Supplementary Information

Multidimensional Characterization of the Conical Intersection Seam in the Normal Mode Space

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SI1. SEVI (slow electron velocity map imaging) spectra via selected S₁ vibrations



SI1-1. Ionization potentials for thioanisole isotopomers

Figure S1. Determination of ionization potentials of two thioanisole isotopomers (upper). 2D images (lower) are shown.

For conducting SEVI experiments (Figure S2), two-color threshold-ionization via each 0_0^0 transition of two isotopomers was done as Figure S1. Ionization potential of each isotopomer is derived from the extrapolation of the measured radial displacement of the ring in VMI images with respect to the two-photon input energies. Ionization potentials of all six thioanisole isotopomers are summarized in Table S1.

Table S1. Experimental S₁ origins, and D₀ origins (ionization potential)

	[‡] C ₆ H ₅ S-CH ₃	[‡] C ₆ H ₅ S-CD ₃	[‡] C ₆ H ₅ S- CH ₂ D ₁	[‡] C ₆ H ₅ S- CH ₁ D ₂	C ₆ D ₅ S-CH ₃	C ₆ D ₅ S-CD ₃
S ₁ origin [cm ⁻¹]	34 504	34 516	34 504, 34 510	34 510, 34 516	34 664	34 677
D ₀ origin [cm ⁻¹]	63 899	63 917	63 904, 63 915	63 909, 63 919	63 896	63 902

* Supporting Ref. 1 * There are two isomers for each partially deuterated Thioanisole-h₂d₁ and Thioanisole-h₁d₂, respectively.





Figure S2. 2 color (1+1') SEVI spectra via selected S_1 vibronic states of (a) C_6D_5S -CD₃ and (b) C_6D_5S -CH₃. Insets show 2D images having comparable ring radius for the peak in each SEVI spectrum whereas the actual reconstruction of images was carried by MEVELER program^{S2} and plotted. Linear polarization axis is shown as red arrow.

 $S_1 \nu_s$ and 7a modes of two isotopomers are assigned by aids of previous reference values^{S3-S6} with SEVI experiments (Figure S2) in this work. Although definitive assignment of the full S_1 normal modes could not be achieved because of limited numbers of SEVI spectra via full S_1 vibronic bands, there are already excellent spectroscopic works for thioanisole molecules.^{S3-S6} and authors did reliable assignment for the $S_1 \nu_s$ and 7a modes for both C_6D_5S -CD₃ and C_6D_5S -CH₃ by getting SEVI spectra. Subsequent comparisons of SEVI peaks (D_0 - S_1 peaks) with calculated values (TDDFT) as well as R2PI peaks engenders the reliable assignments keeping in mind the assumption that the vibronic selection rule ($\Delta \nu = 0$) between D_0 and S_1 is obeyed. Detailed numbers are summarized in Table S2.

SI1-3. Vibrational mode assignments

Table S2. Comparison of calculation and experiment. Normal modes are labelled according to the Varsanyi notation. S_0 , D_0 , S_1 vibrational energies $[cm^{-1}]$ are calculated using the *Gaussian09* package. The (TD) DFT B3LYP level with 6-311++G(3df,3pd) basis set was used. Assignments are done by comparing previous thioanisole references^{S3-S6} with the data of this work (R2PI, TDDFT calculation, and SEVI).

	C	6D5S-CH	I ₃ [cm ⁻¹]				C	C ₆ D ₅ S-CI	D ₃ [cm ⁻¹]		
mode	S ₀	D ₀	S ₁	R2PI	SEVI	mode	S ₀	D ₀	S ₁	R2PI	SEVI
τ	48	77	-59		90	τ	43	68	-55		
10b	159	185	124	123	177	10b	139	107	80		
15	191	199	191			15	176	183	176		
τCH_3	228	147	124			τCD ₃	186	107	90		
$\beta_{\rm s}$	322	326	325			$\beta_{\rm s}$	309	313	312		
16a	359	341	485			16a	358	341			
6a	412	404	410			6a	405	395			
16b	424	386	427			16b	424	386			
6b	605	578				6b	605	578	524		
4	553	515				4	553	515	485		
$v_{\rm s}$	670	660	674	656	669	v _s	656	642	658	642	651
7a	724	689	720	712	711	7a	691	674	693	682	702
11						$v_{s}\tau^{1}$			709	686	727
10a						10a					
y _s CH ₃	972	924	913			$\gamma_{s}CD_{3}$	734	700	688		
17b	760	857	623			17b	760	788	623		
12	957	955	938			12	957	955	937	918	
$\beta_{as}CH_3$	984	981	962			$\beta_{as}CD_3$	773	767	764		
17a	803	827				17a	803	827			
1						1					
5	827	857				5	827	857			
18a						18a					
18b						18b					
9b	856	862				9b	856	862			
9a	885	896	862			9a	885	896	862		
14						14			834		
$\beta_{\rm s} \rm CH_3$	1358	1362	1347			$\beta_{\rm s} {\rm CD}_3$	1032	1034	1025		
3						3					
$\beta_{\rm s} \rm CH_2$	1486	1455	1467			$\beta_{\rm s} {\rm CD}_2$	1076	1046	1056		
19b	842	852	802	768	840	19b	842	855	804		
$\gamma_{as}CH_3$	1469	1459				$\gamma_{as}CD_3$	1062	1054	1049		

 \times 1→9a, 9b↔14 (assignment change from Supporting Ref. 3,4 to this study)





Figure S3. (a), (d) full translational energy (total kinetic energy release; TKER) distributions. TKER distributions of (a) $C_6D_5S-CD_3$ and (d) $C_6D_5S-CH_3$ via selected S_1 vibronic transitions followed by fragmentated into (a) $\cdot CD_3$ and (d) $\cdot CH_3$ detected by (2+1) REMPI via their $3P_z$ state, respectively. Ordinate: normalized population, black-hollow circle: experiment, red: manually fitted \tilde{A} component, blue: gaussian fitted \tilde{X} component, emerald: gaussian fitted two-photon background. Sum of fitted two components (\tilde{A} and \tilde{X}) is drawn in black solid line. \tilde{X}/\tilde{A} branching ratios of (b) $C_6D_5S-CD_3$ and (e) $C_6D_5S-CH_3$ and averaged anisotropy parameters (β_2) of (c) $C_6D_5S-CD_3$ and (f) $C_6D_5S-CH_3$ are shown.

Authors first deconvoluted multiphoton backgrounds from the reconstructed TKER distributions. Considering estimated maximum available TKER for A and X products, ^{S8} most evident X component in TKER distribution via 7a mode was fitted as a single gaussian function. The rest of that TKER distribution was assumed as A products. Authors call this as 'manual deconvolution'. Whereas, to calculate uncertainties, TKER distributions are fitted as a sum of two individual gaussian functions. Error bars for X/A were then calculated from the $\pm 1\sigma$ fitting errors of these two Gaussian functions. Based on the above 'manual' deconvolution concept, corresponding averaged anisotropy parameters were calculated for two components as well as totally (full range of TKER) averaged anisotropies with respect to the S₁ internal energy, Figure S3 (c) and (f). Contrast features of X components of two isotopomers could be recognized. This infers that the C₆D₅S-CH₃ dissociates faster then the C₆D₅S-CD₃ does. Wider vibronic band widths of C₆D₅S-CH₃ than the corresponding narrower widths of C₆D₅S-CD₃ parent R2PI and fragment PHOFEX in Figure 2 strongly support this interpretation.



SI2-2. Three-component $(\tilde{A}_1, \tilde{A}_2, \tilde{X})$ gaussian function-aided deconvolution

Figure S4. (a), (d) full translational energy (total kinetic energy release; TKER) distributions. TKER distributions (upper) of (a) $C_6D_5S-CD_3$ and (d) $C_6D_5S-CH_3$ via S_1 vibronic transitions followed by fragmentated into (a) $\cdot CD_3$ and (d) $\cdot CH_3$ detected by (2+1) REMPI via their $3P_z$ state, respectively. Ordinate: normalized population, black-hollow circle: experiment, red: gaussian fitted \tilde{A}_1 component, orange: gaussian fitted \tilde{A}_2 component, blue: gaussian fitted \tilde{X} component, emerald: two-photon background. Sum of simulated three components (\tilde{A}_1, \tilde{A}_2 , and \tilde{X}) is drawn in black solid line. Branching ratios of (b) $C_6D_5S-CD_3$ and (e) $C_6D_5S-CH_3$ are shown.

From the previously-reported time-resolved work, it has been found that the bifurcation starting from the S_1/S_2 conical intersection ends up with two distinct reaction pathways. These pathways are very different in terms of its X/A product branching ratio as well as the translational energy distribution of the A state. Although the bifurcation ratio should be linearly proportional to the X/A product branching ratio (as above in (5)), they are quantitatively different. This is because that one channel (vibrational predissociation) gives rise to the A state almost exclusively whereas the other channel (electronic predissociation) produces both A and X products. In the former case of vibrational predissociation, the A state is internally highly excited, giving the low translational energy distribution (A₁). On the other hand, the A state from the electronic predissociation is much less excited internally, giving the relatively higher kinetic energy distribution (A₂). Based on these experimental facts, the fit to the experiment is given to extract the bifurcation ratios instead of the final product X/A branching ratios. However, since the current experimental results are not time-resolved, the deconvolution is intrinsically ambiguous to give large uncertainties during the fit.

SI3. Sample information (NMR, Chromatography)

C₆D₅S-CH₃



Instrument:U-3000_2 Sequence:NMA-3-45



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C₆D₅S-CD₃



Instrument:U-3000_2 Sequence:NMA

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njection Nan	ne:	2-NMA-3-49-1			Run Time (min):	20.00	
rial Number:	e.	2 Unknown			Injection Volume:	20.00	
Calibration	evel:	UNKNOWN				254.0	
nstrument M	lethod:	ACN100%-20min-6	0.5	5		n.a.	
Processing N	Nethod:	ACN100%-20min-	1.0		Dilution Factor:	1.0000	
njection Date	e/Time:	14-Oct-16 15:59	14-Oct-16 15:59			1.0000	
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0.0	2.0	4.0 0.0	Tir	me [min]		10,0 10,0	
ntegration	Results	1 e-1	1.50	1	L D-L-K -	Deleter House	
vo. Peak	Name	Retention Time min	Area mAU*min	Height mAU	Relative Area	Relative Height	Amount n.a.
1	den se mente	4 148	246 213	1593 205	97.17	96.80	n.a.
2		4.715	0.450	4 154	0.18	0.25	na
3		4 907	2 315	19 205	0.91	1.17	na
4		4.997	3,000	22 099	1 22	1.40	n.a.
		0.012	0.552	3 704	0.22	0.33	n.a.
5		13 787	0.552	0.124	0.22	0.25	
5		10.101	0.757	2.409	0.30	0.15	n.a.

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Supplementary References

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