

## Supporting Online Material

The supporting on-line material contains:

- A Table with the CC2/cc-pVTZ calculated harmonic wavenumbers compared to the experimental values of tryptamine A. Furthermore, approximate descriptions of the modes and the calculated oscillator strengths in km/mol for all modes are given.
- A Table with the vibrational wavenumbers and Duschinsky matrix for the low frequency vibrations of the A conformer of tryptamine calculated at the CC2/cc-pVTZ level of theory.
- A Figure with the displacement vectors of the normal modes of tryptamine.
- A Figure with the IR-ID spectra of all other conformers.
- A figure with a FC simulation of the R2PI spectrum after pumping IR resonances of the A conformer at  $341.5\text{ cm}^{-1}$ .

Table S1: Harmonic vibrational frequencies and oscillator strengths of tryptamine A calculated at the CC2/cc-pVTZ level of theory, along with the approximate description of the modes and the experimental values. <sup>a</sup> from Ref. [1] <sup>b</sup> this publication

Mode	$\nu$ [cm <sup>-1</sup> ]	I [km/mol]	Description	Exp.	Mode	$\nu$	I [km/mol]	Description	Exp.
1	44.8	1.28123	$\tau_1$	43 <sup>a</sup>	34	1150.0	1.523870	18a	1155 <sup>a</sup>
2	83.6	1.23459	$\beta_1$	81 <sup>a</sup>	35	1171.8	0.487759	9a	1138 <sup>a</sup>
3	135.7	3.18774	$\tau_2$	133 <sup>a</sup>	36	1180.5	0.102449	NH <sub>2</sub> twist	1194 <sup>a</sup>
4	177.9	1.76047	10b	177 <sup>a</sup>	37	1220.2	2.680435	CH <sub>2</sub> twist	1228 <sup>a</sup>
5	219.2	11.05047	10a	215 <sup>a</sup>	38	1244.0	11.262858	13	
6	240.9	19.75197	$N_{inv}/\tau_1/\tau_3$	237 <sup>b</sup>	39	1279.9	4.60635	3	1294 <sup>a</sup>
7	265.7	33.20797	$\gamma NH/\tau_3$	268 <sup>b</sup>	40	1315.7	3.81878	18b	1326 <sup>a</sup>
8	332.3	47.00850	$N_{inv}/\tau_1/\tau_3$	341.5 <sup>b</sup>	41	1340.9	2.16207	C <sub>B</sub> H <sub>2</sub> twist	1334 <sup>a</sup>
9	384.6	22.49646	$N_{inv}/\tau_1/\tau_2$	387 <sup>b</sup>	42	1361.2	1.74642	C <sub>a</sub> H <sub>2</sub> bend	1361 <sup>a</sup>
10	425.1	5.56796	16b	422 <sup>b</sup>	43	1376.1	13.84470	3/ $\nu$ CC	
11	462.1	6.58679	4	454 <sup>b</sup>	44	1380.1	10.10924	C <sub>B</sub> H <sub>2</sub> wag	
12	476.6	19.72806	9b	468 <sup>b</sup>	45	1425.0	46.43047	$\delta$ CH, NH	1421 <sup>a</sup>
13	538.5	2.02159	CCN bend		46	1446.2	5.31395	$\delta$ NH	
14	570.9	1.29528	6a		47	1476.4	5.19797	C <sub>a</sub> H <sub>2</sub> bend	1462 <sup>a</sup>
15	581.6	10.17618	6b	569 <sup>a</sup>	48	1483.5	6.36950	19a	
16	639.7	0.40608	$\gamma NH$		49	1514.7	5.05619	CCC bend	1512 <sup>a</sup>
17	705.9	0.17446	16a		50	1523.5	3.52947	C <sub>B</sub> H <sub>2</sub> bend	1521 <sup>a</sup>
18	740.1	4.53353	18		51	1586.2	4.21056	$\nu$ CC	
19	747.9	73.38435	11		52	1613.4	0.36652	8a	1598 <sup>a</sup>
20	774.4	4.88340	$\gamma CH$		53	1653.8	30.18284	NH <sub>2</sub> bend	1631 <sup>a</sup>
21	799.7	6.46777	1		54	1658.3	2.88088	8b	
22	840.8	1.11881	17a	847 <sup>a</sup>	55	2951.7	44.82044	$\nu_s C_B H$	
23	845.5	49.00812	17b		56	3021.1	24.52387	$\nu_s C_a H$	
24	879.9	6.88880	C <sub>a</sub> H <sub>2</sub> twist		57	3048.7	50.35626	$\nu_{as} C_B H$	
25	892.1	41.24889	5	875 <sup>a</sup>	58	3068.1	19.64459	$\nu_{as} C_a H$	
26	913.1	2.35243	12		59	3163.2	2.61424	$\nu_{as} CH$	
27	919.5	8.50358	NH <sub>2</sub> wag		60	3169.5	1.03134	$\nu_s CH$	
28	947.4	0.06897	C <sub>a</sub> H <sub>2</sub> wag		61	3179.4	20.47852	$\nu_{as} CH$	
29	1025.3	3.03033	18b	1015 <sup>a</sup>	62	3190.5	15.35401	$\nu_s CH$	
30	1036.5	6.66141	CH <sub>2</sub> twist		63	3245.7	0.06675	$\nu_{py} CH$	
31	1093.0	12.22130	NH <sub>2</sub> twist	1091 <sup>a</sup>	64	3487.9	1.52794	$\nu_s NH_2$	
32	1098.7	22.16356	$\delta$ CH, NH		65	3567.6	2.96235	$\nu_{as} NH_2$	
33	1108.5	4.46519	C <sub>a</sub> H <sub>2</sub> wag	1131 <sup>a</sup>	66	3669.8	83.83458	$\nu_{py} NH$	

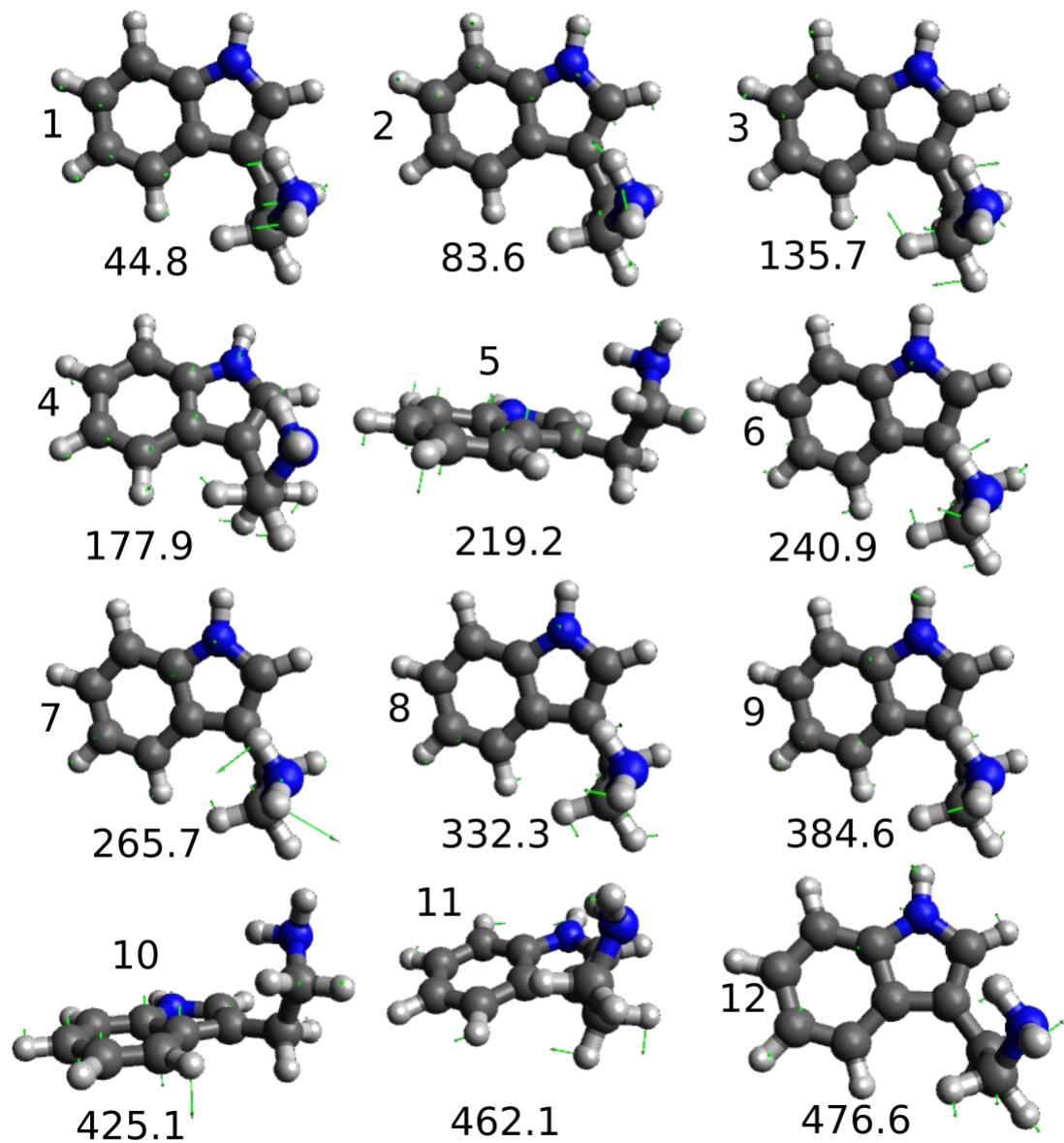


Figure S1: Displacement vectors of the twelve lowest wavenumber vibrations of tryptamine from a normal mode analysis using the CC2/cc-pVTZ optimized geometry. The mode numbering is according to Tables S1 and S2.

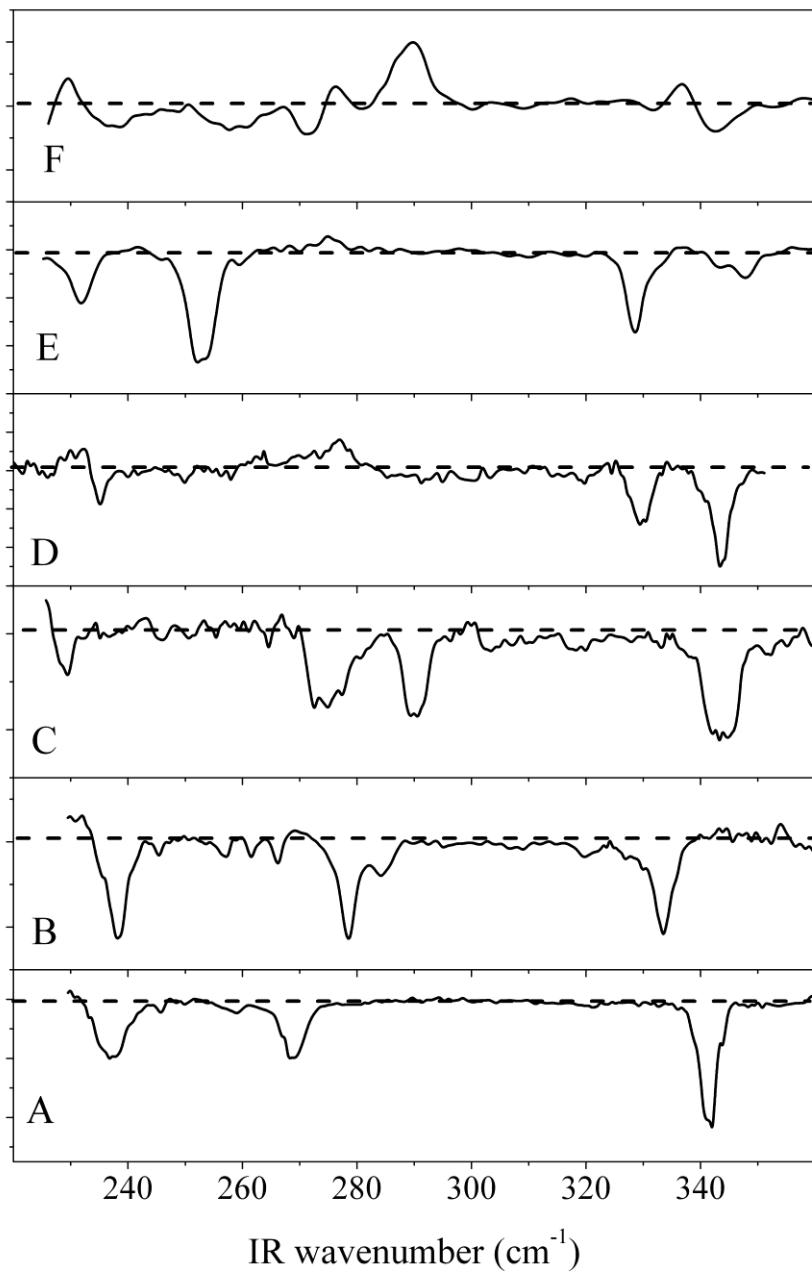


Figure S2: IR-ID spectra of the six conformers (A-F) of tryptamine obtained using a laser desorption source. Solid tryptamine was mixed with graphite powder and applied on a solid graphite bar. A pulsed YAG laser operating at 1064 nm (Polaris Pulsed Nd:YAG Laser, NewWave Research) with a pulse energy of about 1.5 mJ was used to desorb unfragmented tryptamine molecules from the graphite bar. The desorbed molecules are entrained in a supersonic pulsed molecular beam of argon, from a pulsed valve (Jordan) at a backing pressure of 3 bar. In the molecular beam, tryptamine is cooled towards its vibrational ground state and the different conformers of tryptamine are frozen out, depending on their relative energy and the barriers separating the conformer minima. The relative intensity of the conformer origins in the R2PI spectra depend on the temperature and on the relaxation in the beam. For weak conformers, we could enhance the signal by using a laser desorption source. However, all spectra involving only the A conformer in the main paper were taken using the thermal source.

Table S2: Vibrational wavenumbers and Duschinsky matrix for the low frequency vibrations of the A conformer of tryptamine calculated at the CC2/cc-pVTZ level of theory. In the second row the  $S_1$  wavenumber is given, in the second column the  $S_0$  wavenumber. The primed vibrational quantum numbers refer to the excited state vibrations, the doubly primed to the ground state.

$S_0$	$S_1$	$1^t$ 40.8	$2^t$ 78.9	$3^t$ 120.8	$4^t$ 158.7	$5^t$ 163.9	$6^t$ 229.2	$7^t$ 237.6	$8^t$ 279.0	$9^t$ 307.6	$10^t$ 343.9	$11^t$ 378.7	$12^t$ 458.4
$1^{tt}$	44.8	-0.992	0.065	0.053	0.020	-0.002	-0.009	0.032	0.010	0.006	-0.007	0.016	0.011
$2^{tt}$	83.6	-0.065	-0.991	0.046	-0.077	-0.038	0.003	0.002	-0.007	-0.015	0.008	-0.019	-0.000
$3^{tt}$	135.7	0.062	0.011	0.938	0.301	0.113	0.038	0.068	0.009	0.006	0.001	0.017	-0.006
$4^{tt}$	177.9	0.009	0.091	0.313	-0.791	-0.457	-0.214	0.053	0.053	-0.022	-0.027	0.026	-0.029
$5^{tt}$	219.2	0.006	-0.011	-0.042	0.488	-0.835	-0.137	-0.120	-0.032	0.114	0.040	0.075	-0.002
$6^{tt}$	240.9	0.017	0.009	-0.045	-0.023	-0.215	0.661	0.709	-0.012	0.024	-0.032	0.034	0.011
$7^{tt}$	265.7	0.026	-0.023	-0.082	0.144	0.096	-0.635	0.629	0.381	0.048	0.086	-0.042	-0.002
$8^{tt}$	332.3	-0.007	0.007	0.017	-0.045	-0.031	0.280	-0.233	0.723	0.207	0.501	-0.185	-0.043
$9^{tt}$	384.6	-0.014	0.029	0.013	0.009	-0.053	-0.057	0.098	-0.398	-0.393	0.602	-0.532	-0.105
$10^{tt}$	425.1	0.002	-0.006	0.005	-0.088	0.107	-0.069	0.071	-0.392	0.754	0.417	0.221	-0.019
$11^{tt}$	462.1	-0.008	-0.002	-0.016	0.028	0.003	0.014	0.001	0.017	0.071	-0.143	-0.061	-0.948
$12^{tt}$	476.6	-0.001	-0.005	-0.011	-0.007	0.010	0.006	-0.012	0.009	0.085	-0.141	-0.128	-0.109

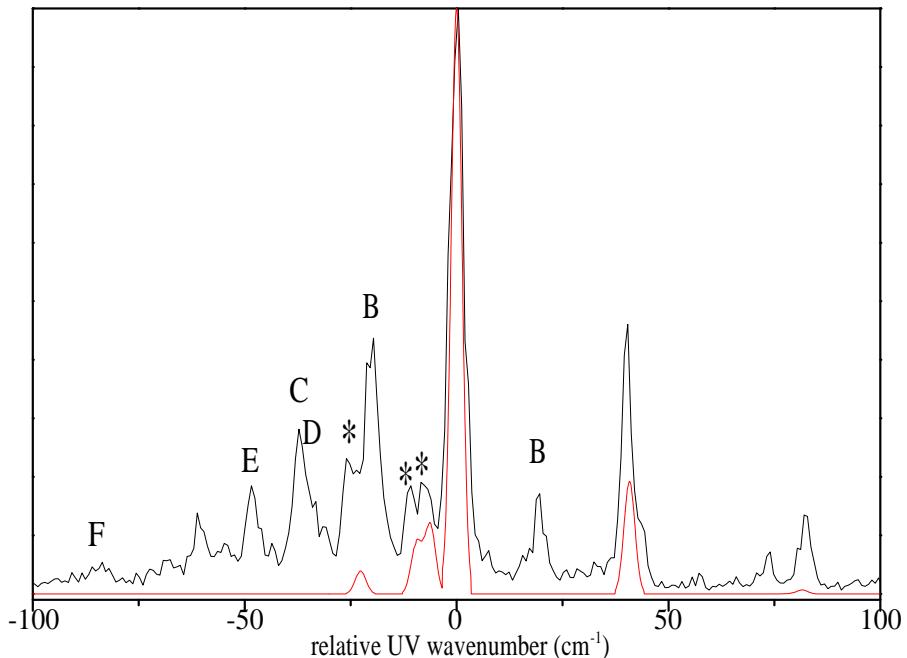


Figure S3: R2PI spectrum of tryptamine after pumping IR resonances of the A conformer at  $341.5\text{ cm}^{-1}$  (black trace) along with a Franck-Condon simulation using the vibrationless origin and the vibration 8 as ground state levels. For details of the simulation see text. The bands marked with \* are those, that appear only upon concerted action of IR and UV radiation. IR pulse energy used here is 0.54 mJ.

## References

- [1] M. Schmitt, K. Feng, M. Böhm, and K. Kleinermanns, *J. Chem. Phys.*, 2006, **125**, 144303(1–9).