho	1	UNK	0.419649
62	H20	1.6270	-1.4800	-5.8110	ho	1	UNK	0.419649
63	H21	1.3530	4.8490	-2.1500	ho	1	UNK	0.420800
64	H22	-1.3910	1.7320	-5.0090	ho	1	UNK	0.420800
65	H23	-4.3040	3.1210	2.4500	ho	1	UNK	0.464000
66	H24	4.2710	-2.6950	-2.8890	ho	1	UNK	0.464000
67	H25	-2.8750	-3.4620	-0.0610	ho	1	UNK	0.415580
68	H26	4.2840	-1.2400	4.9210	ho	1	UNK	0.415580

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@<TRIPOS>SUBSTRUCTURE

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0 **** * 0 ROOT

Hypothetical Compound 9H

lig_00000005.mol2

@<TRIPOS>MOLECULE

UNK

70 75 1 0 0

SMALL

No Charge or Current Charge

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2	C2	1.8810	0.5340	-0.6250	ca	1	UNK	-0.223992
3	C3	-0.4070	4.5270	-0.6250	ca	1	UNK	-0.343795
4	C4	0.4380	0.1780	-4.6030	ca	1	UNK	-0.343795
5	C5	-1.3640	1.8670	-0.6410	ca	1	UNK	0.124944
6	C6	1.3920	0.4300	-1.9550	ca	1	UNK	0.124944
7	C7	-1.9580	2.8280	0.2220	ca	1	UNK	-0.169696
8	C8	1.9890	-0.5130	-2.8330	ca	1	UNK	-0.169696
9	C9	-0.2870	2.2350	-1.4970	cp	1	UNK	-0.110695
10	C10	0.3150	1.2500	-2.3980	cp	1	UNK	-0.110695
11	C11	-3.5450	1.1260	1.0600	ca	1	UNK	-0.364868
12	C12	3.5760	-1.1980	-1.0620	ca	1	UNK	-0.364868
13	C13	-2.9050	0.1860	0.1950	ca	1	UNK	0.213680
14	C14	2.9330	-0.2530	-0.2060	ca	1	UNK	0.213680
15	C15	-1.4520	4.1630	0.2080	ca	1	UNK	0.233604
16	C16	1.4840	-0.6190	-4.1650	ca	1	UNK	0.233604
17	C17	0.1660	3.5550	-1.4800	ca	1	UNK	0.219999
18	C18	-0.1380	1.1140	-3.7110	ca	1	UNK	0.219999
19	C19	-3.0440	2.4340	1.0730	ca	1	UNK	0.302400
20	C20	3.0800	-1.3220	-2.3670	ca	1	UNK	0.302400
21	C21	-4.6780	0.6950	1.8670	c	1	UNK	0.609527
22	C22	4.7180	-1.9580	-0.5680	c	1	UNK	0.609527
23	C23	-5.2330	-0.6750	1.5530	c3	1	UNK	-0.205510
24	C24	5.2140	-1.6050	0.8170	c3	1	UNK	-0.205510
25	C25	-4.0960	-1.6270	1.2060	c3	1	UNK	0.111636
26	C26	4.0620	-1.1370	1.6960	c3	1	UNK	0.111636
27	C27	-6.0910	-1.1980	2.7060	c3	1	UNK	0.139561
28	C28	5.9540	-2.7830	1.4500	c3	1	UNK	0.139561
29	C29	-4.5820	-2.9980	0.7190	c3	1	UNK	0.132825
30	C30	4.5470	-0.5750	3.0390	c3	1	UNK	0.132825
31	O1	-5.1910	1.3930	2.7610	o	1	UNK	-0.590070
32	O2	5.2970	-2.8270	-1.2440	o	1	UNK	-0.590070
33	O3	-3.3230	-1.1250	0.0990	os	1	UNK	-0.336988
34	O4	3.3570	-0.0390	1.0890	os	1	UNK	-0.336988
35	O5	-2.0500	5.0570	1.0540	oh	1	UNK	-0.459593
36	O6	2.0820	-1.5410	-4.9800	oh	1	UNK	-0.459593
37	O7	1.1990	3.8910	-2.3190	oh	1	UNK	-0.477902
38	O8	-1.1710	1.9180	-4.1210	oh	1	UNK	-0.477902
39	O9	-3.5290	3.4300	1.8580	oh	1	UNK	-0.468142
40	O10	3.5650	-2.1860	-3.2940	oh	1	UNK	-0.468142
41	O11	-5.3480	-1.3700	3.8930	oh	1	UNK	-0.606487
42	O12	6.9420	-2.2410	2.3160	oh	1	UNK	-0.606487
43	O13	-5.5180	-2.8890	-0.3300	oh	1	UNK	-0.596740
44	O14	3.4680	-0.3990	3.9280	oh	1	UNK	-0.596740
45	H1	-1.3870	-0.2200	-1.2760	ha	1	UNK	0.164560
46	H2	1.4170	1.2520	0.0670	ha	1	UNK	0.164560
47	H3	-0.0240	5.5560	-0.6330	ha	1	UNK	0.140204
48	H4	0.0560	0.0930	-5.6290	ha	1	UNK	0.140204
49	H5	-5.8990	-0.5730	0.6470	hc	1	UNK	0.110218
50	H6	5.9450	-0.7520	0.6940	hc	1	UNK	0.110218
51	H7	-3.4170	-1.7560	2.0940	h1	1	UNK	0.075068
52	H8	3.3270	-1.9730	1.8660	h1	1	UNK	0.075068
53	H9	-6.4920	-2.2180	2.4600	h1	1	UNK	0.055359
54	H10	-6.9430	-0.4850	2.8740	h1	1	UNK	0.055359
55	H11	6.4400	-3.4080	0.6520	h1	1	UNK	0.055359
56	H12	5.2480	-3.4230	2.0450	h1	1	UNK	0.055359
57	H13	-5.1210	-3.5270	1.5490	h1	1	UNK	0.068788
58	H14	-3.6940	-3.6030	0.3910	h1	1	UNK	0.068788
59	H15	5.0880	0.3930	2.8720	h1	1	UNK	0.068788
60	H16	5.2290	-1.3180	3.5350	h1	1	UNK	0.068788
61	H17	-1.6170	5.9180	0.9430	ho	1	UNK	0.420255
62	H18	1.6530	-1.5020	-5.8490	ho	1	UNK	0.420255
63	H19	1.3980	4.8310	-2.1910	ho	1	UNK	0.421255
64	H20	-1.3620	1.7140	-5.0490	ho	1	UNK	0.421255
65	H21	-4.2580	3.1080	2.4180	ho	1	UNK	0.462860
66	H22	4.3060	-2.7050	-2.9360	ho	1	UNK	0.462860
67	H23	-5.1320	-0.4860	4.2230	ho	1	UNK	0.411952
68	H24	7.5310	-2.9670	2.5510	ho	1	UNK	0.411952
69	H25	-5.0710	-2.4470	-1.0630	ho	1	UNK	0.411638

70 H26 2.8720 0.2480 3.5290 ho 1 UNK 0.411638
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Talaroderxine A (2)

lig_00000006.mol2

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UNK

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SMALL

No Charge or Current Charge

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2	C2	1.8600	0.8320	-0.8980	ca	1	UNK	-0.181185
3	C3	-0.4420	4.8390	-0.8980	ca	1	UNK	-0.335804
4	C4	0.4410	0.5280	-4.8940	ca	1	UNK	-0.335804
5	C5	-1.3690	2.1710	-0.9050	ca	1	UNK	0.098425
6	C6	1.3670	0.7380	-2.2330	ca	1	UNK	0.098425
7	C7	-1.9620	3.1210	-0.0350	ca	1	UNK	-0.146070
8	C8	1.9610	-0.2020	-3.1150	ca	1	UNK	-0.146070
9	C9	-0.3030	2.5490	-1.7700	cp	1	UNK	-0.106387
10	C10	0.3010	1.5710	-2.6760	cp	1	UNK	-0.106387
11	C11	-3.4840	1.3900	0.8260	ca	1	UNK	-0.312707
12	C12	3.4830	-0.9290	-1.3230	ca	1	UNK	-0.312707
13	C13	-2.8820	0.4550	-0.0660	ca	1	UNK	0.011390
14	C14	2.8810	0.0310	-0.4590	ca	1	UNK	0.011390
15	C15	-1.4750	4.4640	-0.0540	ca	1	UNK	0.226417
16	C16	1.4740	-0.2850	-4.4550	ca	1	UNK	0.226417
17	C17	0.1360	3.8740	-1.7560	ca	1	UNK	0.213640
18	C18	-0.1370	1.4560	-3.9970	ca	1	UNK	0.213640
19	C19	-3.0370	2.7110	0.8260	ca	1	UNK	0.281434
20	C20	3.0360	-1.0300	-2.6410	ca	1	UNK	0.281434
21	C21	-4.5710	0.9380	1.6910	c	1	UNK	0.751879
22	C22	4.5700	-1.7570	-0.8070	c	1	UNK	0.751879
23	C23	-3.4210	-0.9310	-0.0770	c3	1	UNK	-0.077832
24	C24	3.4200	0.1490	0.9220	c3	1	UNK	-0.077832
25	C25	-3.8960	-1.3360	1.3110	c3	1	UNK	0.150113
26	C26	3.8960	-1.2040	1.4310	c3	1	UNK	0.150113
27	C27	-4.3380	-5.1570	1.3190	c3	1	UNK	-0.092996
28	C28	4.3410	-0.9200	5.2410	c3	1	UNK	-0.092996
29	C29	-4.6140	-2.6810	1.3100	c3	1	UNK	-0.091266
30	C30	4.6150	-1.1010	2.7720	c3	1	UNK	-0.091266
31	C31	-3.6280	-3.8280	1.3360	c3	1	UNK	-0.080243
32	C32	3.6290	-1.0390	3.9180	c3	1	UNK	-0.080243
33	O1	-5.3450	1.6480	2.3490	o	1	UNK	-0.626078
34	O2	5.3440	-2.4670	-1.4650	o	1	UNK	-0.626078
35	O3	-4.8510	-0.3960	1.8390	os	1	UNK	-0.404076
36	O4	4.8510	-1.8020	0.5340	os	1	UNK	-0.404076
37	O5	-2.0750	5.3470	0.8000	oh	1	UNK	-0.456683
38	O6	2.0740	-1.2040	-5.2710	oh	1	UNK	-0.456683
39	O7	1.1620	4.2190	-2.6000	oh	1	UNK	-0.478251
40	O8	-1.1630	2.2720	-4.4050	oh	1	UNK	-0.478251
41	O9	-3.5430	3.6940	1.6150	oh	1	UNK	-0.470312
42	O10	3.5420	-1.8910	-3.5610	oh	1	UNK	-0.470312
43	H1	-1.4010	0.1040	-1.5840	ha	1	UNK	0.151181
44	H2	1.3990	1.5710	-0.2250	ha	1	UNK	0.151181
45	H3	-0.0740	5.8740	-0.9110	ha	1	UNK	0.141178

46	H4	0.0720	0.4620	-5.9260	ha	1	UNK	0.141178
47	H5	-4.2800	-0.9780	-0.8020	hc	1	UNK	0.074121
48	H6	-2.6430	-1.6600	-0.4240	hc	1	UNK	0.074121
49	H7	4.2790	0.8750	0.9130	hc	1	UNK	0.074121
50	H8	2.6420	0.5490	1.6230	hc	1	UNK	0.074121
51	H9	-3.0230	-1.3560	2.0240	h1	1	UNK	0.066235
52	H10	3.0240	-1.9140	1.5060	h1	1	UNK	0.066235
53	H11	-3.5970	-5.9920	1.3430	hc	1	UNK	0.035741
54	H12	-4.9590	-5.2610	0.3960	hc	1	UNK	0.035741
55	H13	-5.0110	-5.2550	2.2060	hc	1	UNK	0.035741
56	H14	3.6000	-0.8800	6.0760	hc	1	UNK	0.035741
57	H15	4.9600	0.0090	5.2740	hc	1	UNK	0.035741
58	H16	5.0130	-1.7960	5.4070	hc	1	UNK	0.035741
59	H17	-5.2820	-2.7310	2.2120	hc	1	UNK	0.059565
60	H18	-5.2720	-2.7560	0.4060	hc	1	UNK	0.059565
61	H19	5.2830	-1.9960	2.8900	hc	1	UNK	0.059565
62	H20	5.2730	-0.1930	2.7770	hc	1	UNK	0.059565
63	H21	-2.9420	-3.7560	0.4510	hc	1	UNK	0.043355
64	H22	-2.9890	-3.7550	2.2570	hc	1	UNK	0.043355
65	H23	2.9420	-0.1620	3.7790	hc	1	UNK	0.043355
66	H24	2.9910	-1.9620	3.9160	hc	1	UNK	0.043355
67	H25	-1.6670	6.2190	0.6720	ho	1	UNK	0.420592
68	H26	1.6660	-1.1430	-6.1500	ho	1	UNK	0.420592
69	H27	1.3350	5.1670	-2.4930	ho	1	UNK	0.421878
70	H28	-1.3360	2.0930	-5.3420	ho	1	UNK	0.421878
71	H29	-4.2980	3.3590	2.1340	ho	1	UNK	0.464224
72	H30	4.2970	-2.3830	-3.1880	ho	1	UNK	0.464224

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7	C7	-0.2330	-0.7690	-2.4230	cp	1 UNK	-0.041493
8	C8	0.2320	-1.9290	-1.6560	cp	1 UNK	-0.041493
9	C9	-2.4870	0.9800	0.0160	ca	1 UNK	-0.087993
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14	C14	2.8490	-1.4930	1.6780	ca	1 UNK	-0.232493
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24	C24	1.0790	-4.2330	-0.2630	ca	1	UNK	0.250407
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37	O7	1.3450	-1.3590	-4.1050	oh	1	UNK	-0.474793
38	O8	-1.3470	-3.2450	-2.8570	oh	1	UNK	-0.474793
39	O9	-1.3970	2.4180	-4.8640	oh	1	UNK	-0.498393
40	O10	1.3980	-5.4230	0.3210	oh	1	UNK	-0.498393
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42	H2	2.3790	1.7820	0.7610	ha	1	UNK	0.158307
43	H3	-3.9840	4.0350	1.9060	ha	1	UNK	0.159207
44	H4	3.9830	0.1750	4.4580	ha	1	UNK	0.159207
45	H5	0.3850	0.7650	-5.4600	ha	1	UNK	0.162307
46	H6	-0.3860	-5.3240	-1.4320	ha	1	UNK	0.162307
47	H7	-2.3560	1.7120	4.0430	hc	1	UNK	0.054240
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51	H11	3.7180	2.5290	4.2340	hc	1	UNK	0.054240
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53	H13	-3.7800	4.3370	-1.5420	ho	1	UNK	0.459807
54	H14	3.7810	-3.1160	3.3870	ho	1	UNK	0.459807
55	H15	1.7110	-1.0210	-4.9370	ho	1	UNK	0.430307
56	H16	-1.7130	-4.1440	-2.8710	ho	1	UNK	0.430307
57	H17	-2.1480	2.9390	-4.5340	ho	1	UNK	0.463807
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6	C6	1.3790	0.1060	-1.5660	ca	1 UNK	0.124095
7	C7	-1.9720	2.5110	0.6070	ca	1 UNK	-0.168805
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18	C18	-0.1590	0.7800	-3.3190	ca	1	UNK	0.217945
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22	C22	4.7150	-2.2670	-0.1820	c	1	UNK	0.605895
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25	C25	-4.0660	-1.9330	1.6330	c3	1	UNK	0.147095
26	C26	4.0650	-1.4410	2.0800	c3	1	UNK	0.147095
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35	O7	1.1920	3.5630	-1.9320	oh	1	UNK	-0.478305
36	O8	-1.1930	1.5830	-3.7310	oh	1	UNK	-0.478305
37	O9	-3.5480	3.1230	2.2330	oh	1	UNK	-0.470305
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52	H14	3.7040	-0.5560	4.0260	hc	1	UNK	0.052395
53	H15	5.2710	-0.0680	3.2600	hc	1	UNK	0.052395
54	H16	5.0850	-1.7400	3.9570	hc	1	UNK	0.052395
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56	H18	1.6170	-1.8530	-5.4480	ho	1	UNK	0.418795
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23	12	14	ar
24	12	20	ar

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REFERENCES

1. Kalb, S. R.; Smith, T. J.; Moura, H.; Hill, K.; Lou, J.; Geren, I. N.; Garcia-Rodriguez, C.; Marks, J. D.; Smith, L. A.; Pirkle, J. L.; Barr, J.R. The use of Endopep-MS to detect multiple subtypes of botulinum neurotoxins A, B, E, and F. *Int. J. Mass Spectrom.* **2008**, *278*, 101-108.
2. Schmidt, J. J.; Bostian, K. A. Endoproteinase activity of type A botulinum neurotoxin: substrate requirements and activation by serum albumin. *J. Protein Chem.* **1997**, *16*, 19-26.
3. Roxas-Duncan, V.; Enyedy, I.; Montgomery, V. A.; Eccard, V. S.; Carrington, M. A.; Lai, H.; Gul, N.; Yang, D. C.; Smith, L. A. Identification and biochemical characterization of small-molecule inhibitors of *Clostridium botulinum* neurotoxin serotype A. *Antimicrob. Agents. Chemother.* **2009**, *53*, 3478-3486.
4. Koyama, K.; Natori, S. Chaetochromins B, C and D, bis(naphtho-gamma-pyrone) derivatives from *Chaetomium gracile*. *Chem. Pharm. Bull.* **1987**, *35*, 578-584; Koyama, K.; Natori, S.; Iitaka, Y. Absolute configurations of chaetochromin A and related bis(naphtho- γ -pyrone) mold metabolites. *Chem. Pharm. Bull.* **1987**, *35*, 4049-4055.
5. Suzuki, K.; Nozawa, K.; Nakajima, S; Udagawa, S.; Kawai, K. Isolation and structures of antibacterial binaphtho- α -pyrones, talaroderxines A and B, from *Talaromyces derxii*. *Chem. Pharm. Bull.* **1992**, *40*, 1116-1119.
6. Steffens, J. C.; Robeson, D.J. Secalonic acid A, a vivotoxin in pink root-infected onion. *Phytochemistry* **1987**, *26*, 1599-1602. Howard, C.C.; Johnstone, R.A.W. Fungal metabolites. Part I. Stereochemical features and mass spectrometry of secalonic acids. *J. Chem. Soc. Perkin Trans. I* **1973**, 2440-2444.
7. Hegde, V.R.; Miller, J. R.; Patel, M. G.; King, A. H.; Puar, M. S.; Horan, A.; Hart, R.; Yarborough, R.; Gullo, V. SCH 45752--an inhibitor of calmodulin-sensitive cyclic nucleotide phosphodiesterase activity. *J. Antibiot.* **1993**, *46*, 207-213; Kock, I.; Draeger, S.; Schulz, B.; Elsässer, B.; Tibor Kurtán, T.; Kenéz, A.; Antus, S.; Pescitelli, G.; Salvadori, P.; Speakman, J.-B.; Rheinheimer, J.; Krohn, K. Pseudoanguillosporin A and B: two new isochromans isolated from the endophytic fungus *Pseudoanguillospora* sp. *Eur. J. Org. Chem.* **2009**, 1427-1434.
8. Gill, M.; Gimenez, A.; McKenzie, R. W. Pigments of fungi, part 8. Bisanthraquinones from *Dermocybe austroveneta*. *J. Nat. Prod.* **1988**, *51*, 1251-1256. Antonowitz, A.; Gill, M.; Morgan, P. M.; Yu, J. Coupled anthraquinones from the toadstool *Dermocybe icterinoides*. *Phytochemistry* **1994**, *37*, 1679-1683; Koyama, K.; Aida, S.; Natori, S. Supplemental observations on atropisomerism of fungal bis(naphtho- γ -pyrones). *Chem. Pharm. Bull.* **1990**, *38*, 2259-2261.
9. Cardellina, J. H.; Vieira, R. C.; Eccard, V.; Skerry, J.; Montgomery, V.; Campbell, Y.; Roxas-Duncan, V.; Leister, W.; LeClair, C. A.; Maloney, D. J.; Padula, D.; Pescitelli, G.; Khavrutskii, I.; Hu, X.; Wallqvist, A.; Smith, L. A. Separation of Betti reaction product enantiomers: Absolute configuration and inhibition of botulinum neurotoxin A. *ACS Med. Chem. Lett.* **2011**, *2*, 396.
10. Wang, J.; Wolf, R. M.; Caldwell, J. W.; Kollman, P. A.; Case, D. A. Development and testing of a general Amber force field. *J. Comput. Chem.* **2004**, *25*, 1157-1174.
11. Jakalian, A.; Bush, B. L.; Jack, D. B.; Bayly, C. I. Fast, efficient generation of high-quality atomic charges. AM1-BCC model: I. Method. *J. Comput. Chem.* **2000**, *21*, 132-146.
12. Jakalian, A.; Jack, D. B.; Bayly, C. I. Fast, efficient generation of high-quality atomic charges. AM1-BCC model: II. parameterization and validation. *J. Comput. Chem.* **2002**, *23*, 1623-1641.
13. Khavrutskii, I. V.; Wallqvist, A. Computing relative free energies of solvation using Single Reference Thermodynamic Integration augmented with Hamiltonian Replica Exchange. computing relative free

energies of solvation using Single Reference Thermodynamic Integration augmented with Hamiltonian Replica Exchange. *J. Chem. Theory Comput.* **2010**, *6*, 3427-3441.

14. Khavrutskii, I. V.; Wallqvist, A. Improved binding free energy predictions from Single-Reference Thermodynamic Integration augmented with Hamiltonian Replica Exchange. *J. Chem. Theory Comput.* **2011**, *7*, 3001-3011.

15. Trott, O.; Olson, A. J. AutoDock Vina: Improving the speed and accuracy of docking with a new scoring function, efficient optimization, and multithreading. *J. Comput. Chem.* **2010**, *31*, 455-461.

16. Hess, B.; Kutzner, C.; van der Spoel, D.; Lindahl, E. GROMACS 4: Algorithms for Highly Efficient, Load-Balanced, and Scalable Molecular Simulation. *J. Chem. Theory Comput.* **2008**, *4*, 435-447.

17. van der Spoel, D.; Lindahl, E.; Hess, B.; Kutzner, C.; van Buuren, A. R.; Apol, E.; Meulenhoff, P. J.; Tieleman, D. P.; Sijbers, A. L. T. M.; Feenstra, K. A.; Drunen, R. v.; Berendsen, H. J. C., GROMACS User Manual Version 4.0. 4.0 ed.; The GROMACS development team: AG Groningen, The Netherlands, 2006.

18. Clark, A. M.; Labute, P. Two-dimensional depiction of protein-ligand complexes. *J. Chem. Inf. Model.* **2007**, *47*, 1933-1944.