

11 Supplementary Information

11.1 Supplementary figures and tables

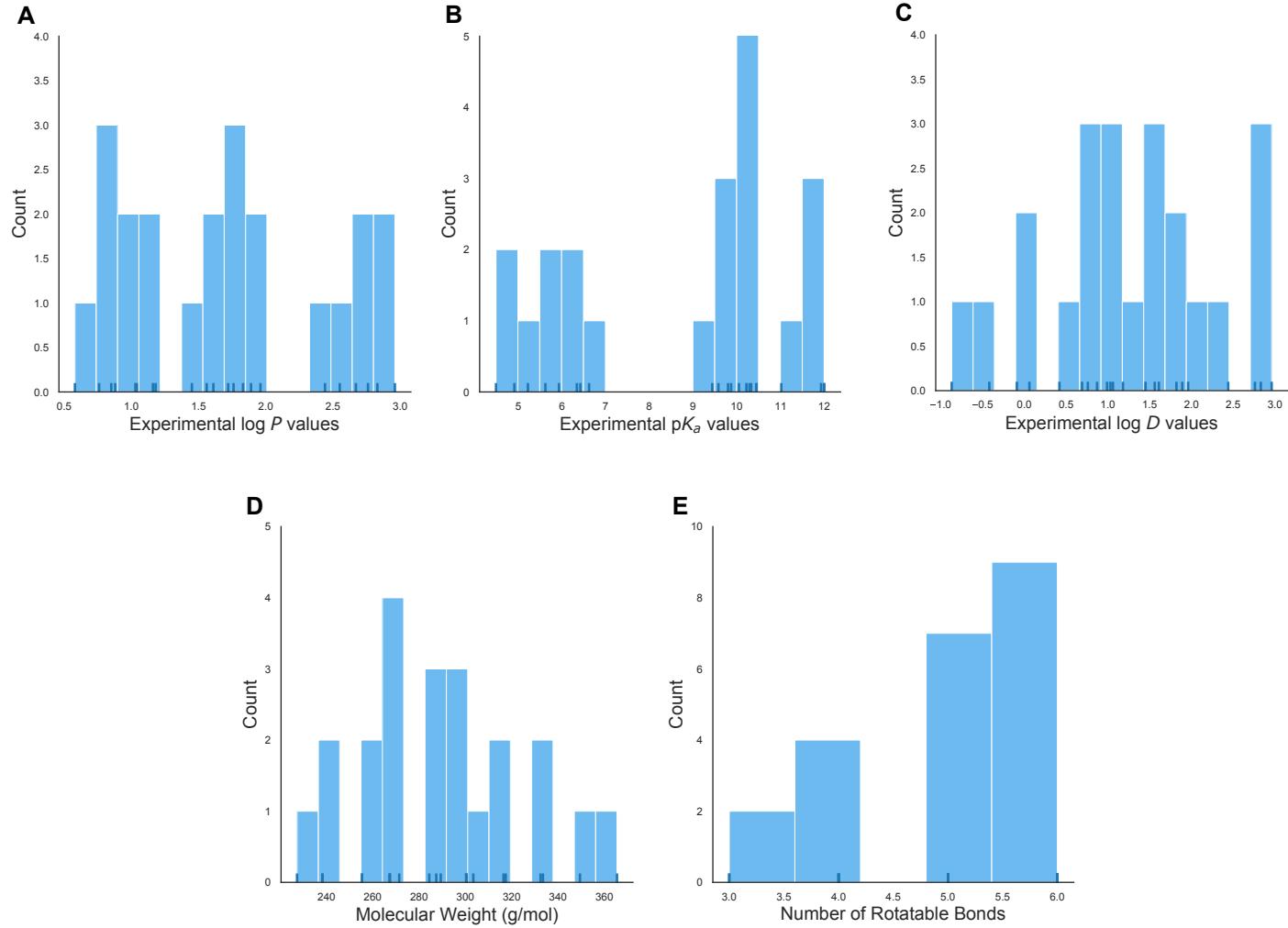


Figure S1. Distribution of molecular properties of the 22 compounds from the SAMPL7 physical property blind challenge. **(A)** Histogram of log P measurements collected with Sirius T3 instrument. The ticks along the x-axis indicate the individual values. Compounds have experimental log P values in the range of 0.58–2.96. **(B)** Histogram of pK_a measurements collected with Sirius T3 instrument. Eight compounds have measured pK_a's in the range of 4.49–6.62 and eleven in the range 9.58–11.93. Two compounds are included here as having pK_a's of 12, but actually had experimental values greater than 12, and were therefore outside of the experimental detection range. **(C)** Histogram of log D measurements between n-octanol and aqueous buffer at pH 7.4 were determined via potentiometric titrations using a Sirius T3 instrument, except for compounds SM27, SM28, SM30–SM34, SM36–SM39 which had log D_{7.4} values determined via shake-flask assay. log D measurements ranged from -0.87–2.96. **(D)** Histogram of molecular weights calculated for the compounds in the SAMPL7 dataset. The molecular weight ranged from 227–365 Da. **(E)** Histogram of the number of rotatable bonds in each molecule. The number of rotatable bonds in challenge molecules ranged from 3–6.

Table S1. Evaluation statistics calculated for all methods in the log P challenge. Submitted predictions are represented by their method name. There are six error metrics reported: the root-mean-squared error (RMSE), mean absolute error (MAE), mean (signed) error (ME), coefficient of determination (R^2), linear regression slope (m), and Kendall's Rank Correlation Coefficient (τ), and error slope (ES). The mean and 95% confidence intervals of each statistic is presented. This table is ranked by increasing RMSE.

Method Name	Category	Submission Type	RMSE	MAE	ME	R^2	m	Kendall's Tau	ES
<i>ClassicalGSG DB2</i>	Empirical	Blind	0.55 [0.38, 0.69]	0.44 [0.31, 0.58]	0.05 [-0.20, 0.26]	0.51 [0.18, 0.82]	0.71 [0.36, 1.06]	0.51 [0.18, 0.78]	0.81 [0.62, 1.03]
<i>TFE MLR</i>	Empirical	Blind	0.58 [0.34, 0.83]	0.41 [0.26, 0.60]	-0.04 [-0.30, 0.19]	0.43 [0.06, 0.80]	0.60 [0.21, 0.95]	0.56 [0.23, 0.82]	1.38 [1.27, 1.45]
<i>ClassicalGSG DB4</i>	Empirical	Blind	0.65 [0.50, 0.78]	0.55 [0.41, 0.69]	0.25 [0.01, 0.50]	0.51 [0.19, 0.76]	0.82 [0.39, 1.22]	0.45 [0.15, 0.71]	0.57 [0.46, 0.85]
<i>Chemprop</i>	Empirical	Blind	0.66 [0.39, 0.88]	0.48 [0.30, 0.68]	-0.17 [-0.44, 0.08]	0.41 [0.11, 0.76]	0.69 [0.31, 1.08]	0.54 [0.25, 0.82]	1.03 [0.79, 1.21]
<i>TFE-SM8-vacuum-opt</i>	Physical (QM)	Blind	0.67 [0.45, 0.86]	0.51 [0.33, 0.69]	0.15 [-0.13, 0.42]	0.45 [0.11, 0.75]	0.80 [0.33, 1.23]	0.50 [0.18, 0.76]	0.99 [0.75, 1.20]
<i>GROVER</i>	Empirical	Blind	0.69 [0.41, 0.96]	0.49 [0.31, 0.71]	-0.21 [-0.50, 0.05]	0.33 [0.04, 0.70]	0.56 [0.18, 0.92]	0.37 [0.05, 0.66]	0.87 [0.62, 1.09]
<i>ClassicalGSG DB1</i>	Empirical	Blind	0.76 [0.56, 0.96]	0.62 [0.45, 0.82]	0.10 [-0.23, 0.40]	0.28 [0.06, 0.60]	0.61 [0.26, 0.99]	0.36 [0.04, 0.63]	0.63 [0.43, 0.85]
<i>ffsampled_deeplearning_cl1</i>	Empirical	Blind	0.77 [0.44, 1.04]	0.51 [0.29, 0.77]	-0.25 [-0.58, 0.04]	0.31 [0.05, 0.70]	0.63 [0.24, 1.05]	0.42 [0.06, 0.74]	0.99 [0.72, 1.19]
<i>ClassicalGSG DB3</i>	Empirical	Blind	0.77 [0.57, 0.96]	0.62 [0.43, 0.82]	-0.15 [-0.46, 0.16]	0.51 [0.18, 0.78]	1.08 [0.55, 1.59]	0.48 [0.15, 0.75]	0.60 [0.42, 0.89]
<i>COSMO-RS</i>	Physical (QM)	Blind	0.78 [0.49, 1.01]	0.57 [0.36, 0.80]	-0.30 [-0.61, -0.01]	0.49 [0.17, 0.79]	0.97 [0.49, 1.45]	0.53 [0.25, 0.78]	0.97 [0.74, 1.18]
<i>TFE_Attentive_FP</i>	Empirical	Blind	0.79 [0.47, 1.07]	0.57 [0.36, 0.82]	-0.18 [-0.53, 0.12]	0.19 [0.00, 0.61]	0.44 [0.04, 0.87]	0.34 [-0.02, 0.69]	0.93 [0.69, 1.13]
<i>ffsampled_deeplearning_cl2</i>	Empirical	Blind	0.82 [0.48, 1.11]	0.56 [0.32, 0.83]	-0.37 [-0.69, -0.08]	0.36 [0.07, 0.72]	0.73 [0.31, 1.16]	0.40 [0.07, 0.69]	0.94 [0.67, 1.15]
<i>TFE-SM12-vacuum-opt</i>	Physical (QM)	Blind	0.82 [0.61, 1.02]	0.66 [0.47, 0.87]	0.28 [-0.06, 0.60]	0.41 [0.08, 0.72]	0.90 [0.36, 1.42]	0.39 [0.05, 0.67]	0.88 [0.65, 1.09]
<i>TFE-SM8-solvent-opt</i>	Physical (QM)	Blind	0.97 [0.71, 1.20]	0.78 [0.55, 1.02]	0.65 [0.35, 0.94]	0.42 [0.10, 0.70]	0.83 [0.35, 1.31]	0.44 [0.13, 0.69]	0.71 [0.47, 0.94]
<i>REF1 ChemAxon</i>	Empirical	Reference	1.00 [0.79, 1.20]	0.85 [0.63, 1.08]	0.46 [0.08, 0.83]	0.39 [0.10, 0.70]	0.98 [0.45, 1.53]	0.40 [0.09, 0.68]	0.13 [-0.00, 0.29]
<i>TFE_IEPPCM MST</i>	Physical (QM)	Blind	1.03 [0.65, 1.41]	0.80 [0.56, 1.10]	-0.07 [-0.53, 0.33]	0.27 [0.01, 0.68]	0.85 [0.12, 1.50]	0.42 [0.10, 0.70]	1.07 [0.88, 1.23]
<i>TFE MD neat oct (GAFF/TIP4P)</i>	Physical (MM)	Blind	1.11 [0.74, 1.43]	0.83 [0.52, 1.15]	-0.74 [-1.10, -0.40]	0.56 [0.24, 0.82]	1.25 [0.64, 1.83]	0.58 [0.27, 0.82]	1.30 [1.19, 1.40]
<i>NULL0 mean clog8 FDA</i>	Empirical	Reference	1.20 [0.94, 1.42]	1.01 [0.73, 1.28]	-0.96 [-1.26, -0.64]	0.00 [0.00, 0.00]	0.00 [-0.00, 0.00]	nan [nan, nan]	0.18 [0.04, 0.32]
<i>NES-1 (GAFF2/OPC3) G</i>	Physical (MM)	Blind	1.21 [0.92, 1.51]	1.03 [0.78, 1.31]	-0.13 [-0.63, 0.37]	0.22 [0.01, 0.59]	0.88 [0.15, 1.59]	0.34 [0.02, 0.63]	1.23 [1.11, 1.33]
<i>NES-1 (GAFF2/OPC3) J</i>	Physical (MM)	Blind	1.28 [0.97, 1.58]	1.08 [0.81, 1.38]	0.01 [-0.54, 0.53]	0.21 [0.01, 0.63]	0.92 [0.09, 1.76]	0.33 [0.00, 0.64]	1.21 [1.08, 1.33]
<i>NES-1 (GAFF2/OPC3) B</i>	Physical (MM)	Blind	1.42 [1.02, 1.81]	1.13 [0.79, 1.51]	-0.51 [-1.08, 0.05]	0.27 [0.02, 0.65]	1.11 [0.30, 1.91]	0.36 [0.05, 0.65]	1.17 [1.01, 1.31]
<i>MD (GAFF/TIP3P)</i>	Physical (MM)	Blind	1.43 [1.15, 1.71]	1.30 [1.06, 1.56]	-1.30 [-1.56, -1.06]	0.48 [0.22, 0.79]	0.77 [0.45, 1.12]	0.55 [0.28, 0.80]	0.94 [0.80, 1.09]
<i>TFE wet oct (GAFF/TIP4P)</i>	Physical (MM)	Blind	1.47 [1.16, 1.77]	1.30 [1.03, 1.60]	-1.30 [-1.60, -1.03]	0.42 [0.10, 0.75]	0.80 [0.30, 1.30]	0.47 [0.14, 0.75]	1.15 [1.03, 1.27]
<i>TFE-NHLBI-TZVP-QM</i>	Physical (QM)	Blind	1.55 [1.19, 1.88]	1.34 [1.02, 1.67]	1.32 [0.00, 1.67]	0.52 [0.19, 0.78]	1.16 [0.59, 1.65]	0.51 [0.19, 0.78]	0.05 [-0.00, 0.17]
<i>0.05 MD (CGenFF/TIP3P)</i>	Physical (MM)	Blind	1.63 [1.25, 1.98]	1.41 [1.08, 1.76]	-1.38 [-1.74, -1.02]	0.54 [0.26, 0.82]	1.26 [0.81, 1.76]	0.52 [0.26, 0.76]	0.90 [0.70, 1.07]
<i>EC_RISM_wet</i>	Physical (QM)	Blind	1.84 [1.31, 2.36]	1.49 [1.07, 1.96]	-1.49 [-1.96, -1.06]	0.29 [0.05, 0.68]	0.96 [0.37, 1.57]	0.38 [0.08, 0.67]	0.67 [0.45, 0.90]
<i>TFE-SMD-vacuum-opt</i>	Physical (QM)	Blind	1.96 [1.60, 2.30]	1.76 [1.42, 2.13]	1.76 [1.42, 2.13]	0.44 [0.12, 0.68]	1.04 [0.46, 1.59]	0.41 [0.03, 0.70]	0.68 [0.50, 0.87]
<i>MD-EE-MCC (GAFF-TIP4P-Ew)</i>	Physical (MM)	Blind	2.06 [1.48, 2.59]	1.61 [1.09, 2.17]	-0.93 [-1.70, -0.17]	0.03 [0.00, 0.28]	0.47 [-0.53, 1.49]	0.11 [-0.16, 0.38]	0.76 [0.51, 1.03]
<i>MD (OPLS-AA/TIP4P)</i>	Physical (MM)	Blind	2.19 [1.69, 2.65]	1.82 [1.31, 2.34]	-1.35 [-2.03, -0.60]	0.28 [0.06, 0.58]	1.47 [0.58, 2.55]	0.36 [0.07, 0.62]	0.73 [0.48, 0.97]
<i>TFE_b3lypd3</i>	Physical (QM)	Blind	2.19 [1.76, 2.57]	1.98 [1.59, 2.37]	1.98 [1.59, 2.37]	0.40 [0.10, 0.67]	1.06 [0.47, 1.64]	0.45 [0.11, 0.72]	0.22 [0.09, 0.41]
<i>MD LigParGen (OPLS-AA/TIP4P)</i>	Physical (MM)	Blind	2.28 [1.80, 2.71]	1.95 [1.46, 2.44]	0.35 [-0.60, 1.29]	0.07 [0.00, 0.37]	0.83 [-0.51, 2.26]	0.19 [-0.14, 0.50]	0.65 [0.42, 0.88]
<i>TFE-SMD-solvent-opt</i>	Physical (QM)	Blind	2.39 [1.97, 2.78]	2.19 [1.79, 2.60]	2.19 [1.79, 2.60]	0.40 [0.09, 0.67]	1.09 [0.45, 1.67]	0.42 [0.09, 0.68]	0.51 [0.34, 0.68]
<i>Ensemble EPI physprop</i>	Empirical	Blind	2.73 [2.27, 3.16]	2.54 [2.13, 2.98]	2.54 [2.13, 2.98]	0.33 [0.04, 0.64]	-0.30 [-0.49, -0.10]	-0.35 [-0.60, -0.03]	-0.00 [-0.00, -0.00]
<i>Ensemble Martel</i>	Empirical	Blind	3.29 [2.89, 3.68]	3.16 [2.78, 3.55]	3.16 [2.78, 3.55]	0.39 [0.05, 0.73]	-0.25 [-0.40, -0.09]	-0.46 [-0.72, -0.14]	-0.00 [-0.00, -0.00]
<i>QSPR_Mordred2D_TPOT_AutoML</i>	Empirical	Blind	3.64 [3.01, 4.24]	3.36 [2.80, 3.96]	3.36 [2.80, 3.96]	0.39 [0.10, 0.71]	-0.72 [-1.12, -0.33]	-0.37 [-0.65, -0.04]	-0.00 [-0.00, -0.00]
<i>TFE-NHLBI-NN-IN</i>	Empirical	Blind	3.97 [3.57, 4.34]	3.85 [3.45, 4.25]	3.85 [3.45, 4.25]	0.00 [0.00, 0.15]	0.02 [-0.30, 0.34]	0.02 [-0.23, 0.27]	0.01 [-0.00, 0.02]

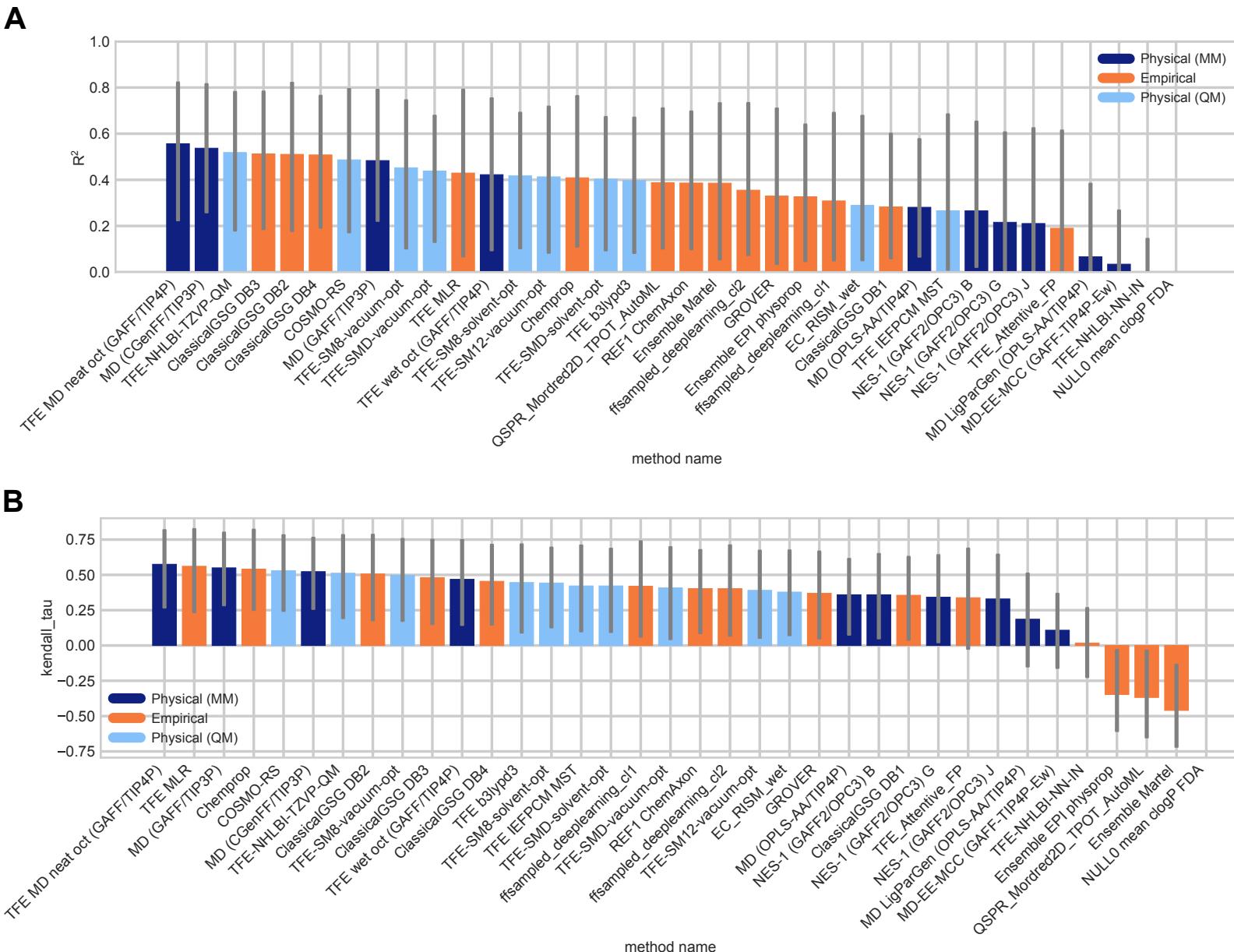


Figure S2. Overall correlation assessment for all methods participating in the SAMPL7 log P challenge show that the uncertainty of each correlation statistic is quite high, not allowing a true ranking based on correlation. Pearson's R^2 and Kendall's Rank Correlation Coefficient Tau (τ) are shown, with error bars denoting 95% confidence intervals obtained by bootstrapping over challenge molecules. Submitted methods are listed in Table 1. The submission REF1 ChemAxon was a reference method included after the blind challenge submission deadline, and NULL0 mean cLogP FDA is the null prediction method; all others refer to blind predictions. Most methods have a statistically indistinguishable performance on ranking because of the small dynamic range of the dataset. Evaluation statistics calculated for all methods are available in Table S1 of the Supplementary Information.

Table S2. Details of MM-based physical methods in the log P prediction challenge. Force fields, water models, and octanol phase choice are reported. A dry octanol phase indicates the octanol phase was composed of only octanol. A wet octanol phase indicates the octanol phase was treated as a mixture of octanol and water. RMSE, MAE, R², and Kendall's Tau values are reported as mean and 95% confidence intervals.

Method Name	Force Field	Water Model	Octanol Phase	RMSE	MAE	R ²	Kendall's Tau
<i>TFE MD neat oct (GAFF/TIP4P)</i>	GAFF	TIP4P	dry	1.11 [0.74, 1.43]	0.83 [0.52, 1.15]	0.56 [0.24, 0.82]	0.58 [0.27, 0.82]
<i>NES-1 (GAFF2/OPC3) G</i>	GAFF2	OPC3	dry	1.21 [0.92, 1.51]	1.03 [0.78, 1.31]	0.22 [0.01, 0.59]	0.34 [0.02, 0.63]
<i>NES-1 (GAFF2/OPC3) J</i>	GAFF2	OPC3	dry	1.28 [0.97, 1.58]	1.08 [0.81, 1.38]	0.21 [0.01, 0.63]	0.33 [0.00, 0.64]
<i>NES-1 (GAFF2/OPC3) B</i>	GAFF2	OPC3	dry	1.42 [1.02, 1.81]	1.13 [0.79, 1.51]	0.27 [0.02, 0.65]	0.36 [0.05, 0.65]
<i>MD (GAFF/TIP3P)</i>	GAFF	TIP3P	dry	1.43 [1.15, 1.71]	1.30 [1.06, 1.56]	0.48 [0.22, 0.79]	0.55 [0.28, 0.80]
<i>TFE wet oct (GAFF/TIP4P)</i>	GAFF	TIP4P	wet	1.47 [1.16, 1.77]	1.30 [1.03, 1.60]	0.42 [0.10, 0.75]	0.47 [0.14, 0.75]
<i>MD (CGenFF/TIP3P)</i>	CGenFF	TIP3P	dry	1.63 [1.25, 1.98]	1.41 [1.08, 1.76]	0.54 [0.26, 0.82]	0.52 [0.26, 0.76]
<i>MD-EE-MCC (GAFF-TIP4P-Ew)</i>	GAFF	TIP4P-eW	dry	2.06 [1.48, 2.59]	1.61 [1.09, 2.17]	0.03 [0.00, 0.28]	0.11 [-0.16, 0.38]
<i>MD (OPLS-AA/TIP4P)</i>	OPLS-AA	TIP4P	dry	2.19 [1.69, 2.65]	1.82 [1.31, 2.34]	0.28 [0.06, 0.58]	0.36 [0.07, 0.62]
<i>MD LigParGen (OPLS-AA/TIP4P)</i>	OPLS-AA	TIP4P	dry	2.28 [1.80, 2.71]	1.95 [1.46, 2.44]	0.07 [0.00, 0.37]	0.19 [-0.14, 0.50]

Table S3. Evaluation statistics calculated for all methods in the pK_a challenge. Submitted predictions are represented by their method name. There are six error metrics reported: the root-mean-squared error (RMSE), mean absolute error (MAE), mean (signed) error (ME), coefficient of determination (R^2), linear regression slope (m), Kendall's Rank Correlation Coefficient (τ), and error slope (ES). The mean and 95% confidence intervals of each statistic is presented. This table is ranked by increasing RMSE.

Method Name	Category	Submission Type	RMSE	MAE	ME	R^2	m	Kendall's Tau	ES
<i>REF00_Chemaxon_Chemicalize</i>	QSPR/ML	Reference	0.71 [0.50, 0.90]	0.56 [0.38, 0.76]	0.09 [-0.23, 0.38]	0.91 [0.86, 0.96]	0.88 [0.72, 1.02]	0.73 [0.51, 0.90]	0.83 [0.58, 1.04]
<i>EC_RISM</i>	QM	Blind	0.72 [0.45, 0.95]	0.53 [0.33, 0.75]	0.20 [-0.10, 0.50]	0.93 [0.87, 0.98]	0.80 [0.72, 0.91]	0.81 [0.63, 0.96]	1.32 [1.19, 1.42]
<i>IEPCM/MST</i>	QM	Blind	1.82 [1.00, 2.69]	1.30 [0.84, 1.92]	0.25 [-0.46, 1.09]	0.56 [0.22, 0.87]	0.86 [0.53, 1.18]	0.52 [0.22, 0.76]	1.00 [0.80, 1.17]
<i>DFT_M05-2X_SMD</i>	QM	Blind	2.90 [2.04, 3.69]	2.28 [1.53, 3.10]	-0.78 [-2.02, 0.41]	0.03 [0.00, 0.37]	0.15 [-0.32, 0.53]	0.17 [-0.22, 0.54]	0.55 [0.31, 0.81]
<i>TZVP-QM</i>	QM	Blind	2.90 [2.52, 3.25]	2.75 [2.34, 3.14]	1.20 [0.02, 2.33]	0.23 [0.03, 0.60]	-0.11 [-0.20, -0.04]	-0.14 [-0.49, 0.23]	-0.00 [-0.00, -0.00]
<i>Standard Gaussian Process</i>	QSPR/ML	Blind	3.49 [2.76, 4.12]	2.91 [2.06, 3.75]	2.47 [1.38, 3.55]	0.30 [0.10, 0.69]	-0.05 [-0.09, -0.02]	-0.42 [-0.70, -0.08]	1.11 [0.96, 1.24]
<i>DFT_M06-2X_SMD_implicit</i>	QM	Blind	4.16 [2.00, 6.38]	2.80 [1.76, 4.33]	-0.07 [-1.61, 1.95]	0.52 [0.39, 0.78]	1.70 [0.80, 2.77]	0.70 [0.48, 0.88]	0.50 [0.30, 0.70]
<i>DFT_M06-2X_SMD_implicit_SAS</i>	QM	Blind	4.16 [2.03, 6.44]	2.81 [1.80, 4.36]	-0.20 [-1.71, 1.85]	0.50 [0.36, 0.77]	1.64 [0.72, 2.72]	0.56 [0.28, 0.81]	0.14 [0.02, 0.31]
<i>DFT_M06-2X_SMD_explicit_water</i>	QM	Blind	5.12 [1.19, 7.92]	2.56 [0.96, 4.76]	-0.35 [-2.62, 1.93]	0.20 [0.00, 0.81]	1.10 [-0.39, 2.50]	0.46 [0.06, 0.78]	0.52 [0.29, 0.77]
<i>Gaussian_corrected</i>	QM+LEC	Blind	5.36 [4.70, 5.95]	5.12 [4.42, 5.79]	5.12 [4.42, 5.79]	0.76 [0.63, 0.88]	0.35 [0.27, 0.45]	0.60 [0.42, 0.76]	0.00 [-0.00, 0.00]

Table S4. Evaluation statistics calculated for all log D estimates. Predictions are represented a name based on method names participants submitted to the and log P challenges. There are six error metrics reported: the root-mean-squared error (RMSE), mean absolute error (MAE), mean (signed) error (ME), coefficient of determination (R^2), linear regression slope (m), Kendall's Rank Correlation Coefficient (τ), and error slope (ES). The mean and 95% confidence intervals of each statistic is presented. This table is ranked by increasing RMSE.

Method Name	Category	Submission Type	RMSE	MAE	ME	R^2	m	Kendall's Tau	ES
<i>REF0 ChemAxon</i>	Empirical	Reference	1.06 [0.82, 1.27]	0.91 [0.68, 1.14]	0.28 [-0.14, 0.70]	0.27 [0.01, 0.58]	0.54 [0.10, 0.90]	0.31 [-0.02, 0.61]	0.12 [-0.00, 0.28]
<i>TFE IEFPCM MST + IEFPCM/MST</i>	Physical (QM)	Standard	1.27 [0.85, 1.64]	0.98 [0.67, 1.33]	0.24 [-0.28, 0.75]	0.55 [0.17, 0.87]	1.31 [0.71, 1.70]	0.57 [0.27, 0.82]	1.16 [0.89, 1.25]
<i>NULL0</i>	Empirical	Reference	1.59 [1.22, 1.93]	1.35 [1.00, 1.71]	1.23 [0.81, 1.65]	0.00 [0.00, 0.00]	0.00 [0.00, 0.00]	nan [nan, nan]	0.65 [0.44, 0.87]
<i>EC_RISM</i>	Physical (QM)	Standard	1.69 [1.30, 2.05]	1.43 [1.07, 1.82]	-1.43 [-1.81, -1.07]	0.53 [0.20, 0.77]	0.95 [0.54, 1.29]	0.51 [0.21, 0.74]	0.84 [0.64, 1.02]
<i>TFE-NHLBI-TZVP-QM + TZVP-QM</i>	Physical (QM)	Standard	1.72 [1.30, 2.12]	1.47 [1.12, 1.86]	1.26 [0.78, 1.75]	0.25 [0.01, 0.64]	0.64 [0.08, 1.25]	0.38 [0.02, 0.70]	0.05 [-0.00, 0.18]
<i>TFE b3lypd3 + DFT_M05-2X_SMD</i>	Physical (QM)	Standard	2.15 [1.56, 2.71]	1.78 [1.31, 2.31]	1.78 [1.31, 2.31]	0.32 [0.04, 0.66]	0.80 [0.27, 1.30]	0.41 [0.05, 0.72]	0.42 [0.27, 0.70]
<i>MD (CGenFF/TIP3P) + Gaussian_corrected</i>	Physical (MM) + QM+LEC	Standard	2.27 [1.97, 2.55]	2.13 [1.80, 2.45]	1.84 [1.21, 2.35]	0.62 [0.35, 0.84]	1.53 [0.93, 2.18]	0.62 [0.36, 0.82]	0.88 [0.75, 1.00]
<i>TFE-SMD-solvent-opt + DFT_M06-2X_SMD_explicit_water</i>	Physical (QM)	Standard	4.54 [2.09, 7.15]	2.92 [1.88, 4.57]	2.88 [1.80, 4.55]	0.25 [0.11, 0.76]	1.92 [0.53, 4.45]	0.55 [0.22, 0.80]	0.55 [0.38, 0.73]

Table S5. Additional info for microscopic pK_a predictions.

Microstate	Total number of relative free energy predictions	Average relative free energy prediction	Average relative free energy prediction STD	Minimum relative free energy prediction	Maximum relative free energy prediction	Number of (+) sign predictions	Number of (-) sign predictions	Number of neutral (0) sign predictions	Shannon entropy (H)
<i>SM25_micro001</i>	9	-0.6	13.2	-15.6	16.3	4	5	0	0.7
<i>SM25_micro002</i>	8	8.8	10.6	-7.5	20.4	6	2	0	0.6
<i>SM25_micro003</i>	8	9.6	2.7	4.5	12.6	8	0	0	0.0
<i>SM25_micro004</i>	2	-8.9	4.5	-12.1	-5.8	0	2	0	0.0
<i>SM25_micro005</i>	2	-0.8	2.1	-2.3	0.7	1	1	0	0.7
<i>SM26_micro001</i>	9	7.3	2.4	3.0	10.7	9	0	0	0.0
<i>SM26_micro002</i>	8	-6.7	20.5	-31.7	22.1	3	5	0	0.7
<i>SM26_micro003</i>	8	20.9	12.0	0.9	32.4	8	0	0	0.0
<i>SM26_micro004</i>	2	4.3	0.7	3.8	4.8	2	0	0	0.0
<i>SM26_micro005</i>	2	8.1	2.6	6.3	10.0	2	0	0	0.0
<i>SM27_micro001</i>	9	13.4	4.9	6.1	19.0	9	0	0	0.0
<i>SM28_micro001</i>	9	-5.7	25.0	-39.0	23.5	4	5	0	0.7
<i>SM28_micro002</i>	8	17.1	8.0	8.2	26.5	8	0	0	0.0
<i>SM28_micro003</i>	8	0.9	8.3	-10.0	12.6	4	4	0	0.7
<i>SM28_micro004</i>	2	25.1	9.1	18.7	31.5	2	0	0	0.0
<i>SM29_micro001</i>	9	12.6	4.3	6.3	18.7	9	0	0	0.0
<i>SM30_micro001</i>	9	12.3	4.2	5.9	17.7	9	0	0	0.0
<i>SM31_micro001</i>	9	13.2	4.4	6.0	18.1	9	0	0	0.0
<i>SM31_micro002</i>	3	-0.6	6.6	-8.1	4.5	2	1	0	0.6
<i>SM32_micro001</i>	9	12.8	4.6	5.9	18.9	9	0	0	0.0
<i>SM33_micro001</i>	9	11.9	3.9	5.2	17.1	9	0	0	0.0
<i>SM34_micro001</i>	9	13.0	4.6	5.7	19.7	9	0	0	0.0
<i>SM34_micro002</i>	3	-0.9	6.4	-8.1	4.4	2	1	0	0.6
<i>SM35_micro001</i>	9	11.7	4.5	3.2	16.2	9	0	0	0.0
<i>SM35_micro002</i>	8	0.2	1.4	-1.9	2.5	5	2	1	0.9
<i>SM35_micro003</i>	8	12.2	5.6	3.2	18.1	8	0	0	0.0
<i>SM36_micro001</i>	9	10.8	3.1	5.2	14.9	9	0	0	0.0
<i>SM36_micro002</i>	8	1.2	1.8	0.0	4.4	4	1	3	1.0
<i>SM36_micro003</i>	8	10.7	3.3	5.2	14.7	8	0	0	0.0
<i>SM37_micro001</i>	9	0.1	9.4	-11.7	13.7	5	4	0	0.7
<i>SM37_micro002</i>	8	9.8	2.9	3.7	12.7	8	0	0	0.0
<i>SM37_micro003</i>	8	0.7	1.8	-1.5	4.2	4	3	1	1.0
<i>SM37_micro004</i>	8	8.9	3.0	3.8	12.4	8	0	0	0.0
<i>SM37_micro005</i>	7	-2.7	7.6	-10.6	11.0	3	4	0	0.7
<i>SM38_micro001</i>	9	11.6	4.6	5.2	17.5	9	0	0	0.0
<i>SM39_micro001</i>	9	10.1	3.1	5.1	14.6	9	0	0	0.0
<i>SM40_micro001</i>	9	10.8	3.3	5.0	15.7	9	0	0	0.0
<i>SM40_micro002</i>	8	-1.8	10.3	-15.5	11.8	4	4	0	0.7
<i>SM41_micro001</i>	9	8.4	3.5	2.2	14.8	9	0	0	0.0
<i>SM41_micro002</i>	8	-0.5	9.9	-12.9	13.9	4	4	0	0.7
<i>SM42_micro001</i>	9	5.5	4.6	0.2	12.3	9	0	0	0.0
<i>SM42_micro002</i>	8	-0.2	8.6	-10.8	14.3	4	4	0	0.7
<i>SM42_micro003</i>	3	-2.0	3.0	-5.1	1.0	1	2	0	0.6
<i>SM43_micro001</i>	9	5.9	4.4	0.5	13.4	9	0	0	0.0
<i>SM43_micro002</i>	8	0.1	9.4	-11.0	11.0	4	4	0	0.7
<i>SM43_micro003</i>	8	-11.6	38.1	-60.9	38.2	4	4	0	0.7
<i>SM43_micro004</i>	2	-3.6	2.2	-5.2	-2.1	0	2	0	0.0
<i>SM43_micro005</i>	2	0.1	0.4	-0.2	0.4	1	1	0	0.7
<i>SM44_micro001</i>	9	9.5	2.9	4.3	12.9	9	0	0	0.0
<i>SM44_micro002</i>	8	-1.1	7.4	-10.3	9.9	4	4	0	0.7
<i>SM45_micro001</i>	9	9.6	3.1	4.4	14.7	9	0	0	0.0
<i>SM45_micro002</i>	8	-1.0	7.8	-11.0	9.6	4	4	0	0.7
<i>SM46_micro001</i>	9	9.9	4.1	4.0	18.4	9	0	0	0.0
<i>SM46_micro002</i>	8	-0.7	7.5	-9.6	10.5	4	4	0	0.7
<i>SM46_micro003</i>	8	-12.2	37.1	-63.5	39.0	4	4	0	0.7
<i>SM46_micro004</i>	3	6.3	4.5	2.4	11.3	3	0	0	0.0

Table S6. SMILES and compound class of SAMPL7 physical property challenge molecules. A view of the compounds and their classes can be found in Figure S3.

SAMPL7 Molecule ID	Compound Class	Isomeric SMILES
SM25	acylsulfonamide	O=C(NS(C1=CC=CC=C1)(=O)=O)CCC2=CC=CC=C2
SM26	acylsulfonamide	O=S(CCC1=CC=CC=C1)(NC(C)=O)=O
SM27	oxetane	O=S(CCC1=CC=CC=C1)(NC2(C)COC2)=O
SM28	thietane-1,1-dioxide	O=S(CC1(NC(C)=O)CCC2=CC=CC=C2)(C1)=O
SM29	oxetane	CS(NC1(COC1)CCC2=CC=CC=C2)(=O)=O
SM30	oxetane	O=S(NC1(COC1)CCC2=CC=CC=C2)(C3=CC=CC=C3)=O
SM31	oxetane	O=S(NC1(COC1)CCC2=CC=CC=C2)(N(C)C)=O
SM32	thietane	CS(NC1(CSC1)CCC2=CC=CC=C2)(=O)=O
SM33	thietane	O=S(NC1(CSC1)CCC2=CC=CC=C2)(C3=CC=CC=C3)=O
SM34	thietane	O=S(NC1(CSC1)CCC2=CC=CC=C2)(N(C)C)=O
SM35	thietane-1-oxide	CS(N[C@@]1(C[S+]([O-])C1)CCC2=CC=CC=C2)(=O)=O
SM36	thietane-1-oxide	O=S(N[C@@]1(C[S+]([O-])C1)CCC2=CC=CC=C2)(C3=CC=CC=C3)=O
SM37	thietane-1-oxide	O=S(N[C@@]1(C[S+]([O-])C1)CCC2=CC=CC=C2)(N(C)C)=O
SM38	thietane-1,1-dioxide	CS(NC1(CS(C1)(=O)=O)CCC2=CC=CC=C2)(=O)=O
SM39	thietane-1,1-dioxide	O=S(NC1(CS(C1)(=O)=O)CCC2=CC=CC=C2)(C3=CC=CC=C3)=O
SM40	thietane-1,1-dioxide	O=S(NC1(CS(C1)(=O)=O)CCC2=CC=CC=C2)(N(C)C)=O
SM41	isoxazole	O=S(NC1=NOC(C2=CC=CC=C2)=C1)(C)=O
SM42	isoxazole	O=S(NC1=NOC(C2=CC=CC=C2)=C1)(C3=CC=CC=C3)=O
SM43	isoxazole	O=S(NC1=NOC(C2=CC=CC=C2)=C1)(N(C)C)=O
SM44	1,2,3-triazole	O=S(NC(N=N1)=CN1C2=CC=CC=C2)(C)=O
SM45	1,2,3-triazole	O=S(NC(N=N1)=CN1C2=CC=CC=C2)(C3=CC=CC=C3)=O
SM46	1,2,3-triazole	O=S(NC(N=N1)=CN1C2=CC=CC=C2)(N(C)C)=O

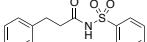
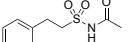
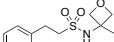
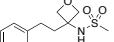
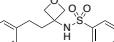
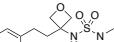
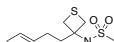
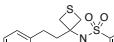
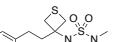
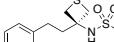
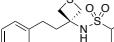
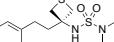
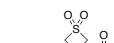
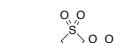
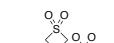
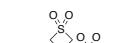
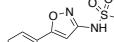
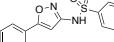
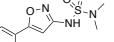
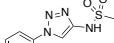
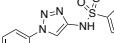
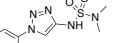
Compound Classes	Structures		
acylsulfonamide	  SM25 SM26		
oxetane	    SM27 SM29 SM30 SM31		
thietane	   SM32 SM33 SM34		
thietane-1-oxide	   SM35 SM36 SM37		
thietane-1,1-dioxide	    SM28 SM38 SM39 SM40		
isoxazole	   SM41 SM42 SM43		
1,2,3-triazole	   SM44 SM45 SM46		

Figure S3. Compound classes and structures of the molecules in the SAMPL7 physical property challenge. SMILES of the compounds are in Table S3.

Table S7. Number of states per charge state for the microstates used in the SAMPL7 pK_a challenge. The total number of microstates (protomers and tautomers) is listed. Some of the molecules have up to 6 microstates, while others have only 2.

	Charge State				
	+2	+1	0	-1	Total #
SM25	0	1	3	2	6
SM26	0	1	3	2	6
SM27	0	0	1	1	2
SM28	0	1	2	2	5
SM29	0	0	1	1	2
SM30	0	0	1	1	2
SM31	0	1	1	1	3
SM32	0	0	1	1	2
SM33	0	0	1	1	2
SM34	0	1	1	1	3
SM35	0	0	2	3	5
SM36	0	0	2	3	5
SM37	0	2	2	2	6
SM38	0	0	1	1	2
SM39	0	0	1	1	2
SM40	0	1	1	1	3
SM41	0	1	1	1	3
SM42	0	1	2	1	4
SM43	1	2	2	1	6
SM44	0	1	1	1	3
SM45	0	1	1	1	3
SM46	1	2	1	1	5