



J | A | C | S
JOURNAL OF THE AMERICAN CHEMICAL SOCIETY

J. Am. Chem. Soc., 1997, 119(52), 12982-12983, DOI: [10.1021/ja971766a](https://doi.org/10.1021/ja971766a)

Terms & Conditions

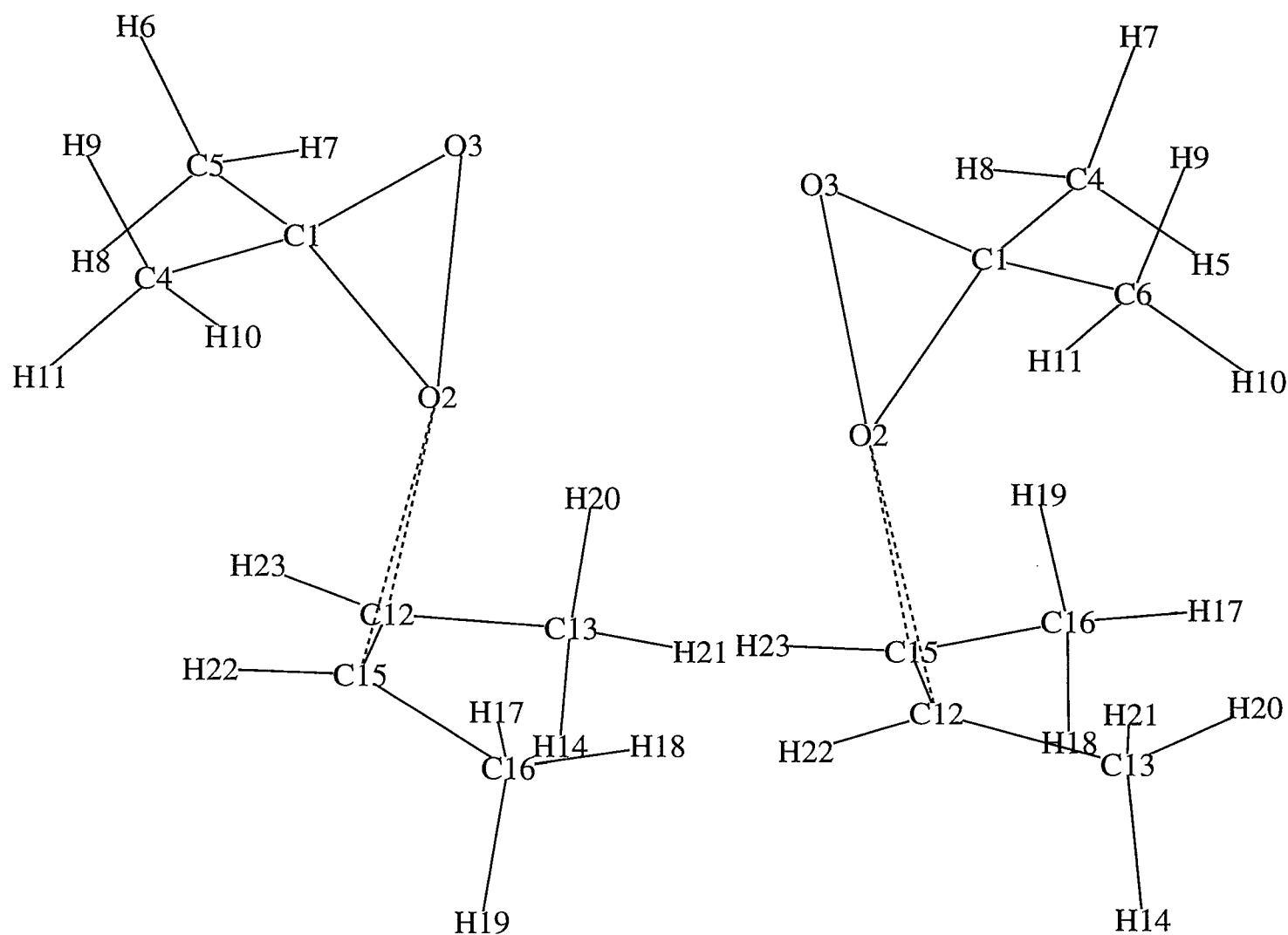
Electronic Supporting Information files are available without a subscription to ACS Web Editions. The American Chemical Society holds a copyright ownership interest in any copyrightable Supporting Information. Files available from the ACS website may be downloaded for personal use only. Users are not otherwise permitted to reproduce, republish, redistribute, or sell any Supporting Information from the ACS website, either in whole or in part, in either machine-readable form or any other form without permission from the American Chemical Society. For permission to reproduce, republish and redistribute this material, requesters must process their own requests via the RightsLink permission system. Information about how to use the RightsLink permission system can be found at <http://pubs.acs.org/page/copyright/permissions.html>



ACS Publications

MOST TRUSTED. MOST CITED. MOST READ.

Copyright © 1997 American Chemical Society



Supplementary Figure 1. Numbering Scheme for *Anti* (left) and *Syn* (right) Systems

Supplementary Table 1a. Non-Bonded Parameters for *Anti* Reactants

Atom	q (e)	σ (Å)	ϵ (kcal/mol)
C1	0.6623	3.50	0.066
O2	-0.2576	2.90	0.140
O3	-0.2576	2.90	0.140
C4	-0.3654	3.50	0.066
C5	-0.3654	3.50	0.066
H6	0.0970	2.50	0.030
H7	0.0979	2.50	0.030
H8	0.0970	2.50	0.030
H9	0.0970	2.50	0.030
H10	0.0979	2.50	0.030
H11	0.0970	2.50	0.030
C12	-0.1856	3.55	0.076
C13	0.1015	3.50	0.066
H14	-0.0056	2.50	0.030
C15	-0.1856	3.55	0.076
C16	0.1018	3.50	0.066
H17	-0.0057	2.50	0.030
H18	0.0024	2.50	0.030
H19	-0.0058	2.50	0.030
H20	-0.0057	2.50	0.030
H21	0.0024	2.50	0.030
H22	0.0929	2.42	0.030
H23	0.0929	2.42	0.030

Supplementary Table 1b. Non-Bonded Parameters for *Anti* Transition Structure

Atom	q (e)	σ (Å)	ϵ (kcal/mol)
C1	0.7983	3.50	0.066
O2	-0.3259	2.90	0.140
O3	-0.5171	2.90	0.140
C4	-0.3738	3.50	0.066
C5	-0.3742	3.50	0.066
H6	0.0842	2.50	0.030
H7	0.0920	2.50	0.030
H8	0.0598	2.50	0.030
H9	0.0840	2.50	0.030
H10	0.0920	2.50	0.030
H11	0.0598	2.50	0.030
C12	0.0040	3.55	0.076
C13	-0.0784	3.50	0.066
H14	0.0348	2.50	0.030
C15	0.0050	3.55	0.076
C16	-0.0796	3.50	0.066
H17	0.0601	2.50	0.030
H18	0.0634	2.50	0.030
H19	0.0349	2.50	0.030
H20	0.0598	2.50	0.030
H21	0.0629	2.50	0.030
H22	0.0769	2.42	0.030
H23	0.0771	2.42	0.030

Supplementary Table 1c. Non-Bonded Parameters for *Syn* Reactants

Atom	q (e)	σ (Å)	ϵ (kcal/mol)
C1	0.6623	3.50	0.066
O2	-0.2577	2.90	0.140
O3	-0.2577	2.90	0.140
C4	-0.3654	3.50	0.066
H5	0.0970	2.50	0.030
C6	-0.3654	3.50	0.066
H7	0.0970	2.50	0.030
H8	0.0979	2.50	0.030
H9	0.0970	2.50	0.030
H10	0.0970	2.50	0.030
H11	0.0979	2.50	0.030
C12	-0.1855	3.55	0.076
C13	0.1015	3.50	0.066
H14	-0.0057	2.50	0.030
C15	-0.1855	3.55	0.076
C16	0.1018	3.50	0.066
H17	0.0024	2.50	0.030
H18	-0.0057	2.50	0.030
H19	-0.0058	2.50	0.030
H20	0.0024	2.50	0.030
H21	-0.0056	2.50	0.030
H22	0.0929	2.42	0.030
H23	0.0929	2.42	0.030

Supplementary Table 1d. Non-Bonded Parameters for *Syn* Transition Structure

Atom	q (e)	σ (Å)	ϵ (kcal/mol)
C1	0.8165	3.50	0.066
O2	-0.3516	2.90	0.140
O3	-0.5249	2.90	0.140
C4	-0.3325	3.50	0.066
H5	0.0429	2.50	0.030
C6	-0.3320	3.50	0.066
H7	0.0702	2.50	0.030
H8	0.0785	2.50	0.030
H9	0.0702	2.50	0.030
H10	0.0426	2.50	0.030
H11	0.0784	2.50	0.030
C12	0.0014	3.55	0.076
C13	-0.0513	3.50	0.066
H14	0.0279	2.50	0.030
C15	0.0002	3.55	0.076
C16	-0.0513	3.50	0.066
H17	0.0460	2.50	0.030
H18	0.0281	2.50	0.030
H19	0.0515	2.50	0.030
H20	0.0465	2.50	0.030
H21	0.0514	2.50	0.030
H22	0.0955	2.42	0.030
H23	0.0958	2.42	0.030

Structure of dimethyldioxirane in PDB format is given below:

```

HEADER
REMARK dimethyldioxirane E = -268.268901 au
HETATM 1 C UNK 1 -0.374 0.258 0.052 0.00 0.00 0
HETATM 2 O UNK 1 -1.683 0.544 -0.367 0.00 0.00 0
HETATM 3 O UNK 1 -1.003 1.307 0.738 0.00 0.00 0
HETATM 4 C UNK 1 0.723 0.663 -0.903 0.00 0.00 0
HETATM 5 H UNK 1 0.870 -0.109 -1.666 0.00 0.00 0
HETATM 6 H UNK 1 0.456 1.604 -1.388 0.00 0.00 0
HETATM 7 H UNK 1 1.670 0.790 -0.365 0.00 0.00 0
HETATM 8 C UNK 1 -0.199 -1.024 0.830 0.00 0.00 0
HETATM 9 H UNK 1 -1.068 -1.182 1.473 0.00 0.00 0
HETATM 10 H UNK 1 -0.095 -1.874 0.147 0.00 0.00 0
HETATM 11 H UNK 1 0.704 -0.975 1.448 0.00 0.00 0
END

```

Structure of *cis*-2-butene in PDB format is given below:

```

HEADER
REMARK cis-2-butene E = -157.2247548 au
HETATM 1 C UNK 1 -0.669 0.783 0.000 0.00 0.00 0
HETATM 2 C UNK 1 0.669 0.783 0.000 0.00 0.00 0
HETATM 3 H UNK 1 -1.170 1.752 0.000 0.00 0.00 0
HETATM 4 H UNK 1 1.170 1.752 0.000 0.00 0.00 0
HETATM 5 C UNK 1 -1.593 -0.403 0.000 0.00 0.00 0
HETATM 6 H UNK 1 -2.252 -0.386 -0.879 0.00 0.00 0
HETATM 7 H UNK 1 -1.061 -1.358 -0.001 0.00 0.00 0
HETATM 8 H UNK 1 -2.250 -0.387 0.881 0.00 0.00 0
HETATM 9 C UNK 1 1.593 -0.403 0.000 0.00 0.00 0
HETATM 10 H UNK 1 2.251 -0.386 0.880 0.00 0.00 0
HETATM 11 H UNK 1 1.061 -1.358 0.001 0.00 0.00 0
HETATM 12 H UNK 1 2.250 -0.387 -0.880 0.00 0.00 0
END

```

Structure of *trans*-2-butene in PDB format is given below:

```

HEADER
REMARK trans-2-butene E = -157.2269605 au
HETATM 1 C UNK 1 0.477 -0.443 0.148 0.00 0.00 0
HETATM 2 C UNK 1 -0.477 0.443 -0.148 0.00 0.00 0
HETATM 3 H UNK 1 -0.193 1.372 -0.648 0.00 0.00 0
HETATM 4 H UNK 1 0.193 -1.372 0.647 0.00 0.00 0
HETATM 5 C UNK 1 1.940 -0.275 -0.150 0.00 0.00 0
HETATM 6 H UNK 1 2.142 0.679 -0.649 0.00 0.00 0
HETATM 7 H UNK 1 2.542 -0.310 0.769 0.00 0.00 0
HETATM 8 H UNK 1 2.311 -1.082 -0.797 0.00 0.00 0
HETATM 9 C UNK 1 -1.940 0.275 0.150 0.00 0.00 0
HETATM 10 H UNK 1 -2.142 -0.679 0.649 0.00 0.00 0
HETATM 11 H UNK 1 -2.311 1.082 0.797 0.00 0.00 0
HETATM 12 H UNK 1 -2.542 0.310 -0.769 0.00 0.00 0
END

```

Structure of the *anti* transition structure for DMD and *cis*-2-butene in PDB format is given below:

```

HEADER
REMARK anti TS cis-2-butene B3LYP/6-31G**/B3LYP/6-31G* E = -425.4715864 au
HETATM 1 C UNK 1 0.822 1.525 0.096 0.00 0.00 0
HETATM 2 O UNK 1 -0.049 2.485 0.378 0.00 0.00 0
HETATM 3 O UNK 1 -0.373 0.642 0.155 0.00 0.00 0
HETATM 4 C UNK 1 -0.336 -1.343 0.560 0.00 0.00 0
HETATM 5 C UNK 1 -0.552 -1.141 -0.788 0.00 0.00 0
HETATM 6 H UNK 1 0.693 -1.496 0.876 0.00 0.00 0
HETATM 7 C UNK 1 -1.376 -1.525 1.619 0.00 0.00 0
HETATM 8 H UNK 1 -2.362 -1.180 1.303 0.00 0.00 0
HETATM 9 H UNK 1 -1.454 -2.587 1.892 0.00 0.00 0
HETATM 10 H UNK 1 -1.094 -0.975 2.523 0.00 0.00 0
HETATM 11 H UNK 1 0.323 -1.150 -1.433 0.00 0.00 0
HETATM 12 C UNK 1 -1.872 -1.060 -1.486 0.00 0.00 0
HETATM 13 H UNK 1 -2.697 -0.862 -0.799 0.00 0.00 0
HETATM 14 H UNK 1 -1.855 -0.265 -2.238 0.00 0.00 0
HETATM 15 H UNK 1 -2.080 -2.004 -2.010 0.00 0.00 0
HETATM 16 C UNK 1 1.830 1.194 1.198 0.00 0.00 0
HETATM 17 H UNK 1 2.559 2.009 1.272 0.00 0.00 0
HETATM 18 H UNK 1 1.306 1.113 2.154 0.00 0.00 0
HETATM 19 H UNK 1 2.385 0.269 1.001 0.00 0.00 0
HETATM 20 C UNK 1 1.430 1.566 -1.307 0.00 0.00 0
HETATM 21 H UNK 1 0.637 1.737 -2.039 0.00 0.00 0
HETATM 22 H UNK 1 2.139 2.400 -1.365 0.00 0.00 0
HETATM 23 H UNK 1 1.975 0.648 -1.560 0.00 0.00 0
END

```

Structure of the *syn* transition structure for DMD and *cis*-2-butene in PDB format is given below:

```

HEADER
REMARK syn TS cis-2-butene B3LYP/6-31G**/B3LYP/6-31G* E = -425.4662337 au
HETATM 1 C UNK 1 0.822 1.525 0.096 0.00 0.00 0
HETATM 2 O UNK 1 -0.049 2.485 0.378 0.00 0.00 0
HETATM 3 O UNK 1 -0.373 0.642 0.155 0.00 0.00 0
HETATM 4 C UNK 1 -0.336 -1.343 0.560 0.00 0.00 0
HETATM 5 C UNK 1 -0.552 -1.141 -0.788 0.00 0.00 0
HETATM 6 H UNK 1 0.693 -1.496 0.876 0.00 0.00 0
HETATM 7 C UNK 1 -1.376 -1.525 1.619 0.00 0.00 0
HETATM 8 H UNK 1 -2.362 -1.180 1.303 0.00 0.00 0
HETATM 9 H UNK 1 -1.454 -2.587 1.892 0.00 0.00 0
HETATM 10 H UNK 1 -1.094 -0.975 2.523 0.00 0.00 0
HETATM 11 H UNK 1 0.323 -1.150 -1.433 0.00 0.00 0
HETATM 12 C UNK 1 -1.872 -1.060 -1.486 0.00 0.00 0
HETATM 13 H UNK 1 -2.697 -0.862 -0.799 0.00 0.00 0
HETATM 14 H UNK 1 -1.855 -0.265 -2.238 0.00 0.00 0
HETATM 15 H UNK 1 -2.080 -2.004 -2.010 0.00 0.00 0
HETATM 16 C UNK 1 1.830 1.194 1.198 0.00 0.00 0
HETATM 17 H UNK 1 2.559 2.009 1.272 0.00 0.00 0
HETATM 18 H UNK 1 1.306 1.113 2.154 0.00 0.00 0
HETATM 19 H UNK 1 2.385 0.269 1.001 0.00 0.00 0
HETATM 20 C UNK 1 1.430 1.566 -1.307 0.00 0.00 0
HETATM 21 H UNK 1 0.637 1.737 -2.039 0.00 0.00 0
HETATM 22 H UNK 1 2.139 2.400 -1.365 0.00 0.00 0
HETATM 23 H UNK 1 1.975 0.648 -1.560 0.00 0.00 0
END

```

Structure of the transition structure for DMD and *trans*-2-butene in PDB format is given below:

```

HEADER
REMARK TS trans-2-butene B3LYP/6-31G**/B3LYP/6-31G* E = -425.4711724 au
HETATM 1 C UNK 1 -0.274 1.588 0.361 0.00 0.00 0
HETATM 2 O UNK 1 0.704 2.478 0.274 0.00 0.00 0
HETATM 3 O UNK 1 0.806 0.615 0.010 0.00 0.00 0
HETATM 4 C UNK 1 0.601 -1.014 -1.014 0.00 0.00 0
HETATM 5 C UNK 1 0.831 -1.509 0.251 0.00 0.00 0
HETATM 6 H UNK 1 -0.430 -0.973 -1.362 0.00 0.00 0
HETATM 7 C UNK 1 1.664 -0.822 -2.053 0.00 0.00 0
HETATM 8 H UNK 1 2.654 -0.748 -1.596 0.00 0.00 0
HETATM 9 H UNK 1 1.667 -1.668 -2.754 0.00 0.00 0
HETATM 10 H UNK 1 1.481 0.091 -2.627 0.00 0.00 0
HETATM 11 H UNK 1 1.865 -1.604 0.577 0.00 0.00 0
HETATM 12 C UNK 1 -1.350 1.665 -0.722 0.00 0.00 0
HETATM 13 H UNK 1 -1.979 2.544 -0.539 0.00 0.00 0
HETATM 14 H UNK 1 -0.874 1.782 -1.700 0.00 0.00 0
HETATM 15 H UNK 1 -2.004 0.785 -0.732 0.00 0.00 0
HETATM 16 C UNK 1 -0.822 1.353 1.768 0.00 0.00 0
HETATM 17 H UNK 1 0.013 1.215 2.459 0.00 0.00 0
HETATM 18 H UNK 1 -1.386 2.240 2.080 0.00 0.00 0
HETATM 19 H UNK 1 -1.495 0.490 1.823 0.00 0.00 0
HETATM 20 C UNK 1 -0.218 -2.015 1.185 0.00 0.00 0
HETATM 21 H UNK 1 -1.229 -1.846 0.802 0.00 0.00 0
HETATM 22 H UNK 1 -0.090 -3.097 1.335 0.00 0.00 0
HETATM 23 H UNK 1 -0.134 -1.548 2.173 0.00 0.00 0
END

```

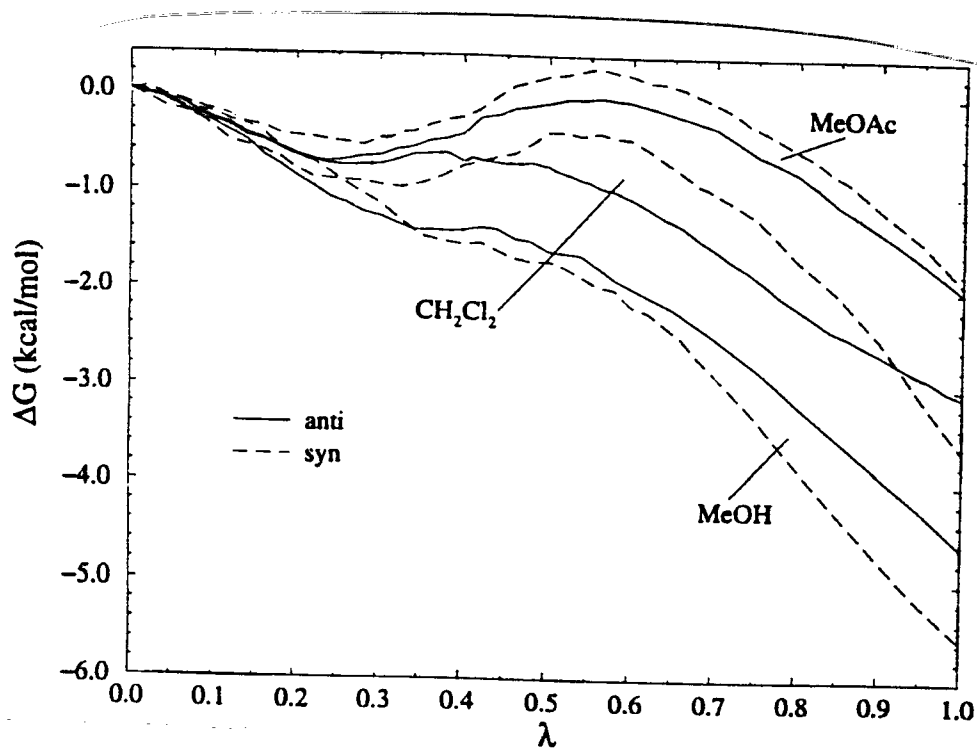


Figure 2. Computed changes in free energies of solvation in CH_3OAc , CH_2Cl_2 , and CH_3OH for the epoxidations of *cis*-2-butene leading from the reactants at $l = 0.0$ to the anti and syn transition structures at $\lambda = 1.0$ (λ is the reaction coordinate used in the FEP calculations).