

Supporting Information

Catalysis by solvation rather than desolvation effect: Exploring the Catalytic power of Methyltransferases

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Figure S1. The two EVB states and atom types used to simulate the methyl transfer from SAM to chloride.

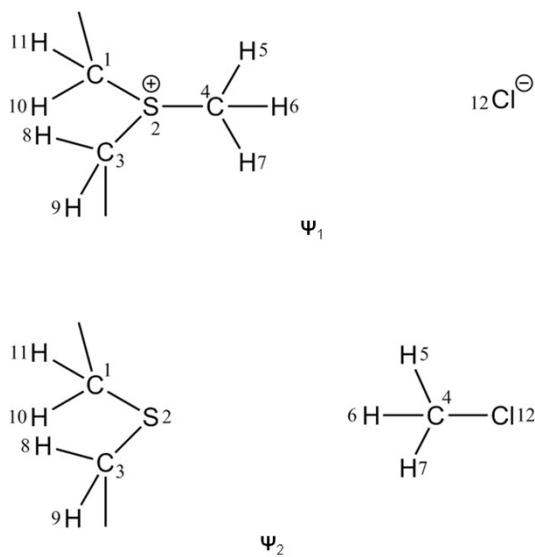


Table S1. Activation and reaction energies (kcal/mol) obtained from HF, M062X and MP2 with set basis 6-31+G(d,p), for reaction involving Cl⁻ and trimethylsulfonium in gaseous and aqueous environment. The ZPVEs were included in the calculations.

	HF		M062X		MP2	
	Gas	Solvent	Gas	Solvent	Gas	Solvent
ΔE^* (kcal/mol)	5.44	30.29	1.39	10.89	3.76	17.37
ΔE^0 (kcal/mol)	-35.30	7.94	-34.85	0.80	-36.50	4.27

Table S2. Key distance calculated at HF, M062X and MP2 with set basis 6-31+G(d,p), level of optimization for Reactants State (RS), Transitions State (TS) and Products State (PS).

	HF			M062X			MP2		
	RS	TS	PS	RS	TS	PS	RS	TS	PS
S-C	1.82	2.18	3.22	1.83	2.05	3.12	1.81	2.08	3.13
C-Cl	3.01	2.52	1.79	2.90	2.52	1.79	2.93	2.47	1.78
Cl-S	4.78	4.70	5.00	4.62	4.58	4.91	4.64	4.55	4.92

Table S3. Key distance calculated at EVB for reaction involving Cl⁻ and trimethylsulfonium, for the Reactants State (RS), the Transitions State (TS) and Products State (PS) on Water, Protein and Mutant (Y70T).

	Water			Protein			Mutant		
	RS	TS	PS	RS	TS	PS	RS	TS	PS
S-C	1.81	1.98	5.58	1.77 (± 0.01)	1.93 (± 0.02)	4.13 (± 0.52)	1.78 (± 0.01)	2.00 (± 0.03)	3.42 (± 0.11)
C-Cl	3.35	2.13	1.73	2.67 (± 0.23)	2.75 (± 0.05)	1.77 (± 0.06)	2.79 (± 0.03)	2.22 (± 0.04)	1.84 (± 0.04)
Cl-S	4.49	4.10	4.81	4.29 (± 0.23)	4.18 (± 0.04)	4.76 (± 0.42)	3.92 (± 0.09)	4.19 (± 0.03)	4.38 (± 0.42)

Table S4. The EVB Parameters for reaction involving Cl⁻ and trimethylsulfonium in solution.

Charges:

Reactant State			Product State	
Atom number	Atom Type	Charge	Atom Type	Charge
1	C0	-0.354179	C0	-0.184985
2	S+	0.314698	S0	-0.361849
3	C0	-0.354179	C0	-0.184985
4	C0	-0.354179	C0	-0.530236
5	H0	0.238665	H0	0.149500
6	H0	0.238665	H0	0.149500
7	H0	0.238665	H0	0.149500
8	H0	0.238665	H0	0.017991
9	H0	0.238665	H0	0.282833
10	H0	0.238665	H0	0.282833
11	H0	0.238665	H0	0.282833
12	L ⁻	-0.922816	L0	-0.131068

$$1 - e^{\{-a(b-b_0)\}_2} \Delta M = D$$

Bonds:

Bond Type	D	b_0	a
C0-S+	70.0	1.80	2.0
C0-S0	70.0	1.80	2.0
C0-L-	120.0	1.40	0.8
C0-L0	120.0	1.35	0.8

$$U_0 = \frac{1}{2} K_\theta (\theta - \theta_0)^2$$

Bond Angles:

Bond Type	θ	K_θ
X-S+-Y	102.5	100.0
X-S0-Y	103.5	100.0
X-C0-Y	109.5	50.0
X-L- -Y	120.0	50.0
X-L0-Y	109.0	50.0

$$U_{nb} = A_i A_j r^{-12} - B_i B_j r^{-6}$$

Nobonded :

Bond Type	A	B
H0	7.00	0.00
S+	1831.8	23.0
S0	1022.5	23.0

	C0	632.0	24.0
	L-	1400.0	24.0
	L0	774.0	24.0
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Nobonded:	$U_{nb} = Ae^{\{-ar\}}$		

	Bond Type	A	a
	C0...S+	19999.0	4.0
	C0...S0	19999.0	4.0

off-diagonal and shift parameters

A=-21.0

α = -140.0

μ =0.0

Table S5. Contributions to Solvation free Energy (kcal/mol) for the Reactant State (RS) and the Transition State (TS) in the water reference reaction. The energies were obtained through LRA.

	Water		Water	
	RS (SAM)	RS (CI)	TS (SAM)	TS (CI)
Intra	-99.04	-63.62	-78.9	-44.56
Solvation/p	0.00	0.00	0.00	0.00
Solvation /w	4.82	-24.11	3.21	-14.36
Elect (total)	-94.21	-87.74	-75.69	-56.92
$\langle U_Q - U_0 \rangle_Q$	-125.55	-155.08	-108.83	-120.01
$\langle U_Q - U_0 \rangle_0$	5.57	10.43	29.06	39.6
ΔG_{solv}	-59.99	-72.33	-39.88	-40.21

Table S5. Contributions to Solvation free Energy (kcal/mol) for the Reactant State (RS) and the Transition State (TS) in the protein native.

	Protein		Protein	
	RS (SAM)	RS (CI)	TS (SAM)	TS (CI)
Intra	-96.42 (± 0.26)	-61.10 (± 0.44)	-78.24 (± 2.74)	-46.80 (± 1.62)
Solvation/p	30.82 (± 0.93)	-60.22 (± 1.02)	27.16 (± 3.89)	-61.10 (± 2.83)
Solvation /w	-16.82 (± 0.43)	16.95 (± 0.57)	-13.43 (± 8.10)	22.85 (± 3.43)
Elect (total)	-82.43 (± 0.24)	-104.38 (± 1.16)	-64.52 (± 1.47)	-85.05 (± 2.22)
$\langle U_Q - U_0 \rangle_Q$	-109.40 (± 1.03)	-130.06 (± 0.09)	-96.99 (± 7.86)	-109.68 (± 2.29)
$\langle U_Q - U_0 \rangle_0$	-3.62 (± 1.46)	-43.51 (± 2.95)	13.56 (± 3.7)	-25.00 (± 0.01)
ΔG_{solv}	-56.51 (± 1.25)	-86.78 (± 1.43)	-41.71 (± 2.08)	-67.34 (± 1.15)

Table S5. Contributions to Solvation free Energy (kcal/mol) for the Reactant State (RS) and the Transition State (TS) in the Mutant.

	Mutant		Mutant	
	RS (SAM)	RS (CI)	TS (SAM)	TS (CI)
Intra	-133.90 (± 37.89)	-68.94 (± 0.06)	-82.85 (± 0.08)	-46.79 (± 0.26)
Solvation/p	30.87 (± 1.82)	-56.84 (± 0.24)	27.75 (± 3.09)	-52.70 (± 2.06)
Solvation /w	-8.93 (± 1.61)	22.22 (± 2.65)	-11.79 (± 0.40)	17.05 (± 1.33)
Elect (total)	-84.97 (± 0.49)	-103.56 (± 2.47)	-66.88 (± 3.41)	-80.93 (± 2.70)
$\langle U_Q - U_0 \rangle_Q$	-133.86 (± 0.30)	-136.62 (± 1.67)	-91.92 (± 0.39)	-112.95 (± 3.5)
$\langle U_Q - U_0 \rangle_0$	-8.84 (± 0.48)	-40.79 (± 2.55)	21.05 (± 12.21)	-11.36 (± 1.9)
ΔG_{solv}	-61.35 (± 0.38)	-88.7 (± 2.11)	-35.43 (± 5.9)	-62.16 (± 3.1)