On the choice of optimal conformation in linear free-energy relationships. Reactivity of 2-[(carboxymethyl)sulfanyl]-4-oxo-4-arylbutanoic acids with diphenyldiazomethane

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Table S1. The second-order rate constants for esterification of 1-11 by DDM (k_2) , their logarithmic values, the Swain-Lupton F and R constants, the backbone RMSD (MD minimums *vs.* same conformations optimized at the semiempirical level), the range of solvent accessible surface area from MD trajectory (Range SA), the same value divided by the molecular weight of compounds (Range SA/M_w).

| Comp- ound | R- | k ₂ (dm ³ mol ⁻¹ s ⁻¹) | logk ₂ | log(k/k ₀) | F | R* | Backbone RMSD | Range SA | Range SA/M _w |
|---------------|------------------|--|-------------------|------------------------|-------|-------|------------------|----------|----------------------------|
| 1 | H- | 0.282 | -0.5498 | / | 0 | 0 | 0.640 | 89.09 | 0.332 |
| 2 | 4-Me- | 0.315 | -0.5017 | 0.9125 | 0.01 | -0.18 | 0.922 | 88.80 | 0.315 |
| 3 | 4-Et- | 0.339 | -0.4698 | 0.8545 | 0 | -0.30 | 0.781 | 104.50 | 0.353 |
| 4 | 4- <i>i</i> -Pr- | 0.346 | -0.4609 | 0.8383 | 0.04 | -0.32 | 0.279 | 109.11 | 0.352 |
| 5 | 4- <i>n</i> -Bu- | 0.323 | -0.4908 | 0.8927 | -0.01 | -0.28 | 0.460 | 135.92 | 0.419 |
| 6 | 4- <i>t</i> -Bu- | 0.278 | -0.5560 | 1.0112 | -0.02 | -0.04 | 0.599 | 108.93 | 0.336 |
| 7 | 4-F- | 0.378 | -0.4225 | 0.7685 | 0.45 | -0.39 | 0.708 | 89.16 | 0.311 |
| 8 | 4-Cl- | 0.329 | -0.4828 | 0.8781 | 0.42 | -0.19 | 0.526 | 91.86 | 0.303 |
| 9 | 4-Br- | 0.351 | -0.4547 | 0.8270 | 0.45 | -0.30 | 0.362 | 93.76 | 0.270 |
| 10 | 3,4-di-Me- | 0.368 | -0.4342 | 0.7897 | 0.02 | -0.36 | 0.540 | 98.29 | 0.332 |
| 11 | 2,5-di-Me- | 0.385 | -0.4145 | 0.7540 | / | / | 0.313 | 102.03 | 0.344 |

* R^+ given in red; R^- given in blue.

Table S2. Values of highest occupied molecular orbitals (HOMO), lowest occupied molecular orbitals (LUMO), and their differences (diff), for compounds 1-11; as obtained by the semiempirical PM6 method, and by single point calculations using the DFT and MP2 methods.

| Compo- | . р | PM6 | | | DFT | MP2 | | | |
|--------|------------------|--------|--------|--------|-----------------|---------|---------|--------|----------|
| und | К- | номо | LUMO | diff | HOMO LUMO | diff | HOMO | LUMO | diff |
| 1 | H- | -9.660 | -0.813 | -8.847 | -6.5413 -1.8955 | -4.6458 | -9.4195 | 2.0343 | -11.4538 |
| 2 | 4-Me- | -9.662 | -0.807 | -8.855 | -6.4592 -1.9154 | -4.5438 | -9.1789 | 1.9464 | -11.1254 |
| 3 | 4-Et- | -9.694 | -0.784 | -8.910 | -6.5506 -1.8931 | -4.6575 | -9.1716 | 2.0147 | -11.1863 |
| 4 | 4- <i>i</i> -Pr- | -9.731 | -0.836 | -8.895 | -6.7321 -1.7453 | -4.9868 | -9.1452 | 2.1598 | -11.3050 |
| 5 | 4- <i>n</i> -Bu- | -9.692 | -0.787 | -8.905 | -6.6102 -1.7565 | -4.8537 | -9.0932 | 2.1304 | -11.2236 |
| 6 | 4- <i>t</i> -Bu- | -9.445 | -1.031 | -8.414 | -6.2385 -2.1195 | -4.1190 | -9.1147 | 1.7783 | -10.8930 |
| 7 | 4-F- | -9.898 | -0.939 | -8.959 | -6.7985 -1.7374 | -5.0610 | -9.5066 | 2.2036 | -11.7102 |
| 8 | 4-Cl- | -9.605 | -1.004 | -8.601 | -6.3647 -1.8406 | -4.5242 | -9.3809 | 2.0332 | -11.4141 |
| 9 | 4-Br- | -9.776 | -1.038 | -8.738 | -6.6404 -1.8904 | -4.7500 | -9.3966 | 1.9649 | -11.3616 |
| 10 | 3,4-di-Me- | -9.583 | -0.872 | -8.711 | -6.5473 -1.6351 | -4.9122 | -8.9721 | 2.2833 | -11.2554 |
| 11 | 2,5-di-Me- | -9.428 | -0.654 | -8.774 | -6.3084 -1.1902 | -5.1182 | -8.7974 | 3.0801 | -11.8775 |

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| surfac | Irrace area (PSA) derived from MD trajectories. | | | | | | | |
|--------|---|--------------------|-------------------------|--------------------|------------------------|------------------------|------------------------|----------------------------------|
| q | Experimental | | | | | | | |
| unodu | | | | Swain- Lupton | PM6 | DFT | MP2 | |
| Con | R- | log k ₂ | log(k/k ₀)* | log k ₂ | log(k/k ₀) | log(k/k ₀) | log(k/k ₀) | Average PSA (Å ²) |
| 1 | H- | -0.5498 | / | -0.5592 | / | / | / | 223.525 |
| 2 | 4-Me- | -0.5017 | 0.9125 | -0.5043 | 0.8863 | 0.9008 | 0.9007 | 224.776 |
| 3 | 4-Et- | -0.4698 | 0.8545 | -0.4683 | 0.8697 | 0.8746 | 0.8868 | 223.043 |
| 4 | 4- <i>i</i> -Pr- | -0.4609 | 0.8383 | -0.4606 | 0.8504 | 0.7987 | 0.8596 | 223.108 |
| 5 | 4- <i>n</i> -Bu- | -0.4908 | 0.8927 | -0.4747 | 0.8707 | 0.8294 | 0.8782 | 220.633 |
| 6 | 4- <i>t</i> -Bu- | -0.5560 | 1.0112 | -0.5479 | 0.9991 | 0.9987 | 0.9540 | 223.395 |
| 7 | 4-F- | -0.4225 | 0.7685 | -0.4232 | 0.7636 | 0.7816 | 0.7667 | 221.638 |
| 8 | 4-Cl- | -0.4828 | 0.8781 | -0.4850 | 0.9159 | 0.9053 | 0.8346 | 223.084 |
| 9 | 4-Br- | -0.4547 | 0.8270 | -0.4505 | 0.8271 | 0.8533 | 0.8466 | 220.873 |
| 10 | 3,4-di-Me- | -0.4342 | 0.7897 | -0.4493 | / | 0.8159 | 0.8709 | 223.695 |
| 11 | 2,5-di-Me- | -0.4145 | 0.7540 | / | / | 0.7684 | 0.7284 | 218.968 |

Table S3. The experimental and predicted rate constants, as obtained by the Swain-Lupton approach; and by the semiempirical (PM6), DFT, and MP2 calculations. The average polar surface area (PSA) derived from MD trajectories.

* $\overline{k_0}$ – refer to the second order rate constant for unsubstituted derivative (1)

| Compound | R- | pK _{a1} | pK _{a2} |
|----------|------------------|------------------|------------------|
| 1 | H- | 3.27 | 4.62 |
| 2 | 4-Me- | 3.28 | 4.63 |
| 3 | 4-Et- | 3.31 | 4.67 |
| 4 | 4- <i>i</i> -Pr- | 3.33 | 4.69 |
| 5 | 4- <i>n</i> -Bu- | 3.35 | 4.73 |
| 6 | 4- <i>t</i> -Bu- | 3.34 | 4.72 |
| 7 | 4-F- | 3.24 | 4.57 |
| 8 | 4-Cl- | 3.17 | 4.44 |
| 9 | 4-Br- | 3.25 | 4.54 |
| 10 | 3,4-di-Me- | 3.28 | 4.63 |
| 11 | 2,5-di-Me- | 3.27 | 4.62 |

Table S4. Estimated pK_a values for compounds 1-11, see Figure S9.



Figure S1. Experimental vs. calculated log k₂ for compounds 1-10.



Figure S2. The two-dimensional free energy landscape of compound **11**, obtained by the ABF calculation. X-axis shows distance between centroids defined on the phenyl, and on the distal carboxyl (collective variable 1). Y-axis show radius of gyration of heavy atoms (collective variable 2). See Figure S3.

Supplementary data



Figure S3. a) Collective variable 1; distance is given as a red line. c) Collective variable 2; radius of gyration is measured for all heavy atoms, except for those included in the first collective variable. Exemplified for compound **11**.



Figure S4. Root-mean-square deviation (rmsd) of MD minimum *vs*. the same conformation optimized on semiempirical level, for heavy atoms of backbone (excluding substituents), ending on carboxyl carbons (<u>Ph-C(O)-CH₂-CH(S-CH₂-COOH)-C</u>OOH). The main contribution to the rmsd were Ph-C(O) torsion. The initial MD minimum is given in blue, the same conformation optimized on the semiempirical level is given in red. Exemplified for compound **3**.



Figure S5. Experimental $log(k/k_0)$ vs. highest occupied molecular orbital (HOMO) energy, obtained by the semiempirical calculations, for compounds 1-11.



Figure S6. Experimental $log(k/k_0)$ vs. energy difference between highest occupied (HOMO) and lowest unoccupied (LUMO) molecular orbitals, obtained by the DFT calculations, for compounds 2-11.



Figure S7. Experimental $log(k/k_0)$ *vs*. energy difference between highest occupied (HOMO) and lowest unoccupied (LUMO) molecular orbitals, obtained by the MP2 calculations, for compounds **2-11**.



Figure S8. Experimental $log(k/k_0)$ vs. conformation (3D) dependent polar surface area (PSA) of compounds 2-11. Trend is assigned by colored line.



Figure S9. Estimated pH profile for compound 1.



a) b) Figure S10. Experimentally obtained $log(k/k_0) vs$. estimated pK_{a1} (a) and pK_{a2} (b) for compounds 2-11. Compound 8 is outlier. Trend is assigned by colored line.