

Supporting Information

Probing the Radical and Base Dual Properties of Peptide Sulfinyl Radicals via Mass Spectrometry

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Materials

Synthesis of Disulfide Precursor

The amine was dissolved in water at a concentration of 1.0 mg/mL, and the pH of the solution was adjusted to 8 by adding 0.1 M NaOH. Freshly prepared DSP or d₈-DSP (10 mM in DMSO) was then added at a molar ratio of 5:1 (DSP: amine). The mixture was allowed to react at room temperature for 30 min and then subjected to RP-HPLC separation (Agilent 1200 series, Agilent Technologies, Santa Clara, CA). The collected eluent was vacuum dried using a centrifrap concentrator (Labconco, Kansas City, MO) and redissolved in water/methanol/acetic acid (v/v/v=50:49:1) at 10 μ M for MS analysis. To prepare the sodium-ion adduct, the sample was reconstituted in water/methonal (v/v=1:1) at 10 μ M, and 10 mM NaOH was added to adjust the pH to 8. Acetylation was achieved by mixing the tetrapeptide disulfide-linked dimer solution (100 μ L, approximately 0.2 mg/mL in water, pH = 5) and 10 μ L of acetylation solution (acetic anhydride/methanol (v/v) = 1:3). Notably, the pH of the solution was adjusted to acidic conditions to avoid acetylation at the ϵ -NH₂ from the Lys side chain.^{S1}

Theoretical calculations

Optimization for [^{SO}•Cys-Arg + H]⁺

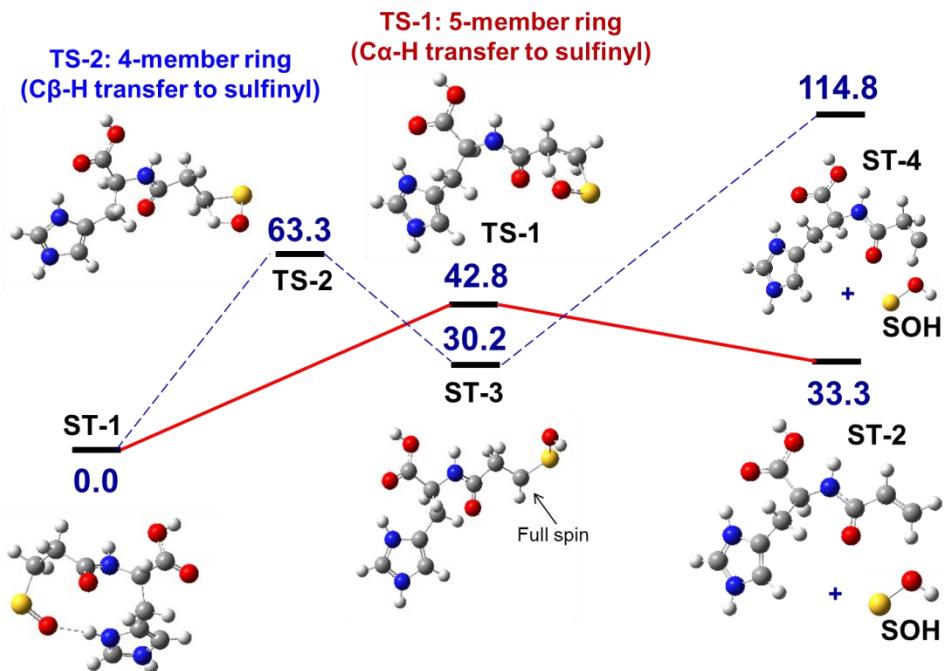
Exhaustive search of the conformational space of the dipeptide sulfinyl radical [^{SO}•Cys-Arg + H]⁺ using the ConformSearch engine^{S2} is very challenging because the force fields used in the molecular dynamics step have not been optimized for sulfinyl radicals. Therefore, an indirect approach was adopted, using the [Asn-Arg + H]⁺ ion protonated at Arg as a proxy of analogous hydrogen bonding properties for the Asn amide and SO groups. Replica exchange molecular dynamics^{S3} with CHARMM^{S4}/NAMD^{S5} produced 800,000 structures from which 8000 were sampled for full geometry optimization with PM6.^{S6} Over 100 lowest-energy [Asn-Arg + H]⁺ structures were selected and converted to sulfinyl radical analogues by deleting the amide group (-NH₂) from the Asn side chain and changing the carbonyl C=O to S=O. These cation-radical constructs were then fully optimized with B3LYP^{S7} and M06-2X^{S8}/6-31+G(d,p), the structures were characterized as local energy minima by harmonic frequency analysis (all frequencies real), and their relative energies were determined by single point calculations with B3LYP, M06-2X and MP2(frozen core)^{S9} and the 6-311++G(2d,p) basis set.

Benchmark Calculation for Proton Affinity

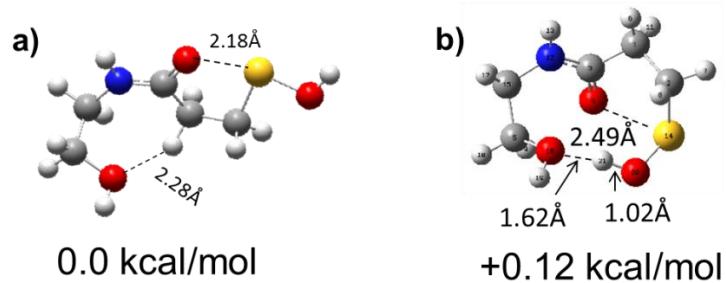
Proton affinity is calculated as the enthalpy difference between protonated sulfinyl radical and the neutral structure. This method has been successfully applied in predicting the proton affinity of small molecules and the results agree well with experimental values.^{S10} Three model molecules, methonal, propylamine, and pyridine, were used to calibrate the calculation method. The result is listed in SI Table 1, which yields r.m.s error of 1.6 kcal mol⁻¹.

SI Table 1. Benchmark calculation of proton affinity by three model compounds. Level of calculation CCSD(T)//B3LYP/6-31G(d) + ΔZPE

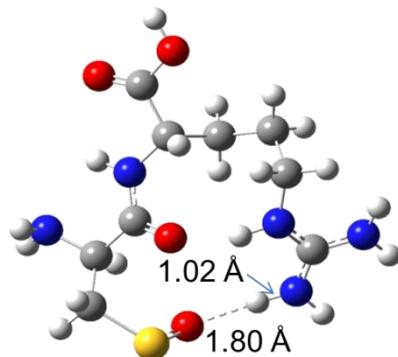
Name	Calc. (kcal mol ⁻¹)	Exp. (kcal mol ⁻¹) ^{S11}	Δ(Calc.-Exp.) (kcal mol ⁻¹)
Methanol	181.0	180.3	0.7
Propylamine	221.1	219.4	1.7
Pyridine	224.5	222.3	2.2



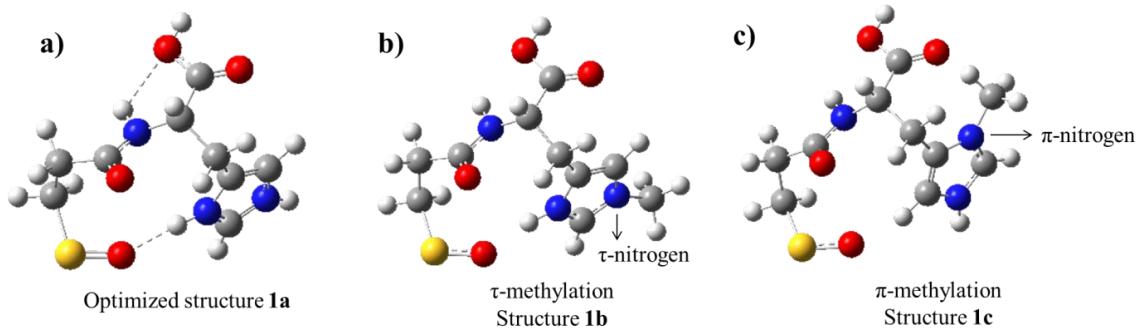
SI Scheme 1. Energy diagram for radical induced SOH loss comparing involvement of C α -H vs. C β -H, using [n-³⁴S-Cys-His + H]⁺ as the model system. For the mechanism involving the C α -H, the sulfinyl radical oxygen grabs the C α -H of the cysteine residue through a five-member ring transition state, followed by the leaving of SOH to form a dehydroalanine motif. The transition state and the products are 42.8 kcal mol⁻¹ and 33.3 kcal/mol higher in enthalpy relative to the initial structure, which are quite reasonable for low energy CID conditions. However, if C β -H were involved, the fragmentation should have gone through a high energy four-member ring transition state (63.3 kcal mol⁻¹ higher relative to the initial structure), and the resulting motif from the SOH loss does not obey the octet rule and is thus unstable (product ions with energy of 114.8 kcal mol⁻¹ higher relative to the initial structure). It shows that the 5-member ring transition state to abstract C α -H is more favorable than the 4-member ring transition state to abstract the C β -H. Level of calculation: CCSD(T)//B3LYP/6-31G(d) + Δ ZPE.



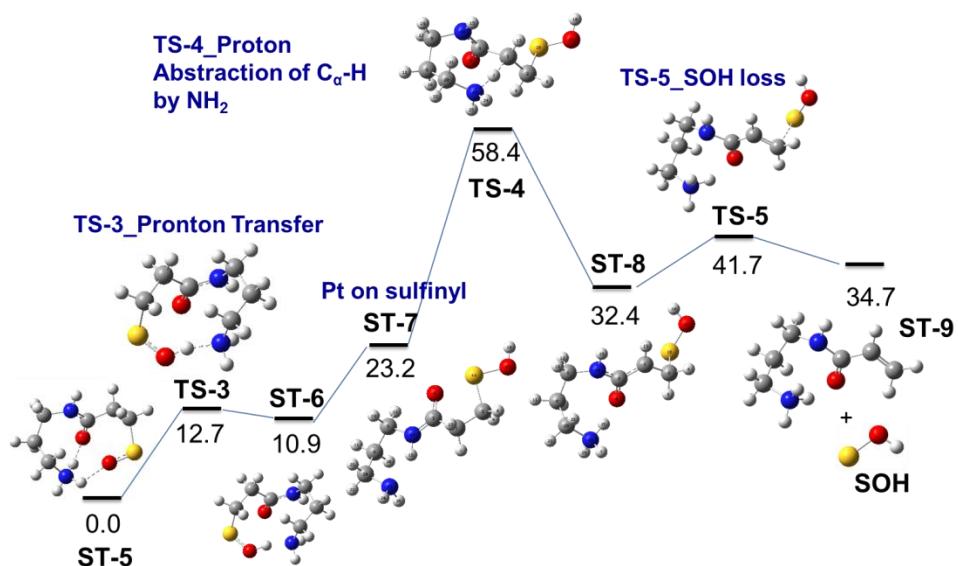
SI Scheme 2. Geometry optimization of protonated **4a**. The relative energy of two low energy structures are listed, with the protonation on sulfinyl radical being the most stable one. Level of calculation: B3LYP/6-31G(d) + ΔZPE.



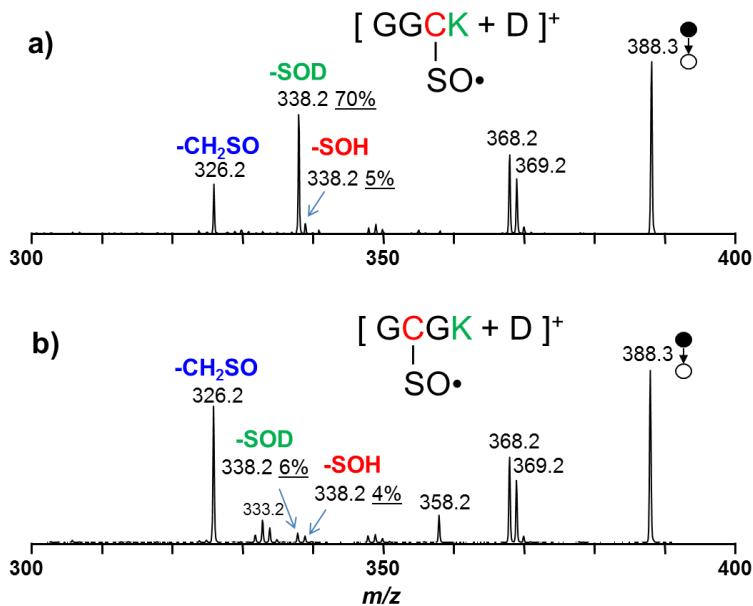
SI Scheme 3. Geometry optimization of protonated ^{SO}•Cys-Arg.



SI Scheme 4. 3D structure for compound **1a-c**. Structure **1a** is optimized on level of B3LYP/6-31G(d), while structure **1b** and **1c** are not optimized. It can be seen from the model that the proton bridging between sulfinyl radical and π -nitrogen can be retained in case of τ -methylation (b). However, it is disrupted in π -methylation (scheme c). No matter how we rotate the bonds, it is difficult for the proton on τ -nitrogen to access sulfinyl radical or other backbone atoms.

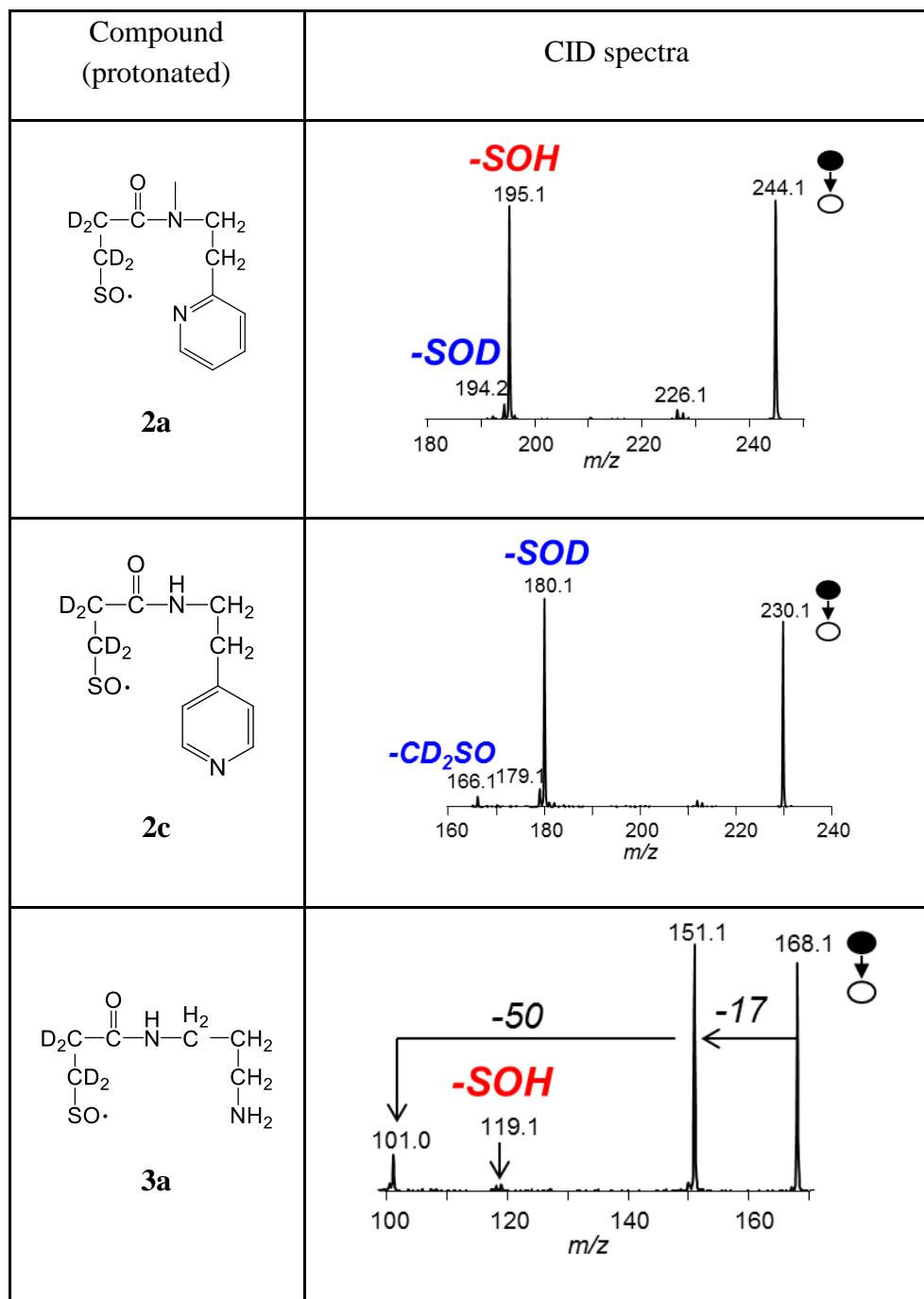


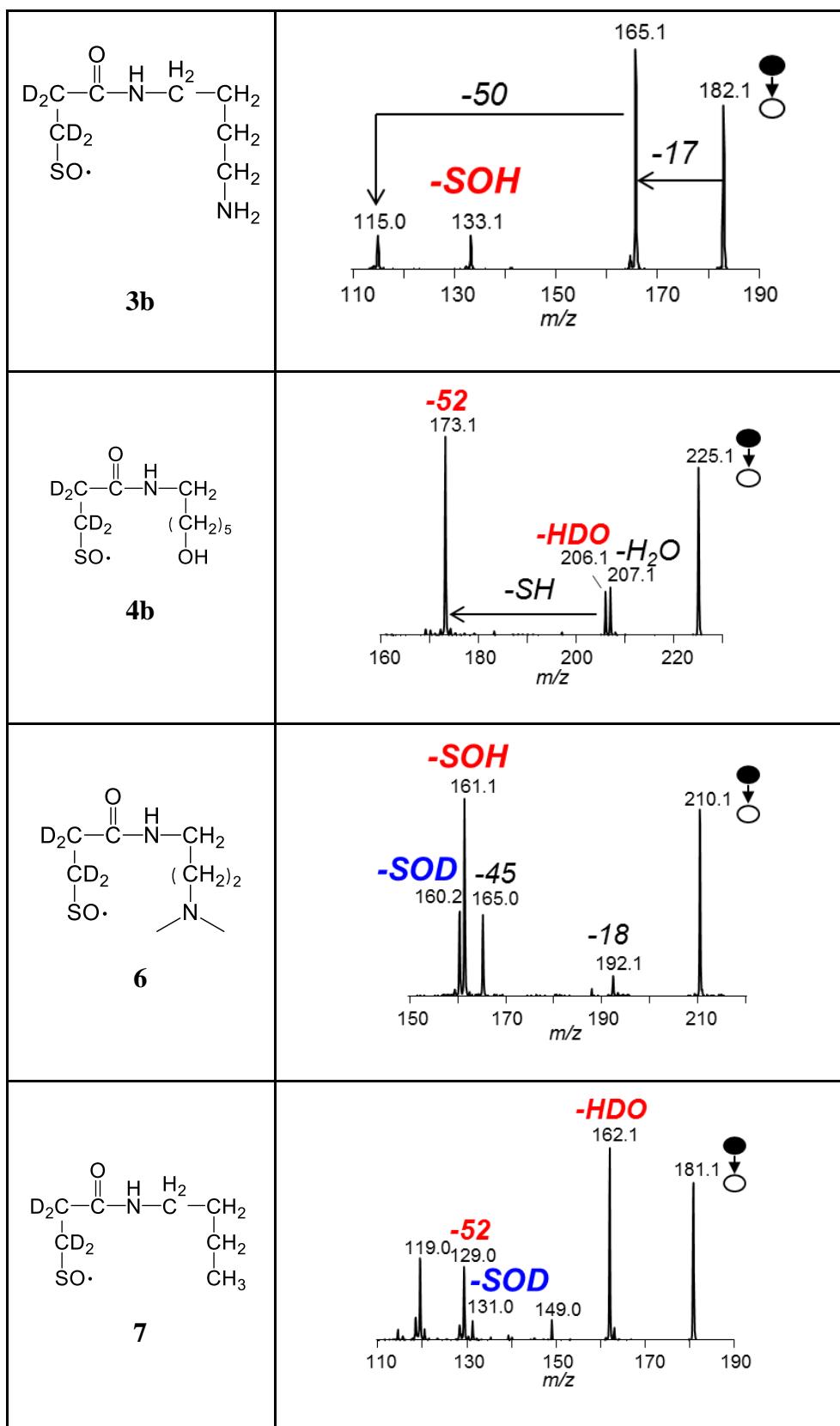
SI Scheme 5. Energy diagram for proton induced SOH loss, using protonated **3b** as the model system. Level of calculation: CCSD(T)//B3LYP/6-31G(d) + Δ ZPE.



SI Figure 1. MS² ion trap CID of tetrapeptide sulfinyl radical after solution phase hydrogen/deuterium exchange: a) $[\text{GG}^{\text{SO}\bullet} \text{CK} + \text{D}]^+$; b) $[\text{G}^{\text{SO}\bullet} \text{CGK} + \text{D}]^+$. The degree of deuterium exchange is about 95%.

SI Table 2. CID spectra of d₄- n-SO[•]Cys-X, the number of compound is indicated below each structure.





SI Table 3. Electronic energies (kcal mol⁻¹) for key optimized structures. Level of theory: CCSD(T)//B3LYP/6-31G(d)

Name	CCSD (T)	Δ ZPE/B3LYP	CCSD (T) + Δ ZPE
SI Scheme 1			
ST-1	0.0	0	0.0
ST-2 + SOH	36.3	-3.0	33.3
ST-3	32.1	-1.9	30.2
ST-4 + SOH	120.4	-5.7	114.8
TS-1	46.3	-3.5	42.8
TS-2	67.5	-4.2	63.3
SI Scheme 2			
ST-5	0.0	0.0	0.0
ST-6	11.4	-0.5	10.9
ST-7	24.9	-1.7	23.2
ST-8	34.5	-2.1	32.4
ST-9 + SOH	38.2	-3.5	34.7
TS-3	13.5	-0.8	12.7
TS-4	63.6	-5.2	58.4
TS-5	44.4	-2.7	41.7

Gaussian archive entries for B3LYP/6-31G(d) calculation for results shown in SI Scheme 1 and 4

ST-1

```
1\1\GINC-CARTER-A076\FOpt\UB3LYP\6-31G(d)\C9H13N3O4S1(1+,2)\FRANCISC\2
6-Aug-2013\0\#p b3lyp/6-31g* scfcyc=300 trans=iabc opt=calcfc\nCH 25
9 sulfinyl, proton bridging between pi-N and sulfinyl_2\1,2\C,0.26998
80697,-0.0461315607,-2.0093475802\C,0.8368339503,-0.0472550568,-3.4446
443598\C,-0.4291211469,1.2612409322,-1.6492249062\O,0.2099127787,2.249
5437835,-1.3106018597\C,-2.5651133638,2.4276272038,-1.2993547713\C,-4.
0592172401,2.104422737,-1.2984179169\O,-4.9305169065,2.914517666,-1.51
27127335\H,-0.7017972211,3.0159127658,-4.30777813\C,-2.2921129548,3.75
08204647,-2.0522074792\H,-0.3969059664,-0.9110967997,-1.9108933677\H,1
.4674706301,-0.9311601079,-3.594212028\H,0.039919024,-0.067304986,-4.1
95730945\H,-1.2419563596,4.0046837216,-1.8947353368\H,-2.9026762617,4.
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 3.5161471732\C,-3.7293585129,4.0779953391,-4.2025062512\H,-4.68628438,
 4.4195225128,-3.8415162845\N,-1.6704571465,3.3498991052,-4.479111494\N
 ,-3.4554791516,3.9079794741,-5.5470234953\H,-4.0946234812,4.0998628907
 ,-6.3090614962\C,-2.1998417333,3.4669149893,-5.6951427147\H,-1.7040033
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 917\H,-5.2629310767,0.6826710557,-0.9719248823\N,-1.7948376964,1.28805
 947,-1.7996633925\H,-2.2893898355,0.4056441913,-1.8426910195\S,1.89665
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ST-2

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 5,-2.2941451889\C,-3.3761723977,2.4774781716,-0.3318578753\H,0.4389836
 909,-1.3819189565,0.3221450326\H,2.1627703707,-0.7357074837,0.59209233
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ST-3

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6-Aug-2013\0\#p b3lyp/6-31g* scfcyc=300 trans=iabc opt=calcfc\nCH --
 -> hoS loss (4 center transition state, grab belta-carbon hydrogen) h
 ydrogen transfer to so group\1,2\C,2.8025108381,0.4720746926,-0.59627
 93389|C,3.6918074206,-0.729834587,-0.6757038988|C,1.3273089654,0.09915
 73973,-0.5804627577|O,0.8660015987,-0.9087634129,-1.1035080162|C,-0.93
 60404637,0.8211359962,-0.0692743305|C,-1.6995123634,2.112514304,0.2032
 546012|O,-2.9004211431,2.1576772149,0.4221612736|H,-3.9523096812,0.670
 6938087,0.3149775509|C,-1.4400070872,-0.3436690429,0.840976225|H,3.064
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 509|H,3.3556224708,-1.6088962841,-1.2166946959|H,-1.5596233108,0.01141
 26181,1.8712565274|H,-0.6501464324,-1.0954952938,0.8361414117|H,-1.154
 5526582,0.5402267377,-1.1110488807|H,2.9494846284,1.1247843678,-1.4776
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 267059229,0.0637314005|H,-2.2931811441,-3.1871296177,0.0570770685|N,-3
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ST-4

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 0.898167436,-1.1018695232|O,-1.0724461138,-0.0389131079,-1.3188440045|\br/>
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 547,3.4882765068,-2.8265128056|C,-3.1625088562,2.1911805589,-0.9433499
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 06,0.4310541595|H,-3.2890635809,3.064000626,-0.2921589078|H,-2.8544054
 975,1.3519052641,-0.3189040494|H,-2.1073807471,1.7246392894,-2.7735267
 24|H,1.6836132168,0.1474281197,-1.2807188948|C,-4.4719300545,1.8166053
 727,-1.5662856364|C,-5.2627787777,0.7133131708,-1.389573627|H,-5.12587
 324,-0.1617001136,-0.7735820657|N,-5.1421523446,2.6186313025,-2.478996
 2629|N,-6.3772587665,0.8840694056,-2.1914740271|H,-7.1491334446,0.2325

477949,-2.2758665773\|C,-6.2848753491,2.0452772763,-2.8505327377\|H,-7.0
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 O3)]\\@

ST-5

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 9-Jan-2014\0\\#p b3lyp/6-31g* scfcyc=300 trans=iabc opt=readfc geom=ch
 eck guess=read\\h+nh2(ch2)3nhc(o)ch2ch2so radical cation protonation a
 t nh2 site Lei-Lei structure\\1,2\|C,1.3883048434,0.7266371198,0.332502
 6053\|C,2.6435992351,0.8178356447,-0.5621105398\|C,0.243934683,1.6468788
 368,-0.0873479831\|O,0.0444257216,2.7151921161,0.5131030985\|C,-1.844320
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 7\|H,3.4273747184,0.1626287472,-0.1648558786\|H,2.4389736396,0.514085157
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 \|Dipole=-0.944255,0.3727687,-0.8554475\|Quadrupole=1.2161249,9.6259462,
 -10.8420711,-6.3990633,6.9792141,-3.3752338\|PG=C01 [X(C6H14N2O2S1)]\\@

ST-6

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 ead\\h+nh2(ch2)3ch2nhc(o)ch2ch2so radical cation, minimum structure f
 ollowing transfer yupaq3.lnh2tsqst folded\\1,2\|C,-1.0002512445,-1.62744
 9272,-1.0908772012\|C,-1.4311376725,-0.304341141,-1.7586514865\|S,-2.409
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 23621453\|C,-0.4616268869,-1.49280506,0.3345289256\|O,-1.1635485787,-1.7

605679404,1.2923571326|N,0.826503685,-1.0057366671,0.5396075109|H,1.11
 81677545,-1.210934747,1.4927053301|H,-1.8661689065,-2.2862737475,-0.99
 94984547|H,-0.5800531433,0.2924551496,-2.1046556548|H,-2.0639229742,-0
 .5146054271,-2.628375498|O,-1.3628357727,1.5523679321,0.1741359028|C,1
 .9108868345,-0.8950824881,-0.4348163479|C,2.8832431826,0.2277861253,-0
 .0437337981|H,1.4826558583,-0.6863399173,-1.4194073273|H,2.4749371171,
 -1.833580686,-0.5228218157|H,3.2952860918,0.0333660522,0.9568432532|H,
 3.7421348184,0.2003255616,-0.7246865087|C,2.3361527334,1.6554142736,-0
 .1049085045|H,1.9185739128,1.8934978736,-1.0869639835|H,3.1300394212,2
 .3736640431,0.1123451831|H,1.0573219828,2.9321913134,0.9924417928|H,0.
 3037977555,1.5074319403,0.5967397725|N,1.2239911218,1.9244595736,0.891
 8533691|H,1.4500653193,1.5573637452,1.8216879984\\Version=EM64L-G09Rev
 C.01\\State=2-A\\HF=-894.9492718\\S2=0.753688\\S2-1=0.|\\S2A=0.75001\\RMSD=9.
 582e-09\\RMSF=2.753e-05\\Dipole=2.7326225,1.8191357,-0.5136184\\Quadrupol
 e=5.1501283,-0.5746264,-4.575502,5.6129247,9.7259151,7.7972654\\PG=C01
 [X(C6H14N2O2S1)]\\@

ST-7

1\\1\\GINC-CARTER-A269\\FOpt\\UB3LYP\\6-31G(d)\\C6H14N2O2S1(1+,2)\\ROOT\\04-Fe
 b-2014\\0\\#p b3lyp/6-31g* scfcyc=300 trans=iabc opt=readfc geom=check
 guess=read\\nh2(ch2)3nhc(o)ch2ch2so-h+ radical cation protonation at s
 -o site Structure before grabbing alpha-carbon hydrogen with amine, pt
 at sulfinyl\\1,2|C,-2.9187486804,2.0676305069,2.3069342889|C,-3.22477
 20888,0.5865632519,2.5051744926|C,-2.8212300522,2.344123177,0.81679343
 56|O,-2.4708152184,1.3676027714,0.0828865257|C,-2.9161744179,3.9175902
 529,-1.0761859149|C,-3.7975570195,5.1309455765,-1.3969984986|H,-3.6858
 178925,2.6808928489,2.788513559|H,-2.9215308068,0.2136828022,3.4860919
 664|H,-4.2850324319,0.3556804159,2.3567888741|H,-4.8562865172,4.857124
 8296,-1.2951985334|H,-3.6426393814,5.3838270156,-2.4518169203|H,-3.186
 7727394,3.0526798488,-1.6874168287|H,-1.9554412472,2.3272769906,2.7636
 616568|N,-3.0520466167,3.5474151292,0.343708735|H,-3.3121824584,4.3522
 755189,0.9535154797|S,-2.3217606188,-0.4289626356,1.2661221561|O,-2.85
 05810797,-1.8439218528,1.9965728096|H,-2.4924211149,-2.6095271706,1.50
 24763339|H,-1.8620089209,4.1466267562,-1.2787388956|C,-3.4973974259,6.
 3695826261,-0.5404762824|H,-2.4247849898,6.5930100144,-0.5790646881|H,
 -4.0207631587,7.2406415604,-0.957201519|N,-3.8512498271,6.1257039874,0
 .8818870097|H,-3.4483620657,6.8534905922,1.47081731|H,-4.8622395798,6.
 1992311862,0.9991425781\\Version=EM64L-G09RevC.01\\State=2-A\\HF=-894.93
 66179\\S2=0.753621\\S2-1=0.|\\S2A=0.750011\\RMSD=6.216e-09\\RMSF=7.085e-06\\D
 ipole=-0.170326,-0.1883176,0.9016281\\Quadrupole=-15.0242755,24.9287905
 ,-9.904515,-7.620105,1.0519533,-7.1962102\\PG=C01 [X(C6H14N2O2S1)]\\@

ST-8

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1\1\GINC-CARTER-A192\FOpt\UB3LYP\6-31G(d)\C6H14N2O2S1(1+,2)\ROOT\04-Fe
b-2014\0\#p b3lyp/6-31g* scfcyc=300 trans=iabc opt=readfc geom=check
guess=read\nh2(ch2)3nhc(o)ch2ch2so-h+ radical cation protonation at s
-o site Structure after grabbing alpha-carbon hydrogen with amine\\1,2
|C,-2.3972476315,1.3888257263,1.897305464|C,-3.3475934624,0.6196047137
,2.7085265228|C,-2.8603940241,2.3929527799,0.9732358198|O,-4.092162915
9,2.6538879036,0.8487333782|C,-2.0773315377,3.7509060444,-1.0301286405
|C,-3.2832808554,4.6871979034,-1.2300334048|H,-4.600175595,5.661716347
,1.7304677947|H,-2.957260469,0.2791137223,3.6696248352|H,-4.301536908,
1.1344190076,2.8430450735|H,-4.1906941596,4.1003636958,-1.4050590361|H
,-3.0920860627,5.231827182,-2.1606414292|H,-2.1150261247,2.9909316196,
-1.822072651|H,-1.3466500793,1.1121572508,1.9100896139|N,-1.9098749487
,3.0946517861,0.2773823813|H,-0.9551917705,2.8035377547,0.4522818111|S
,-3.762330791,-0.9248530791,1.7387648367|O,-2.5179077817,-1.9301455701
,2.2498475226|H,-1.9264620362,-2.0783749875,1.4911665347|H,1.15921369
7,4.3263683917,-1.1824500425|C,-3.5370282942,5.7106320133,-0.111744787
|H,-2.6459523509,5.8815223351,0.4966859516|H,-3.8778930863,6.671047306
5,-0.5027861406|N,-4.6080251025,5.1916455651,0.8211112124|H,-5.5413335
06,5.3178148674,0.4178049625|H,-4.4499969695,4.1149661004,0.9387717767
\\Version=EM64L-G09RevC.01\State=2-A\HF=-894.9235213\S2=0.755789\S2-1=
0.\$2A=0.750022\RMSD=5.355e-09\RMSF=1.913e-05\Di pole=0.1457089,3.81990
32,-0.823492\Quadrupole=-5.278813,15.5752128,-10.2963998,-12.709051,-2
.0903397,-3.4080688\PG=C01 [X(C6H14N2O2S1)]\\@

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ST-9

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1\1\GINC-CARTER-A505\FOpt\RB3LYP\6-31G(d)\C6H13N2O1(1+)\ROOT\12-Feb-20
14\0\#p b3lyp/6-31g* scfcyc=300 opt=readfc geom=check guess=read\\str
ucture following soh loss from yupa3 structure Lei-Lei structure\\1,1\
C,-2.3921786599,1.4325675794,1.989975034|C,-3.2460507297,0.7886578568,
2.7922628222|C,-2.8790594234,2.4208888452,1.002607437|O,-4.0945918347,
2.7116033343,0.9062815437|C,-2.0779600624,3.7365370245,-1.0334598905|C
,-3.2910693664,4.6643281993,-1.2153197815|H,-4.6570197417,5.7580915956
,1.6930616042|H,-2.9019762128,0.0487571423,3.5076147445|H,-4.312796564
1,0.9842379342,2.7394624152|H,-4.1984156029,4.0720021295,-1.3683816687
|H,-3.1155746831,5.1942039804,-2.1574815011|H,-2.0935693432,3.01251638
23,-1.8579799876|H,-1.3244281533,1.2289202943,2.029188238|N,-1.9247539
992,3.0085959061,0.2377738705|H,-0.9807349272,2.6757554529,0.395283316
3|H,-1.1627787835,4.3267571789,-1.1451789169|C,-3.5284712298,5.7099544
298,-0.1115464578|H,-2.6460862236,5.8536744312,0.5166218047|H,-3.82093
47448,6.6790440939,-0.5196787989|N,-4.6410945678,5.2498025489,0.804182
0268|H,-5.5580556955,5.370261202,0.362782723|H,-4.4983490009,4.1889324
782,0.972079873\\Version=EM64L-G09RevC.01\State=1-A\HF=-420.9664378\RM

```

SD=4.164e-09|RMSF=4.529e-06|Dipole=-0.2963946,2.0198718,-0.0097212|Qua
 drupole=-1.1394991,4.9463867,-3.8068876,-9.5434703,-1.9155284,-5.13413
 25|PG=C01 [X(C6H13N2O1)]\\@

TS-1

1\1\GINC-CARTER-A213\FTS\UB3LYP\6-31G(d)\C9H13N3O4S1(1+,2)\FRANCISC\27
 -Aug-2013\0\#p b3lyp/6-31g* scfcyc=300 trans=iabc opt=(ts,calcf,nc,oei
 gen) optcyc=100\\nCH ---> hoS loss (5 center transition state)\\1,2\\
 C,0.7747058658,0.7459522578,0.0786425915|C,1.2519458043,-0.5736240297,
 0.1935881632|C,-0.6107327193,0.9334901898,-0.3931111095|O,-1.473106759
 7,0.0522494083,-0.388651134|C,-2.2495181818,2.4199943047,-1.3955006014
 |C,-2.2927999464,3.6563016006,-2.2855463873|O,-3.3246081409,4.20234860
 61,-2.6478436746|H,-4.9532429933,3.4985454955,-2.2784589122|C,-3.34911
 67565,2.4703921017,-0.287543739|H,0.5519296134,0.9405405471,1.57460218
 6|H,0.5928657987,-1.3890218706,-0.0884587102|H,2.3151988952,-0.7749338
 828,0.1127482929|H,-3.3618656934,3.4616548377,0.1806171609|H,-3.043613
 6488,1.7467274182,0.4685842622|H,-2.4785372426,1.5587526308,-2.0429782
 5|H,1.493311874,1.5460649016,-0.0971689371|C,-4.7277556804,2.090654842
 1,-0.7298339128|C,-5.5744361674,1.1113469135,-0.2845499337|H,-5.446391
 799,0.3637403965,0.4824225644|N,-5.4148646747,2.7530276091,-1.73679380
 41|N,-6.7371260497,1.215103397,-1.0278681886|H,-7.5556431354,0.6259545
 455,-0.9290034237|C,-6.6202877254,2.2153371591,-1.909817495|H,-7.36485
 60354,2.5220553007,-2.6283817189|O,-1.0890739896,4.088390098,-2.663704
 5427|H,-1.2105166112,4.8648691892,-3.2463051011|N,-0.9445116306,2.2150
 578253,-0.8077767601|H,-0.2156032427,2.8865596397,-1.0051386544|S,1.08
 80183661,-0.9168116516,2.514348091|O,0.5475400869,0.566096789,2.646746
 4584\\Version=EM64L-G09RevC.01\\State=2-A\\HF=-1213.8427471\\S2=0.754229\\
 S2-1=0.\\S2A=0.750014|RMSD=7.717e-09|RMSF=7.203e-06|Dipole=-3.9927632,1
 .0095485,-1.8926168|Quadrupole=37.6145405,-15.5072926,-22.1072479,-1.5
 414438,9.9311982,-10.6732013|PG=C01 [X(C9H13N3O4S1)]\\@

TS-2

1\1\GINC-CARTER-A233\FTS\UB3LYP\6-31G(d)\C9H13N3O4S1(1+,2)\FRANCISC\26
 -Aug-2013\0\#p b3lyp/6-31g* scfcyc=300 trans=iabc opt=(ts,calcf,nc,oei
 gen)\\nCH ---> hoS loss (4 center transition state, grab belta-carbon
 hydrogen)\\1,2|C,0.6396154534,-0.2028306644,-0.4130676148|C,1.3212251
 692,0.5937307119,0.6881987799|C,-0.7470519091,0.308188546,-0.808063520
 1|O,-1.5484539369,-0.3771773559,-1.4338561284|C,-2.2630126302,2.197763
 118,-0.9479987225|C,-2.2139548735,3.7208509317,-0.886032392|O,-3.19813
 73618,4.4376640155,-0.9916845995|H,-4.8382916669,3.7574969749,-1.37332
 58454|C,-3.5186311486,1.6339864745,-0.2092199146|H,1.2373609111,-0.177

1088544,-1.3344155431\H,1.8669882639,2.0407347576,0.8251059059\H,0.896
 2277948,0.4151618814,1.6833755254\H,-3.6424283756,2.1389482652,0.75596
 18737\H,-3.3066674632,0.5823880276,-0.0132501359\H,-2.362229145,1.9291
 483614,-2.0112895394\H,0.4997811419,-1.2636013571,-0.170257222\C,-4.79
 6283702,1.687523782,-0.9876325822\C,-5.6601600604,0.6934329717,-1.3611
 362466\H,-5.6214391736,-0.3724169053,-1.1993409845\N,-5.3458268481,2.8
 659664711,-1.4718469227\N,-6.6966285444,1.2982713083,-2.0501591935\H,
 7.4950250241,0.8216396743,-2.4532124825\C,-6.48624637,2.618747457,-2.1
 125495053\H,-7.1224827204,3.3436067991,-2.5969936925\O,-0.9859476164,4
 .2111133189,-0.7243970134\H,-1.0410694397,5.1879389476,-0.7160752115\N
 ,-1.0499871836,1.5948483402,-0.4440325289\H,-0.3274313632,2.1633279061
 ,-0.0185393495\S,3.088754938,0.5722378738,0.8035378267\O,3.046172014,2
 .1743133113,1.0510207797\\Version=EM64L-G09RevC.01\State=2-A\HF=-1213.
 8037362\\$2=0.753438\\$2-1=0.\\$2A=0.750008\RMSD=8.772e-09\RMSF=5.411e-06
 \Dipole=-5.1836165,0.7741064,-1.1820774\Quadrupole=37.6017893,-16.1186
 546,-21.4831347,-5.0346828,19.263363,-5.3365828\PG=C01 [X(C9H13N3O4S1)
]\\@\n

TS-3

1\\1\GINC-CARTER-A018\FTS\UB3LYP\6-31G(d)\C6H14N2O2S1(1+,2)\ROOT\13-Feb
 -2014\0\\#p b3lyp/6-31g* opt=(ts,readfc,noeigen) geom=check guess=read
 \\h+-nh2(ch2)3ch2nhc(o)ch2ch2so radical cation, ts to transfer proton
 from amine to sulfinyl\\1,2\C,-0.9172476677,-1.4322970651,-1.005315241
 9\C,-1.4001775446,-0.1649199208,-1.7431336369\S,-2.4195084365,1.013705
 2376,-0.7639473633\H,-0.2249273494,-1.9666181642,-1.6636460805\C,-0.33
 93011847,-1.2017025791,0.3896655299\O,-1.0792331044,-0.9857076067,1.34
 01641621\N,1.0289678191,-1.2575115753,0.5883929345\H,1.2758240184,-1.4
 266658829,1.5599816228\H,-1.7736184563,-2.0918921166,-0.8361570717\H,
 0.5696058539,0.4312864932,-2.137495951\H,-2.0264057842,-0.4578589423,-
 2.5935458268\O,-1.4143726858,1.9693545269,-0.0991596235\C,2.1250771712
 ,-1.0874216963,-0.361431897\C,2.9003284738,0.2235035324,-0.1125869759\
 H,1.7134001992,-1.0888693281,-1.3743187683\H,2.8166282394,-1.934427144
 7,-0.2957082772\H,3.4017307991,0.1862688036,0.8640178903\H,3.701514662
 3,0.3010633053,-0.8570962936\C,2.0624153187,1.501036942,-0.2201991178\
 H,1.4589855052,1.5081862045,-1.1313317943\H,2.6963665068,2.3890802853,
 -0.2344108595\H,1.3840069469,2.3909730117,1.6041051485\H,0.1215888944,
 1.9327243708,0.5676420798\N,1.0785572334,1.6809859453,0.9325053623\H,0
 .9347578676,0.7954798706,1.4331863984\\Version=EM64L-G09RevC.01\State=
 2-A\HF=-894.9471039\\$2=0.75365\\$2-1=0.\\$2A=0.75001\RMSD=9.337e-09\RMSF
 =4.476e-06\Dipole=2.523962,0.6893604,-0.3308093\Quadrupole=5.3650241,-
 2.0349943,-3.3300298,4.3215244,9.1140359,5.6592333\PG=C01 [X(C6H14N2O2
 S1)]\\@\n

TS-4

```
1\1\GINC-CARTER-A220\FTS\UB3LYP\6-31G(d)\C6H14N2O2S1(1+,2)\FRANCISC\04
-Feb-2014\0\#p b3lyp/6-31g* scfcyc=300 trans=iabc opt=(ts,readfc,noeigen)
geom=check guess=read\TS to grab alpha-carbon hydrogen with amine
e\1,2\C,-2.9668014707,2.1254176952,1.5701888385\O,-3.9739272246,1.124
4531787,2.051602057\O,-2.8269878316,2.2227335516,0.0798503164\O,-3.670
429064,1.7302858269,-0.6650143946\O,-2.1704915098,4.0119778299,-1.4464
590358\O,-3.4185892687,4.8721202751,-1.1407379358\H,-3.5595584714,3.40
0093001,1.5729049931\H,-4.1598961414,1.1318759107,3.1273553725\H,-4.90
64473572,1.1588273388,1.4869933416\H,-4.321968556,4.2741474286,-1.3110
859458\H,-3.4571610854,5.6936146306,-1.8644309944\H,-2.3327203056,3.44
04897202,-2.3627246128\H,-2.054479085,2.1985494861,2.16706496\N,-1.859
9165007,3.0940222844,-0.3475356672\H,-1.2183037098,3.4396777201,0.3550
261618\S,-3.1901934585,-0.4917747303,1.6546133625\O,-2.6228948754,-0.8
647761347,3.1693590561\H,-1.7647865501,-1.3226298534,3.076107835\H,-1.
2906719529,4.6427643779,-1.5993803869\O,-3.4672842491,5.4827245553,0.2
731058075\H,-2.4678111982,5.7149751769,0.6537948467\H,-4.0353022275,6.
4186594438,0.2556603049\N,-4.1269650626,4.5940456884,1.2993216745\H,-4
.20514697,5.0837850308,2.1942476752\H,-5.0787170036,4.3624813575,1.001
1453298\Version=EM64L-G09RevC.01\State=2-A\HF=-894.8811024\S2=0.75433
7\S2-1=0.\$2A=0.750014\RMSD=3.940e-09\RMSF=6.333e-06\Dipole=0.1214968,
1.6872558,1.5680794\Quadrupole=-6.3882214,15.3124311,-8.9242097,-7.463
0333,-0.5499465,-8.3621939\PG=C01 [X(C6H14N2O2S1)]\@\@
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TS-5

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1\1\GINC-CARTER-A472\FTS\UB3LYP\6-31G(d)\C6H14N2O2S1(1+,2)\FRANCISC\13
-Feb-2014\0\#p b3lyp/6-31g* opt=(ts,readfc,noeigen) geom=c
heck guess=read\nh2(ch2)3nhc(o)ch2ch2so-h+ radical cation protonation
at s-o site Structure after grabbing alpha-carbon hydrogen with amine
breaking the C-S bond\1,2\C,-2.3028065316,1.6236053622,2.0080052821\
C,-3.1718474181,0.9366972041,2.8225904747\O,-2.8002765333,2.6112116706
,1.0585363727\O,-4.0347685408,2.8245494263,0.9133770921\O,-2.032335290
3,4.0150179161,-0.9224727113\O,-3.2691903196,4.9098929747,-1.122125458
2\H,-4.6990515248,5.7940519518,1.8171910694\H,-2.7969033066,0.29235556
25,3.611090758\H,-4.2014306682,1.265604535,2.909533096\H,-4.150849745,
4.2932990247,-1.3245618181\H,-3.082810335,5.4792336045,-2.0388147814\H
,-2.0248098893,3.2776219024,-1.7358340654\H,-1.241672287,1.3891809651,
2.0136298602\N,-1.8649983297,3.3242693575,0.3668939438\H,-0.9043355222
,3.0644408396,0.5569644342\S,-3.7447268141,-0.986251276,1.4804598977\O
,-2.874772302,-2.1648419927,2.2677661006\H,-2.220206008,-2.5295029562,
```

1.6435210267\H,-1.1339499472,4.6288211197,-1.0394882535\C,-3.580125178
8,5.9019998026,0.010481629\H,-2.7106904668,6.0828049908,0.6468198493\H
, -3.9356143218,6.8615553413,-0.3694284022\N,-4.6614888501,5.3355888852
,0.9026073411\H,-5.5854133615,5.4332944604,0.4707861255\H,-4.466097978
2,4.2658840377,1.0130414568\\Version=EM64L-G09RevC.01\State=2-A\HF=-89
4.9110188\S2=0.767459\S2-1=0.\S2A=0.75008\RMSD=6.564e-09\RMSF=4.689e-0
6\Di pole=0.1829882,3.6941324,-0.6714356\Quadrupole=-6.7055154,19.33588
94,-12.630374,-14.0653529,-1.1333077,-4.4451462\PG=C01 [X(C6H14N2O2S1)
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