

**Supporting Information: Does water relay play an important role in phosphoryl transfer reactions? Insights from theoretical study of a model reaction in water and *tert*-butanol**

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**1. Test for the MM model of *tert*-butanol**

As shown in Table S1, a few key properties (density, diffusion constant and heat of vaporization) are well reproduced by the MM model for the *tert*-butanol.

**2. Benchmark calculations for SCC-DFTBPR for pseudo-rotation barriers**

As pointed by York et al.,<sup>1</sup> conventional semiempirical methods (MNDO, AM1, PM3 and MNDO/d methods are tested) are not reliable for pseudo-rotation and that generally the difference between the semi-empirical and DFT results exceed 50% of the actual barrier at the DFT level. Therefore, it is crucial to test the performance of SCC-DFTBPR in this context to show that the corresponding PMFs that we calculate are meaningful. Accordingly, we have selected representative pseudo-rotation processes from the QCRNA database<sup>2</sup> (10 neutral plus 10 negatively charged cases) and compared the SCC-DFTBPR barriers to high-level DFT data in that database; for further comparison, we have also included results from the second-order SCC-DFTB and a

number of standard semi-empirical methods (see Table S2, S3 and S4). in the **Supporting Materials**).

If the B3LYP/6-31++G(d,p) structures are used and only single point energy calculations are performed, SCC-DFTBPR gives consistently better results than the popular semi-empirical methods. For example, for the 10 neutral systems (5 acyclic and 5 cyclic), SCC-DFTBPR has a RMSE of 2.09 kcal/mol, while the values for AM1, PM3, MNDO are 3.20, 5.50, 3.85 kcal/mol, respectively; the second order SCC-DFTB has a similar RMSE of 2.50 kcal/mol. For the 10 negatively charged cases, the performance becomes worse for all methods; the RMSE for SCC-DFTBPR is 3.72 kcal/mol, and the values for AM1, PM3 and MDNO are 7.51, 5.01 and 5.73 kcal/mol, respectively. The exception is that the second-order SCC-DFTB remains close to the high-level DFT results with a RMSE of 1.6 kcal/mol, even smaller than for the neutral cases.

Besides the single-point calculations, geometry optimizations also indicate that SCC-DFTBPR is a reasonable method for describing pseudo-rotation. For example, for the P(O)(OH)(-O-sugar-O-)(OCH3)\_PR case from QCRNA data base, which is structurally similar to the model UNP studied here, the SCC-DFTBPR barrier after geometry optimization is 5.76 kcal/mol, which is only 0.10 kcal/mol higher than the B3LYP/6-311++G(3df,2p)//B3LYP/6-31++G(d,p) reference. For more test cases, see Table S4.

These gas-phase benchmark calculations indicate that SCC-DFTBPR can be used to meaningfully compare the energetics of pseudo-rotation for the UNP system in different solutions.

### **3. van der Waals parameters for the SCC-DFTBPR atoms**

As shown in Fig. S1, the radial distributions of the solvent (water) near the solute (model UNP) do not change in any major way when the QM region expands from solute-only to include all water molecules within 6 Å from the phosphorus atom or the oxygen in the ribose ring; the results also do not change when different van der Waals parameters (i.e., using the CHARMM parameters vs. optimized values for the key oxygen atoms:  $\epsilon = -0.1397$  kcal/mol;  $r_0 = 1.698$  Å) are used for the solute-QM simulations. This suggests that the QM/MM results reported in the main text are not expected to depend sensitively on the van der Waals parameters for the QM atoms.

#### **4. Reaction coordinate for the pseudo-rotation PMF calculations**

As shown in Fig. S2, there is clear linear correlations between the approximate reaction coordinate ( $d(O_7-O_{12})-d(O_8-O_{11})$ ) and the relevant angles that characterize the pseudo-rotation process in both the aqueous and *tert*-butanol simulations. Therefore, the computed PMF profiles can be meaningful compared to characterize the energetic differences between the two solution conditions.

#### **5. Full Reference for CHARMM 22 force field (Ref. 40 in main text)**

MacKerell, A. D.; Bashford, D.; Bellott, M.; Dunbrack, R. L.; Evanseck, J. D.; Field, M. J.; Fischer, S.; Gao, J.; Guo, H.; Ha, S.; Joseph-McCarthy, D.; Kuchnir, L.; Kuczera, K.; Lau, F. T. K.; Mattos, C.; Michnick, S.; Ngo, T.; Nguyen, D. T.; Prodhom, B.; Reiher, W. E.; Roux, B.; Schlenkrich, M.; Smith, J. C.; Stote, R.; Straub, J.; Watanabe, M.; Wiorkiewicz-Kuczera, J.; Yin, D.; Karplus, M. *J. Phys. Chem. B* **1998**, 102, 3586–3616.

## References

1. (a) C. S. Lopez, O. N. Faza, B. A. Gregersen, X. Lopez, A. R. de Lera and D. M. York, *ChemPhysChem*. 5, 1045-1049 (2004) (b) C. S. Lopez, O. N. Faza, A. R. de Lera and D. M. York, *Chem. Eur. J.* 11, 2081-2093 (2005)
2. T. J. Giese et al., and D. M. York, *J. Mol. Graph. Model.* 25, 423-433 (2006)
3. D. Riccardi, G. Li and Q. Cui, *J. Phys. Chem. B* 108, 6467-6478 (2004)

## Figure Captions

### Figure S1.

Radial distribution functions,  $g(r)$  (solid lines) and running coordinate numbers,  $rcn$  (dashed lines), of water oxygen around different oxygen atoms of the model UNP in SCC-DFTBPR/MM simulations. Black indicates simulations in which the QM region includes the model UNP and the first solvation shell (within 6 Å from the phosphorus atom or the oxygen atom in the ribose ring; the partitioning of the QM/MM water molecules is updated on the fly). Red/green indicates simulations with only the solute treated as QM; the van der Waals parameters in those simulations are either the standard CHARMM parameters (red) or optimized based on standard protocols (green).<sup>3</sup>

### Figure S2.

Correlation between the defined reaction coordinate ( $d(O_7-O_{12})-d(O_8-O_{11})$ ) and the relevant angles that characterize the pseudo-rotation process in the aqueous solution (left) and *tert*-butanol (right) PMF simulations. Each data point indicates the average value from a given window. For the atomic labels, see Fig. 1 in the main text.

Table S1. Comparison of computed physical properties to experimental data  
for *tert*-butanol

Properties	Experiment <sup>a</sup>	Computation
Number density ( $\text{\AA}^{-3}$ )	0.006342	0.006395
Self-diffusion coefficient ( $\text{cm}^2/\text{s}$ )	N/A	$0.77054 \times 10^{-5}$
Heat of vaporization (kcal/mol)	10.20	11.86

a. The value refers to 298.15K; from CRC Handbook.

Table S2. The performance of several semi-empirical methods for the pseudo-rotation barrier of ten neutral phosphates in the QCRNA database <sup>a</sup>

Reactions	Ring Type	DFT Reference <sup>b</sup>	SCC-DFTB <sup>c</sup>	SCC-DFTBPR <sup>c</sup>	AM1 <sup>c</sup>	PM3 <sup>c</sup>	MNDO <sup>c</sup>
P(OH)(OCH3)(-O-CH2CH2-O-)(OCH3)_PR	-CH2CH2-	7.86	-3.95	-3.09	-3.24	-6.23	-5.11
P(OH)(OH)(-O-CH2CH2-O-)(OCH3)_PR	-CH2CH2-	7.76	-3.17	-3.57	-3.51	-6.85	-5.48
P(OH)(OH)(-O-CH2CH2-O-)(OH)_PR	-CH2CH2-	5.15	-2.22	-0.19	-1.50	-4.58	-3.30
P(OH)(OH)(-O-sugar-O-)(OCH3)_PR	-ribose-	6.75	-3.03	-3.22	-2.97	-5.88	-4.00
P(OH)(OH)(-O-sugar-O-)(OH)_PR	-ribose-	4.06	-1.68	0.50	-0.84	-3.66	-2.20
P(OCH3)(OCH3)(OCH3)(OCH3)(OCH3)_PR	acyclic	4.33	-0.48	0.33	-2.28	-3.41	-1.72
P(OH)(OCH3)(OCH3)(OCH3)(OCH3)_PR	acyclic	7.60	-2.30	-1.17	-3.91	-5.72	-4.50
P(OH)(OH)(OCH3)(OCH3)(OCH3)_PR	acyclic	8.47	-2.80	-2.22	-4.61	-6.82	-5.13
P(OH)(OH)(OCH3)(OH)(OCH3)_PR	acyclic	6.36	-2.35	-1.01	-3.95	-5.90	-2.89
P(OH)(OH)(OCH3)(OH)(OH)_PR	acyclic	5.81	-0.99	-1.80	-3.22	-4.74	-1.69
MAXE <sup>d</sup>			-3.95/-3.95/-2.8	-3.57/-3.57/-2.22	-4.61/-3.51/-4.61	-6.85/-6.85/-6.82	-5.48/-5.48/-5.13
RMSE <sup>d</sup>			2.50/2.91/1.99	2.09/2.57/1.46	3.20/2.63/3.68	5.50/5.56/5.44	3.85/4.19/3.49
MUE <sup>d</sup>			2.30/2.81/1.78	1.71/2.11/1.31	3.00/2.41/3.59	5.38/5.44/5.32	3.60/-4.02/3.19
MSE <sup>d</sup>			-2.30/-2.81/-1.78	-1.54/-1.91/-1.17	-3.00/-2.41/-3.59	-5.38/-5.44/-5.32	-3.60/4.02/-3.19

- a) <http://theory.chem.umn.edu/QCRNA/>. The notations follow the QCRNA convention. For the semi-empirical methods including SCC-DFTB(PR), the values are single point energies at the B3LYP geometries.
- b) Gas phase potential energy barrier at the level of B3LYP/6-311++G(3df,2p)//B3LYP/6-31++G(d,p) in kcal/mol.
- c) Barrier difference from the DFT reference, i.e.,  $E_{\text{Semiempirical}} - E_{\text{DFT}}$ ; for AM1, PM3, MNDO, reaction enthalpy is used. “SCC-DFTB” is the second-order method, “SCC-DFTBPR” is the 3<sup>rd</sup>-order parameterization for phosphate chemistry (Ref. 29 in the main text).
- d) MAXE: largest absolute error; RMSE: root-mean-square error; MUE: mean unsigned error; MSE: mean signed error. Before the slash refers to the overall statistics including both the cyclic and acyclic cases; between the slash refers to cyclic cases; after the slash refers to acyclic cases.

Table S3. The performance of several semi-empirical methods for the pseudo-rotation barrier of ten anionic phosphates in the QCRNA database <sup>a</sup>

Reactions	Ring Type	DFT Reference	SCC-DFTB	SCC-DFTBPR	AM1	PM3	MNDO
P(O)(OCH3)(-O-CH2CH2-O-)(OCH3)_PR	-CH2CH2-	0.99	-0.47	-0.50	-4.07	-3.03	-2.09
P(O)(OH)(-O-CH2CH2-O-)(OCH3)_PR	-CH2CH2-	4.70	-0.95	-1.71	-10.51	-4.98	-8.36
P(O)(OH)(-O-CH2CH2-O-)(OH)_PR	-CH2CH2-	2.01	-1.59	1.64	-10.96	-7.83	-6.68
P(O)(OH)(-O-sugar-O-)(OCH3)_PR	-ribose-	5.66	-0.15	-1.80	-5.37	-4.72	-8.15
P(O)(OH)(-O-sugar-O-)(OH)_PR	-ribose-	1.38	-0.44	2.04	-3.78	-3.08	0.00
P(O)(OCH3)(OCH3)(OCH3)(OCH3)_PR	acyclic	1.75	1.01	1.80	-4.99	-3.64	-0.71
P(O)(OH)(OCH3)(OCH3)(OCH3)_PR	acyclic	6.65	-1.32	-3.62	-5.61	-2.17	-4.50
P(O)(OH)(OCH3)(OH)(OCH3)_PR	acyclic	4.15	-1.31	-0.66	-8.57	-6.63	-3.09
P(O)(OH)(OCH3)(OH)(OH)_PR	acyclic	8.40	-3.49	-6.03	-8.69	-7.05	-7.07
P(O)(OH)(OH)(OCH3)(OCH3)_PR	acyclic	12.36	-2.31	-8.48	-8.24	-3.47	-7.90
MAXE			-3.49/-1.59/-3.49	-8.48/2.04/-8.48	-10.96/-10.96/-8.69	-7.83/-7.83/-7.05	-8.36/-8.36/-7.90
RMSE			1.61/0.88/2.10	3.72/1.63/5.00	7.51/7.62/7.39	5.01/5.04/4.97	5.73/6.09/5.34
MUE			1.30/0.72/1.89	2.83/1.54/4.12	7.08/6.94/7.22	4.66/4.73/4.59	4.86/5.06/4.65
MSE			-1.10/-0.72/-1.48	-1.73/-0.07/-3.40	-7.08/-6.94/-7.22	-4.66/-4.73/-4.59	-4.86/-5.06/-4.65

a. refer to Table S2 for notations.



Table S4. The performance of several semi-empirical methods with geometry optimization for the pseudo-rotation barrier of selected phosphates in the QCRNA database <sup>a</sup>

Reactions	Charge	DFT Reference	SCC-DFTB	SCC-DFTBPR	AM1	PM3	MNDO
P(O)(OH)(-O-sugar-O-)(OCH3)_PR	-1	5.66	-0.92 <sup>e</sup>	0.10	-1.77	-1.80	-0.77
P(O)(OH)(-O-sugar-O-)(OH)_PR	-1	1.38	0.37	3.71	-1.50	1.50	-0.88
P(OH)(OH)(-O-sugar-O-)(OCH3)_PR	0	6.75	-2.86	-3.13	-6.01	-5.73	-3.64
P(OH)(OH)(-O-sugar-O-)(OH)_PR	0	4.06	-1.74	-0.22	-0.22	-3.00	-1.20

a. Please refer to Table S2 for notation. Optimizations with SCC-DFTB(PR) are done with CHARMM, while those for AM1/PM3/MNDO are done with Gaussian03.

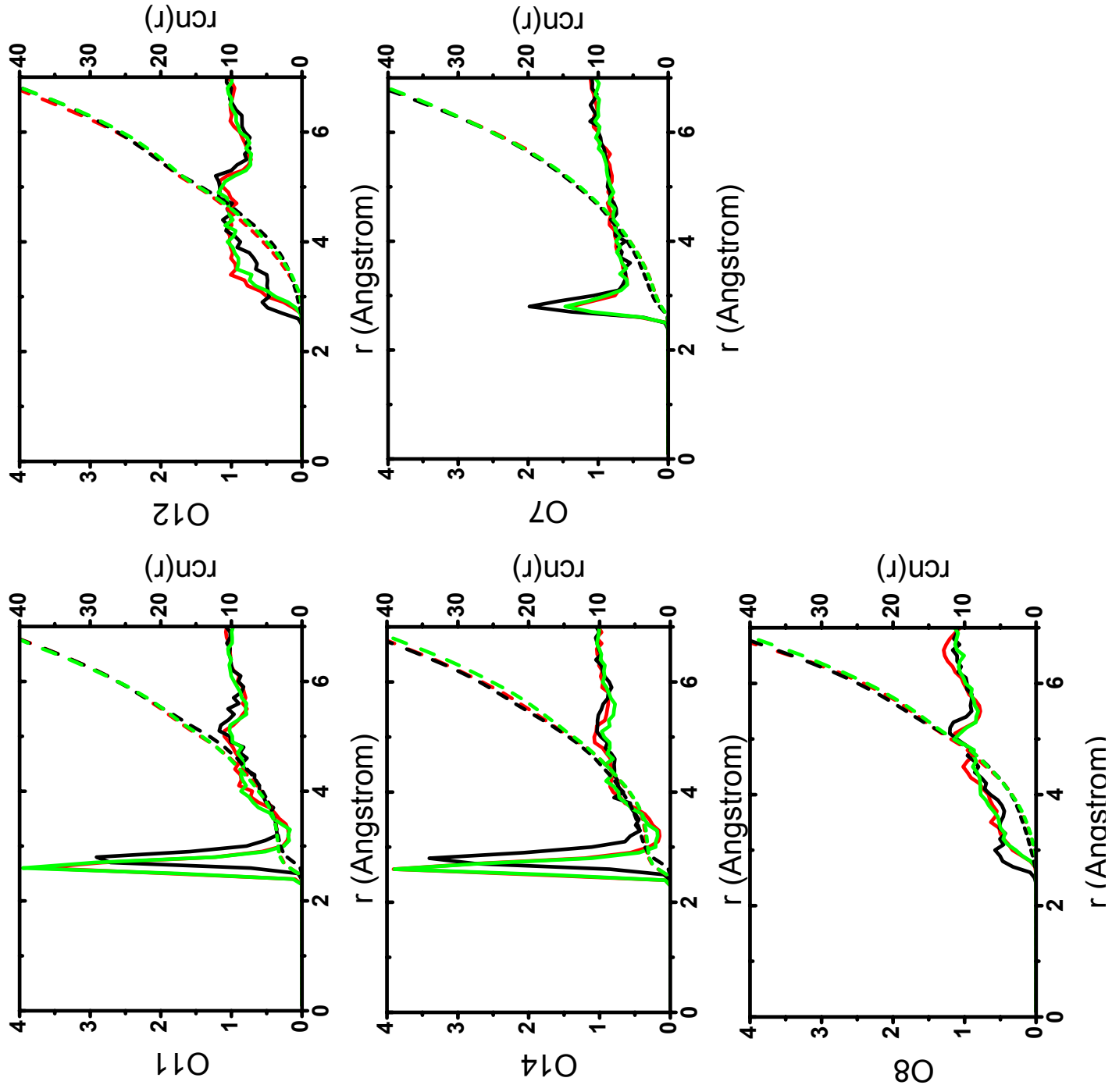


Figure S1

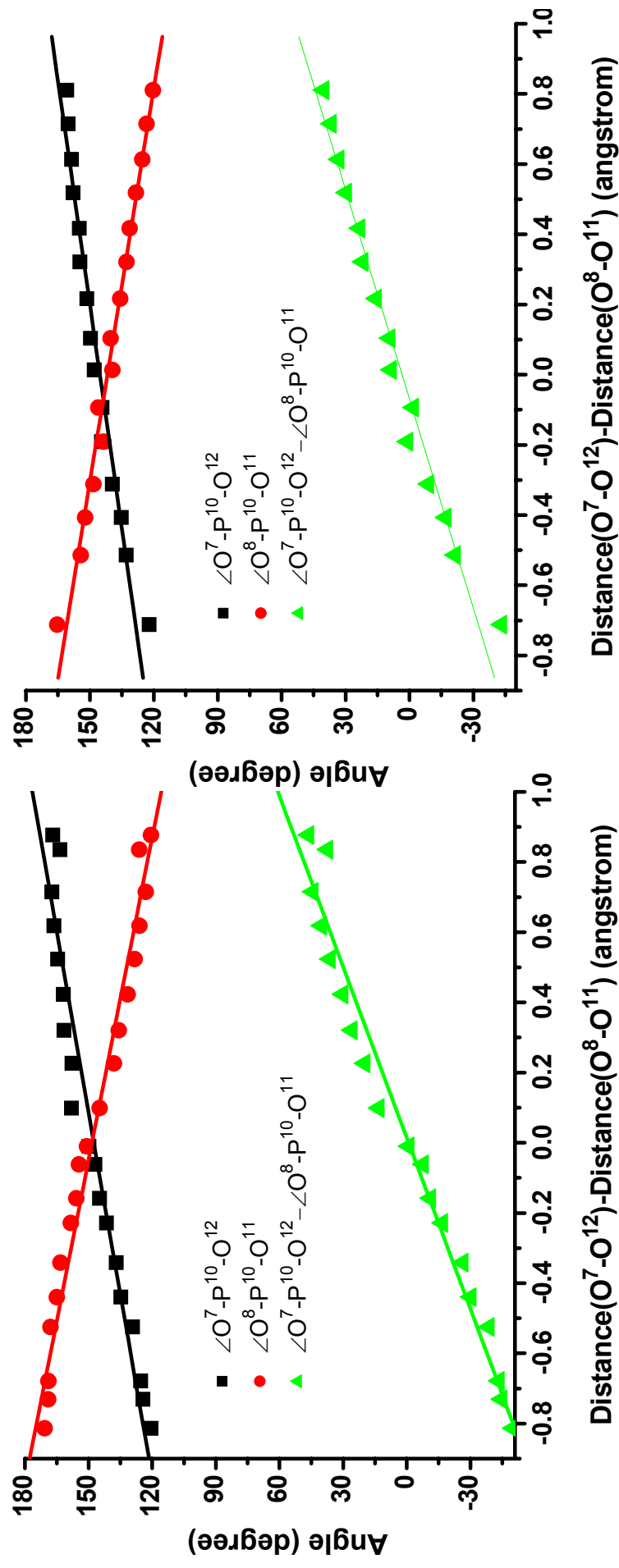


Figure S2