

Electronic supplementary information for “Interaction in the indole...imidazole heterodimer: Structure, Franck-Condon analysis and energy decomposition”.

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Table SI. SAPT2+3/aug-cc-pVDZ contributions (kcal mol⁻¹) to the total interaction energy for four proposed structures of the indole...imidazole heterodimer. All ground electronic state geometries optimized at the M05-2X/6-31+G(d) level.

Structure	$E_{\text{electrostatic}}$	E_{exchange}	$E_{\text{induction}}$	$E_{\text{dispersion}}$
HB	-13.07	14.32	-4.69	-6.36
T1	-6.31	8.79	-2.43	-8.15
T2	-6.23	9.08	-2.67	-7.97
Syn-PD	-8.17	11.37	-1.95	-9.37

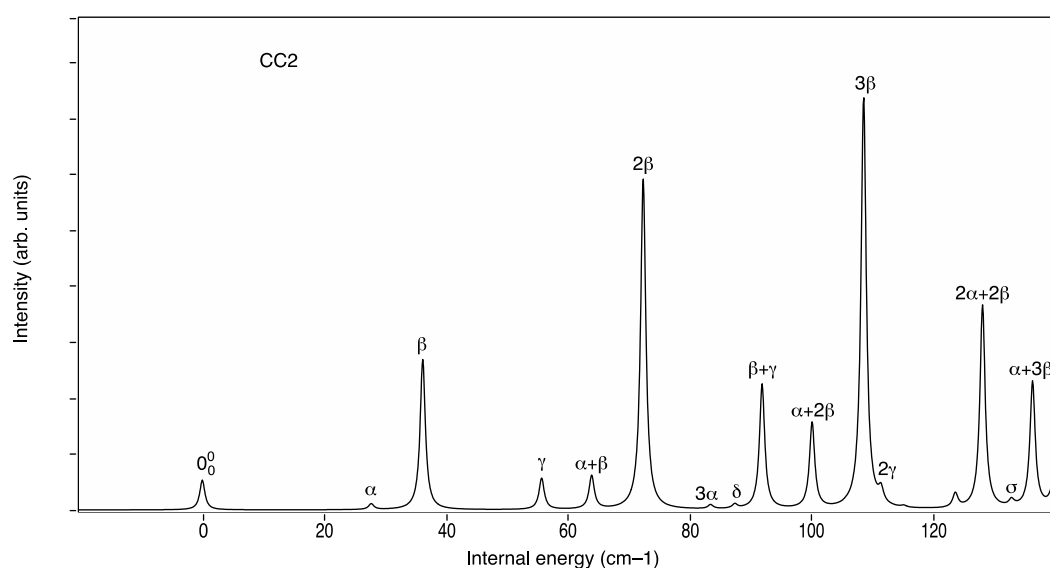


Fig. S1. Franck-Condon simulated electronic spectrum for the HB indole...imidazole structure obtained from calculations at the CC2/def2-TZVPP level of theory.

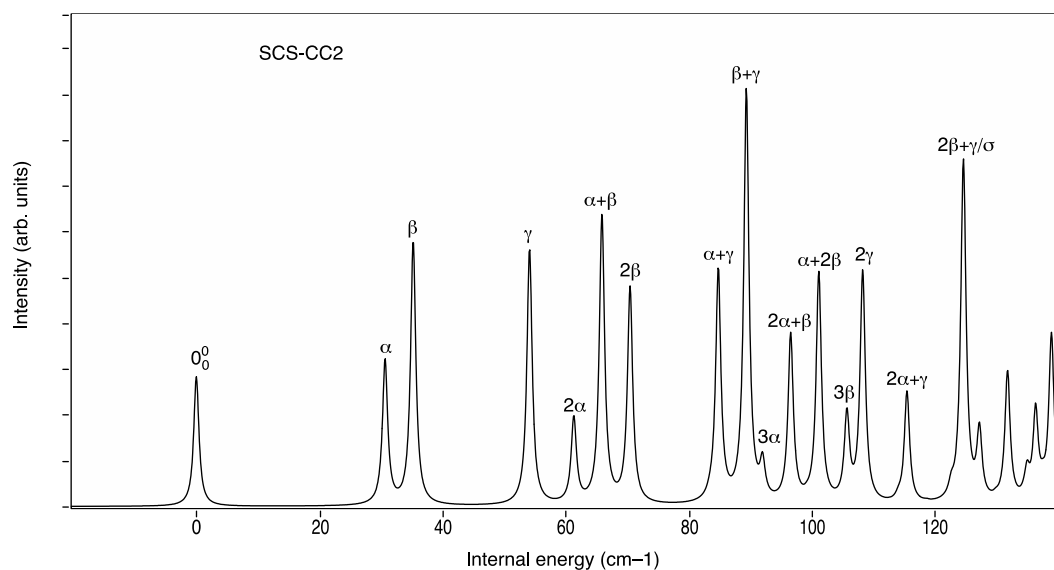


Fig. S2. Franck-Condon simulated electronic spectrum for the HB indole...imidazole structure obtained from calculations at the SCS-CC2/def2-TZVPP level of theory.

OPTIMIZED GEOMETRIES

HB structure

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S0 structure for HB indole...imidazole M05-2X/6-31+G*

C	-1.250493	0.135229	-0.320761
C	-2.499175	0.567452	0.192973
C	-3.512097	-0.383648	0.402822
C	-3.259699	-1.711840	0.105660
C	-2.009438	-2.119312	-0.405869
C	-0.991575	-1.206660	-0.627727
C	-1.132555	2.344621	-0.005185
C	-2.396311	1.986818	0.383518
H	-4.476959	-0.079661	0.791524
H	-4.033484	-2.453425	0.262562
H	-1.846011	-3.165258	-0.635761
H	-0.034865	-1.512176	-1.034307
H	0.537729	1.208940	-0.706140
H	-0.662801	3.315437	-0.012370
H	-3.159148	2.651357	0.756642
N	-0.441232	1.238957	-0.435886
C	4.420947	-0.460629	0.066574
C	3.730038	0.297395	-0.839805
C	2.324107	-0.208641	0.697834
N	3.506252	-0.776378	1.043757
H	3.678471	-1.328418	1.867350
H	5.443793	-0.792854	0.104012
H	4.092394	0.740052	-1.752732
H	1.427351	-0.303581	1.291079
N	2.425269	0.447612	-0.436254

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S1 structure for HB indole...imidazole TD-M05-2X/6-31+G*

C	-1.267185	0.090100	-0.249883
C	-2.518336	0.588171	0.151494
C	-3.604554	-0.301415	0.395751
C	-3.350037	-1.687115	0.175984
C	-2.113725	-2.150736	-0.221680
C	-0.984875	-1.265686	-0.435582
C	-1.042415	2.334175	-0.137899
C	-2.390481	1.997284	0.212826
H	-4.571536	0.058506	0.716016
H	-4.153431	-2.397739	0.327535
H	-1.971972	-3.212578	-0.381785
H	-0.047300	-1.620390	-0.838727
H	0.593756	1.100210	-0.610278
H	-0.551086	3.291190	-0.185053
H	-3.156147	2.712288	0.472295
N	-0.409865	1.192647	-0.408253
C	4.486992	-0.273641	0.039224
C	3.711589	0.575084	-0.702888
C	2.365930	-0.459777	0.613015
N	3.610590	-0.926655	0.873870
H	3.844899	-1.630227	1.554264
H	5.545531	-0.466514	0.047475
H	4.022013	1.261276	-1.473144
H	1.479022	-0.817858	1.115332
N	2.392852	0.449950	-0.337229

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Fitted S1 structure for HB indole...imidazole

C	-1.25475	0.110198	-0.242225
C	-2.50874	0.574995	0.193060
C	-3.60276	-0.328284	0.340665
C	-3.35300	-1.68718	-0.011485
C	-2.11374	-2.11778	-0.435364
C	-0.97717	-1.22424	-0.543397
C	-1.02025	2.33076	0.090096
C	-2.37322	1.97074	0.397205
H	-4.57317	0.006467	0.679030
H	-4.16548	-2.40103	0.056740
H	-1.97753	-3.15830	-0.698634
H	-0.028661	-1.54915	-0.946877
H	0.609976	1.14135	-0.506418
H	-0.523559	3.28556	0.133992
H	-3.13546	2.66330	0.719689
N	-0.393525	1.21733	-0.286617
C	4.47861	-0.347251	-0.017466
C	3.70998	0.534629	-0.729859
C	2.37032	-0.465891	0.630336
N	3.60898	-0.978956	0.844239
H	3.83757	-1.69989	1.51194
H	5.53089	-0.576895	-0.049346
H	4.02115	1.22000	-1.50781
H	1.48767	-0.800177	1.16371
N	2.39972	0.452004	-0.315660

T1 structure

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S0 structure for T1 indole...imidazole M05-2X/6-31+G*

C	-1.415724	-0.191099	0.745852
C	-1.675443	-0.168843	-0.647508
C	-1.665384	1.066061	-1.319636
C	-1.402666	2.220439	-0.598850
C	-1.153022	2.172070	0.790352
C	-1.156245	0.971350	1.479665
C	-1.745226	-2.303196	0.064712
C	-1.884195	-1.530502	-1.055186
H	-1.858438	1.112230	-2.384567
H	-1.394598	3.178087	-1.104230
H	-0.951646	3.091687	1.325164
H	-0.951749	0.934239	2.542494
H	-1.278226	-1.832717	2.080079
H	-1.826192	-3.371964	0.181260
H	-2.110313	-1.885865	-2.047533
N	-1.467027	-1.503215	1.149345
C	1.805970	-0.924898	-0.399447
C	3.104928	-1.156894	-0.023753
C	2.872795	0.964422	-0.040412
N	1.675495	0.440936	-0.409330
H	0.830422	0.954760	-0.615064
H	0.988253	-1.577796	-0.655969
H	3.602576	-2.105651	0.095729
H	3.040206	2.026972	0.036667
N	3.762244	0.028361	0.198731

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S1 structure for T1 indole...imidazole M05-2X/6-31+G*

C	-0.946005	0.016517	0.801286
C	-1.316537	0.709061	-0.396188
C	-0.561796	1.858496	-0.811213
C	0.467156	2.276044	0.059223
C	0.808212	1.575199	1.229703
C	0.072497	0.451592	1.682112
C	-2.733590	-1.034792	-0.096764
c	-2.441127	0.025574	-0.933700
H	-0.894845	2.478209	-1.633264
H	1.025341	3.175496	-0.179446
H	1.615311	1.956313	1.843912
H	0.324748	-0.080035	2.588193
H	-1.862749	-1.641756	1.740695
H	-3.508158	-1.781583	-0.157327
H	-2.982964	0.282002	-1.830951
N	-1.817240	-1.044081	0.933461
C	0.624374	-1.288414	-0.904854
C	1.414473	-1.856062	0.134451
C	2.433495	-0.120380	-0.531362
N	1.355273	-0.241524	-1.366490
H	0.900045	0.572166	-1.799851
H	-0.288857	-1.626279	-1.368822
H	1.190281	-2.744467	0.702858
H	3.170752	0.654631	-0.651048
N	2.516850	-1.131781	0.340012

T2 structure

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S0 structure for T2 indole...imidazole M05-2X/6-31+G*

C	1.341749	0.342454	0.761821
C	1.573426	0.516533	-0.623831
C	2.024548	-0.579412	-1.380992
C	2.229357	-1.790841	-0.745619
C	1.993832	-1.938883	0.638312
C	1.547501	-0.880819	1.409739
C	0.832747	2.465455	0.237373
C	1.240432	1.881822	-0.933701
H	2.206122	-0.474276	-2.443702
H	2.575271	-2.643692	-1.315833
H	2.160089	-2.902211	1.103652
H	1.354772	-0.998750	2.468894
H	0.599308	1.709954	2.203942
H	0.508438	3.475566	0.430045
H	1.315353	2.369768	-1.892476
N	0.895693	1.546244	1.257278
C	-3.034857	0.746219	-0.271477
C	-3.739019	-0.362938	0.122200
C	-1.737682	-1.030234	-0.210671
N	-1.753008	0.300862	-0.481428
H	-0.957544	0.855186	-0.767248
H	-3.320507	1.773398	-0.421318
H	-4.784021	-0.429113	0.378604
H	-0.839563	-1.622189	-0.295295
N	-2.921247	-1.465645	0.157265

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S1 structure for T2 indole...imidazole M05-2X/6-31+G*

C	0.900715	0.926177	0.444038
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C	0.787538	0.534261	-0.903647
C	1.629251	-0.497954	-1.415810
C	2.623955	-1.006736	-0.528080
C	2.735391	-0.566555	0.778877
C	1.864584	0.428251	1.331860
C	-0.865866	2.062584	-0.411680
C	-0.320946	1.238791	-1.443233
H	1.610760	-0.771115	-2.462383
H	3.317261	-1.756808	-0.889576
H	3.512162	-0.982642	1.409309
H	1.953620	0.768780	2.353691
H	-0.316629	2.257458	1.588837
H	-1.693790	2.750289	-0.427482
H	-0.682476	1.209506	-2.459533
N	-0.122779	1.856101	0.684844
C	-1.920270	-1.115850	-0.880476
C	-2.725858	-0.384510	-0.011324
C	-1.133593	-1.171213	1.159439
N	-0.941937	-1.653026	-0.102696
H	-0.041725	-1.968143	-0.465760
H	-1.994328	-1.314057	-1.936233
H	-3.636971	0.142872	-0.243814
H	-0.448315	-1.388212	1.962344
N	-2.225137	-0.436132	1.253685

Syn-PD structure

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S0 structure for Syn-PD indole...imidazole M05-2X/6-31+G*

C	0.950318	0.018452	-0.709022
C	1.607119	0.946978	0.134938
C	2.773528	0.545046	0.808458
C	3.234915	-0.748845	0.638349
C	2.558370	-1.659283	-0.200458
C	1.412418	-1.290512	-0.885133
C	-0.208643	1.934572	-0.759852
C	0.846280	2.166257	0.080551
H	3.302097	1.239194	1.451101
H	4.132149	-1.070543	1.152427
H	2.945798	-2.664145	-0.314823
H	0.892856	-1.986105	-1.533563
H	-0.884655	0.191241	-1.753905
H	-1.012728	2.588293	-1.057156
H	1.065208	3.094142	0.584563
N	-0.142825	0.652580	-1.251009
C	-1.747946	-1.348760	0.520307
C	-3.377287	-0.184950	-0.227727
C	-2.826683	0.550207	0.789671
N	-1.788088	-0.212528	1.262341
H	-1.103807	0.068636	1.946149
H	-1.004588	-2.113315	0.678325
H	-4.228966	0.061554	-0.840865
H	-3.069629	1.512819	1.205434
N	-2.693097	-1.368157	-0.391071

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S1 structure for Syn-PD indole...imidazole M05-2X/6-31+G*

C	0.892201	0.369685	-0.905393
C	1.031225	1.015622	0.335923
C	1.997047	0.548408	1.276480
C	2.815546	-0.539728	0.858892
C	2.673510	-1.138893	-0.378988
C	1.672841	-0.712907	-1.323720
C	-0.813326	1.867232	-0.760605
C	-0.031504	1.956051	0.430392
H	2.153136	1.047085	2.222644
H	3.580542	-0.907439	1.532528
H	3.326998	-1.959456	-0.648601
H	1.563279	-1.176524	-2.294140
H	-0.592173	0.633435	-2.425935
H	-1.667689	2.435573	-1.085513
H	-0.232958	2.633100	1.246584
N	-0.241693	0.923490	-1.526051
C	-1.588924	-1.550001	-0.373069
C	-2.798988	-0.115070	0.630641
C	-1.640370	-0.255100	1.392758
N	-0.903847	-1.194807	0.745724
H	0.062531	-1.424686	0.950571
H	-1.198765	-2.264624	-1.079326
H	-3.644830	0.522475	0.830828
H	-1.316842	0.194859	2.315224
N	-2.752858	-0.929480	-0.457491