

Supporting Information

$[\text{Mo}_3\text{S}_{13}]^{2-}$ as a Model System for Hydrogen Evolution Catalysis by MoS_x : Probing Protonation Sites in the Gas Phase by Infrared Multiple Photon Dissociation Spectroscopy

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Abstract: Molybdenum sulfide based materials are known as efficient hydrogen evolution reaction (HER) catalysts. As the binding site for H atoms on molybdenum sulfide for the catalytic process is under debate, $[\text{HMo}_3\text{S}_{13}]^-$ is an interesting molecular model system. Herein, we probe the $[\text{HMo}_3\text{S}_{13}]^-$ cluster in the gas phase by coupling Fourier transform ion cyclotron resonance mass spectrometry (FT-ICR MS) with infrared multiple photon dissociation (IRMPD) spectroscopy. Our investigations show one distinct S-H stretching vibration at 2450 cm^{-1} . Thermochemical arguments based on DFT calculations strongly suggest a terminal disulfide unit as the H adsorption site.

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Experimental Section

The experimental setup has been described in detail previously.^[1,2] Photodissociation spectroscopy was performed in ultrahigh vacuum ($\approx 10^{-9}$ mbar) inside the ICR cell of a commercial Bruker Apex Qe FT-ICR MS, equipped with a combined electrospray ionization (ESI) and matrix assisted laser desorption ionization (MALDI) Apollo II Dual ESI/MALDI ion source along with a 9.4 T superconducting magnet. Tuneable IR laser light from optical parametric oscillators (OPO) EKSPLA NT277 and NT273, operated at a 1 kHz repetition rate, is guided through a CaF₂ window at the rear end of the vacuum chamber into the ICR cell.^[1,2] Electrospray ionization of (NH₄)₂[⁹²Mo₃S₁₃] dissolved in water-methanol (1:1) yields mainly [Mo₃S₁₃]²⁻ and [HMo₃S₁₃]. Subsequently the protonated clusters [HMo₃S₁₃]⁻ were preselected by a quadrupole mass filter and guided into the ICR cell, where they were trapped for irradiation. The IR photodissociation ratio was obtained by recording a mass spectrum at each set wavelength, yielding an IR action spectrum. The wavelength of the OPO was calibrated with an accuracy of <2 nm by a laser spectrum analyser Ångstrom LSA IRIII L (HighFinesse). (NH₄)₂[⁹²Mo₃S₁₃] was prepared from isotopically enriched ⁹²MoO₃ (STB Isotope Germany), following a procedure of Müller et al.,^[3] as described previously.^[4] Quantum chemical calculations were performed using the ω B97XD functional, with the vibrational scaling factor of 0.957 (see the SI for benchmark calculations). The Gaussian software was used for calculations.^[5]

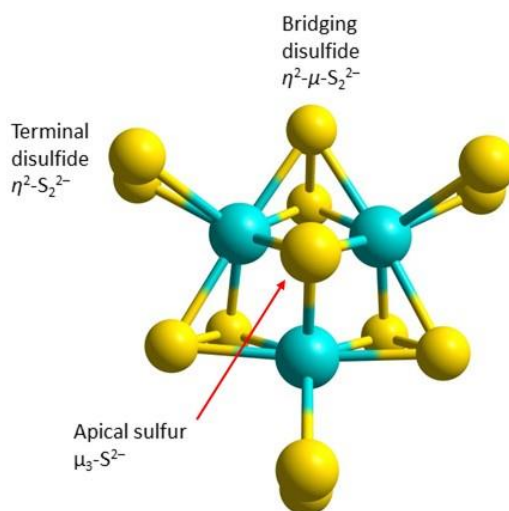
[Mo₃S₁₃]²⁻ Geometry

Figure S1. Different sulfur moieties of the [Mo₃S₁₃]²⁻ cluster.

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Additional Experimental Data

Table S1. Gauss functions used to model the experimental data inFigure 3b. Fitting model: $G(x) = \frac{A}{w\sqrt{\pi/2}} e^{-2\left(\frac{x-x_c}{w}\right)^2}$

	x_c / cm^{-1}	w / cm^{-1}	$A / 10^{-19} \text{ cm}$	R^2
G	2451.7 ± 0.1	10.2 ± 0.2	5.6 ± 0.1	0.974
G1	2449 ± 5	10 ± 3	3.1 ± 3.4	0.984
G2	2454 ± 1	7 ± 2	2.5 ± 3.4	

Additional Computational Data

Table S2 shows calculations of S–H vibrations for several molecules containing S–H bonds. Based on this calculations, we can conclude that scaling factors of 0.968 and 0.957 for B3LYP and ω B97XD from NIST^[6] lie within 1% of the values calculated for the tested molecules and are therefore used also in our calculations. For the M06, we pick the factor of 0.967.

Table S2. Unscaled S–H vibrational frequencies in several molecules along with experiment (in cm^{-1}). The aug-cc-pVTZ basis set was used for all calculations.

	CCSD	B3LYP	M06	ω B97XD	exp.
H ₂ S	2727.9; 2742.8	2679.0; 2692.1	2695.7; 2705.9	2743.7; 2756.2	2614.6; 2626 ^[7]
H ₂ S ₂	2696.2; 2697.8	2627.5; 2629.6	2639.2; 2642.2	2689.4; 2691.2	2553.8; 2556.6 ^[8]
CH ₃ SH	2742.8	2674.3	2693.1	2735.5	2597 ^[9]

Table S3. Relative energies ΔE (in eV) and vibrational frequencies ν (in cm^{-1}) calculated for various [HMO₃S₁₃]⁻ isomers using the aug-cc-pVTZ(-PP) basis set. Scaling factors of 0.968 (B3LYP), 0.967 (M06) and 0.957 (ω B97XD) were used.

Isomer	ΔE			ν		
	B3LYP	M06	ω B97XD	B3LYP	M06	ω B97XD
A	1.26	1.31	1.31	2477	2506	2552
T1	0.00	0.00	0.00	2483	2478	2513
T2	0.10	0.09	0.09	2479	2489	2509
B	1.27	1.39	1.51	2409	2411	2442

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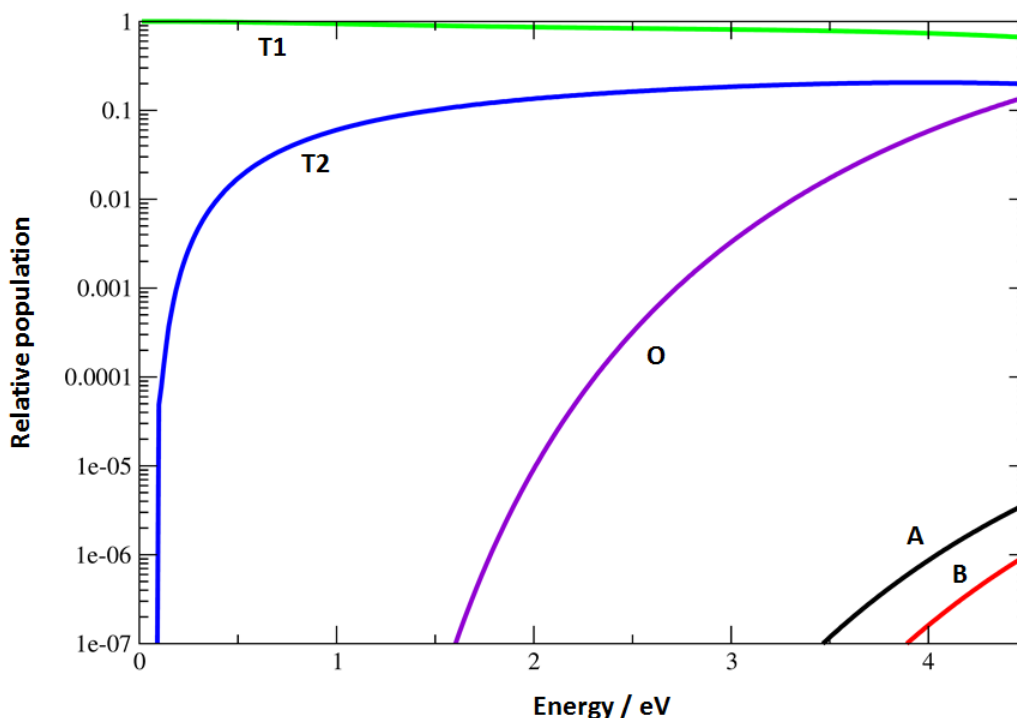


Figure S2. Relative population of various $[\text{HMo}_3\text{S}_{13}]^-$ isomers calculated through the density of states at the B3LYP/aug-cc-pVTZ(-PP) level of theory.

Cartesian coordinates (in Å) and electronic energies (in a.u.) of various $\text{HMo}_3\text{S}_{13}^-$ structures optimized using various DFT functionals; see methods for more computational details

Isomer A, wB97XD

E= -5381.897509

S 0.000239 -0.000415 1.776810
 Mo -1.498590 -0.530315 0.031039
 S -1.440442 1.229234 -1.625026
 S -2.199108 1.878991 0.138971
 Mo 0.289787 1.562598 0.030637
 S 2.726452 0.965191 0.136853
 S 1.784089 0.631809 -1.626596
 Mo 1.209018 -1.032334 0.030321
 S -0.345025 -1.859484 -1.625758
 S -0.527213 -2.844038 0.136658
 S 0.652128 3.510800 1.391252
 S 0.718950 3.878458 -0.640770
 S 3.000020 -2.562659 -0.640232
 S 2.715182 -2.318846 1.391729
 S -3.719000 -1.316373 -0.641235
 S -3.367037 -1.192476 1.390858
 H 0.003216 -0.000961 3.119923

Isomer T2, wB97XD

E= -5381.942227

S 0.022032 -0.002060 1.807039
 Mo -1.529411 -0.004816 -0.004760
 S -0.963314 1.672129 -1.640744
 S -1.440509 2.477777 0.165665
 Mo 0.799332 1.350548 0.011270
 S 2.885296 0.009220 0.086197
 S 1.902615 0.006634 -1.691818
 Mo 0.808950 -1.345660 0.010942
 S -0.948032 -1.674478 -1.644105
 S -1.425198 -2.489506 0.160282

S 1.768567 2.981633 1.469582
 S 2.008564 3.424358 -0.538861
 S 2.032135 -3.412205 -0.537404
 S 1.787558 -2.969824 1.470690
 S -3.976551 -0.024272 -0.640639
 S -3.628344 0.076936 1.380163
 H -3.709697 -1.224538 1.730308

Isomer T1, wB97XD

E= -5381.945514

S -0.015327 -0.005581 1.799838
 Mo 1.516065 -0.249743 0.001894
 S 0.662319 -1.788840 -1.665087
 S 1.016218 -2.698581 0.125139
 Mo -1.006593 -1.208731 0.006213
 S -2.844432 0.453379 0.112355
 S -1.899505 0.301988 -1.678890
 Mo -0.578755 1.451464 0.009572
 S 1.186283 1.478425 -1.665070
 S 1.821417 2.231330 0.123141
 S -2.215245 -2.668318 1.469364
 S -2.548537 -3.046655 -0.538547
 S -1.466978 3.681871 -0.530626
 S -1.264005 3.212545 1.475858
 S 4.019650 -0.631938 -0.505112
 S 3.457588 -0.542330 1.476205
 H 4.358726 0.657651 -0.719579

Isomer B, wB97XD

E= -5381.890017

S -0.097476 0.026825 -1.832585
 Mo 1.334098 -0.728703 -0.046263

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S 2.183395 0.582482 1.968813
 S 2.366707 1.568415 0.152389
 Mo -0.129773 1.598307 -0.096350
 S -2.585381 1.379862 0.114525
 S -1.502179 0.872628 1.782901
 Mo -1.359851 -0.795413 0.017056
 S 0.048520 -2.003518 1.570928
 S 0.039681 -2.846679 -0.274823
 S 0.054921 3.437477 -1.503972
 S -0.206453 3.971986 0.443113
 S -3.386479 -2.040499 0.633040
 S -3.079937 -1.692072 -1.385710
 S 3.464691 -1.984489 0.224630
 S 2.896234 -1.465106 -1.687672
 H 3.392206 -0.033019 1.984196

Isomer O, wB97XD

E= -5381.913185
 S 0.858281 -1.395436 -0.718545
 Mo 3.008797 -0.550678 0.119527
 S 4.623850 0.824643 -1.015524
 S 3.312940 -0.241253 -2.200892
 S 1.470298 -0.087296 1.888680
 S 3.106984 1.155919 1.794572
 Mo -0.449842 0.340210 0.050553
 S -0.201980 2.630660 0.695121
 S 0.907161 2.317005 -1.017846
 Mo -3.109485 -0.478276 -0.116447
 S -4.975365 0.226454 1.167853
 S -5.016297 0.734748 -0.846613
 S -3.232335 -2.509812 -0.636081
 S -1.755162 -0.333420 1.769982
 S -1.834749 0.593512 -1.729721
 S 4.048167 -2.246891 0.736250
 H 2.133606 2.225902 -0.448403

Isomer A, B3LYP

E= -5382.158739
 S -0.000813 0.000336 1.779419
 Mo 1.582756 -0.348045 0.032644
 S 0.579995 -1.827340 -1.645113
 S 0.886290 -2.794592 0.130294
 Mo -1.092920 -1.196298 0.031758
 S -2.863531 0.629859 0.127196
 S -1.871056 0.411382 -1.647303
 Mo -0.489954 1.544327 0.031472
 S 1.293163 1.414924 -1.645704
 S 1.977112 2.164895 0.129449
 S -2.430317 -2.659455 1.411814
 S -2.692151 -2.947578 -0.637188
 S -1.206661 3.805298 -0.636789
 S -1.089575 3.433262 1.412112
 S 3.898640 -0.857422 -0.637418
 S 3.519581 -0.773718 1.411767
 H -0.005853 0.003049 3.132677

Isomer T2, B3LYP

E= -5382.201574
 S 0.025147 -0.002012 1.817437
 Mo -1.546751 -0.003071 -0.004156
 S -0.978737 1.693441 -1.656476
 S -1.460243 2.510795 0.163558
 Mo 0.816299 1.371517 0.013169
 S 2.919666 0.004937 0.068192
 S 1.920768 0.003953 -1.722035
 Mo 0.822356 -1.368809 0.013201
 S -0.966082 -1.692437 -1.660018
 S -1.453761 -2.517120 0.157649
 S 1.789822 3.000821 1.488828
 S 2.037385 3.458552 -0.534071
 S 2.050718 -3.452552 -0.532067

S 1.800198 -2.993931 1.490446
 S -4.022354 -0.020433 -0.643017
 S -3.668888 0.083305 1.393807
 H -3.758197 -1.221869 1.751280

Isomer T1, B3LYP

E= -5382.205143
 S -0.018181 -0.004990 1.808468
 Mo 1.535165 -0.237003 0.001789
 S 0.695113 -1.801111 -1.684208
 S 1.058523 -2.723642 0.118246
 Mo -1.012709 -1.238719 0.006389
 S -2.880225 0.431287 0.097591
 S -1.922844 0.285542 -1.709168
 Mo -0.604909 1.470664 0.009973
 S 1.190073 1.504916 -1.683103
 S 1.828237 2.277064 0.115839
 S -2.205651 -2.708422 1.490965
 S -2.548275 -3.105484 -0.530596
 S -1.520977 3.709233 -0.523194
 S -1.305884 3.223473 1.496895
 S 4.084436 -0.604057 -0.495879
 S 3.485578 -0.513732 1.494660
 H 4.424236 0.691209 -0.706591

Isomer B, B3LYP

E= -5382.158636
 S -0.108046 0.028767 -1.842675
 Mo 1.343798 -0.757260 -0.053766
 S 2.267408 0.555926 2.000018
 S 2.391425 1.569775 0.162104
 Mo -0.113901 1.619036 -0.098360
 S -2.601582 1.422986 0.109654
 S -1.514116 0.910991 1.797332
 Mo -1.393125 -0.795179 0.019626
 S 0.031976 -2.014433 1.603217
 S 0.007616 -2.890039 -0.248411
 S 0.107572 3.482611 -1.511293
 S -0.173504 4.023293 0.443548
 S -3.447637 -2.026234 0.636786
 S -3.123303 -1.679883 -1.398647
 S 3.485151 -2.050265 0.194007
 S 2.888613 -1.503280 -1.720098
 H 3.470413 -0.080512 1.956333

Isomer A, M06

E= -5381.696964
 S -0.000653 0.000098 1.789804
 Mo 1.564838 -0.368530 0.036271
 S 0.549567 -1.824603 -1.636541
 S 0.838156 -2.786084 0.129654
 Mo -1.101625 -1.170715 0.035527
 S -2.832177 0.667167 0.128729
 S -1.855062 0.436855 -1.637494
 Mo -0.463310 1.539172 0.035473
 S 1.306410 1.387899 -1.636516
 S 1.993534 2.119286 0.129950
 S -2.459851 -2.614190 1.393038
 S -2.712634 -2.884420 -0.641348
 S -1.140375 3.791589 -0.641254
 S -1.035169 3.436918 1.393263
 S 3.853734 -0.907043 -0.642031
 S 3.494995 -0.823366 1.392685
 H -0.003530 0.001381 3.143574

Isomer T2, M06

E= -5381.741559
 S 0.025315 -0.001964 1.822606
 Mo -1.538601 -0.006140 0.000253
 S -0.979896 1.681682 -1.647697
 S -1.450800 2.489307 0.163457

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Mo 0.806555 1.364497 0.013407
S 2.901555 0.010839 0.069190
S 1.908837 0.007633 -1.708731
Mo 0.817950 -1.358591 0.013529
S -0.960777 -1.685005 -1.651304
S -1.433281 -2.501711 0.157981
S 1.775256 2.994148 1.472773
S 2.014696 3.437189 -0.536299
S 2.041784 -3.423032 -0.534098
S 1.797771 -2.979959 1.474563
S -3.987514 -0.028894 -0.641637
S -3.646926 0.077614 1.380031
H -3.704297 -1.235748 1.724694

Isomer T1, M06

E= -5381.744950

S -0.017125 -0.005137 1.814650
Mo 1.522674 -0.269766 0.005812
S 0.654845 -1.808080 -1.673025
S 0.987993 -2.724397 0.122154
Mo -1.030647 -1.209219 0.007645
S -2.852947 0.489559 0.099173
S -1.904995 0.324678 -1.695409
Mo -0.566928 1.472524 0.010582
S 1.222348 1.470679 -1.672251
S 1.856086 2.218472 0.122327
S -2.254967 -2.651977 1.474344
S -2.594112 -3.027445 -0.533924
S -1.428926 3.711737 -0.527082
S -1.230353 3.239317 1.479748
S 4.017658 -0.675742 -0.507090
S 3.469753 -0.583871 1.478532
H 4.341721 0.626678 -0.723991

Isomer B, M06

E= -5381.693750

S -0.099624 0.028849 -1.848734
Mo 1.303686 -0.807251 -0.047416
S 2.212883 0.498612 1.965989
S 2.462739 1.436001 0.123553
Mo -0.043074 1.613956 -0.102558
S -2.521924 1.517642 0.096005
S -1.471719 0.967620 1.777242
Mo -1.414049 -0.733594 0.013916
S -0.055202 -2.003516 1.598056
S -0.111273 -2.866873 -0.243807
S 0.250867 3.484968 -1.485959
S 0.005911 3.980859 0.468240
S -3.502139 -1.872091 0.628182
S -3.170929 -1.548862 -1.391904
S 3.366342 -2.167448 0.232217
S 2.825251 -1.635004 -1.686255
H 3.385488 -0.202731 1.989200

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Author Contributions

AB: Experimental planning and investigation (lead), data analysis (lead), results discussion (equal), preparation of the manuscript (lead).

MP: Computations (supporting), results discussion (equal).

MO: Computations (lead), results discussion (equal), manuscript editing (lead).

MLG: Chemical synthesis (lead), results discussion (equal).

PK: Project supervision and funding acquisition (lead), results discussion (equal), manuscript editing (supporting).

MKB: Conception of original idea, project supervision and funding acquisition (lead), data validation (lead), results discussion (equal), manuscript editing (lead).