

Table S-I. Molecular properties predictors calculated by the MolSoft platform [52, 91].

| Comp. | TPSA (Å ²) ^a | Volume (Å ³) ^b | LogP | LogS (in mg/L) | Number of HBA | Number of HBD | %ABS =109-(0.345 x TPSA) ^c | Drug-Likeness Model Score (DLS) | BBB Score |
|-------|-------------------------------------|---------------------------------------|------------|----------------|---------------|---------------|---------------------------------------|---------------------------------|-----------|
| 1a | 24.31 | 403.94 | 5.80 (> 5) | 0.07 | 2 | 1 | 100.61 | -1.36 | 4.64 |
| 1b | 26.89 | 302.71 | 2.68 | 238.66 | 4 | 1 | 99.72 | -0.72 | 5.54 |
| 1c | 44.60 | 382.88 | 3.87 | 17.56 | 2 | 3 | 93.1 | -1.64 | 4.11 |
| 1d | 44.42 | 379.48 | 3.48 | 61.76 | 2 | 3 | 93.8 | -1.42 | 4.07 |
| 1e | 39.31 | 559.95 | 7.80 (> 5) | 0.42 | 4 | 1 | 95.44 | -1.22 | 3.93 |
| 1f | 68.44 | 271.94 | 0.06 | 6010.37 | 4 | 3 | 85.39 | -0.69 | 3.53 |
| 1g | 24.86 | 306.30 | 3.48 | 53.22 | 2 | 1 | 100.42 | -1.28 | 5.57 |
| 1h | 26.89 | 336.64 | 3.94 | 19.86 | 4 | 1 | 99.72 | -1.05 | 5.53 |
| 1i | 26.89 | 350.95 | 4.07 | 23.47 | 4 | 1 | 99.72 | -0.74 | 5.55 |
| 1j | 24.86 | 318.13 | 3.61 | 36.11 | 2 | 1 | 100.42 | -1.34 | 5.57 |
| 2a | 28.62 | 151.60 | 2.22 | 522.96 | 2 | 1 | 99.13 | -1.17 | 4.26 |
| 2b | 28.35 | 200.43 | 3.19 | 68.87 | 2 | 1 | 99.22 | -1.34 | 4.86 |
| 2c | 61.72 | 174.88 | 1.27 | 3541.88 | 4 | 3 | 87.7 | -0.35 | 2.80 |
| 3a | 28.29 | 550.36 | 7.93 (> 5) | 0.13 | 2 | 0 | 99.24 | -0.75 | 3.43 |
| 3b | 28.56 | 501.54 | 6.57 (> 5) | 0.12 | 2 | 0 | 99.15 | -0.75 | 3.81 |
| 3c | 28.01 | 599.18 | 8.89 (> 5) | 0.10 | 2 | 0 | 99.34 | -0.79 | 3.27 |
| 3d | 28.83 | 464.54 | 5.74 (> 5) | 0.60 | 2 | 0 | 99.05 | -0.56 | 4.40 |
| 3e | 30.59 | 497.95 | 5.77 (> 5) | 0.23 | 4 | 0 | 98.45 | -0.55 | 3.87 |
| 3f | 28.56 | 513.37 | 6.70 (> 5) | 0.12 | 2 | 0 | 99.15 | -0.35 | 3.72 |
| 3g | 28.83 | 452.72 | 5.61 (> 5) | 0.62 | 2 | 0 | 99.05 | -0.76 | 4.49 |
| 3h | 61.38 | 573.63 | 6.97 (> 5) | 0.18 | 4 | 2 | 87.82 | 0.08 | 2.45 |
| 1k | 12.86 | 394.39 | 6.86 (> 5) | 0.03 | 1 | 0 | 104.56 | -0.90 | 3.98 |
| 1l | 21.94 | 299.34 | 3.91 | 14.25 | 2 | 0 | 101.43 | -1.31 | 5.17 |
| 1m | 21.40 | 396.98 | 6.23 (> 5) | 0.05 | 2 | 0 | 101.62 | -1.39 | 4.10 |
| 1n | 12.49 | 413.39 | 7.22 (> 5) | 0.04 | 2 | 0 | 104.69 | -1.36 | 4.22 |

| | | | | | | | | | |
|------------------------|-------|--------|------------|--------|---|---|--------|-------|------|
| 1o | 11.71 | 313.93 | 4.18 | 11.10 | 2 | 1 | 104.96 | -1.38 | 5.74 |
| 1p | 11.17 | 411.57 | 6.50 (>5) | 0.03 | 2 | 1 | 105.15 | -1.47 | 4.81 |
| 1q | 39.78 | 372.71 | 5.24 (> 5) | 0.22 | 4 | 1 | 95.8 | -1.23 | 4.12 |
| Curcumin (Keto) | 73.83 | 393.60 | 2.83 | 428.21 | 6 | 2 | 83.53 | -0.82 | 2.83 |
| Curcumin (Enol) | 77.11 | 401.76 | 3.29 | 380.63 | 6 | 3 | 82.4 | -0.61 | 2.78 |

^aTopological polar surface area, ^bMolecular volume, ^cPercentage absorption

Number of HBA:Number of hydrogen bond acceptors (n-ON)

Number of HBD:Number of hydrogen bond donors (n-OH/NH)

Data were calculated by the **MolSoftplatform** (<http://molsoft.com/mprop/>). The percentage of absorption (%ABS) was calculated by using $\%ABS = 109 - (0,345 \times TPSA)$ and is referred to the degree of absorption.

All molecular property predictors are calculated using fragment-based contributions.

LogP (octanol/water partition coefficient)

LogS (water solubility)

Molecular Polar Surface Area (PSA) and Volume

PSA is defined as sum of surfaces of oxygens, nitrogens and attached hydrogens.