

Supporting Information
for
Thermodynamics and spectroscopy of halogen- and hydrogen-bonded complexes of haloforms with aromatic and aliphatic amines

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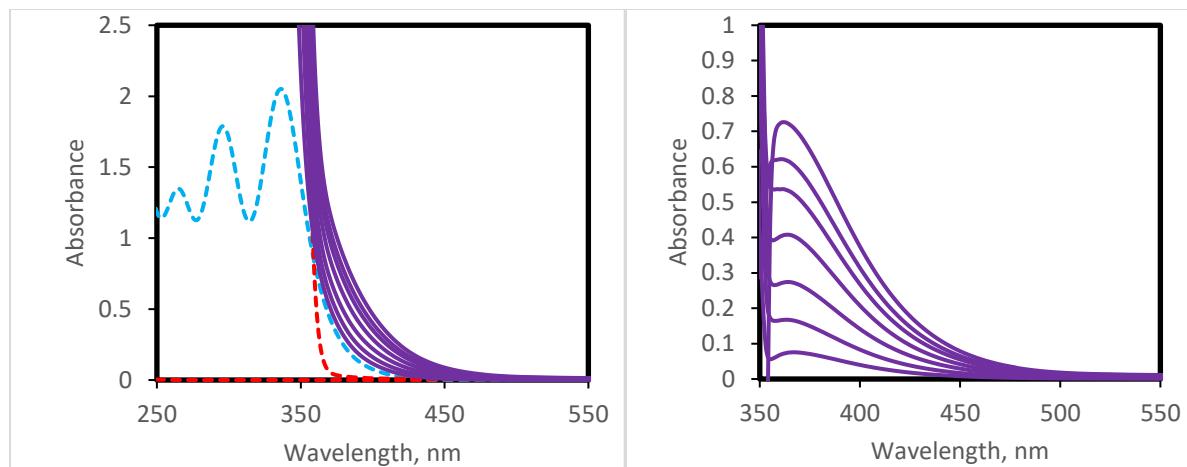


Figure S1. (Left) Spectra of the solutions with constant concentration of CHI_3 and various concentrations of p-methoxy-N,N-dimethylaniline (pMeODMA). Spectra of the solutions of individual reactants are shown as dashed blue (CHI_3) or red (pMeODMA) lines. (Right): Spectra of the complexes obtained by subtraction of the absorption of components from the spectra of their mixtures

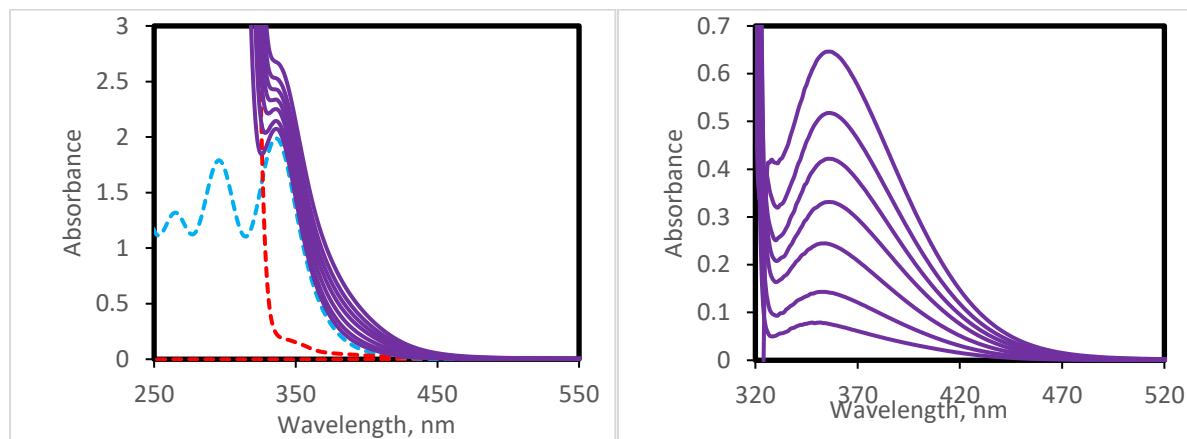


Figure S2. (Left) Spectra of the solutions with constant concentration of CHI_3 and various concentrations of m-methoxy-N,N-dimethylaniline (mMeODMA). Spectra of the solutions of individual reactants are shown as dashed blue (CHI_3) or red (mMeODMA) lines. (Right): Spectra of the complexes obtained by subtraction of the absorption of components from the spectra of their mixtures

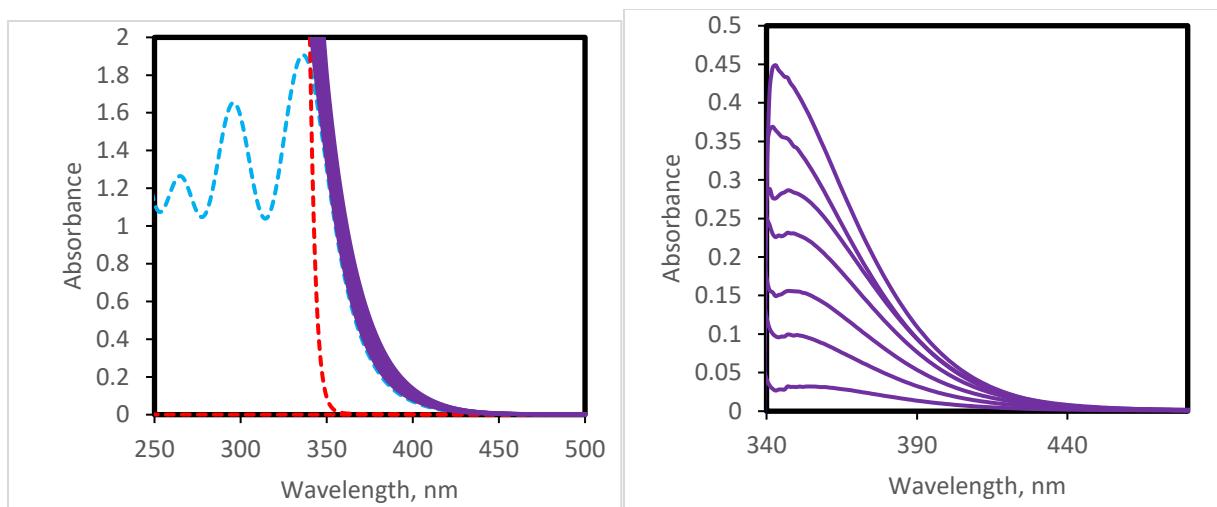


Figure S3. (Left) Spectra of the solutions with constant concentration of CHI₃ and various concentrations of p-cyano-N,N-dimethylaniline (DMACN). Spectra of the solutions of individual reactants are shown as dashed blue (CHI₃) or red (DMACN) lines. (Right): Spectra of the complexes obtained by subtraction of the absorption of components from the spectra of their mixtures

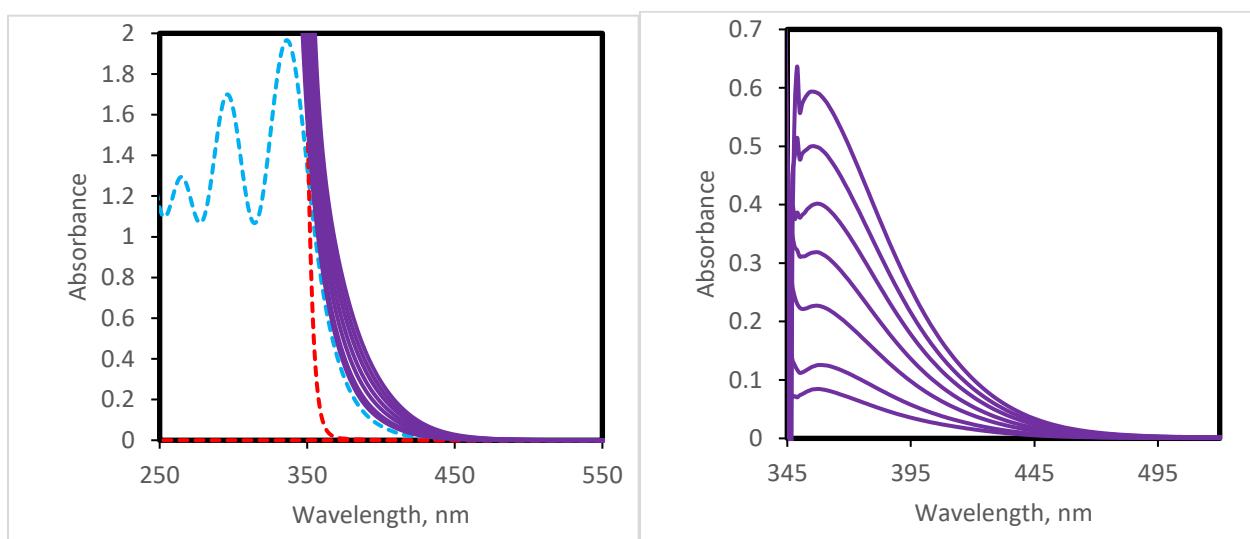


Figure S4. (Left) Spectra of the solutions with constant concentration of CHI₃ and various concentrations of p-bromo-N,N-dimethylaniline (DMABr). Spectra of the solutions of individual reactants are shown as dashed blue (CHI₃) or red (DMABr) lines. (Right): Spectra of the complexes obtained by subtraction of the absorption of components from the spectra of their mixtures

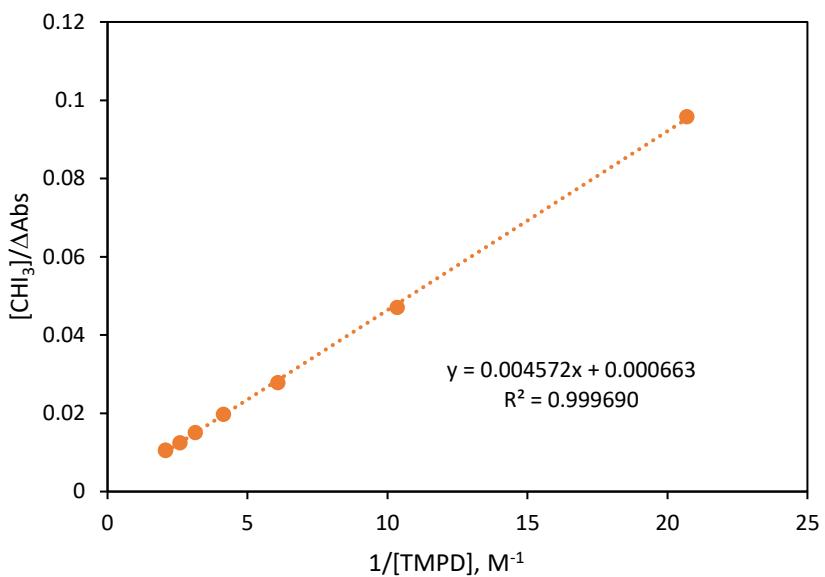
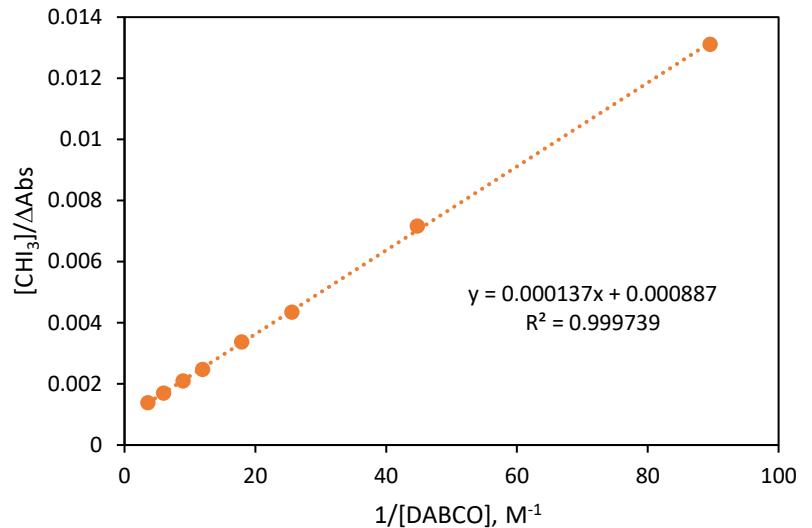


Figure S5. Benesi-Hilderbrand plots $C_D^0 / \text{Abs} = 1/(\varepsilon l) + \{1/(K_{\text{eff}} \varepsilon l)\} \times 1/[C_A^0]$ (where C_D^0 and C_A^0 are initial concentrations of CHI_3 and amine, respectively, ε and l are extinction coefficient of the complex and the length of the cell which was used in the UV-Vis measurements, K_{eff} is the effective formation constant of the complex and ΔAbs is the absorbance of the complex at certain wavelength (obtained by subtraction of the absorption of the components from the spectra of the mixtures), based on the treatments of spectra measured in solution of CHI_3 with DABCO (top) and TMPD (bottom).

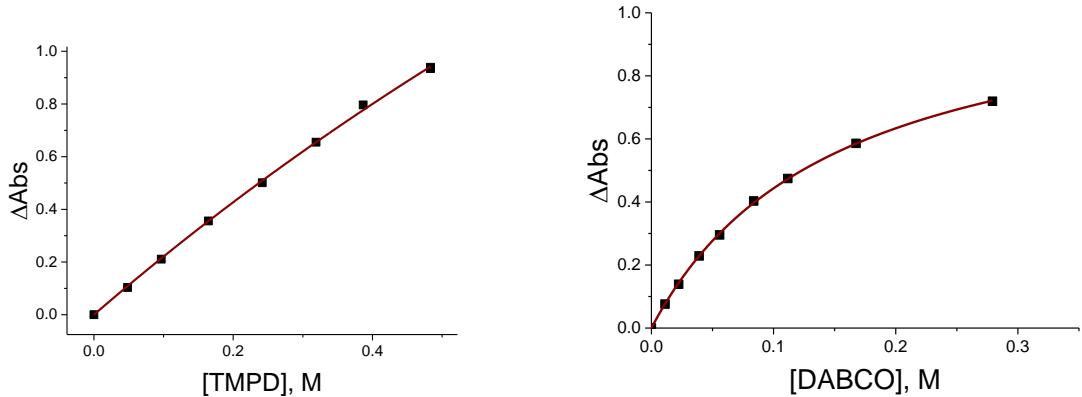


Figure S6. The fit (using Origin 16) of the UV-Vis spectral data of the solutions containing constant concentrations of CHI_3 and variable concentrations of TMPD (left) or DABCO (right) to 1:1 binding isotherm as $\Delta\text{Abs} = \varepsilon l \times C_{\text{com}} = \varepsilon l \times \{(C^{\circ}_{\text{A}} + C^{\circ}_{\text{D}} + 1/K_{\text{eff}}) - ((C^{\circ}_{\text{A}} + C^{\circ}_{\text{D}} + 1/K_{\text{eff}})^2 - 4C^{\circ}_{\text{A}}C^{\circ}_{\text{D}})^{0.5}\}/2$, where ΔAbs is the absorbance of the complex at certain wavelength (obtained by subtraction of the absorption of the components from the spectra of the mixtures), C_{com} is the concentration of the complex, and C°_{D} and C°_{A} are initial concentrations of CHX_3 and amine, ε and l are extinction coefficient of the complex and the length of the cell which was used in the UV-Vis measurements, and K_{eff} is the effective formation constant of the complex.

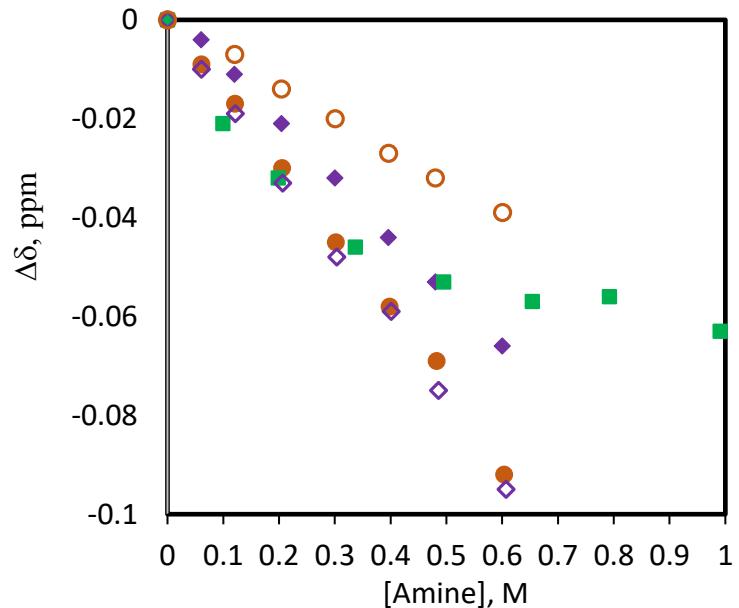


Figure S7. Dependencies of the chemical shifts of the protons of CHI_3 (as compared to that in the corresponding isolated molecules) on the concentration of added trimethylamine (TEA) (■), DMA (●), DMACN (○), DMABr (◆) or mMeODMA (◊) (in CD_3CN , 22°C).

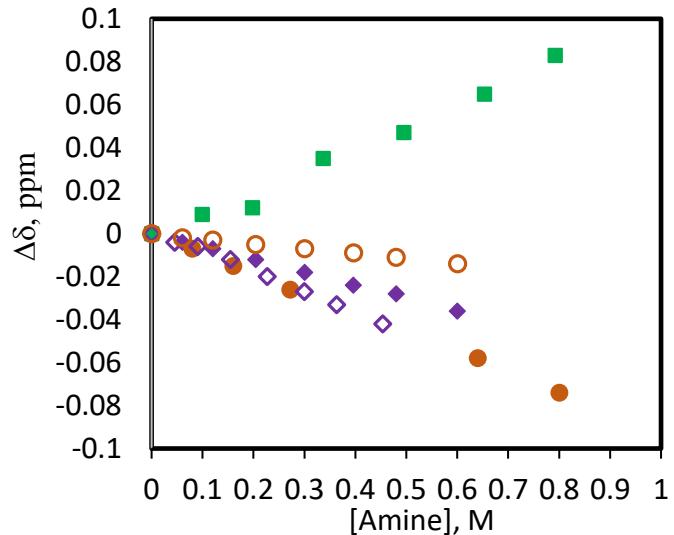


Figure S8. Dependencies of the chemical shifts of the protons of CHBr_3 as compared to that in the corresponding isolated molecules) on the concentration of added TEA (■), DMA (●), DMACN (○), DMABr (♦) or mMeODMA (◊) (in CD_3CN , 22°C).

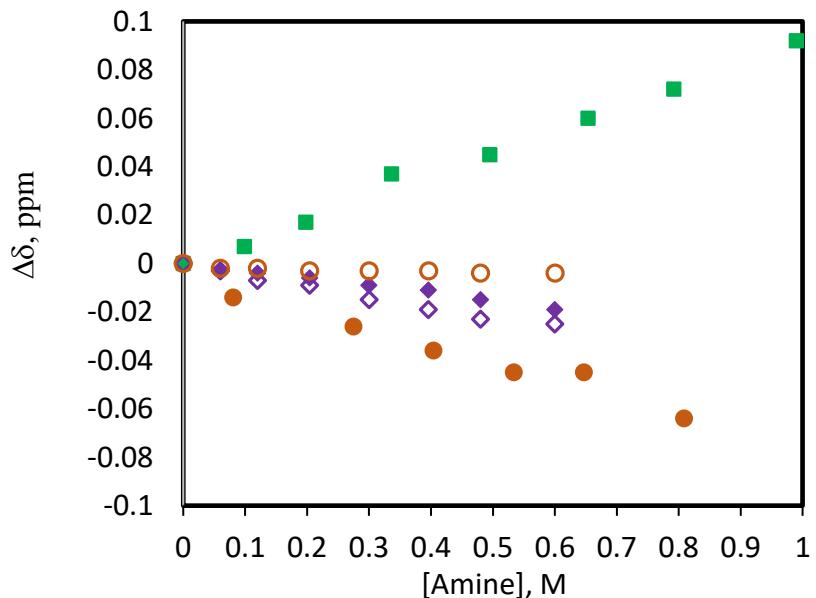


Figure S9. Dependencies of the chemical shifts of the protons of CHCl_3 as compared to that in the corresponding isolated molecules) on the concentration of added TEA (■), DMA (●), DMACN (○), DMABr (♦) or mMeODMA (◊) (in CD_3CN , 22°C).

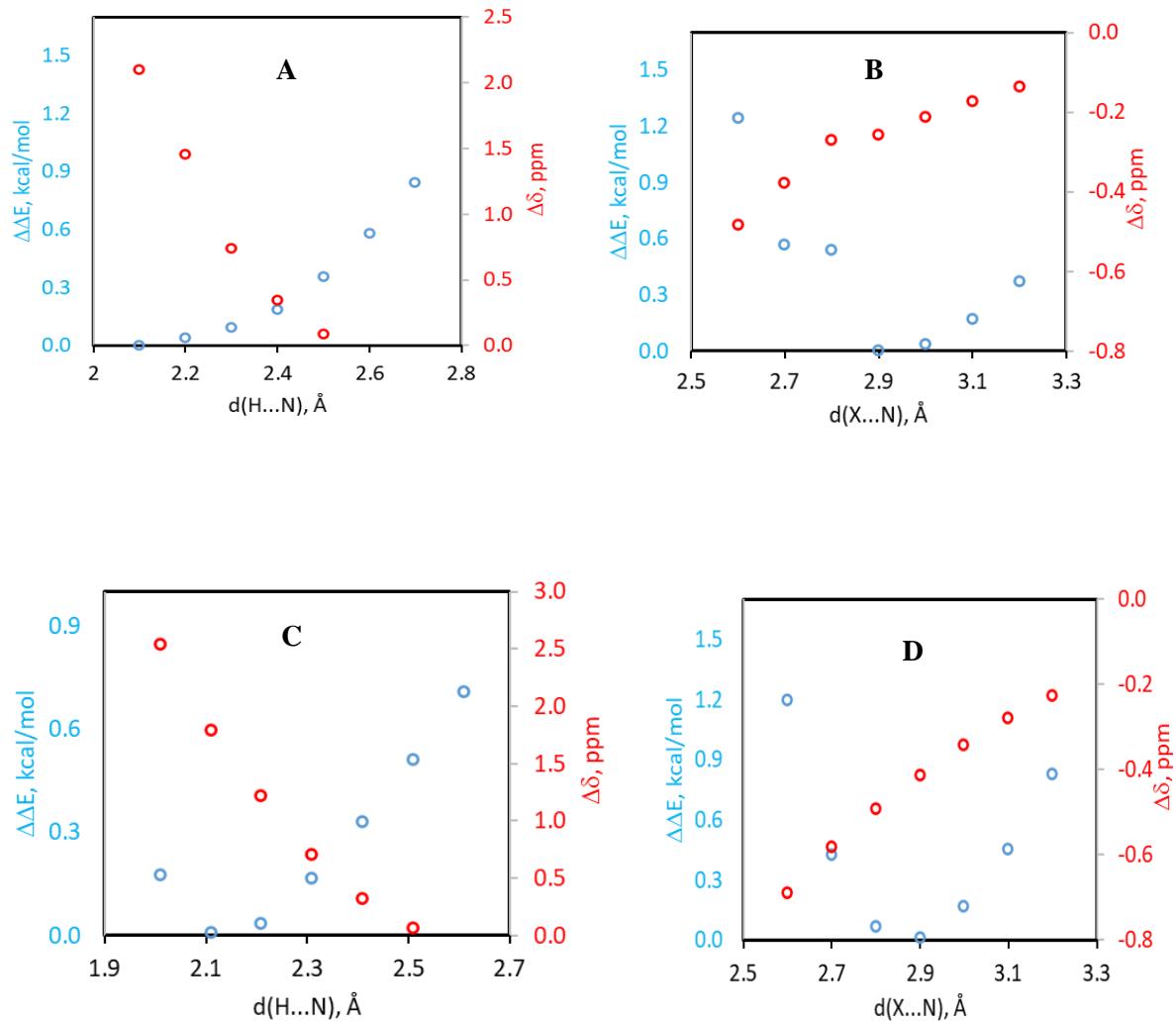


Figure S10. Effect of variations of interatomic H...N and X...N separations on the energies of HyB and HaB complexes (blue circle) relative to that of the energy minimum and the shift of the proton signal in their NMR spectra (red circles) relative to that in the individual CHX₃ for: A) HyB complexes of CHCl₃ with DABCO, B) HaB complexes of CHCl₃ with DABCO, C) HyB complexes of CHBr₃ with TMPD, D) HaB complexes of CHBr₃ with TMPD.

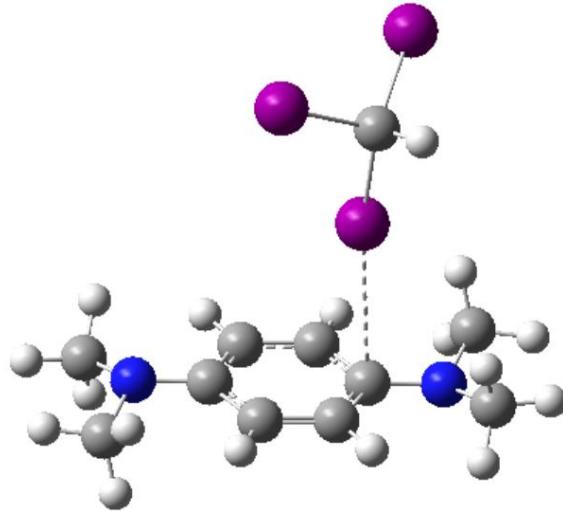


Figure S11. Alternative structure of the HaB complex between CHI_3 and TMPD showing I...C bonding ($d_{\text{I} \dots \text{C}} = 3.35 \text{ \AA}$, $\Delta E = -4.3 \text{ kcal/mol}$, chemical shift relative to that of the individual CHI_3 $\Delta\Delta\delta = -0.55 \text{ ppm}$ and absorption band with $\lambda_{\text{max}} = 392 \text{ nm}$ and $\varepsilon = 2900 \text{ M}^{-1} \text{ cm}^{-1}$).

Table S1. Values of V_{max} on the surfaces of halogen (V^X) and hydrogen (V^H) atoms (at 0.001 a.u. electron density) in the individual haloforms and in the molecules polarized by the presence of electron rich center nearby halogen (in HaB complex) or hydrogen (in HyB complex) atoms.

Molecule	V_{max}^H , a.u.	V_{max}^X , a.u.
CHI_3 (Ind)	0.0541	0.0425
CHI_3 (HaB complex)	0.0434	0.124
CHI_3 (HyB complex)	0.108	0.0334
CHBr_3 (Ind)	0.0611	0.0302
CHBr_3 (HaB complex)	0.0528	0.0828
CHBr_3 (HyB complex)	0.108	0.0181
CHCl_3 (Ind)	0.0624	0.02
CHCl_3 (HaB complex)	0.055	0.055
CHCl_3 (HyB complex)	0.103	0.0141

Table S2. Crystallographic, data collection and refinement details.

	CHI ₃ ·DABCO	2CHI ₃ ·TMPD	CHI ₃ ·TMPD
Chemical formula	C ₆ H ₁₂ N ₂ ·CHI ₃	C ₁₀ H ₁₆ N ₂ ·2(CHI ₃)	C ₁₃ H ₂₀ I ₃ N ₃
M _r	505.89	951.68	599.02
Crystal system, space group	Orthorhombic, <i>Pnma</i>	Triclinic, <i>P</i> ̄ 1	Orthorhombic, <i>Pnma</i>
Temperature (K)	150	150	150
<i>a</i> , <i>b</i> , <i>c</i> (Å)	16.4635 (13), 10.6079 (9), 7.3574 (6)	6.6114 (3), 9.6487 (5), 9.8861 (5)	10.5562 (5), 15.1118 (6), 11.6277 (5)
α, β, γ (°)		63.578 (2), 83.094 (3), 74.633 (3)	1854.89 (14)
<i>V</i> (Å ³)	1284.92 (18)	544.58 (5)	4
Z	4	1	Mo <i>Kα</i>
Radiation type	Mo <i>Kα</i>	Mo <i>Kα</i>	5.05
μ (mm ⁻¹)	7.26	8.55	0.32 × 0.25 × 0.16
Crystal size (mm)	0.25 × 0.23 × 0.18	0.21 × 0.20 × 0.13	C ₁₃ H ₂₀ I ₃ N ₃
Diffractometer	Bruker AXS D8 Quest CMOS diffractometer	Bruker AXS D8 Quest diffractometer with PhotonII charge-integrating pixel array detector (CPAD)	Bruker AXS D8 Quest diffractometer with PhotonII charge-integrating pixel array detector (CPAD)
Absorption correction	Multi-scan <i>SADABS</i> 2016/2: Krause, L., Herbst-Irmer, R., Sheldrick G.M. & Stalke D., <i>J. Appl. Cryst.</i> 48 (2015) 3-10	Multi-scan <i>SADABS</i> 2016/2: Krause, L., Herbst-Irmer, R., Sheldrick G.M. & Stalke D., <i>J. Appl. Cryst.</i> 48 (2015) 3-10	Multi-scan <i>SADABS</i> 2016/2: Krause, L., Herbst-Irmer, R., Sheldrick G.M. & Stalke D., <i>J. Appl. Cryst.</i> 48 (2015) 3-10
<i>T</i> _{min} , <i>T</i> _{max}	0.432, 0.747	0.500, 0.747	0.539, 0.747
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	13583, 3209, 2702	13429, 4038, 3603	52270, 3646, 3302
<i>R</i> _{int}	0.040	0.037	0.043
(sin θ/λ) _{max} (Å ⁻¹)	0.833	0.771	0.769
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.037, 0.082, 1.06	0.022, 0.046, 1.05	0.013, 0.028, 1.06
No. of reflections	3209	4038	3646
No. of parameters	65	94	98
H-atom treatment	H-atom parameters constrained	H-atom parameters constrained	H-atom parameters constrained
Δ <i>ρ</i> _{max} , Δ <i>ρ</i> _{min} (e Å ⁻³)	2.58, -2.25	1.79, -1.73	0.50, -0.43

Computer programs: Apex3 v2016.9-0 (Bruker, 2016), SAINT V8.37A (Bruker, 2016), *SHELXS*97 (Sheldrick, 2008), *SHELXL*2017/1 (Sheldrick, 2015, 2016), *SHELXLE* Rev714 (Hübschle *et al.*, 2011).

Table S3. Energies of the HyB and HaB complexes between CHX₃ and amines resulting from the M06-2X/def-TZVPP computations (acetonitrile).

CHX ₃	Amine	Energy + ZPE, Hartree		BSSE, Hartree		ΔE, kcal/mol ^a	
		HaB complex	HyB complex	HaB complex	HyB complex	HaB	HyB
CHI ₃	DABCO	-1276.702449	-1276.698772	0.000676	0.000626	-7.0	-4.7
	TMPD	-1431.471322	-1431.471276	0.000619	0.000635	-5.5	-5.5
CHBr ₃	DABCO	-8106.386286	-8106.387885	0.000667	0.000793	-3.5	-4.4
	TMPD	-8261.157135	-8261.159697	0.000649	0.000902	-3.3	-4.7
CHCl ₃	DABCO	-1764.400973	-1764.404661	0.000553	0.000832	-1.8	-4.0
	TMPD	-1919.172264	-1919.176305	0.000583	0.000940	-1.9	-4.2

a) $\Delta E = E_{\text{comp}} - (E_{\text{CHX}_3} + E_{\text{D}}) + \text{BSSE}$ where E_{comp} , E_{CHX_3} and E_{D} are sums of the electronic and ZPE of the complex, CHX₃ and amine and BSSE is a basis set superposition error. Energies of the individual molecules

Amines	Energy+ZPE, Hartree	CHX ₃	Energy+ZPE, Hartree
TMPD	-499.89700337	CHI ₃	-931.564856
DABCO	-345.125783	CHBr ₃	-7761.254232
		CHCl ₃	-1419.271718

Table S4. Atomic coordinates of the optimized HyB and HaB complexes between CHX₃ and amines (from the M06-2X/def-TZVPP computations in acetonitrile)

CHI ₃ -TMPD (HyB)				CHI ₃ -TMPD(HaB)			
C	-2.50128400	2.11661400	0.00105700	C	4.03883200	-1.62919400	-0.19174400
C	-1.57644800	1.66757400	0.93943400	C	3.08456700	-1.09818500	0.67069100
C	-1.84988200	0.44981200	1.56934800	C	3.22299400	0.24679500	1.02451500
C	-2.98718700	-0.28073700	1.28251700	C	4.27124500	1.01682000	0.55981500
C	-3.92871500	0.17240400	0.34032300	C	5.25125200	0.48036800	-0.29605500
C	-3.64249300	1.38379000	-0.30202000	C	5.09028700	-0.85828700	-0.67424200
H	-2.33789800	3.04428000	-0.52837300	H	3.97917900	-2.65897200	-0.51215900
H	-1.14420900	0.04616100	2.28556800	H	2.48246800	0.71350100	1.66084500
H	-3.13230400	-1.22035500	1.79379500	H	4.31687700	2.05175200	0.86281100
H	-4.31236000	1.77723500	-1.05140100	H	5.79099000	-1.32516000	-1.34950300
N	-5.08760500	-0.54463100	0.07146100	N	6.32508800	1.24121100	-0.73356300
N	-0.35376200	2.33474300	1.20942500	N	1.96788000	-1.84137200	1.14817600
C	-5.12405000	-1.93910000	0.46701900	C	6.23505500	2.68233100	-0.60525900
H	-6.07235200	-2.36821000	0.15593600	H	7.13594900	3.12864900	-1.01659600
H	-4.31083000	-2.52405900	0.01907800	H	5.36694700	3.09795600	-1.13153100
H	-5.05743100	-2.03788800	1.55001500	H	6.17043200	2.97594700	0.44216400
C	-5.83127600	-0.19567400	-1.12220800	C	7.10536200	0.73429900	-1.84404600
H	-6.69302600	-0.85196500	-1.20493100	H	7.89146300	1.44694600	-2.07677800
H	-6.20004900	0.82791300	-1.06264700	H	7.58196000	-0.21031900	-1.58280100
H	-5.22940600	-0.28969500	-2.03518200	H	6.49974000	0.57386700	-2.74481900
C	-0.12423600	2.66694500	2.61081600	C	1.81076300	-1.83606200	2.60140800
H	0.91328100	2.97331000	2.73885500	H	0.85120600	-2.28568000	2.85384100
H	-0.77488100	3.48689500	2.94285700	H	2.60881300	-2.40746400	3.09043400
H	-0.30513500	1.80404700	3.24553000	H	1.82115700	-0.82053000	2.98603300

C	-0.03044500	3.44703500	0.33726400	C	1.83287300	-3.18625400	0.61413700
H	0.98395200	3.77629800	0.55671800	H	0.89298900	-3.60580000	0.96921300
H	-0.07136600	3.12860500	-0.70432600	H	1.80648300	-3.15617000	-0.47398400
H	-0.70612600	4.30326900	0.47122800	H	2.64946000	-3.84617800	0.93265000
C	1.20460200	-0.17745600	0.00075000	C	-2.13490500	0.38666700	-0.61688200
H	0.64941700	0.50756900	0.63144500	H	-1.89526400	0.77196100	-1.59787300
I	1.33638000	-2.03381500	1.03579800	I	-0.36771600	-0.54943300	0.16219700
I	3.10101600	0.75706700	-0.27202300	I	-3.73270600	-0.99934600	-0.91601500
I	0.09541100	-0.32290200	-1.81124900	I	-2.74583400	2.07936500	0.53366200
CHBr ₃ -TMPD (HyB)				CHBr ₃ -TMPD (HaB)			
C	-1.98135200	1.92281000	-0.57393700	C	-3.31674500	1.63856500	-0.30699700
C	-1.09686600	1.72540100	0.48213500	C	-2.32496900	1.16970200	0.55126700
C	-1.39242300	0.69308200	1.37631400	C	-2.47435100	-0.13210100	1.04071100
C	-2.51342400	-0.10091900	1.22824300	C	-3.56359300	-0.91465800	0.70967500
C	-3.41394300	0.09738000	0.16553300	C	-4.57820700	-0.43809400	-0.14054200
C	-3.10571000	1.12275100	-0.73775700	C	-4.40936700	0.85225300	-0.65563300
H	-1.79854800	2.69787900	-1.30434300	H	-3.25350000	2.63142200	-0.72812000
H	-0.71357500	0.48438500	2.19429000	H	-1.70856400	-0.55598900	1.67692500
H	-2.67668900	-0.89186400	1.94447900	H	-3.61387200	-1.91422100	1.11416800
H	-3.74402800	1.31419300	-1.58686300	H	-5.13673400	1.27059800	-1.33465700
N	-4.55525100	-0.68210100	0.02837100	N	-5.69440700	-1.20960000	-0.44247200
N	0.11241200	2.45485400	0.63603000	N	-1.17374300	1.92272400	0.88742900
C	-4.60345100	-1.93951600	0.74893100	C	-5.61521700	-2.63627300	-0.19723100
H	-5.53449600	-2.44581700	0.51023700	H	-6.54688500	-3.09883900	-0.51084700
H	-3.76789400	-2.60284600	0.49228900	H	-4.78833500	-3.11181000	-0.73953200
H	-4.58638300	-1.77184000	1.82533800	H	-5.48757900	-2.84138500	0.86518500
C	-5.24445100	-0.64540900	-1.24585100	C	-6.51261600	-0.79552600	-1.56431000
H	-6.09704900	-1.31738500	-1.20465600	H	-7.32724500	-1.50385000	-1.68654400
H	-5.62222700	0.35546500	-1.45265500	H	-6.95181000	0.18486000	-1.38162700
H	-4.59966900	-0.94705200	-2.08117900	H	-5.94641500	-0.74654700	-2.50324400
C	0.25485900	3.13996500	1.91643200	C	-0.89514700	2.04019800	2.31307500
H	1.27936700	3.49498400	2.01966400	H	0.10391300	2.45450400	2.44528700
H	-0.42296300	4.00084000	1.98903600	H	-1.61697400	2.69889100	2.81333400
H	0.04335000	2.46324500	2.73941200	H	-0.92220800	1.06686100	2.79412300
C	0.47246200	3.32038900	-0.47107800	C	-1.01660400	3.19188400	0.20594200
H	1.47268600	3.71236000	-0.29433500	H	-0.03756000	3.59907900	0.45439300
H	0.48260900	2.75222500	-1.40101700	H	-1.06407800	3.04745900	-0.87245800
H	-0.21326000	4.17108200	-0.58607700	H	-1.77900200	3.92740100	0.49673200
C	1.59416800	-0.31693800	-0.01764700	C	2.63916100	-0.60291600	-0.58444300
H	1.15017200	0.57930300	0.40462700	H	2.35454800	-1.18304800	-1.44966200
Br	1.58185300	-1.69123200	1.33638400	Br	3.22268200	-1.85380000	0.76837100
Br	0.52726300	-0.81774300	-1.54374200	Br	1.09766900	0.40751800	-0.00231000
Br	3.39457700	0.14976400	-0.53798300	Br	4.08551300	0.55612700	-1.13169600
CHCl ₃ -TMPD (HyB)				CHCl ₃ -TMPD (HaB)			
C	-1.08545100	1.70398500	-0.79175900	C	-2.15301000	1.63525800	-0.54977700
C	-0.21799500	1.50518600	0.27829300	C	-1.11957400	1.26555200	0.30987800

C	-0.61950500	0.59976600	1.26397100	C	-1.28189000	0.06365500	1.00908700
C	-1.82722100	-0.06792600	1.19265000	C	-2.41677000	-0.71270200	0.87225300
C	-2.71259900	0.13680000	0.11849000	C	-3.46913600	-0.33333100	0.02009800
C	-2.29707600	1.02886000	-0.87855700	C	-3.29158000	0.85160100	-0.70236100
H	-0.82252800	2.38081700	-1.59204300	H	-2.08659900	2.54653400	-1.12602900
H	0.04098500	0.38906600	2.09645100	H	-0.49192100	-0.29175900	1.65692400
H	-2.07246800	-0.76483900	1.97963500	H	-2.47318700	-1.63270500	1.43418300
H	-2.91885600	1.21315200	-1.74141500	H	-4.04812200	1.19051000	-1.39376100
N	-3.94050300	-0.50873600	0.05845500	N	-4.63036100	-1.09422900	-0.08590000
N	1.06674300	2.10451200	0.36606100	N	0.07191300	2.01039500	0.44255900
C	-4.13484700	-1.67149600	0.90227100	C	-4.57303700	-2.47129300	0.36442700
H	-5.12484900	-2.07862600	0.71711800	H	-5.53788900	-2.93890000	0.18841700
H	-3.39213000	-2.45674300	0.71363800	H	-3.80165400	-3.05327100	-0.15587300
H	-4.08235300	-1.39983900	1.95615900	H	-4.37353500	-2.52224200	1.43420500
C	-4.63935800	-0.50911200	-1.21092600	C	-5.49923700	-0.83081000	-1.21517700
H	-5.56505600	-1.06743400	-1.10339600	H	-6.34583100	-1.51064800	-1.17444300
H	-4.89861900	0.50607900	-1.51021600	H	-5.88990500	0.18546400	-1.17375900
H	-4.04768300	-0.96240800	-2.01639400	H	-4.98989800	-0.96201700	-2.17872000
C	1.29193000	2.88939200	1.57539100	C	0.50658400	2.28605000	1.80372300
H	2.34984100	3.13780900	1.64851100	H	1.52302200	2.67740200	1.77764300
H	0.71354000	3.82280500	1.56198100	H	-0.13981300	3.02467300	2.29646900
H	1.01218300	2.32244200	2.45874700	H	0.51386300	1.37964400	2.40179600
C	1.51324500	2.81592000	-0.81660900	C	0.21008700	3.16777100	-0.41566500
H	2.55323400	3.10579700	-0.67579400	H	1.21722800	3.56556400	-0.30116700
H	1.44979000	2.16658800	-1.68932500	H	0.07183000	2.88312900	-1.45778700
H	0.92831100	3.72531100	-1.01195900	H	-0.50460100	3.96701300	-0.17356100
C	2.16194200	-0.89248500	-0.07565700	C	3.47654500	-0.97417700	-0.63478600
H	1.88788300	0.08208100	0.31756000	H	3.24265800	-1.46755300	-1.56772000
Cl	3.86229400	-0.82264900	-0.54963600	Cl	2.18866200	0.17183100	-0.26614300
Cl	1.91136300	-2.11030500	1.17661700	Cl	3.59362000	-2.21257200	0.61934600
Cl	1.13850300	-1.21562400	-1.47558200	Cl	5.02298900	-0.14393400	-0.83282700
CHBr₃-DABCO (HyB)				CHBr₃-DABCO (HaB)			
C	-1.22279200	0.00045700	0.08015600	C	2.18262800	-0.00010400	-0.56732600
N	1.85197500	0.08602100	0.40700900	N	-2.52276100	-0.01376500	-0.02562200
C	2.16656300	0.60389700	-0.93149800	C	-3.08030000	1.03518400	-0.88396400
C	2.59312000	0.86822800	1.40397700	C	-3.06829800	-1.31072500	-0.43556200
C	2.28103500	-1.31689800	0.48571800	C	-2.90274200	0.25035000	1.36489500
N	4.36400500	-0.09053400	-0.03407200	N	-5.06516800	0.01369500	0.18441200
H	1.52899200	0.09715100	-1.65833700	H	-2.60108700	1.98093500	-0.62736100
H	1.91307700	1.66475500	-0.94720500	H	-2.82119600	0.79697800	-1.91659800
H	2.19489000	1.88330500	1.41418700	H	-2.70598400	-1.53460900	-1.43980000
H	2.40865900	0.42466000	2.38305200	H	-2.67370100	-2.07022400	0.24079500
H	1.94565000	-1.72928800	1.43807900	H	-2.41065400	-0.48542100	2.00227600
H	1.77757900	-1.86450500	-0.31264000	H	-2.52276800	1.23648400	1.63545000
C	3.67112900	0.37650800	-1.23961500	C	-4.62014500	1.10589300	-0.68733100
H	4.14530700	1.29723300	-1.58046500	H	-5.14237500	1.02310000	-1.64092200
H	3.80344200	-0.37704400	-2.01692200	H	-4.91388900	2.04859200	-0.22433500

C	4.10646900	0.85692400	1.05565000	C	-4.62082800	-1.26097000	-0.38887100
H	4.70584300	0.56689800	1.91905100	H	-5.02183400	-2.07287500	0.21849200
H	4.44158400	1.84248300	0.73024400	H	-5.04659600	-1.34794900	-1.38911600
C	3.82423800	-1.40108500	0.34260500	C	-4.44891000	0.18017200	1.50468800
H	4.10880600	-2.12816800	-0.41846600	H	-4.84414200	1.08891300	1.95950100
H	4.29024000	-1.69936000	1.28249400	H	-4.74886400	-0.66428200	2.12619200
H	-0.18913200	0.00662700	0.42780500	Br	2.99463900	1.60008400	0.15434800
Br	-1.33098000	-1.32002100	-1.32633200	Br	3.01291500	-1.58934800	0.15759300
Br	-1.58508300	1.77391800	-0.59059300	Br	0.26495700	-0.01067700	-0.30813100
Br	-2.37266100	-0.45050700	1.56111400	H	2.38227300	-0.00001100	-1.62861600
CHCl₃-DABCO (HyB)				CHCl₃-DABCO (HaB)			
C	2.03809100	0.00013900	0.10152100	C	3.08696500	-0.00121800	-0.50344800
N	-1.05816600	0.05327600	0.36557000	N	-1.58488400	-0.29863800	-0.39288100
C	-1.52148100	-1.34038800	0.38763200	C	-1.82258800	1.14122000	-0.53013200
C	-1.82733200	0.82698200	1.34765300	C	-2.61855000	-1.02498900	-1.13660400
C	-1.29961900	0.61114800	-0.97164600	C	-1.67592500	-0.66055600	1.02491000
N	-3.55155600	-0.05294700	-0.19332700	N	-3.90046400	0.29819900	0.51715900
H	-0.99370600	-1.87967000	-0.40043800	H	-1.00315200	1.67322400	-0.04420000
H	-1.24121800	-1.78461200	1.34337300	H	-1.79715700	1.38672700	-1.59284800
H	-1.69672700	0.35564100	2.32243300	H	-2.49234400	-0.81537000	-2.19934800
H	-1.40865300	1.83244700	1.40169100	H	-2.45648500	-2.09268600	-0.98331200
H	-1.02062800	1.66540600	-0.94944200	H	-1.41365600	-1.71422800	1.12984600
H	-0.64222000	0.10666100	-1.68231500	H	-0.93235500	-0.07554800	1.56858000
C	-3.05809700	-1.38520900	0.17091400	C	-3.19199000	1.51359300	0.10211300
H	-3.57411800	-1.69812500	1.07944300	H	-3.81928400	2.05474300	-0.60707900
H	-3.32264800	-2.08450000	-0.62276800	H	-3.06049100	2.14410700	0.98243000
C	-3.32346000	0.85865400	0.93287500	C	-4.02690100	-0.59320200	-0.64063200
H	-3.62330400	1.85991600	0.62115500	H	-4.62360100	-1.45798500	-0.34837300
H	-3.96688100	0.55827300	1.76035600	H	-4.57111200	-0.05813800	-1.41995800
C	-2.79378600	0.42985100	-1.35273800	C	-3.11046200	-0.37919500	1.55078200
H	-2.90897300	-0.29809100	-2.15674200	H	-3.08759300	0.25329200	2.43895300
H	-3.23016300	1.37135000	-1.68750800	H	-3.62223500	-1.30619800	1.81264900
H	1.03095200	0.00459200	0.51621500	H	3.46990900	-0.00160700	-1.51431100
Cl	2.29464100	1.56403300	-0.67893500	Cl	1.33974300	-0.21832700	-0.56115200
Cl	2.11062500	-1.30552500	-1.08766700	Cl	3.85998300	-1.32897000	0.36894300
Cl	3.21711300	-0.25960500	1.38728300	Cl	3.50402100	1.54958000	0.23368500
CHI₃-DABCO (HyB)				CHI₃-DABCO (HaB)			
C	-0.79508700	-0.03847900	-0.02185900	C	1.72078000	-0.00017000	-0.61980300
N	2.26804100	0.09817400	0.50236500	N	-3.09669500	-0.00628500	-0.02812100
C	2.72131600	0.35538500	-0.87118600	C	-3.66314700	1.23231400	-0.57776400
C	2.88736600	1.08290300	1.39768600	C	-3.69298700	-1.15873900	-0.71629100
C	2.70130500	-1.24782000	0.90023300	C	-3.40531300	-0.08570800	1.40570800
N	4.81497200	-0.07987200	0.37032200	N	-5.62176000	0.00611700	0.30863500
H	2.16827300	-0.29968600	-1.54749200	H	-3.14412500	2.07710000	-0.12384800
H	2.46375400	1.38563500	-1.12029000	H	-3.45787300	1.24646300	-1.64871000
H	2.47654000	2.06676600	1.16833500	H	-3.37822100	-1.13467200	-1.75999500

H	2.60480600	0.83045100	2.42041500	H	-3.28820400	-2.06411300	-0.26273300
H	2.26586100	-1.47968400	1.87303000	H	-2.90097200	-0.96097800	1.81644800
H	2.29689400	-1.95556800	0.17475500	H	-2.98819900	0.80002600	1.88623100
C	4.25173400	0.11559200	-0.96995100	C	-5.18827600	1.27321000	-0.28689300
H	4.75167400	0.96337800	-1.43921300	H	-5.75521000	1.43955000	-1.20276100
H	4.47336100	-0.77383900	-1.56102100	H	-5.43109600	2.07568800	0.40989200
C	4.42932800	1.05569700	1.21528000	C	-5.23927400	-1.09323900	-0.58211800
H	4.93692700	0.96415600	2.17593000	H	-5.63472100	-2.02410700	-0.17590100
H	4.78368300	1.96824000	0.73434800	H	-5.70812400	-0.92282600	-1.55145700
C	4.25180800	-1.30680200	0.94358600	C	-4.94455900	-0.16923300	1.59654200
H	4.62815100	-2.16082500	0.37976700	H	-5.29447500	0.60246700	2.28204700
H	4.61322500	-1.39599500	1.96875500	H	-5.23534900	-1.13791600	2.00336700
H	0.22370400	0.00004400	0.36515800	H	1.92864200	0.00009000	-1.68096000
I	-0.89900500	-1.81078800	-1.20219900	I	-0.42196800	-0.00685600	-0.34561400
I	-1.00768800	1.73364200	-1.18744300	I	2.64937600	-1.76858300	0.14289000
I	-2.11086600	-0.08420700	1.65216700	I	2.63662500	1.77569900	0.14099800