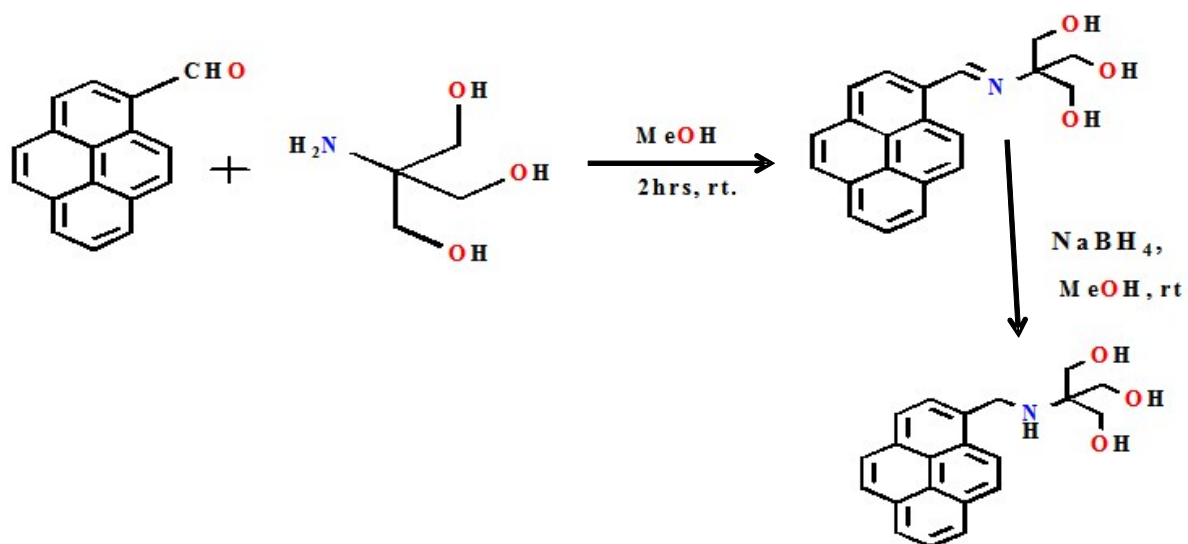


**Interaction of a Synthesized Pyrene Based Fluorescent Probe with CT-DNA:  
Spectroscopic, Thermodynamic and Molecular modeling studies**

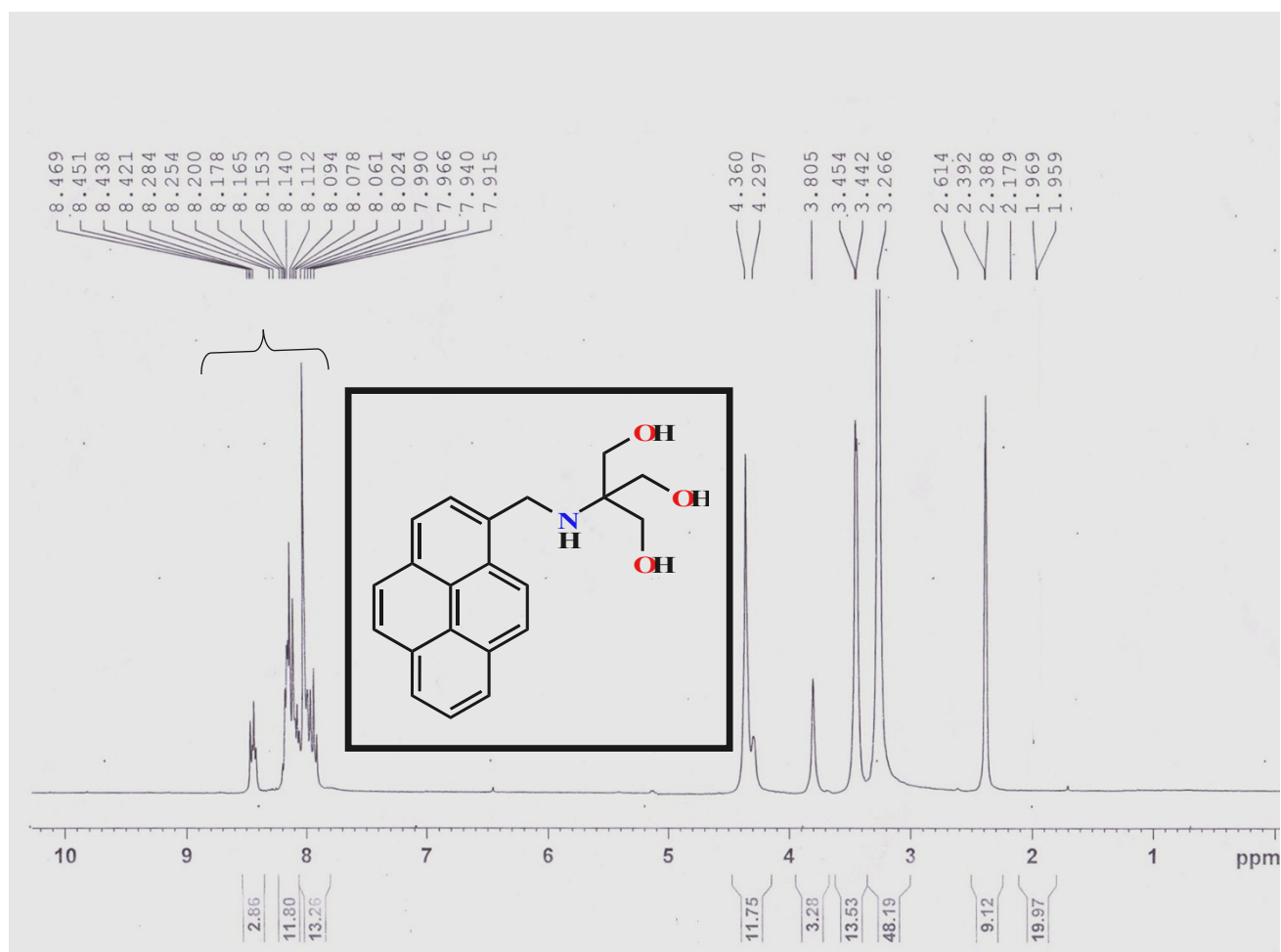
Soumen Ghosh<sup>a</sup>, Abdulla Al Masum<sup>b</sup>, Aniruddha Ganguly<sup>a</sup>, Md. Akhtarul Alam<sup>b</sup>, Md. Maidul Islam<sup>b\*</sup> and Nikhil Guchhait<sup>a\*</sup>

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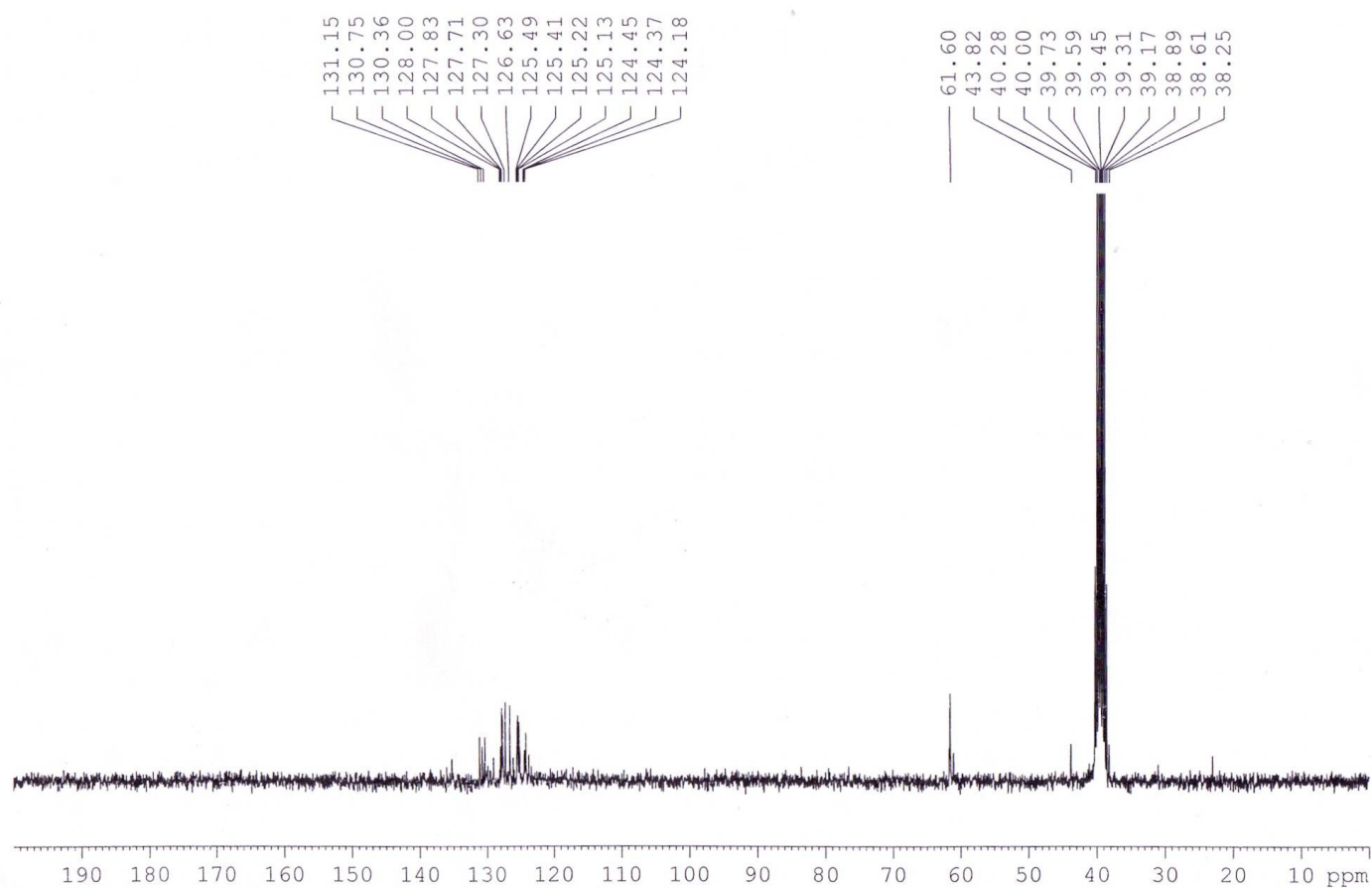
<sup>b</sup>*Department of Chemistry, Aliah University, Action Area IIA/27, Kolkata 700156.*



**Scheme S1.** Synthetic route of compound 1.



**Figure S1.** <sup>1</sup>H NMR (300 MHz) spectrum of **1** in DMSO-d<sub>6</sub> at 20 °C



**Figure S2.** <sup>13</sup>C NMR (75.5 MHz) spectrum of **1** in *d*<sub>6</sub>-DMSO at 20 °C

**Table S1. Four DNA decamer sequences**

| S.No. | DNA | Sequence                                |
|-------|-----|-----------------------------------------|
| 1     | S1  | 5'-d(GAT <u>GGCC</u> ATC) <sub>2</sub>  |
| 2     | S2  | 5'-d(GAT <u>CCGG</u> ATC) <sub>2</sub>  |
| 3     | S3  | 5'-d(GGCA <u>AATT</u> GCC) <sub>2</sub> |
| 4     | S4  | 5'-d(GGCT <u>TAAG</u> CC) <sub>2</sub>  |

| S1-pytris docking table |           |           |            |            |
|-------------------------|-----------|-----------|------------|------------|
| S                       | rmsd      | E_conf    | E_place    | E_score1   |
| -6.4546618              | 7.179431  | 0         | -668.33075 | -6.4546618 |
| -6.9276376              | 7.1241064 | 1.8063836 | -674.39392 | -6.9276376 |
| -7.7952886              | 7.105413  | 0         | -669.61407 | -7.7952886 |
| -8.4600554              | 7.0959425 | 2.0509024 | -679.12665 | -8.4600554 |
| -6.4867949              | 7.0909152 | 0         | -670.79437 | -6.4867949 |
| -6.3626819              | 7.0853705 | 0         | -672.44769 | -6.3626819 |
| -6.3328075              | 7.0820994 | 0         | -669.20404 | -6.3328075 |
| -7.3279901              | 7.0621223 | 0         | -665.3371  | -7.3279901 |
| -6.3158188              | 7.0573668 | 0         | -671.45886 | -6.3158188 |
| -7.9114366              | 7.0262923 | 0         | -672.1687  | -7.9114366 |
| -8.5631628              | 7.0082469 | 2.0513055 | -679.55054 | -8.5631628 |
| -7.6667867              | 6.9982057 | 0         | -671.11646 | -7.6667867 |
| -6.3457031              | 6.9916921 | 0         | -671.67682 | -6.3457031 |
| -8.8124399              | 6.9690537 | 2.2563903 | -687.34387 | -8.8124399 |
| -8.4452066              | 6.9598751 | 2.2561491 | -686.63971 | -8.4452066 |
| -8.7948046              | 6.9373326 | 1.998136  | -678.63818 | -8.7948046 |
| -8.9760923              | 2.7277198 | 1.5962512 | -785.32898 | -8.9760923 |
| -9.3244419              | 2.6845901 | 1.59613   | -779.87915 | -9.3244419 |
| -10.555079              | 2.6388922 | 1.5963793 | -778.24445 | -10.555079 |
| -10.474515              | 2.6342528 | 1.5963116 | -778.34509 | -10.474515 |
| -9.4319153              | 2.5617173 | 1.6351231 | -779.85431 | -9.4319153 |

|            |           |           |            |            |
|------------|-----------|-----------|------------|------------|
| -9.9312782 | 2.5442975 | 1.7154114 | -775.35364 | -9.9312782 |
| -8.7853031 | 2.5160537 | 1.9945502 | -767.75281 | -8.7853031 |
| -9.915637  | 2.5110061 | 1.6817863 | -779.30719 | -9.915637  |
| -10.51836  | 2.5022755 | 1.6352916 | -778.41888 | -10.51836  |
| -10.186988 | 2.4944973 | 1.7154398 | -769.52301 | -10.186988 |
| -10.191694 | 2.4622648 | 1.6818099 | -773.41003 | -10.191694 |
| -9.4528351 | 2.432831  | 1.7154876 | -778.23621 | -9.4528351 |
| -9.4138355 | 2.3993895 | 1.6818545 | -782.16907 | -9.4138355 |
| -9.7803926 | 2.3774307 | 2.2563462 | -762.89679 | -9.7803926 |

| S2-pytris docking table |           |           |            |             |
|-------------------------|-----------|-----------|------------|-------------|
| S                       | rmsd      | E_conf    | E_place    | E_score1    |
| -10.034606              | 6.8968883 | 1.6975729 | -622.12256 | -10.034606  |
| -9.7919636              | 6.8823619 | 1.5182332 | -612.32928 | -9.7919636  |
| -10.204964              | 6.8735442 | 1.3769805 | -645.50311 | -10.204964  |
| -9.5849695              | 6.8422637 | 1.7013464 | -610.50824 | -9.5849695  |
| -9.8206396              | 6.7628417 | 1.3769788 | -643.96552 | -9.8206396  |
| -10.059402              | 6.7492332 | 1.7419789 | -658.20728 | -10.059402  |
| -10.583174              | 6.7233791 | 1.6980435 | -658.53894 | -10.583174  |
| -9.9027309              | 6.5937419 | 1.7420311 | -659.10406 | -9.9027309  |
| -10.530219              | 6.5864015 | 1.6980658 | -658.25964 | -10.530219  |
| -1.4828229              | 6.4732137 | 1.5182403 | -587.99823 | -1.4828229  |
| -1.4340451              | 6.3915849 | 1.5181806 | -588.30975 | -1.4340451  |
| -1.6817904              | 6.3449621 | 1.7888666 | -603.5791  | -1.6817904  |
| -2.0954382              | 6.3367147 | 1.7013464 | -586.98895 | -2.0954382  |
| -1.6723944              | 6.2518449 | 1.7888476 | -604.22632 | -1.6723944  |
| -0.85963154             | 6.1890893 | 1.7013148 | -590.99493 | -0.85963154 |
| -13.053338              | 2.2384264 | 0         | -667.19012 | -13.053338  |
| -12.721235              | 2.1680107 | 0         | -665.71368 | -12.721235  |
| -11.335208              | 1.9214787 | 0         | -663.76135 | -11.335208  |
| -10.992325              | 1.8861812 | 0         | -670.3316  | -10.992325  |
| -11.538211              | 1.864049  | 0         | -664.17596 | -11.538211  |
| -10.89166               | 1.8328191 | 0         | -669.31128 | -10.89166   |



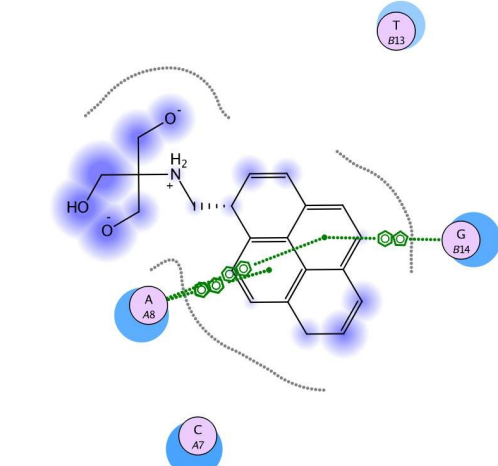
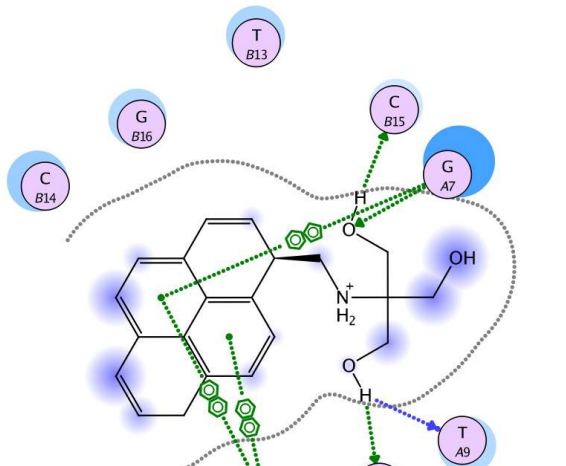
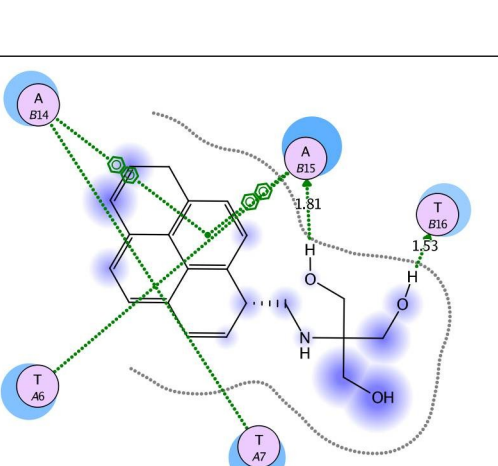
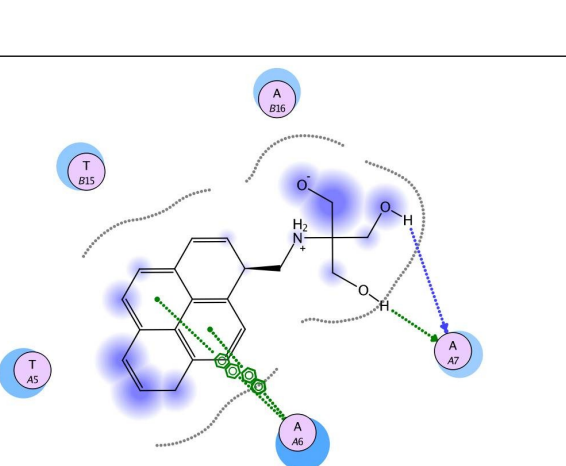
|            |           |   |            |            |
|------------|-----------|---|------------|------------|
| -11.922042 | 1.794799  | 0 | -668.28589 | -11.922042 |
| -11.601603 | 1.7777536 | 0 | -673.83838 | -11.601603 |
| -11.743905 | 1.5099301 | 0 | -671.711   | -11.743905 |
| -11.77538  | 1.5027348 | 0 | -671.58643 | -11.77538  |
| -11.587506 | 1.4267851 | 0 | -661.77747 | -11.587506 |
| -11.458681 | 1.4265821 | 0 | -660.94049 | -11.458681 |
| -11.599536 | 1.4146892 | 0 | -673.94769 | -11.599536 |
| -11.59997  | 1.4130015 | 0 | -670.08167 | -11.59997  |
| -10.963889 | 1.2992067 | 0 | -659.75641 | -10.963889 |

| S3-pytris dockin table |           |             |            |            |
|------------------------|-----------|-------------|------------|------------|
| S                      | rmsd      | E_conf      | E_place    | E_score1   |
| -10.141603             | 6.9137802 | 1.4802929   | -689.4436  | -10.141603 |
| -10.009198             | 6.8776774 | 1.4803308   | -688.50928 | -10.009198 |
| -10.30027              | 6.8583522 | 1.4592804   | -690.04401 | -10.30027  |
| -9.5048351             | 6.8468351 | 1.4802989   | -689.27686 | -9.5048351 |
| -10.197693             | 6.8467264 | 1.4593091   | -688.90265 | -10.197693 |
| -9.6938133             | 6.8164277 | 1.459278    | -689.66986 | -9.6938133 |
| -9.4734163             | 6.6056628 | 0.043115843 | -599.25256 | -9.4734163 |
| -9.3524256             | 6.5624537 | 0           | -594.47003 | -9.3524256 |
| -9.4462204             | 6.5423951 | 0           | -592.16217 | -9.4462204 |
| -9.5140982             | 6.521142  | 0.043113727 | -599.92206 | -9.5140982 |
| -9.8149042             | 6.4616256 | 0           | -594.17664 | -9.8149042 |
| -9.7813759             | 6.4401097 | 0.043112118 | -598.76245 | -9.7813759 |
| -10.877054             | 6.2290454 | 0.15002993  | -687.83417 | -10.877054 |
| -11.144661             | 6.1475258 | 0.15003026  | -687.5954  | -11.144661 |
| -10.676573             | 6.1318712 | 0.17941804  | -691.75067 | -10.676573 |
| -10.932964             | 6.0480628 | 0.17941876  | -691.47571 | -10.932964 |
| -9.775733              | 6.036365  | 0.15003374  | -685.01965 | -9.775733  |
| -11.619782             | 4.1934342 | 0.60000002  | -672.43811 | -11.619782 |
| -9.4616261             | 4.1088805 | 0.60000002  | -672.32452 | -9.4616261 |
| -9.8311424             | 3.448595  | 0.15045162  | -713.87164 | -9.8311424 |
| -9.4071321             | 3.4408824 | 0.15045534  | -713.95142 | -9.4071321 |

|            |           |             |            |            |
|------------|-----------|-------------|------------|------------|
| -9.5048923 | 3.4151387 | 0.17983626  | -713.97797 | -9.5048923 |
| -10.924271 | 3.1644521 | 1.4802989   | -724.12    | -10.924271 |
| -9.3308001 | 3.156287  | 1.4802929   | -724.15997 | -9.3308001 |
| -9.9733047 | 2.0062029 | 0.17983767  | -707.28589 | -9.9733047 |
| -11.489121 | 1.9539614 | 0.15045534  | -707.49316 | -11.489121 |
| -9.8017664 | 1.9114515 | 0           | -670.58667 | -9.8017664 |
| -9.7758608 | 1.8978614 | 0.043115843 | -671.67792 | -9.7758608 |
| -9.6469202 | 1.7481135 | 0           | -671.013   | -9.6469202 |
| -11.684961 | 1.5173985 | 0           | -657.56439 | -11.684961 |

| S4-pytris docking table |           |            |            |            |
|-------------------------|-----------|------------|------------|------------|
| S                       | rmsd      | E_conf     | E_place    | E_score1   |
| -7.3543439              | 6.8449535 | 2.3977594  | -657.85327 | -7.3543439 |
| -12.003634              | 3.3526378 | 1.1591526  | -660.59705 | -12.003634 |
| -10.874158              | 3.3524058 | 1.0730581  | -662.64423 | -10.874158 |
| -10.375343              | 3.3506923 | 0.60000002 | -630.40167 | -10.375343 |
| -10.834942              | 3.3495309 | 1.1591526  | -660.01318 | -10.834942 |
| -11.720356              | 3.3073516 | 1.0730581  | -661.09021 | -11.720356 |
| -10.579964              | 3.305783  | 1.0730581  | -660.51135 | -10.579964 |
| -8.6411629              | 3.2964044 | 1.0052307  | -654.98016 | -8.6411629 |
| -9.2367783              | 3.27966   | 1.2830637  | -662.41882 | -9.2367783 |
| -10.802885              | 3.2787411 | 1.3731635  | -661.63263 | -10.802885 |
| -10.890466              | 3.2746527 | 1.389603   | -703.6015  | -10.890466 |
| -11.897175              | 3.269079  | 1.3731608  | -662.84094 | -11.897175 |
| -10.923158              | 3.2482934 | 1.4475715  | -702.07471 | -10.923158 |
| -11.496552              | 3.239063  | 1.3731623  | -661.72498 | -11.496552 |
| -11.37088               | 3.237813  | 1.3731633  | -659.97534 | -11.37088  |
| -11.86519               | 3.2318439 | 1.389603   | -704.34729 | -11.86519  |
| -11.99263               | 3.2079558 | 1.4475714  | -702.9317  | -11.99263  |
| -10.59781               | 3.207428  | 1.3731633  | -658.87494 | -10.59781  |
| -11.323713              | 3.1977823 | 1.389603   | -702.5542  | -11.323713 |
| -9.0633535              | 3.1944628 | 1.4544754  | -658.95349 | -9.0633535 |
| -10.585618              | 3.1799052 | 1.0052327  | -656.53918 | -10.585618 |

|            |           |           |            |            |
|------------|-----------|-----------|------------|------------|
| -11.866062 | 3.1755123 | 1.4475713 | -701.39337 | -11.866062 |
| -11.360275 | 3.1715798 | 1.4475706 | -701.09546 | -11.360275 |
| -10.136869 | 3.1696568 | 1.2830637 | -668.43518 | -10.136869 |
| -10.083429 | 3.1487181 | 1.45446   | -659.94318 | -10.083429 |
| -12.155    | 3.1273401 | 1.3731612 | -667.68359 | -12.155    |
| -11.586825 | 3.0943685 | 1.3731633 | -664.87238 | -11.586825 |
| -10.735434 | 3.0727437 | 1.45444   | -661.01099 | -10.735434 |
| -12.086841 | 2.7308867 | 2.3977592 | -703.77454 | -12.086841 |
| -11.935977 | 2.7161479 | 2.3977597 | -703.52759 | -11.935977 |

|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |  |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   |
|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
|  <p> <span style="color: blue;">○</span> polar    <span style="color: green;">→</span> sidechain acceptor    <span style="color: grey;">○</span> solvent residue    <span style="color: green;">⊗</span> arene-arene<br/> <span style="color: red;">○</span> acidic    <span style="color: green;">←</span> sidechain donor    <span style="color: grey;">⊗</span> metal complex    <span style="color: green;">⊕</span> arene-cation<br/> <span style="color: blue;">○</span> basic    <span style="color: blue;">→</span> backbone acceptor    <span style="color: grey;">—</span> solvent contact    <span style="color: green;">—</span> metal contact<br/> <span style="color: green;">○</span> greasy    <span style="color: blue;">→</span> backbone donor    <span style="color: grey;">—</span> metal contact<br/> <span style="color: grey;">○</span> proximity    <span style="color: blue;">○</span> ligand exposure    <span style="color: blue;">○</span> receptor contact<br/> <span style="color: grey;">○</span> contour </p> |  |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| S1- pyrene 2D interaction                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |  | S2- pyrene 2D interaction                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |  |  <p> <span style="color: blue;">○</span> polar    <span style="color: green;">→</span> sidechain acceptor    <span style="color: grey;">○</span> solvent residue    <span style="color: green;">⊗</span> arene-arene<br/> <span style="color: red;">○</span> acidic    <span style="color: green;">←</span> sidechain donor    <span style="color: grey;">⊗</span> metal complex    <span style="color: green;">⊕</span> arene-cation<br/> <span style="color: blue;">○</span> basic    <span style="color: blue;">→</span> backbone acceptor    <span style="color: grey;">—</span> solvent contact    <span style="color: green;">—</span> metal contact<br/> <span style="color: green;">○</span> greasy    <span style="color: blue;">→</span> backbone donor    <span style="color: grey;">—</span> metal contact<br/> <span style="color: grey;">○</span> proximity    <span style="color: blue;">○</span> ligand exposure    <span style="color: blue;">○</span> receptor contact<br/> <span style="color: grey;">○</span> contour </p> |
| S3- pyrene 2D interaction                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |  | S4- pyrene 2D interaction                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |