

Supplementary Materials

Exploring the solubility limits of Edaravone in neat solvents and binary mixtures: experimental and machine learning study

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S1. New edaravone solubility data

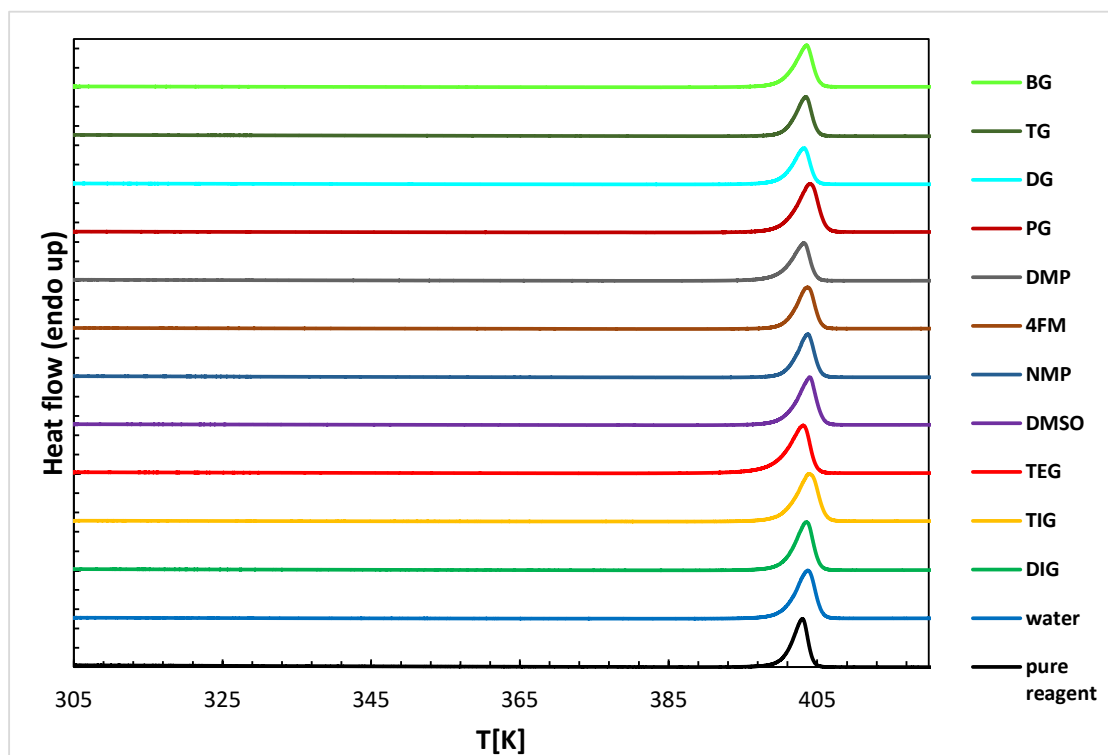
Table S1.1. The experimentally determined mole fraction solubility of Edaravone (x_{EDA} , $\cdot 10^2$) in diglyme (DIG), triglyme (TIG), tetraglyme (TEG), dimethyl sulfoxide (DMSO), 1-methyl-2-pyrrolidone, (NMP), 4-formylmorpholine (4FM), and 2,4-dimethylphenol (DMP) at various temperatures. Standard deviation values are given in parentheses.

| solvent | Edaravone solubility (x_{EDA} , $\cdot 10^2$) | | | |
|---------|--|----------------------|----------------------|----------------------|
| | 25 °C | 30 °C | 35 °C | 40 °C |
| DMSO | 7.58 (± 0.32) | 12.25 (± 0.16) | 20.1 (± 0.15) | 29.81 (± 1.14) |
| TEG | 4.63 (± 0.05) | 8.63 (± 0.22) | 13.05 (± 0.19) | 21.21 (± 0.64) |
| NMP | 3.83 (± 0.01) | 4.96 (± 0.49) | 6.94 (± 0.19) | 8.61 (± 0.21) |
| DIG | 2.05 (± 0.02) | 2.69 (± 0.04) | 3.33 (± 0.11) | 4.16 (± 0.06) |
| DMP | 1.88 (± 0.05) | 2.82 (± 0.58) | 4.08 (± 0.21) | 5.65 (± 0.25) |
| TIG | 1.53 (± 0.07) | 2.20 (± 0.04) | 3.11 (± 0.17) | 4.57 (± 0.39) |
| 4FM | 1.11 (± 0.01) | 1.46 (± 0.08) | 1.76 (± 0.19) | 2.17 (± 0.01) |

Table S1.2. The experimentally determined mole fraction solubility of Edaravone (x_{EDA} , $\cdot 10^2$) in 1,2-propanediol (PG), diethylene glycol (DG), triethylene glycol (TG), and 1,3-butanediol (BG), as well as in their aqueous binary mixtures at 25 °C. In the first column, x_2^* denotes the mole fraction of the organic solvent in solute-free solutions. Standard deviation values are given in parentheses.

| x_2^* | Edaravone solubility (x_{EDA} , $\cdot 10^2$) | | | |
|---------|--|-----------------------|-----------------------|-----------------------|
| | PG | DG | TG | BG |
| 0.0 | 0.02 ($< \pm 0.01$) | 0.02 ($< \pm 0.01$) | 0.02 ($< \pm 0.01$) | 0.02 ($< \pm 0.01$) |
| 0.1 | 0.06 (± 0.01) | 0.09 (± 0.01) | 0.31 (± 0.04) | 0.12 (± 0.02) |
| 0.2 | 0.15 (± 0.01) | 0.25 (± 0.01) | 0.78 (± 0.03) | 0.29 (± 0.04) |
| 0.3 | 0.26 (± 0.02) | 0.48 (± 0.04) | 1.28 (± 0.07) | 0.53 (± 0.03) |
| 0.4 | 0.40 (± 0.02) | 0.74 (± 0.03) | 1.72 (± 0.06) | 0.83 (± 0.04) |
| 0.5 | 0.54 (± 0.02) | 1.09 (± 0.01) | 2.11 (± 0.08) | 1.07 (± 0.05) |
| 0.6 | 0.71 (± 0.02) | 1.39 (± 0.02) | 2.54 (± 0.08) | 1.27 (± 0.03) |
| 0.7 | 0.86 (± 0.02) | 1.65 (± 0.05) | 3.03 (± 0.09) | 1.42 (± 0.05) |
| 0.8 | 0.99 (± 0.02) | 1.87 (± 0.04) | 3.51 (± 0.09) | 1.50 (± 0.03) |
| 0.9 | 1.00 (± 0.02) | 1.98 (± 0.02) | 3.58 (± 0.06) | 1.51 (± 0.04) |
| 1.0 | 0.85 (± 0.01) | 2.04 (± 0.06) | 2.75 (± 0.08) | 1.43 (± 0.04) |

A



B

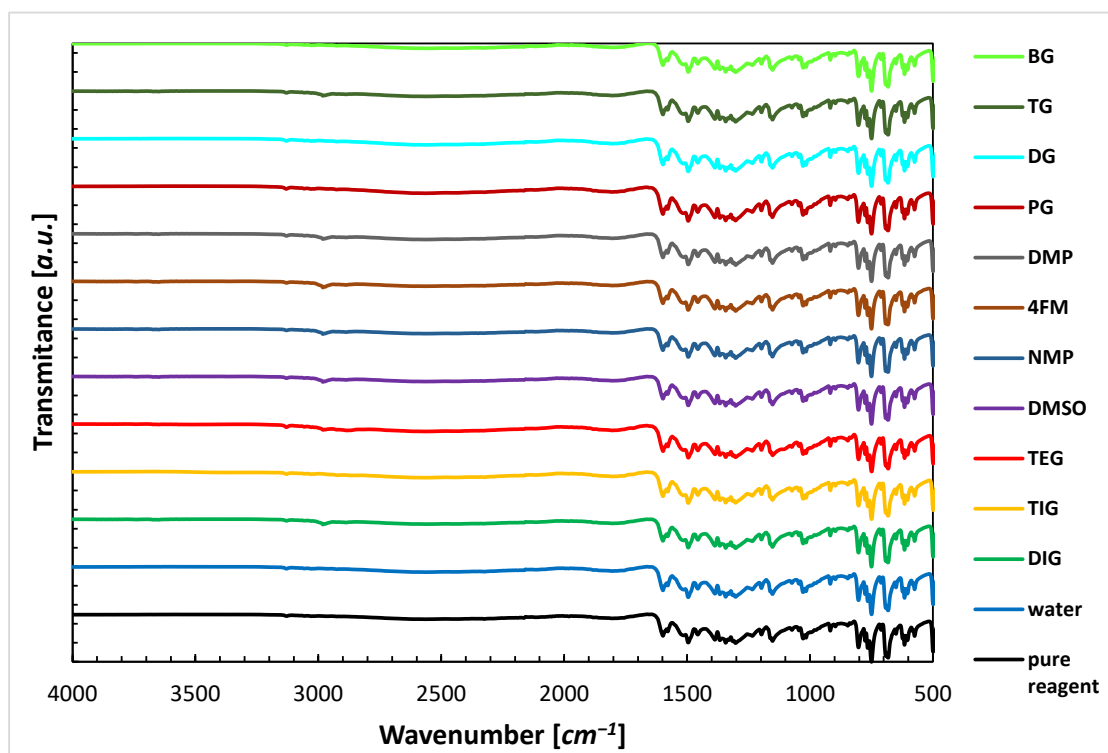


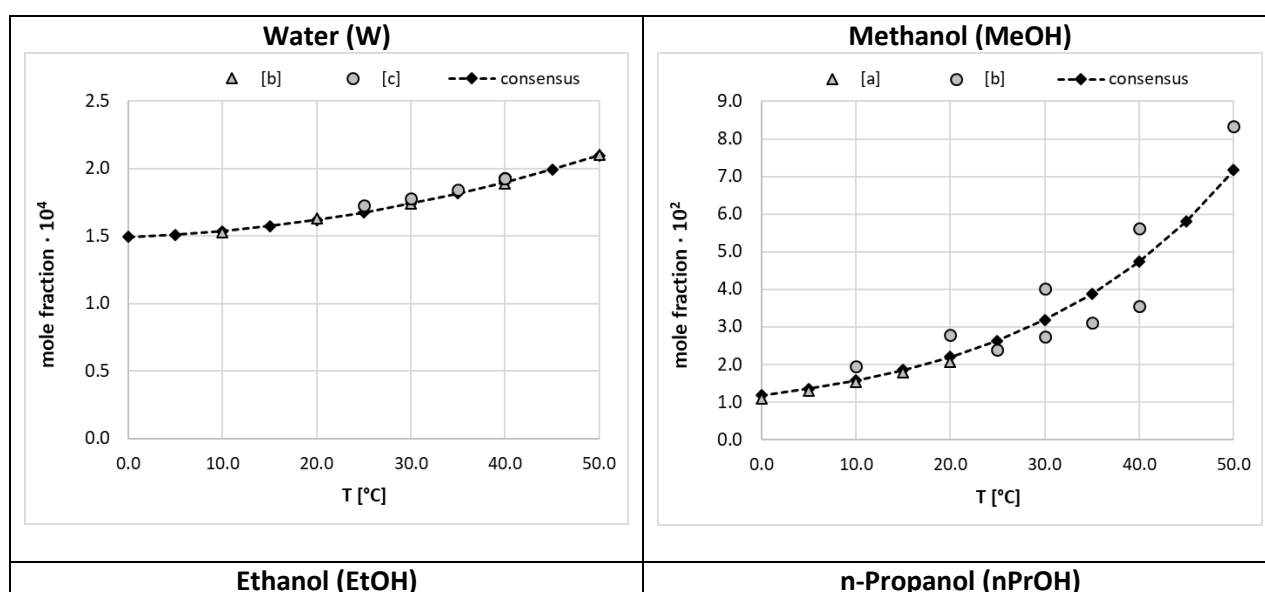
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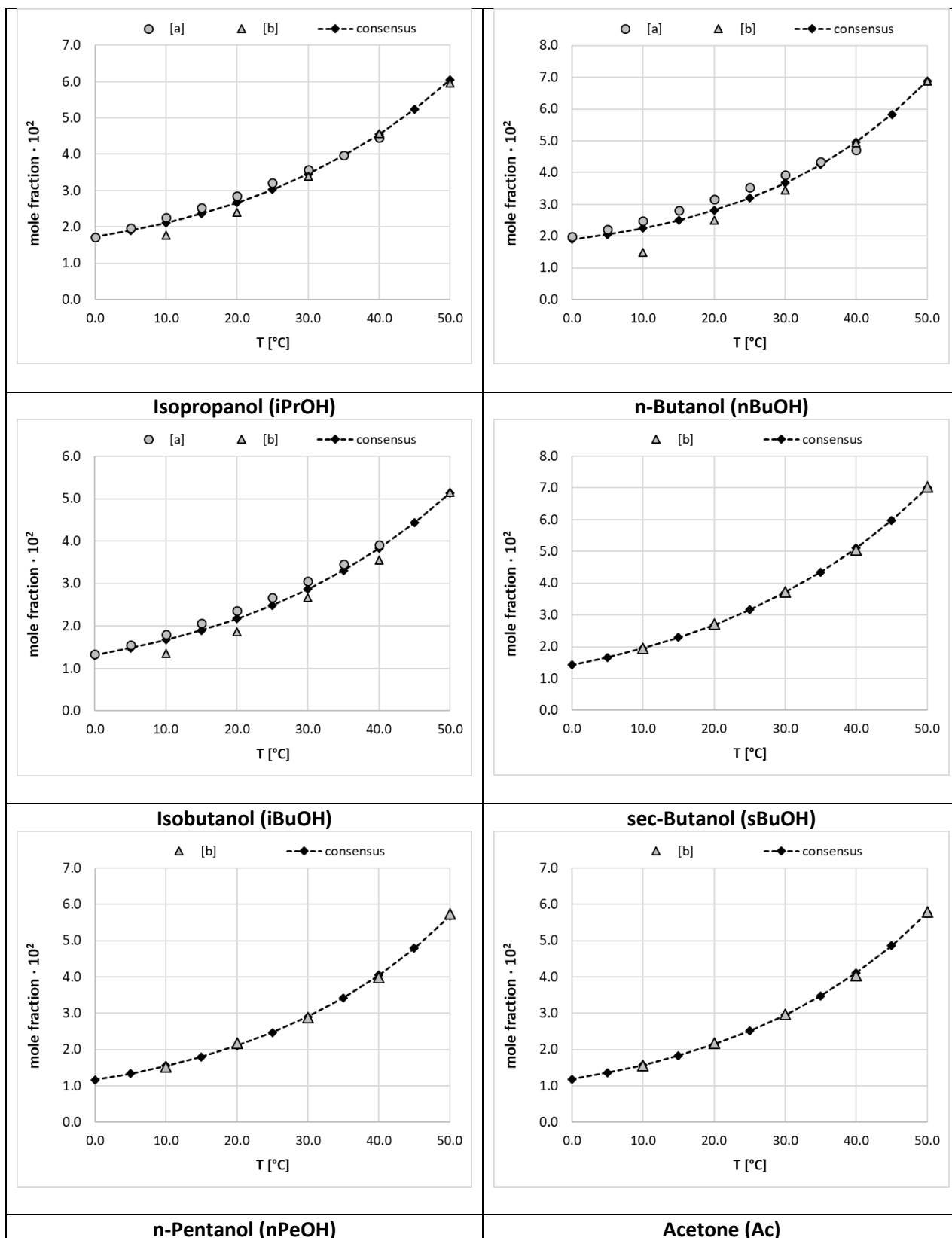
S2. Solubility data curation

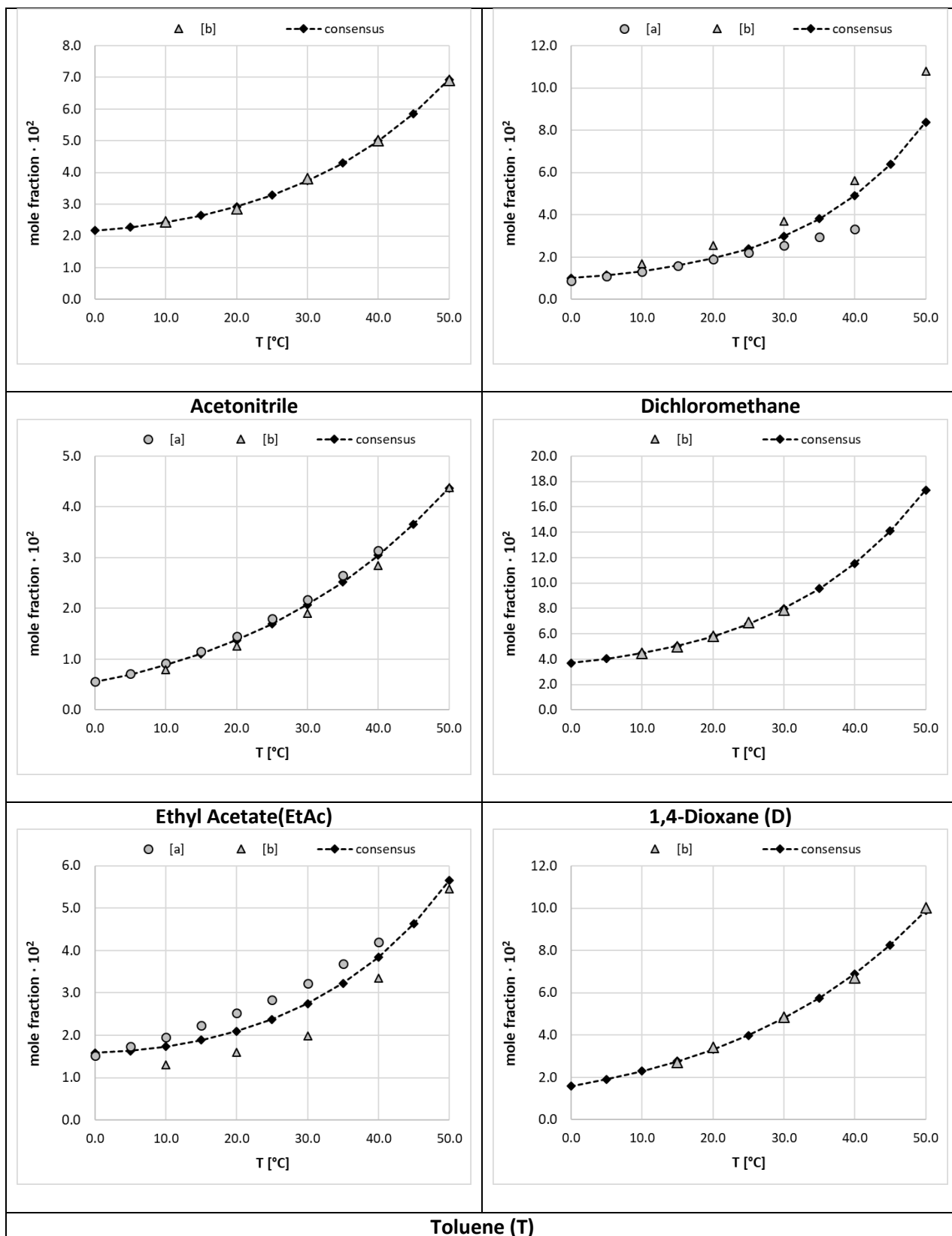
S2.1. Collection of EDA solubility in neat solvents

Table S2.1. Optimized values of model parameters used for data curation along with statistical measure of the fitting accuracy.

| solvent | acronim | A·10 ³ | B·10 ³ | C·10 ³ | RMSD·10 ³ | MAPE |
|------------------------|---------|-------------------|-------------------|-------------------|----------------------|------|
| water | W | 3.111 | -6.398 | 0.858 | 4.03 | 0.04 |
| methanol | MeOH | 31.984 | -17.949 | 2.185 | 165.47 | 4.19 |
| ethanol | EtOH | 20.389 | -11.949 | 1.439 | 68.09 | 1.33 |
| n-propanol | nPrOH | 29.274 | -17.128 | 2.198 | 133.95 | 2.51 |
| isopropanol | iPrOH | 18.599 | -10.825 | 1.246 | 90.26 | 1.90 |
| n-butanol | nBuOH | 15.486 | -8.444 | 0.834 | 8.89 | 0.24 |
| isobutanol | iBuOH | 21.045 | -11.898 | 1.348 | 18.43 | 0.45 |
| sec-butanol | sBuOH | 20.873 | -11.790 | 1.332 | 11.53 | 0.30 |
| n-pentanol | nPeOH | 34.206 | -20.257 | 2.695 | 14.61 | 0.34 |
| acetone | AT | 49.838 | -28.045 | 3.600 | 192.27 | 4.65 |
| acetonitrile | AE | 8.189 | -3.658 | 0.000 | 58.89 | 1.22 |
| dichloromethane | DC | 40.161 | -22.691 | 2.956 | 11.54 | 0.36 |
| ethyl acetate | EA | 44.601 | -26.411 | 3.577 | 178.67 | 4.16 |
| 1,4-dioxane | D | 17.497 | -9.079 | 0.865 | 21.92 | 0.68 |
| toluene | T | 3.251 | -1.145 | -0.296 | 7.08 | 0.16 |
| 2,4-dimethylphenol | DMP | -16.508 | 14.856 | -3.315 | 3.76 | 0.10 |
| 1-methyl-2-pyrrolidone | NMP | 10.517 | -3.004 | -0.330 | 23.14 | 0.73 |
| diglyme | DIG | -12.269 | 9.697 | -2.146 | 7.41 | 0.18 |
| triglyme | TIG | 62.116 | -33.404 | 4.066 | 5.93 | 0.14 |
| tetraglyme | TEG | -31.442 | 27.130 | -5.565 | 32.42 | 1.16 |
| DMSO | DMSO | -9.106 | 13.021 | -3.302 | 14.09 | 0.74 |
| 4FM | 4FM | 9.308 | -4.112 | 0.000 | 15.91 | 0.31 |







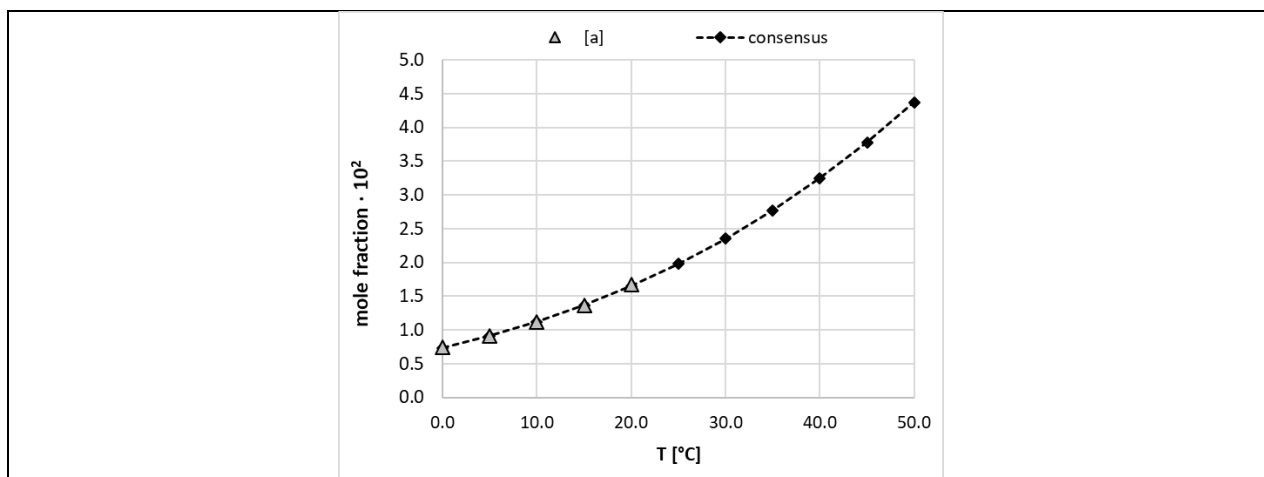
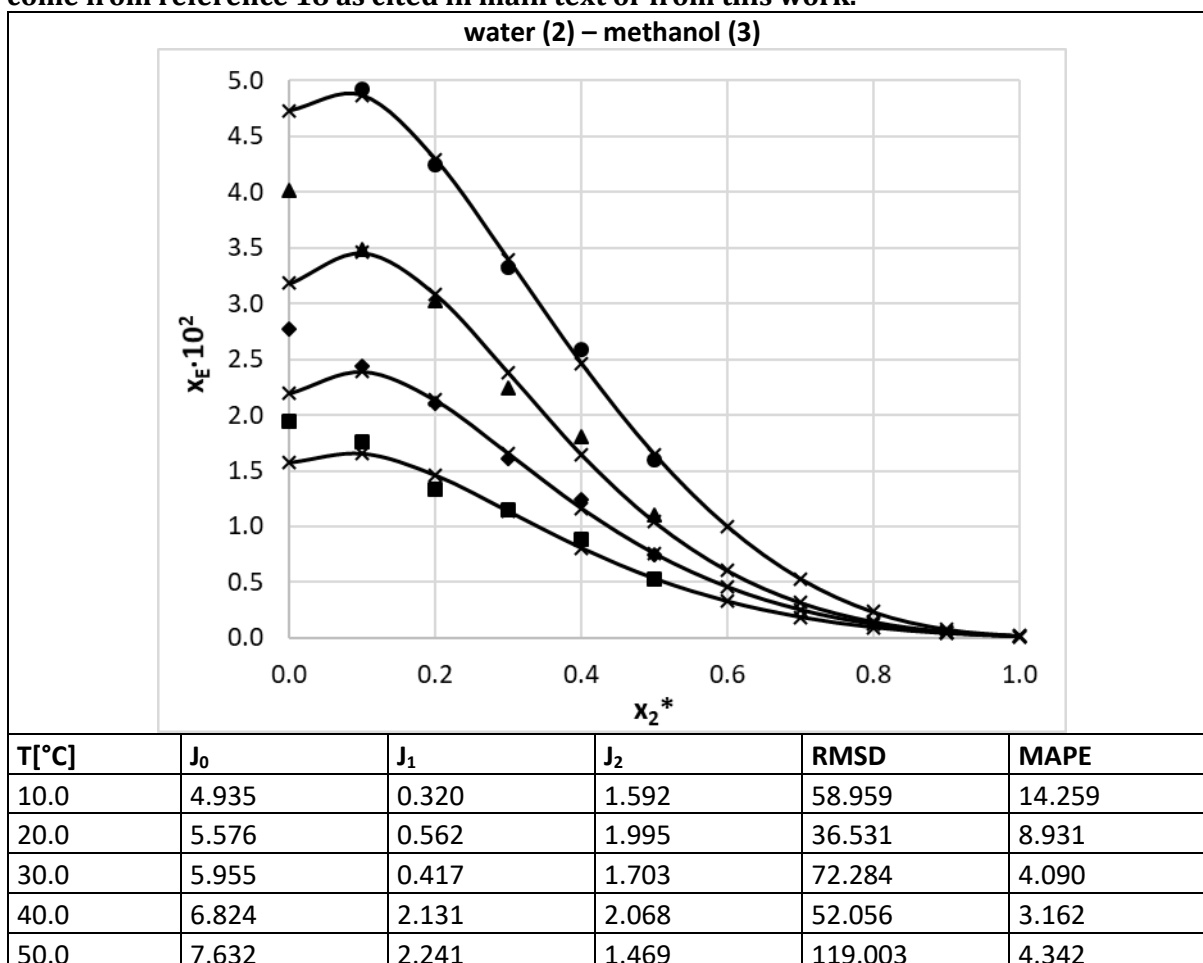


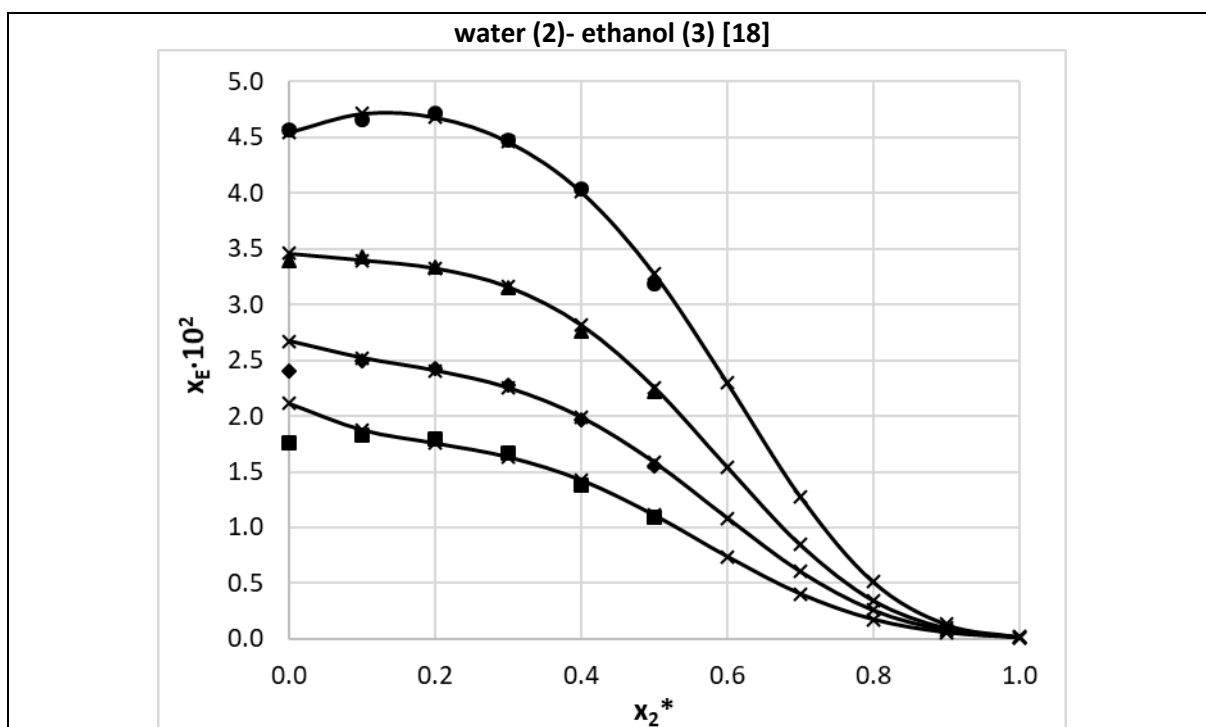
Figure S2.1. Graphical illustration of the solubility data curation of edaravone in neat solvents. The black diamonds represent curated values constituting the solubility dataset, while gray symbols characterize experimental points ([a], [b], [c] taken from references 18, 16, 17, respectively as cited in main text).

S2.2. Binary solvents mixtures

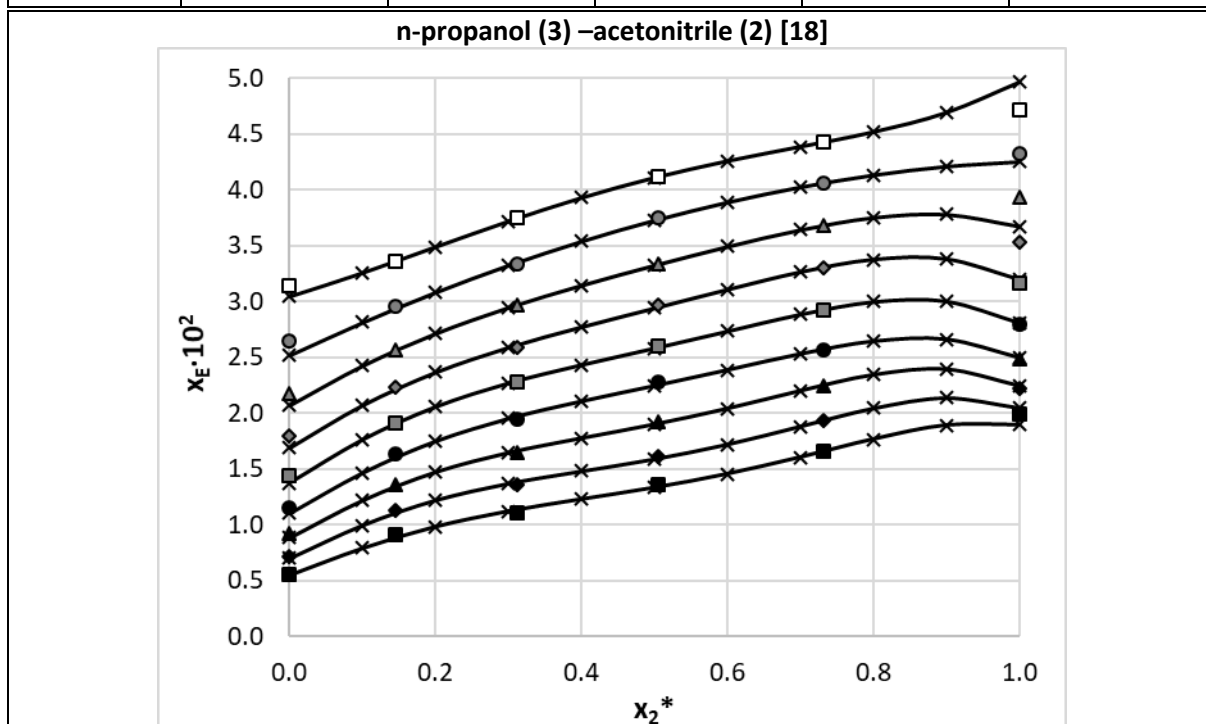
$$\ln x_{w,T} = w_1 \ln x_{1,T} + w_2 \ln x_{2,T} + \frac{w_1 w_2}{T/K} \sum_{i=0}^2 J_i (w_1 - w_2)^i$$

Table S2.2. Solubility data curation of edaravone in binary solvents. All experimental data come from reference 18 as cited in main text or from this work.



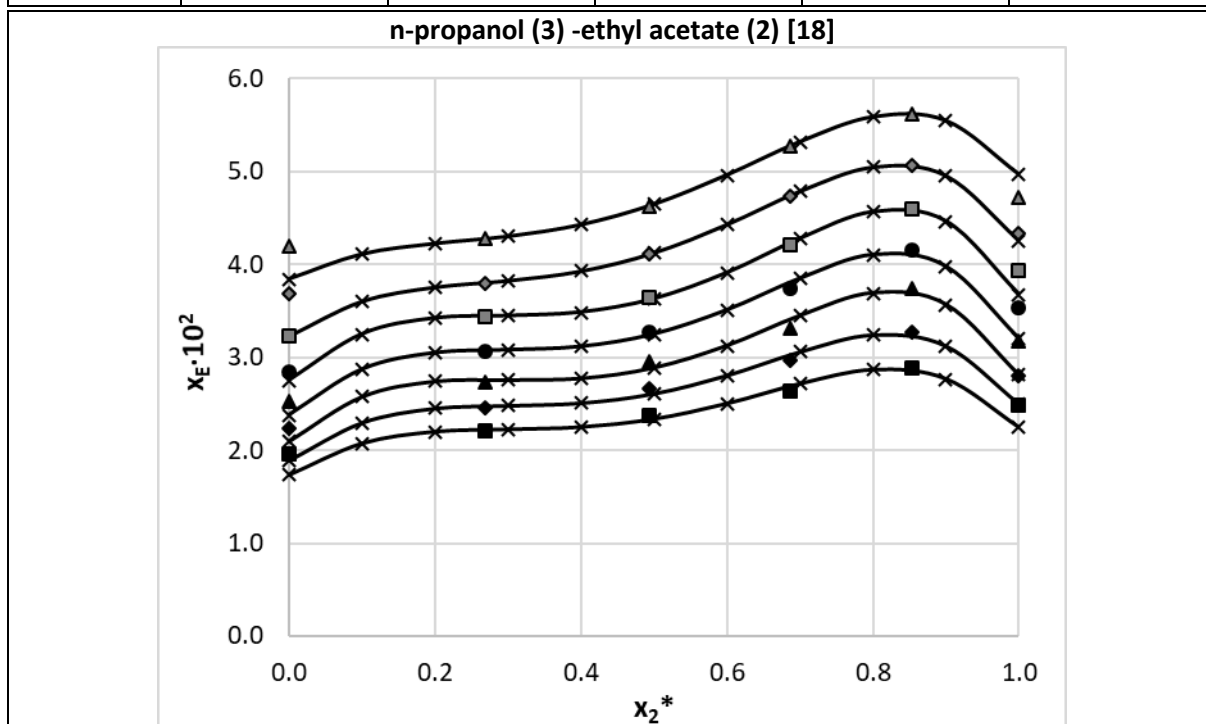


| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|--------|-------|--------|--------|-------|
| 10.0 | 7.281 | 3.449 | -0.574 | 32.544 | 6.264 |
| 20.0 | 8.122 | 4.338 | 0.587 | 30.913 | 5.251 |
| 30.0 | 8.877 | 4.757 | 0.964 | 49.154 | 5.418 |
| 40.0 | 9.646 | 5.615 | 2.082 | 58.181 | 5.770 |
| 50.0 | 10.520 | 6.562 | 3.602 | 87.403 | 6.526 |

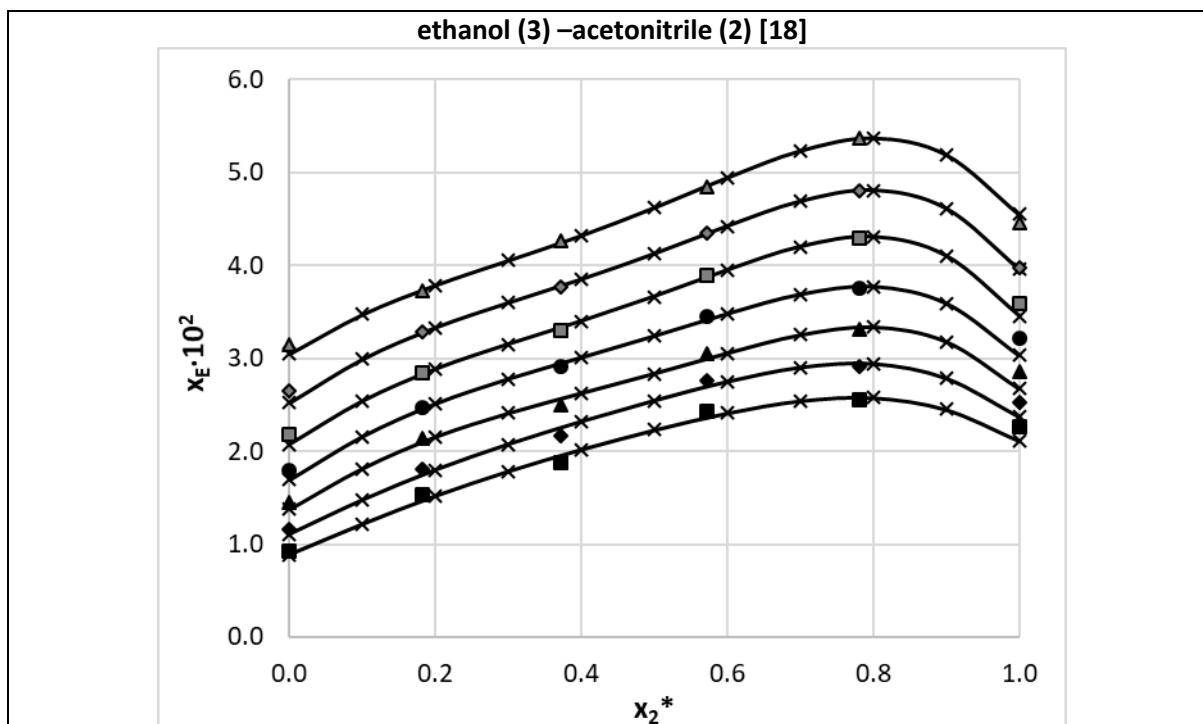


| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|-------|--------|-------|--------|-------|
| 0.0 | 1.069 | -0.816 | 1.467 | 15.902 | 1.065 |
| 5.0 | 1.131 | -0.666 | 1.658 | 12.582 | 0.691 |

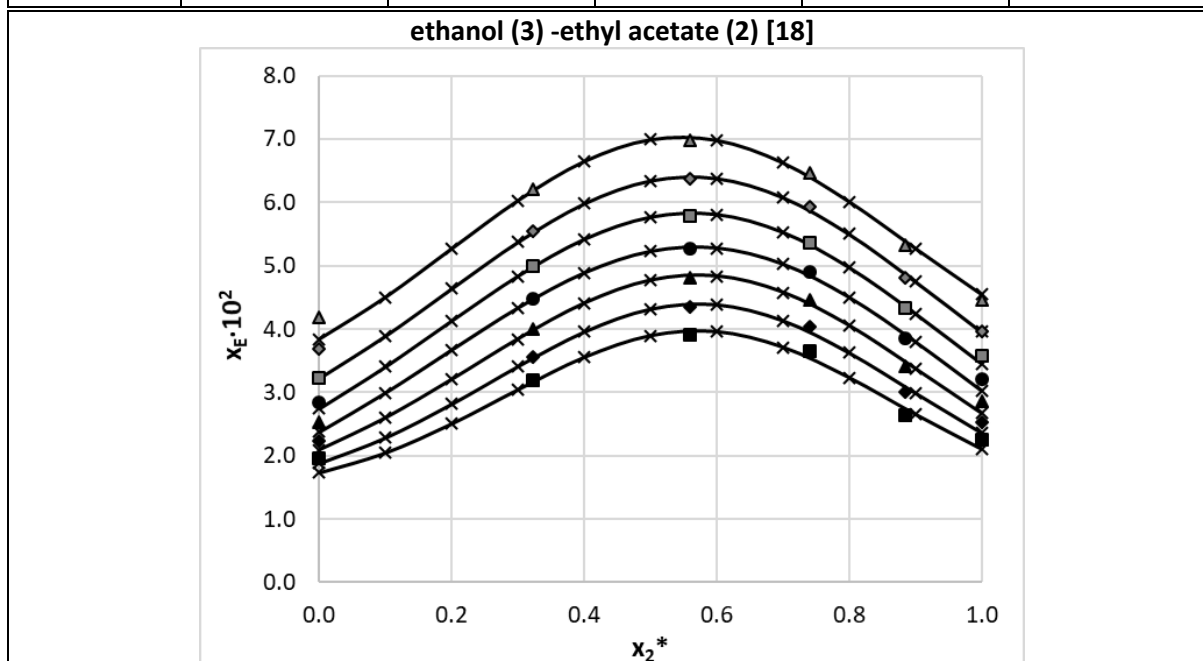
| | | | | | |
|------|-------|--------|--------|--------|-------|
| 10.0 | 1.198 | -0.494 | 1.470 | 13.761 | 0.629 |
| 15.0 | 1.204 | -0.398 | 1.128 | 19.602 | 0.756 |
| 20.0 | 1.092 | -0.275 | 1.030 | 6.988 | 0.228 |
| 25.0 | 0.936 | -0.138 | 0.778 | 13.604 | 0.392 |
| 30.0 | 0.746 | -0.106 | 0.453 | 3.783 | 0.096 |
| 35.0 | 0.521 | -0.109 | 0.060 | 13.166 | 0.294 |
| 40.0 | 0.223 | -0.180 | -0.249 | 1.361 | 0.027 |



| T[°C] | J ₀ | J ₁ | J ₂ | RMSD | MAPE |
|-------|----------------|----------------|----------------|--------|-------|
| 0.0 | 0.348 | 0.736 | 0.694 | 2.541 | 0.094 |
| 5.0 | 0.523 | 0.591 | 1.736 | 17.786 | 0.608 |
| 10.0 | 0.674 | 0.563 | 2.305 | 32.779 | 0.989 |
| 15.0 | 0.737 | 0.579 | 2.493 | 41.052 | 1.104 |
| 20.0 | 0.693 | 0.629 | 2.765 | 54.727 | 1.314 |
| 25.0 | 0.655 | 0.618 | 2.495 | 34.850 | 0.716 |
| 30.0 | 0.531 | 0.592 | 2.309 | 15.065 | 0.290 |
| 35.0 | 0.427 | 0.671 | 1.604 | 5.045 | 0.087 |
| 40.0 | 0.246 | 0.652 | 1.153 | 2.607 | 0.040 |

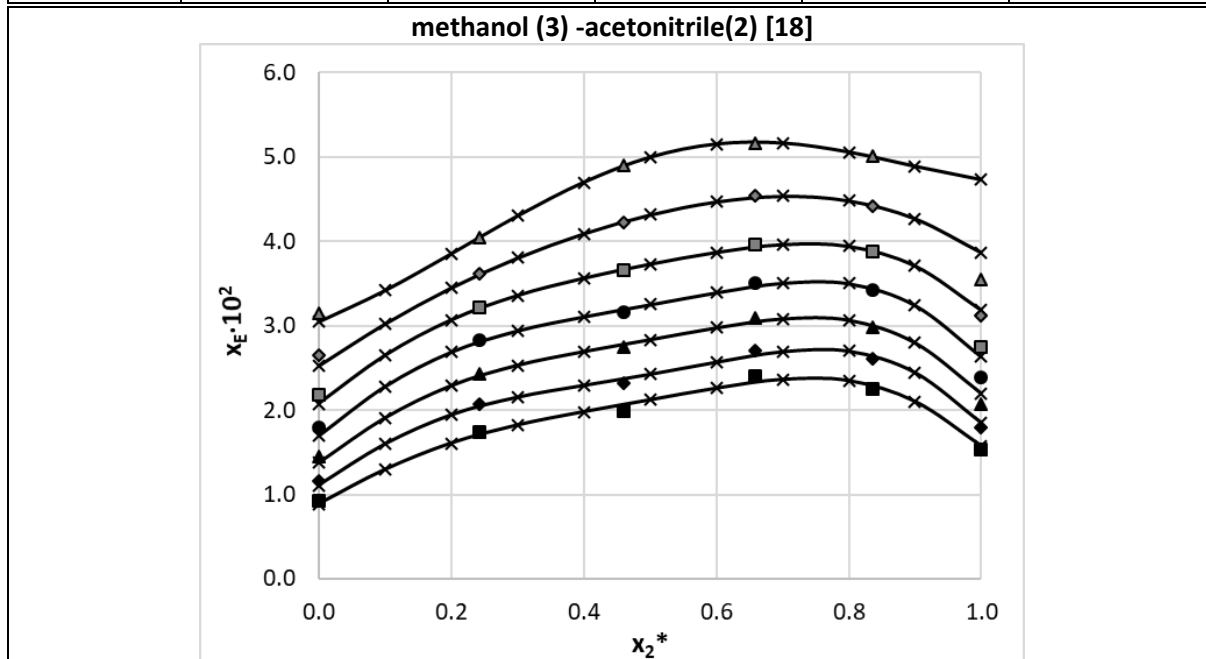


| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|-------|--------|-------|--------|-------|
| 0.0 | 2.244 | -0.142 | 0.369 | 16.928 | 0.823 |
| 5.0 | 2.130 | -0.042 | 0.741 | 27.080 | 1.264 |
| 10.0 | 1.959 | 0.039 | 0.986 | 49.030 | 1.951 |
| 15.0 | 1.810 | 0.182 | 1.075 | 50.854 | 1.740 |
| 20.0 | 1.570 | 0.198 | 1.470 | 39.716 | 1.175 |
| 25.0 | 1.433 | 0.297 | 1.346 | 25.193 | 0.621 |
| 30.0 | 1.255 | 0.491 | 1.274 | 18.085 | 0.407 |
| 35.0 | 1.066 | 0.501 | 1.135 | 4.698 | 0.093 |
| 40.0 | 0.865 | 0.566 | 0.927 | 10.274 | 0.180 |

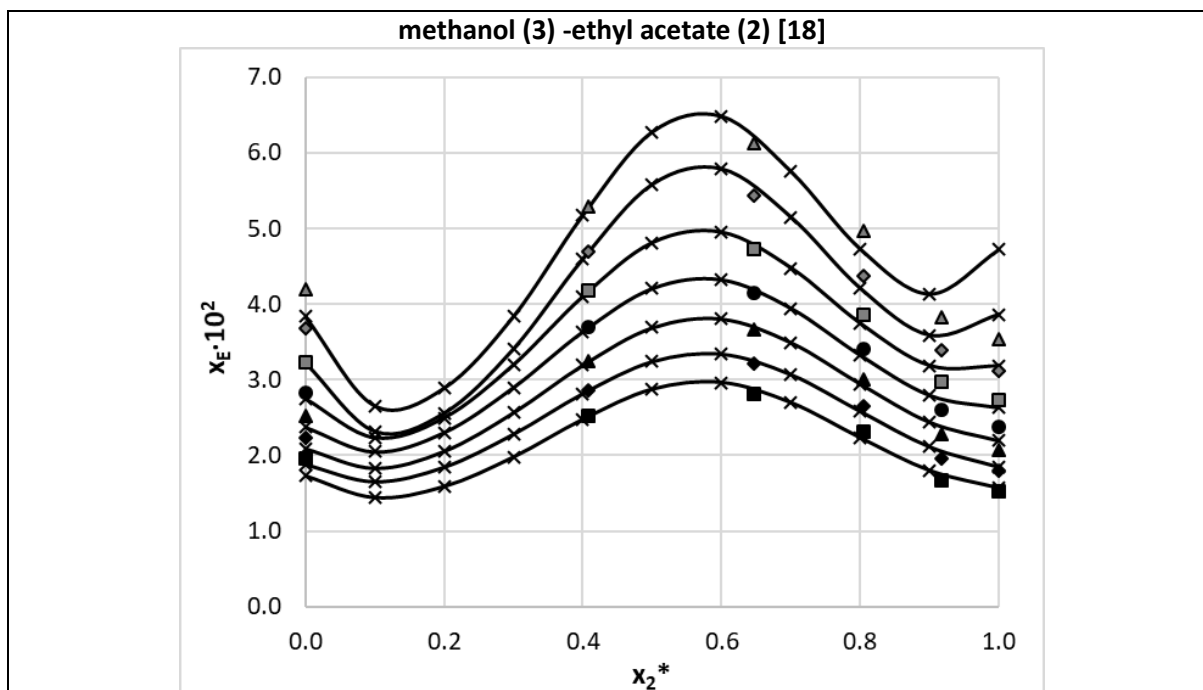


| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|-------|-------|--------|---------|-------|
| 0.0 | 2.559 | 0.951 | -1.637 | 102.143 | 2.974 |

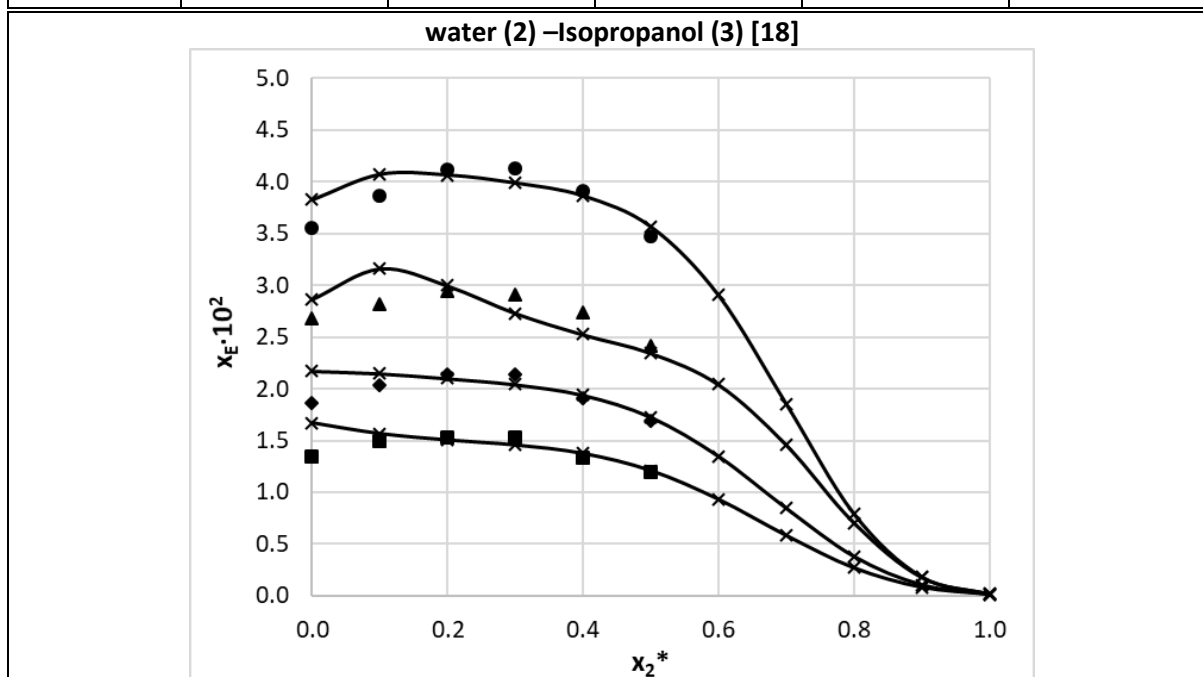
| | | | | | |
|------|-------|-------|--------|--------|-------|
| 5.0 | 2.792 | 0.817 | -1.586 | 91.718 | 2.223 |
| 10.0 | 2.843 | 0.714 | -0.996 | 68.643 | 1.564 |
| 15.0 | 2.860 | 0.601 | -0.737 | 55.115 | 1.117 |
| 20.0 | 2.805 | 0.463 | -0.471 | 47.899 | 0.867 |
| 25.0 | 2.665 | 0.301 | -0.192 | 35.629 | 0.579 |
| 30.0 | 2.503 | 0.246 | -0.261 | 27.505 | 0.393 |
| 35.0 | 2.296 | 0.247 | -0.361 | 37.908 | 0.502 |
| 40.0 | 2.058 | 0.159 | -0.546 | 47.431 | 0.570 |



| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|-------|--------|--------|--------|-------|
| 0.0 | 2.752 | -0.076 | 1.068 | 24.958 | 1.287 |
| 5.0 | 2.534 | 0.047 | 1.954 | 40.614 | 1.634 |
| 10.0 | 2.350 | 0.179 | 2.151 | 49.757 | 1.839 |
| 15.0 | 2.113 | 0.109 | 2.314 | 36.525 | 1.162 |
| 20.0 | 1.960 | 0.051 | 1.891 | 26.510 | 0.726 |
| 25.0 | 1.726 | -0.005 | 1.693 | 21.156 | 0.504 |
| 30.0 | 1.493 | -0.034 | 1.118 | 12.033 | 0.242 |
| 35.0 | 1.293 | 0.030 | 0.405 | 9.733 | 0.178 |
| 40.0 | 1.098 | 0.043 | -0.434 | 6.343 | 0.102 |



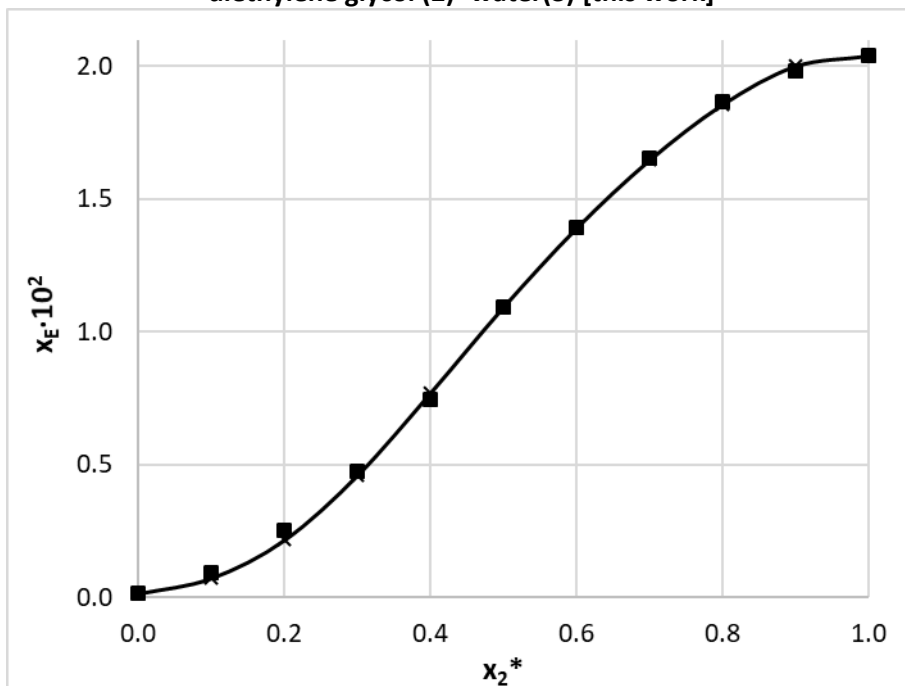
| T[°C] | J ₀ | J ₁ | J ₂ | RMSD | MAPE |
|-------|----------------|----------------|----------------|---------|-------|
| 0.0 | 1.765 | 1.825 | -2.584 | 61.288 | 2.725 |
| 5.0 | 2.022 | 1.807 | -2.824 | 52.796 | 2.027 |
| 10.0 | 2.220 | 2.071 | -3.857 | 55.394 | 1.836 |
| 15.0 | 2.209 | 1.805 | -3.397 | 55.023 | 1.589 |
| 20.0 | 2.172 | 1.724 | -3.630 | 57.723 | 1.448 |
| 25.0 | 2.079 | 1.590 | -4.015 | 71.032 | 1.563 |
| 30.0 | 1.948 | 1.661 | -4.848 | 92.465 | 1.790 |
| 35.0 | 1.832 | 2.029 | -6.378 | 117.226 | 2.005 |
| 40.0 | 1.551 | 1.919 | -6.782 | 176.555 | 2.651 |



| T[°C] | J ₀ | J ₁ | J ₂ | RMSD | MAPE |
|-------|----------------|----------------|----------------|--------|-------|
| 10.0 | 8.087 | 5.722 | 1.542 | 42.478 | 4.391 |

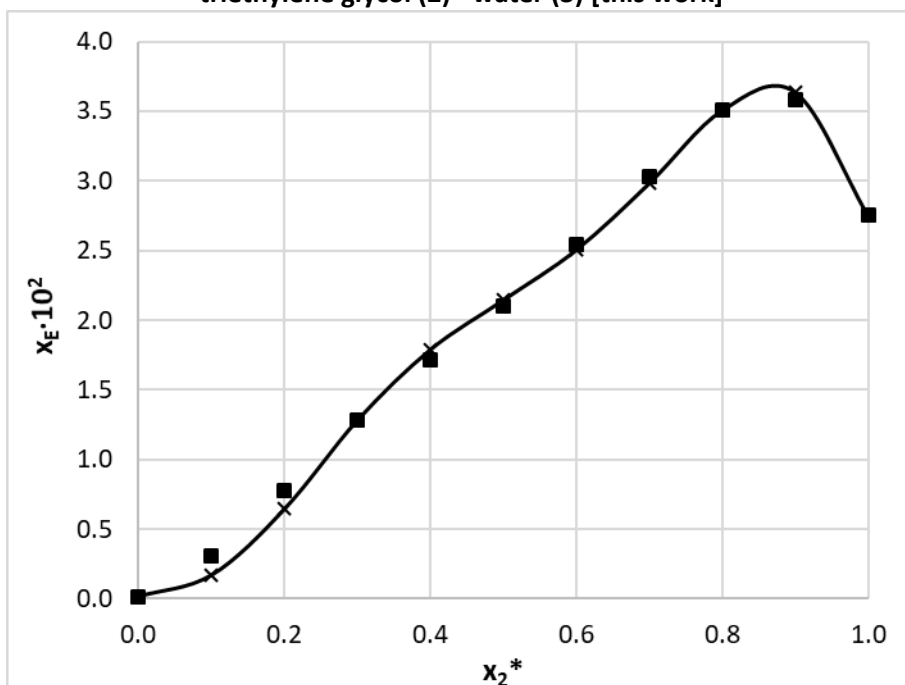
| | | | | | |
|------|--------|-------|-------|---------|-------|
| 20.0 | 8.881 | 6.437 | 2.481 | 63.409 | 6.063 |
| 30.0 | 9.406 | 8.426 | 6.408 | 149.800 | 5.715 |
| 40.0 | 10.331 | 8.081 | 4.253 | 137.453 | 8.299 |
| 50.0 | 11.214 | 8.878 | 5.335 | 218.016 | 9.134 |

diethylene glycol (2) -water(3) [this work]

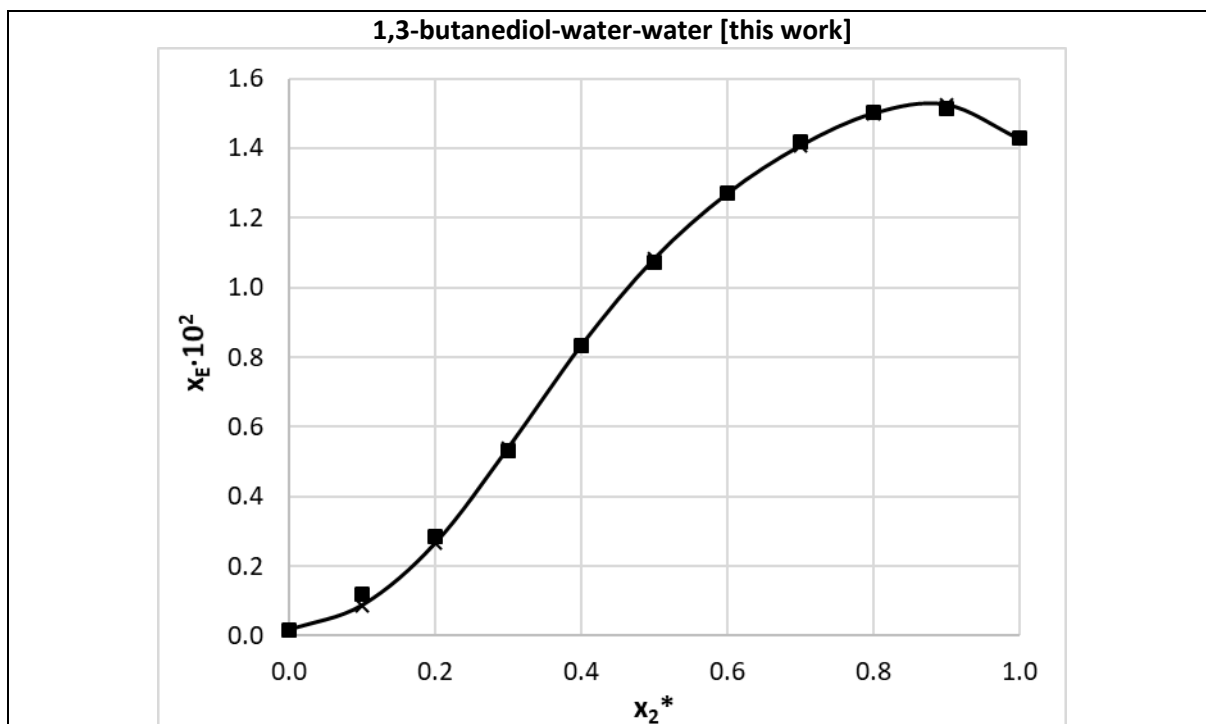


| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|-------|--------|-------|--------|-------|
| 25.0 | 7.112 | -3.865 | 1.720 | 16.725 | 3.815 |

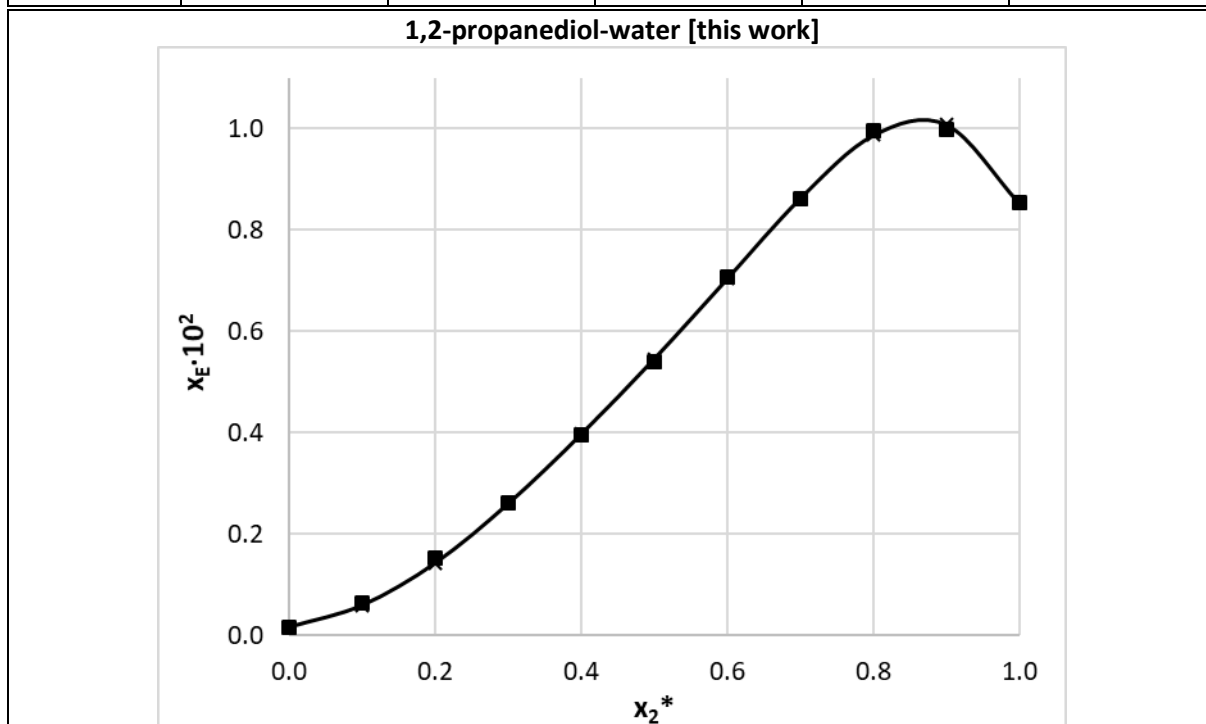
triethylene glycol (2) -water (3) [this work]



| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|-------|--------|-------|--------|-------|
| 25.0 | 9.207 | -7.124 | 8.221 | 67.058 | 6.571 |



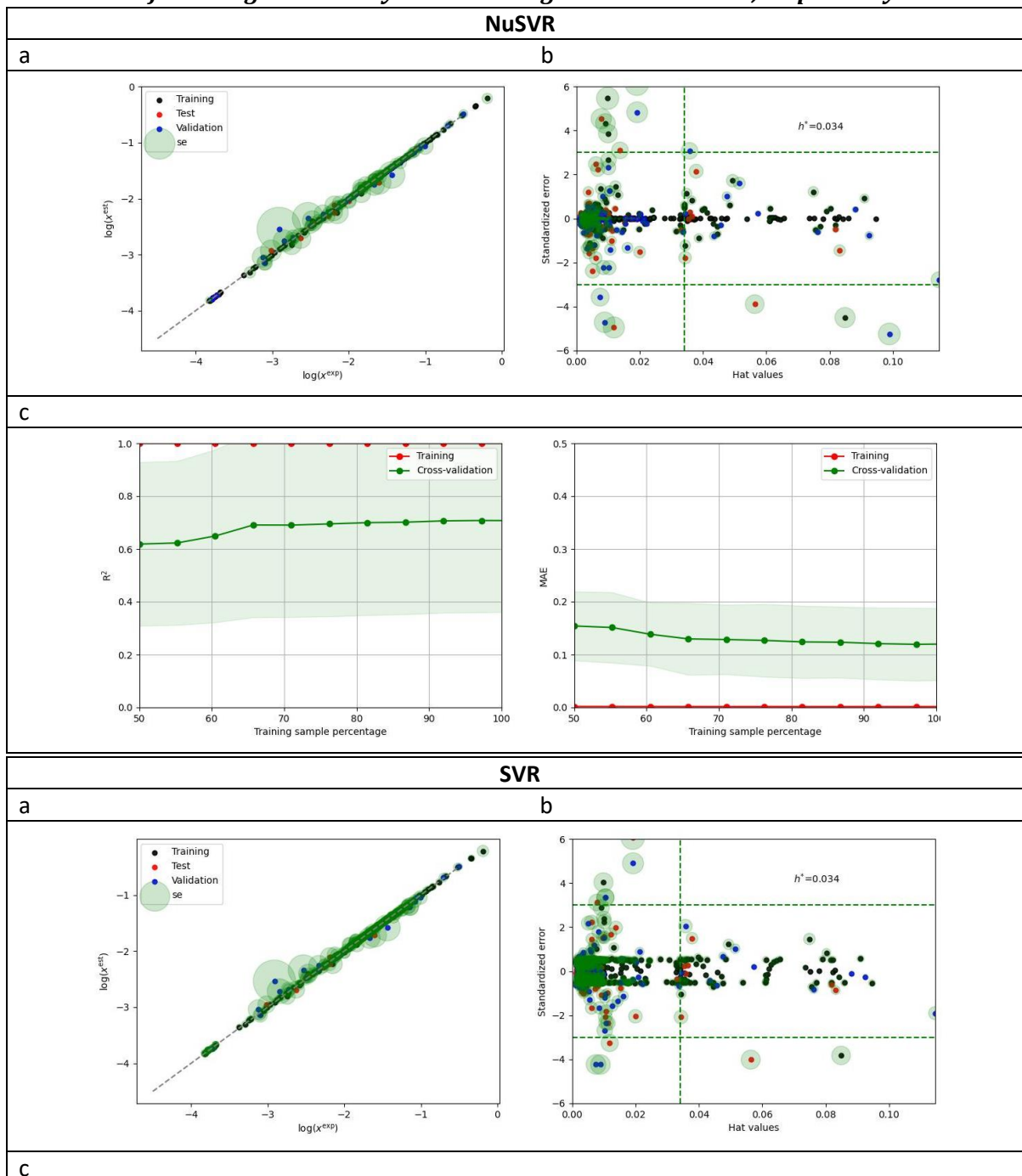
| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|-------|--------|-------|--------|-------|
| 25.0 | 7.787 | -4.889 | 2.818 | 13.222 | 3.480 |

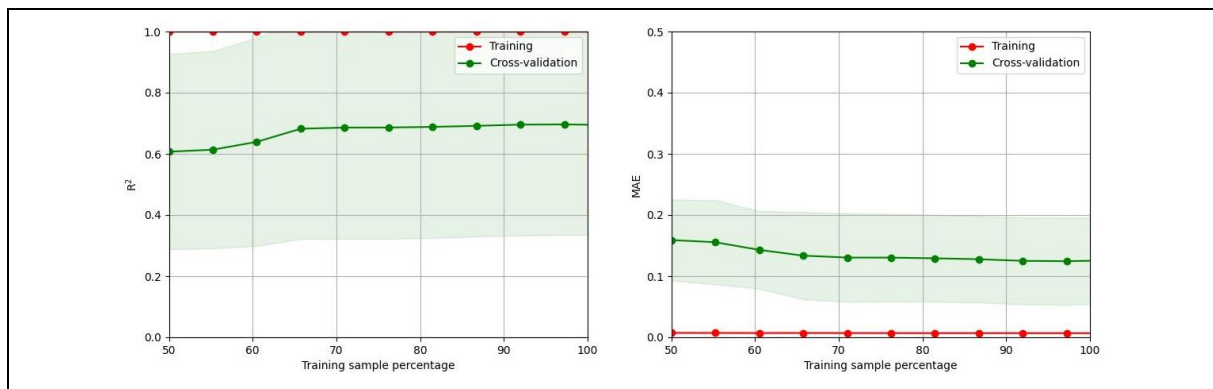


| T[°C] | J_0 | J_1 | J_2 | RMSD | MAPE |
|-------|-------|--------|-------|-------|-------|
| 25.0 | 6.077 | -2.239 | 3.021 | 5.422 | 1.480 |

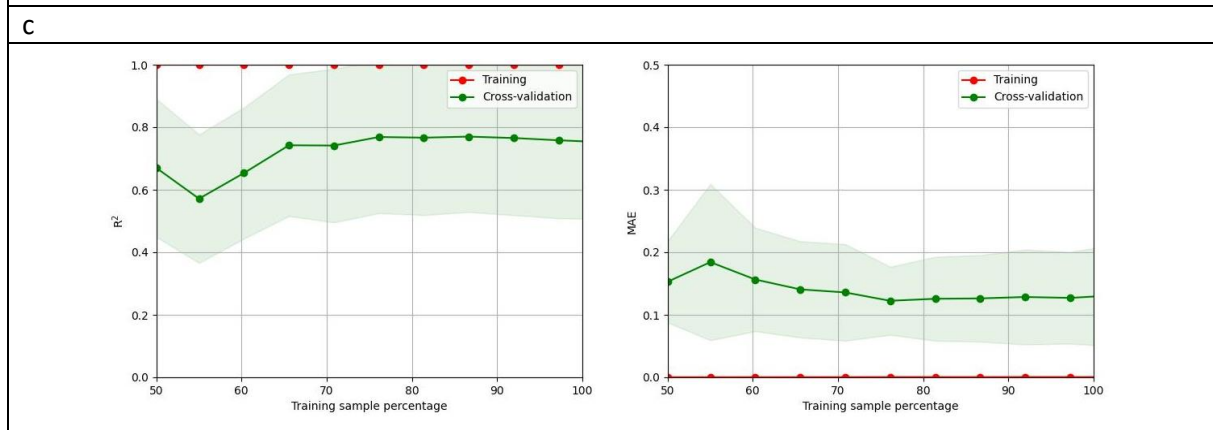
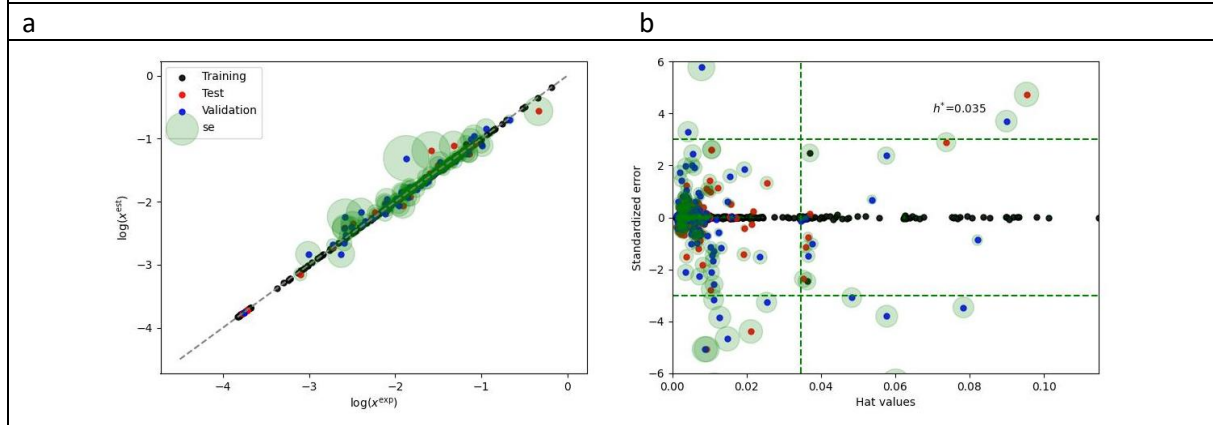
S3. Regressor models characteristics

Table S3.1. Graphical illustration of the of the regression model performance belonging to set A. The panels a,b, and c document the correlation between computed and consensus solubility values with annotation of the standard deviation as circles radius, applicability domain plots, and results of learning curve analysis concerning both R^2 and MAE, respectively.

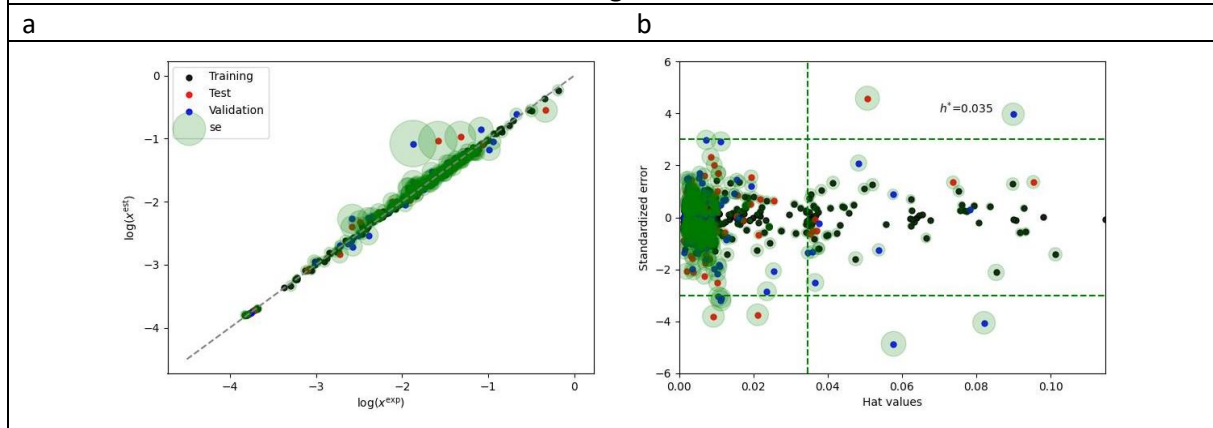




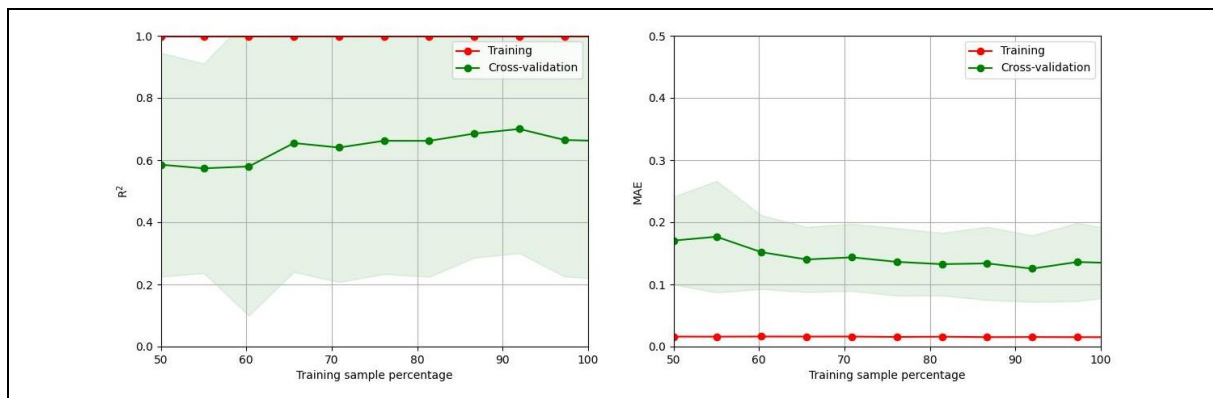
CatBoostRegressor



XGBRegressor

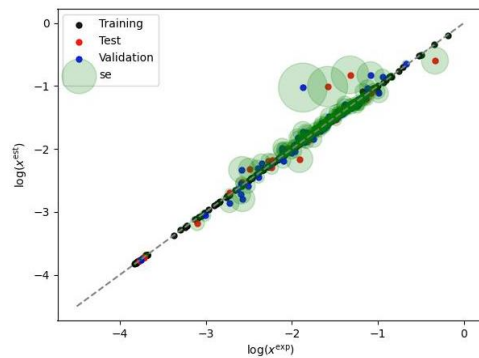


C

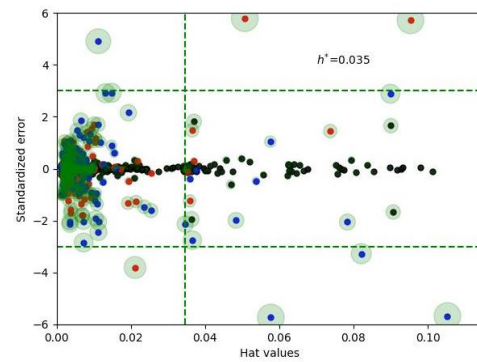


HistGradientBoostingRegressor

a



b



c

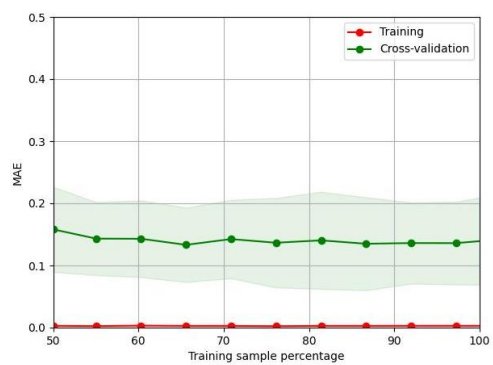
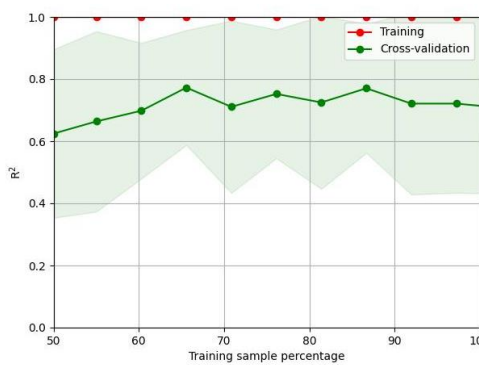
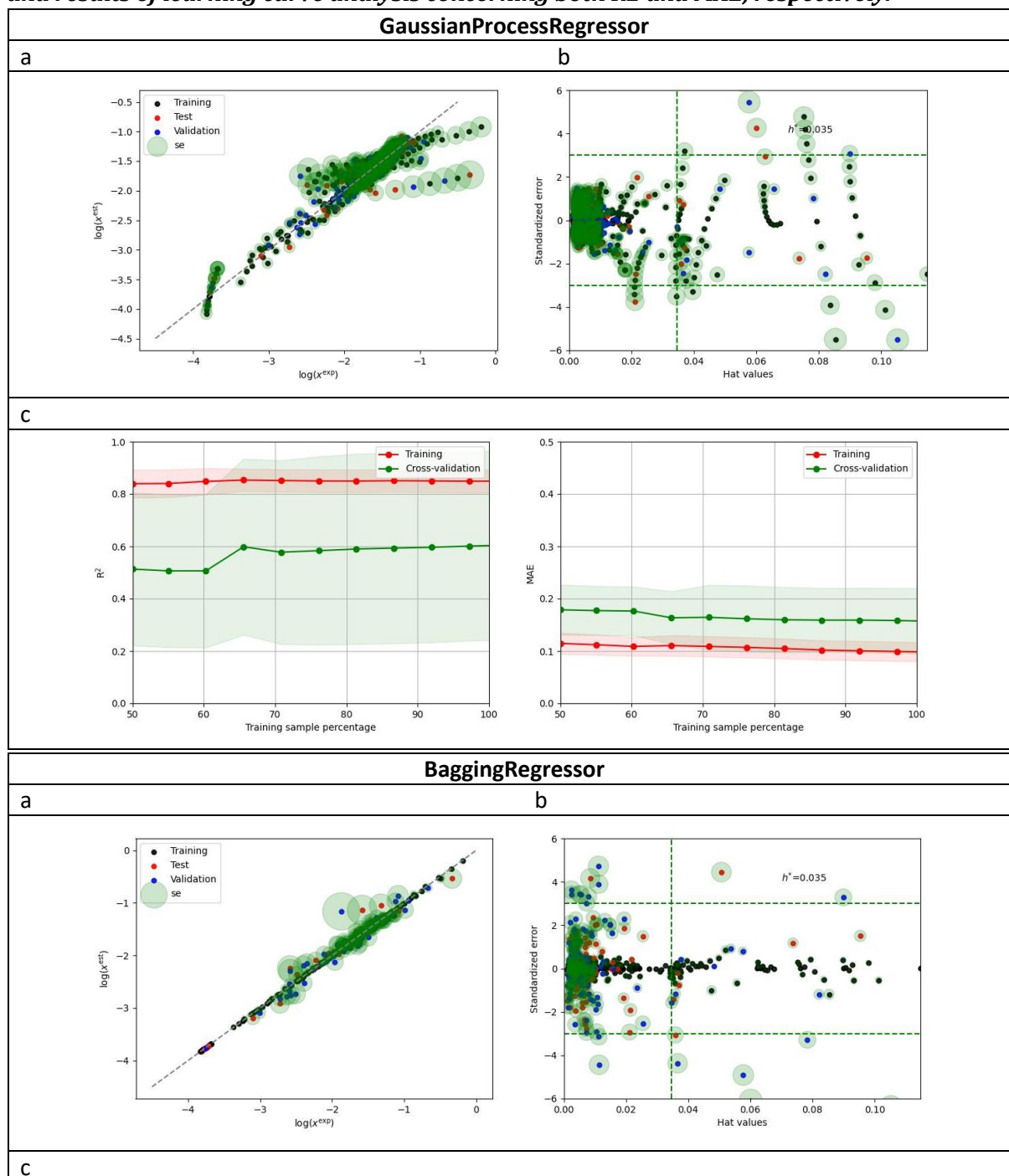
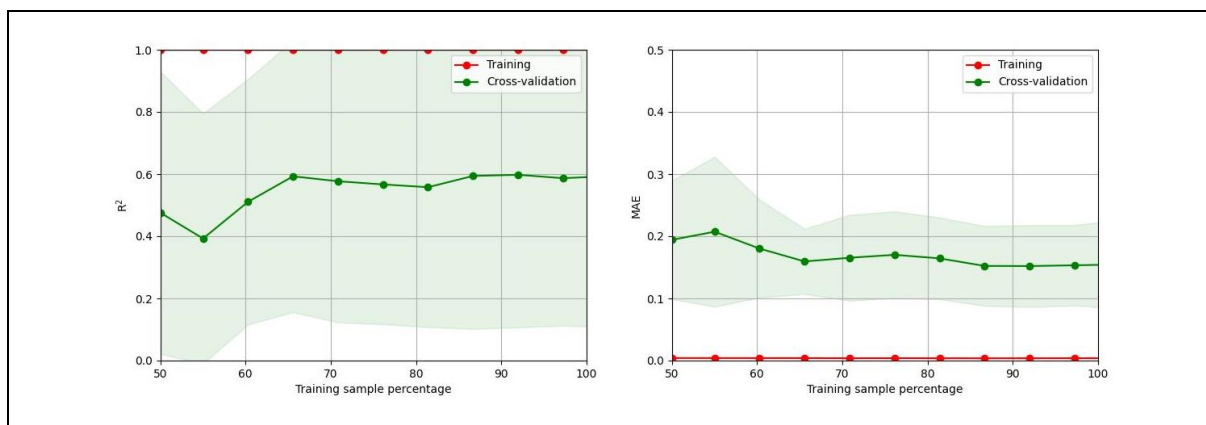
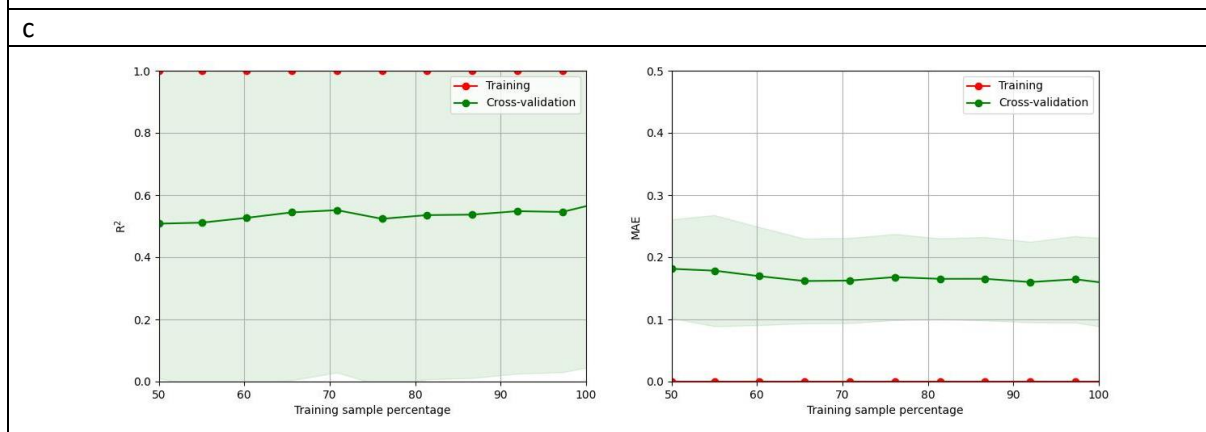
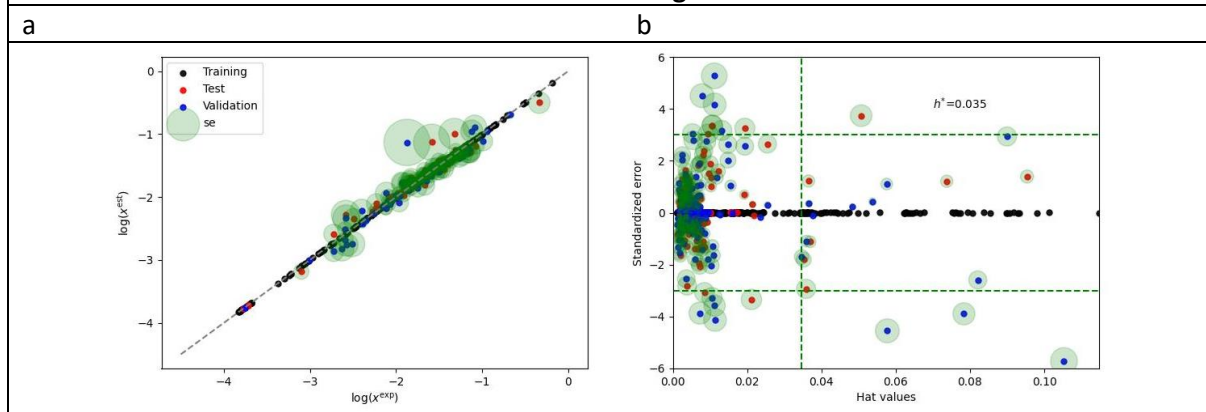


Table S3.2. Graphical illustration of the of the regression model performance belonging to set B. The panels a,b, and c document the correlation between computed and consensus solubility values with annotation of the standard deviation as circles radius, applicability domain plots, and results of learning curve analysis concerning both R² and MAE, respectively.

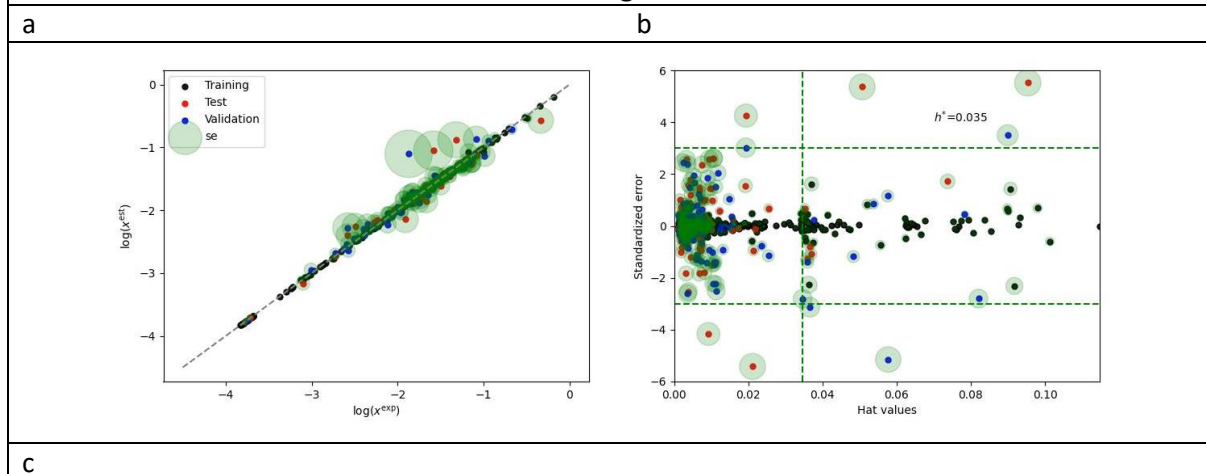


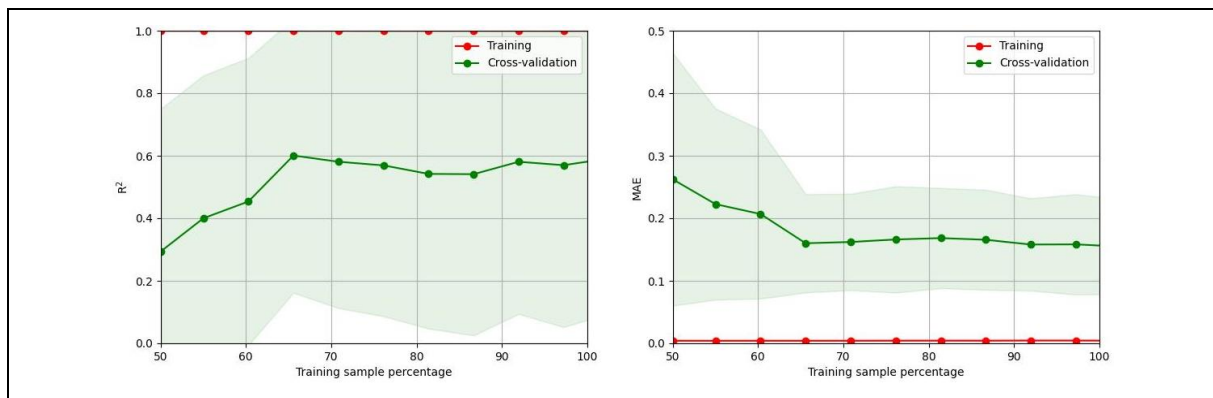


RandomForestRegressor

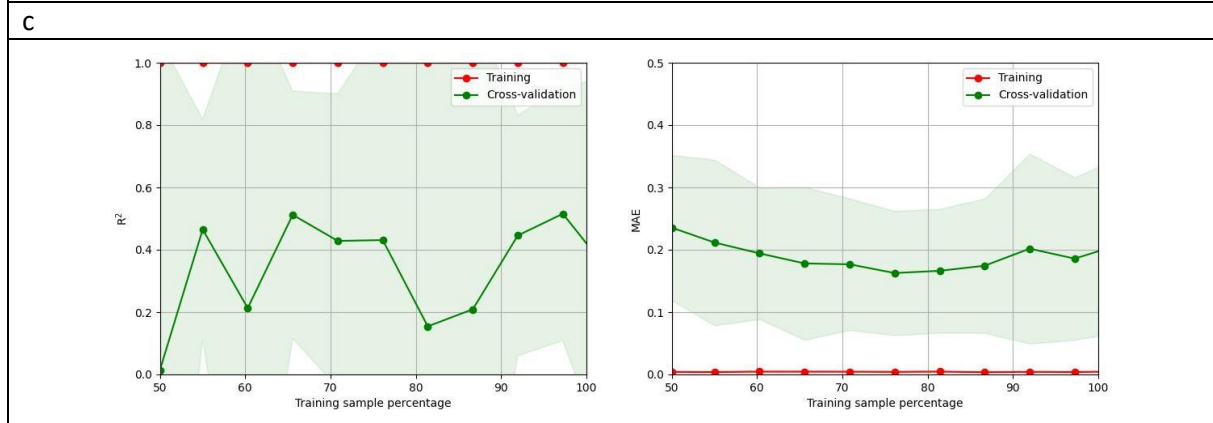
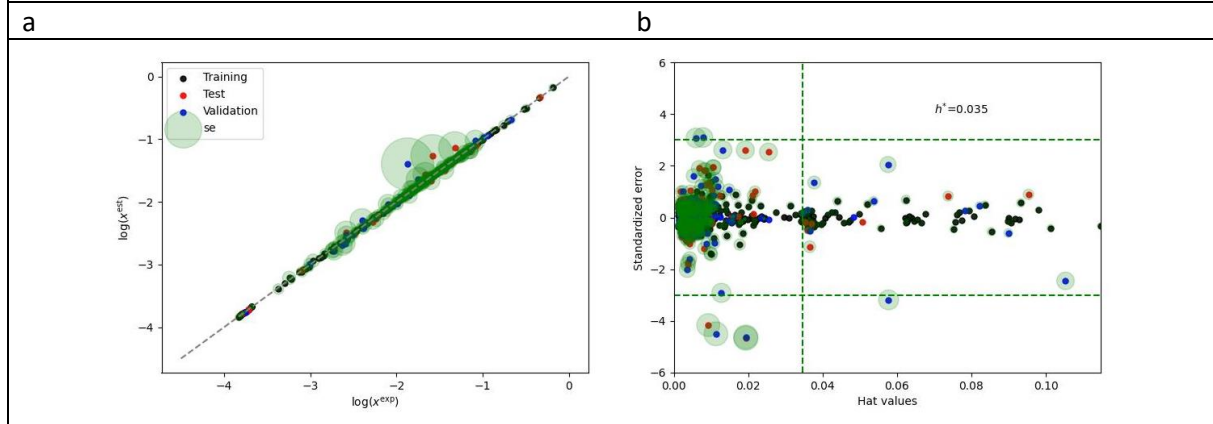


LGBMRegressor

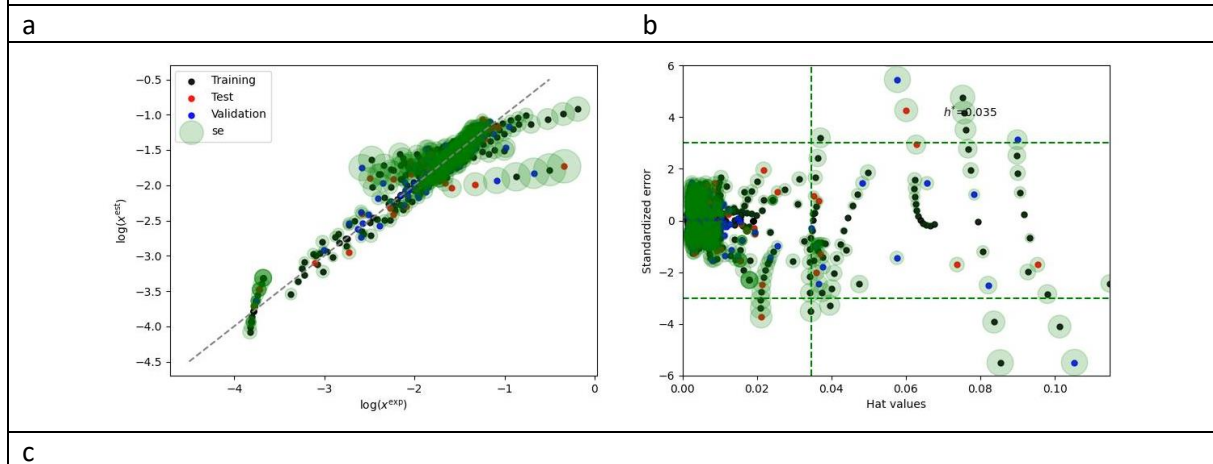


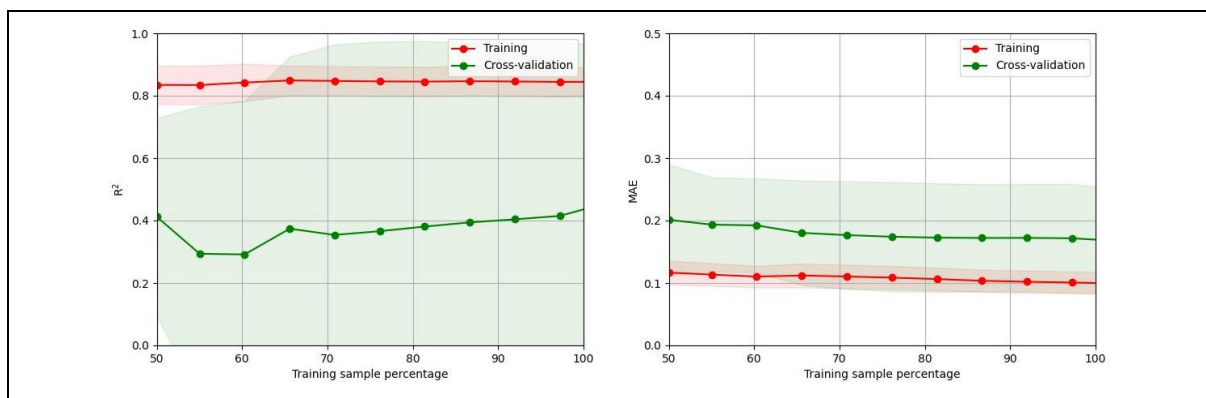


MLPregressor



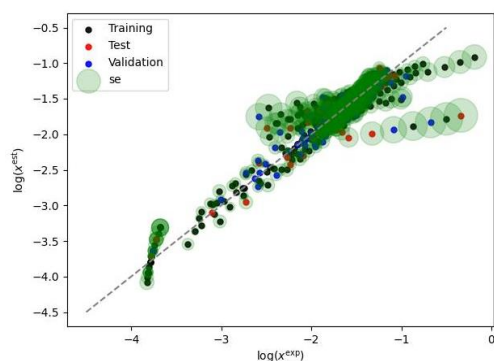
Lassolars



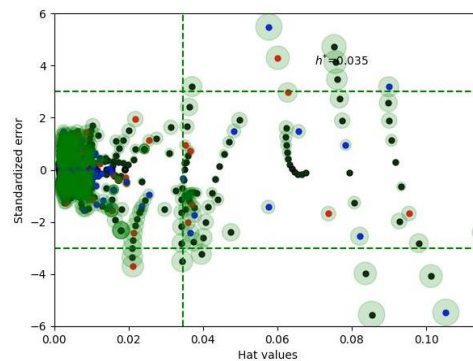


LassoLarsCV

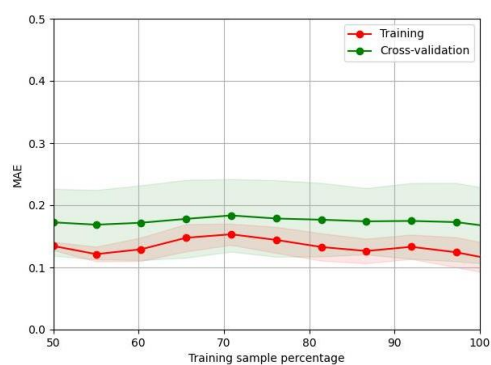
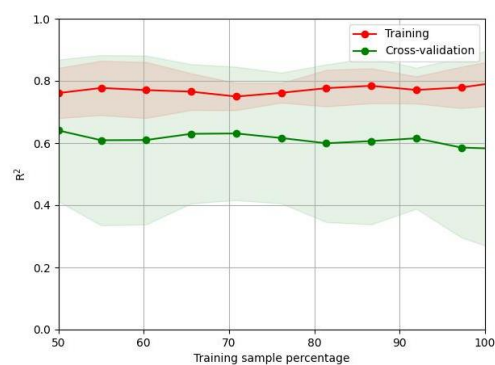
a



b

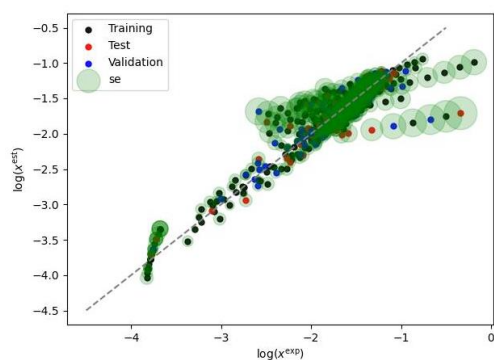


c

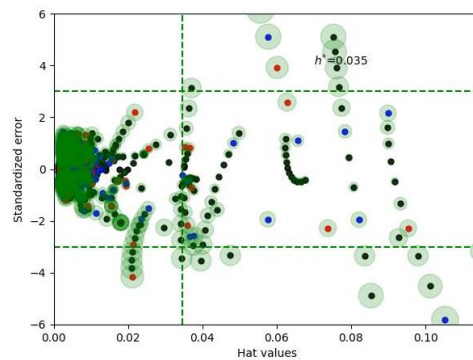


Ridge

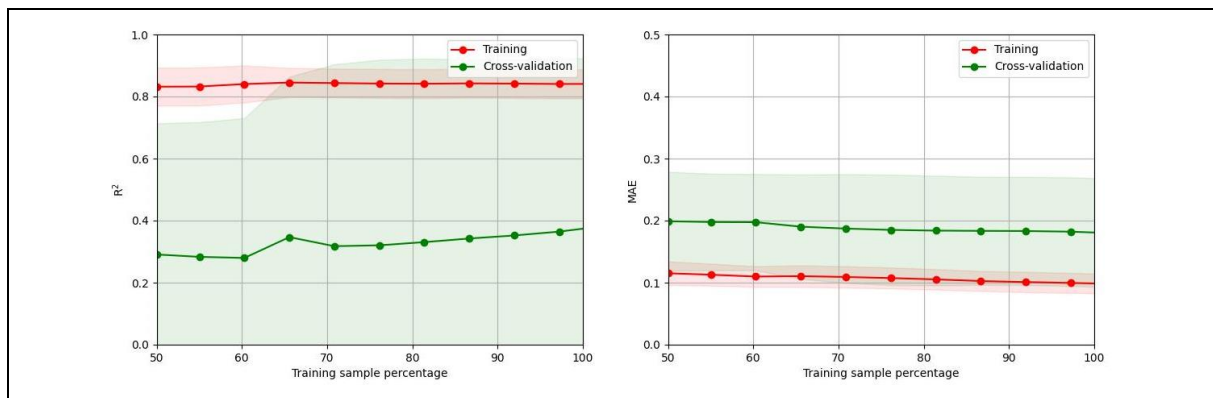
a



b

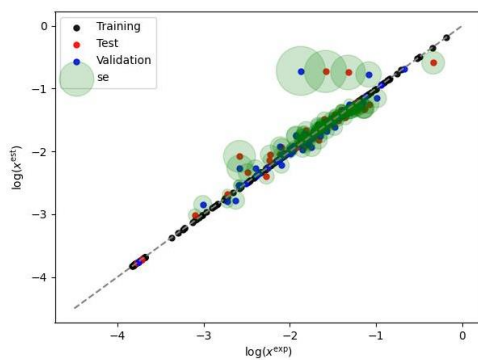


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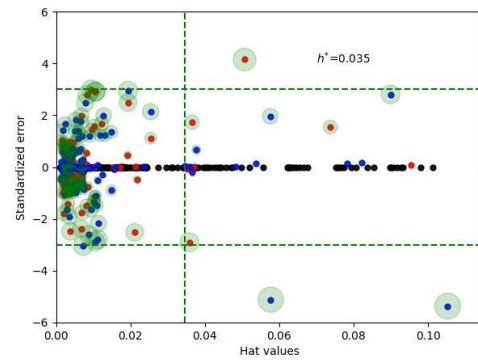


KNeighborsRegressor

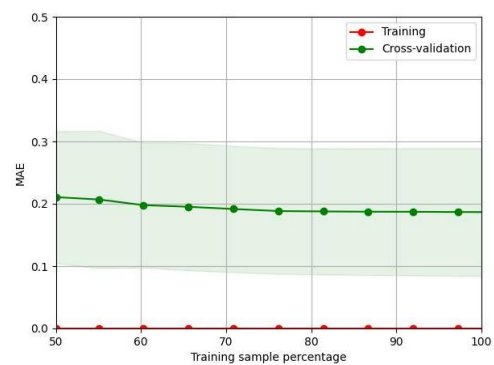
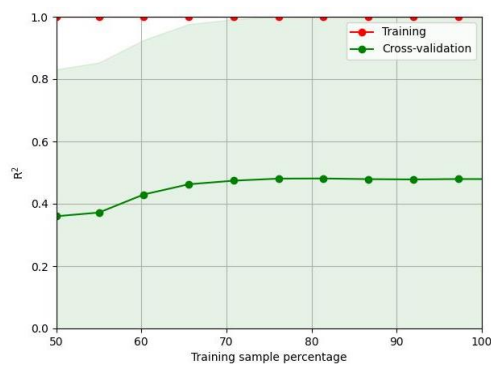
a



b

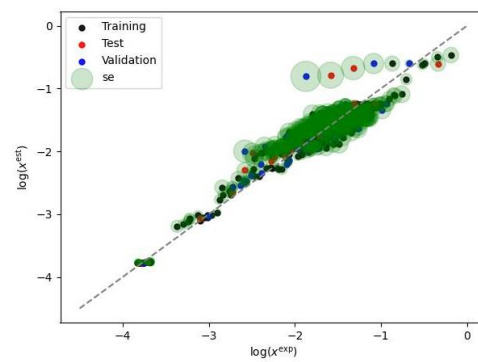


c

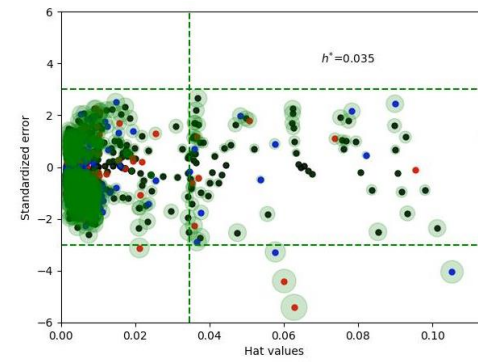


AdaBoostRegressor

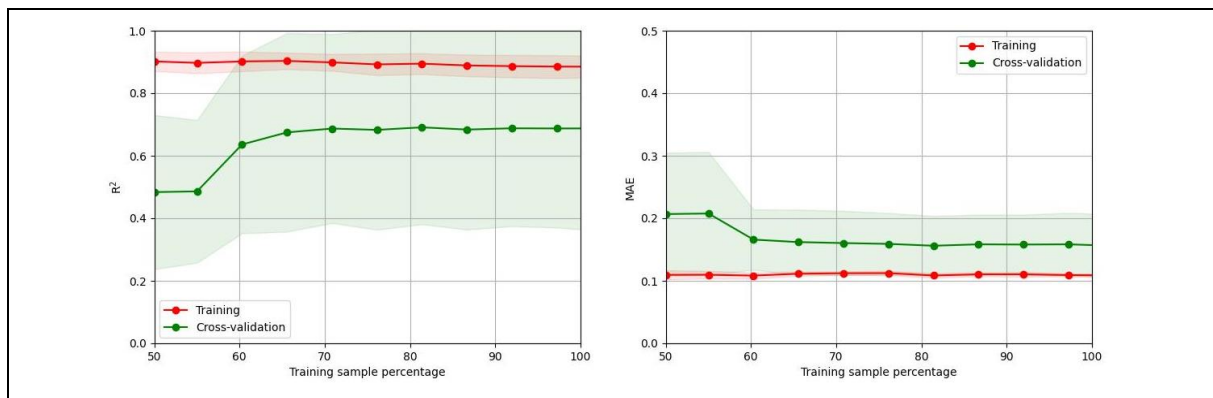
a



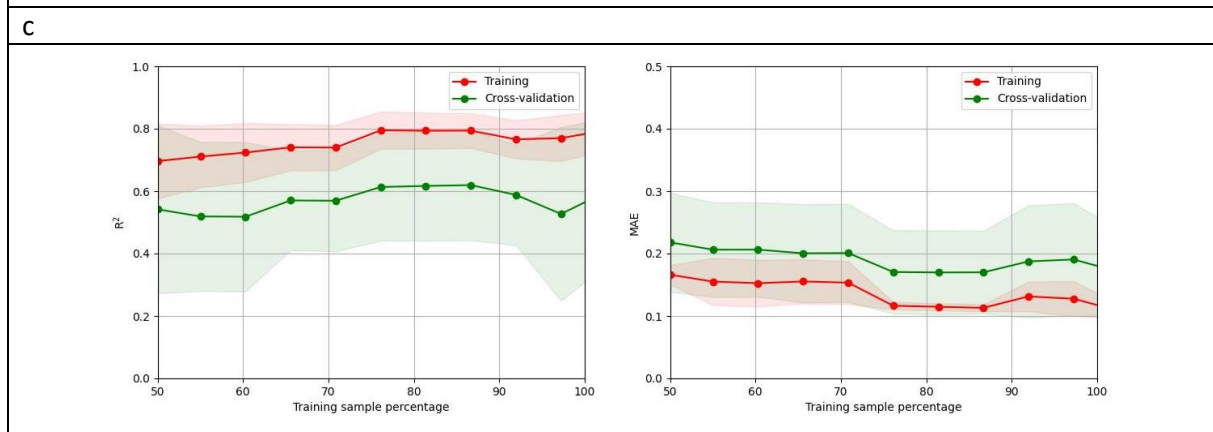
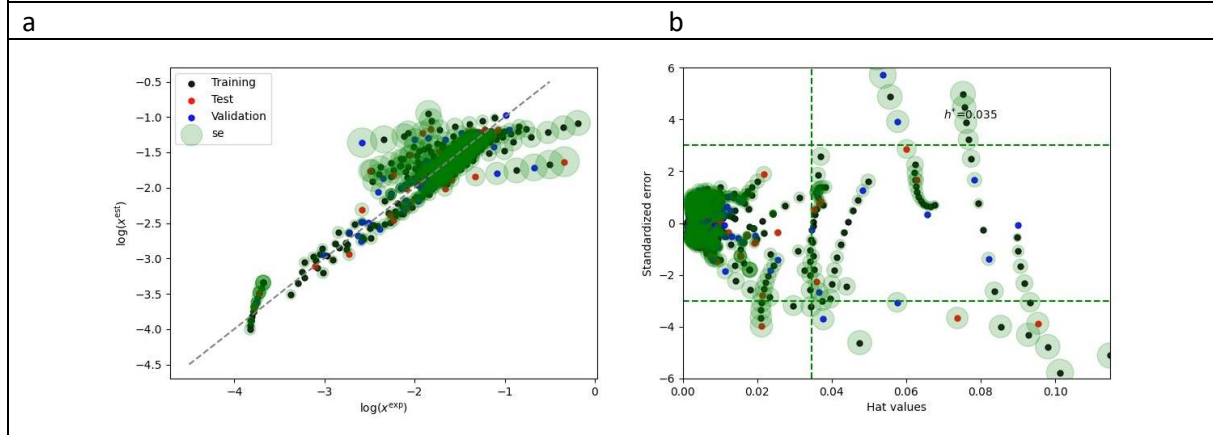
b



c



OrthogonalMatchingPursuitCV



TransformedTargetRegressor

