Electronic Supplementary Information

Lysine triggered ratiometric conversion of dynamic to static excimer of a pyrene derivative: aggregation-induced emission, nanomolar detection and human breast cancer cell (MCF7) imaging

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1. Materials and methods

High-purity HEPES, pyrene-1-carbaldehyde and phenylpropanolamine are purchased from Sigma Aldrich (India). Amino acids are purchased from Merck (India). Solvents used are of spectroscopic grade. Other chemicals are of analytical reagent grade and has been used without further purification except when specified. Mili-Q Milipore 18.2 M Ω cm⁻¹ water has been used throughout all the experiments. A Shimadzu Multi Spec 2450 spectrophotometer is used for recording UV-Vis spectra. FTIR spectra are recorded on a Shimadzu FTIR (model IR Prestige 21 CE) spectrophotometer. Mass spectra have been recorded using a QTOF 60 Micro YA 263 mass spectrometer in ES positive mode. ¹H NMR titration in CD₃OD-D₂O has been performed using a Bruker Avance 500 (500 MHz).¹³C NMR spectrum of A3 is recorded using a Bruker Advance 400 (100 MHz). ¹H NMR spectra of A3 with other representative amino acids are recorded using a Bruker Avance 300 (300 MHz). Elemental analyses are performed on a Perkin Elmer 2400 CHN analyzer. The steady state emission and excitation spectra are recorded with a Hitachi F-4500 spectrofluorimeter. Systronics digital pH meter (model 335) is used for measurement of solution pH. Dilute HCl or NaOH (50 μ M) are used for pH adjustment. The imaging system is composed of an inverted fluorescence microscope (Dewinter, Italy). Images are captured through an attached CCD camera equipped with BioWizard 4.2 software. The microscope is equipped with a 50 W mercury arc lamp. All spectra are recorded at room temperature.

2. General method of UV-Vis and fluorescence titration

Path length of the cells used for absorption and emission studies is 1 cm. Stock solutions of A3 and Lys are prepared in CH₃OH/ water (4/1, v/v). Working solutions of A3 and Lys are prepared from their respective stock solutions. Fluorescence measurements are performed using 2.5 nm x 2.5 nm slit width.

3. Determination of binding constant

The binding constant of A3 for Lys is determined using a modified Benesi–Hildebrand equation¹: $(I_{max} - I_0)/(I_x - I_0) = 1 + (1/K) (1/[C]^n)$ where I_{max} , I_0 and I_x are emission intensity values for A3 in the presence of Lys at saturation, in the absence of Lys and at any intermediate Lys concentrations, respectively. A plot of $(I_{max} - I_0)/(I_x - I_0) = 1 + (1/K) (1/[C]^n)$ (here n = 0.5) yields the binding constant value from the slope.

4. In vitro cell imaging

Human breast cancer cell line MCF7 is grown in DMEM (Sigma, St. Louis, USA) supplemented with 10% fetal bovine serum (Sigma, St. Louis, USA), 2 mM glutamine, 100 U m/L penicillin-streptomycin solution (Gibco, Invitrogen, USA) in the presence of 5% CO₂ at 37 °C. For *in vitro* imaging studies, cells are seeded in 6 well culture plates with a seeding density of 10^5 cells per well. After reaching 60–70% confluence, the previous media is replaced with serum free media, supplemented by Lys and A3 at 50 µM and 20 µM, and incubated for 2 h to facilitate the Lys or A3 uptake by the cells. Then the cells were placed under an inverted microscope (Dewinter, Italy) at different magnifications to examine any adverse effect on cellular morphology.A3 treated cells are then incubated with Lys for 15–30 min and observed under an inverted fluorescence microscope at different magnifications with a blue filter.

Synthesis of (E)-2-((pyren-3-yl)methyleneamino)-1-phenylpropan-1-ol (A3)

A methanol solution of pyrene 1-carboxaldehyde (230 mg, 1.0 mmol) is added drop wise under stirring condition to a methanol solution of 2-amino-1-phenylpropan-1-ol hydrochloride (187.6 mg, 1.0 mmol) in presence of NaHCO₃ (Scheme 1). Stirring is continued for further 30 min followed by reflux for 8 h. Upon removal of excess NaHCO₃ through filtration, an orange solution is obtained which on slow evaporation gives an orange residue with a yield of 93%. The residue is then recrystallized to get orange crystals suitable for SC-XRD analysis. ESI-MS (m/z) (Fig.S1, ESI): $[M + H]^+$ 364.2, $[M + Na]^+$ 386.2. ¹H NMR (CD₃OD, 500 MHz) (Fig.S2, ESI): δ 1.56 (d, J= 5 Hz, 3H), δ 3.85 (m, J= 5 Hz, 1H), 4.77 (t, 1H), 7.20 (t, 1H, benzene), 7.31 (t, J = 5 Hz, 2H, benzene), 7.42 (d, J = 10 Hz, 2H, benzene), 8.04 (m, J = 5 Hz, 3H, pyrene), 8.12 (m, J = 5 Hz, 3H, pyrene), 8.21 (m, J = 5 Hz, 3H, pyrene), 9.10 (s, 1H, CH=N) ppm. ¹³C NMR (CDCl₃, 400 MHz) (Fig. S3, ESI): δ 16.48, 71.99, 77.26, 122.30-132.72, 141.16, 159.28 ppm. Elemental analysis for C₂₆H₂₁NO (%): calculated, C 85.92, H 5.82, N 3.85, O 4.40; found C 85.84, H 5.91, N 3.90, O 4.32.











Fig. S2a Aromatic region of the ¹H NMR spectrum of A3







Fig. S4 FTIR spectrum of A3



Fig. S5 Crystal packing (i), molecular structure (ii) and H-bonding network (iii) of A3



Fig. S6 (i) Emission spectra and (ii) UV light exposed colors of A3 (10 μ M) in absence and presence of Lys (20 μ M) in different solvents, $\lambda_{ex} = 360$ nm



Fig.S7 Effect of pH on the emission intensities of A3 (10 μ M) before and after addition of Lys (2 equiv.), $\lambda_{ex} = 360$ nm



Fig.S8 Emission spectra of A3 (10 μ M) in presence of different amino acids (20 μ M) in HEPES buffered (0.1 M; MeOH: H₂O, 4:1 v/v, pH 7.4; $\lambda_{ex} = 360$ nm) solution



Fig. S9 Changes in $I_{505 nm} / I_{404 nm}$ value of A3 (10 µM) in presence of different amino acids (20 µM) in HEPES buffered (0.1 M; MeOH: H₂O, 4:1 v/v; pH 7.4; λ_{ex} = 360 nm) solution



Fig S10 UV light exposed colors of A3 (10 μ M) in presence of different amino acids (20 μ M) in HEPES buffered (0.1 M; MeOH: H₂O, 4:1 v/v; pH 7.4)



Fig S11 Benesi-Hildebrand plot for determination of the association constant of A3 with Lys



Fig S12 Emission intensities of A3 (10 μ M) as a function of externally added [Lys] in HEPES buffered (0.1 M; MeOH: H₂O, 4:1 v/v; pH 7.4; $\lambda_{ex} = 360$ nm, $\lambda_{em} = 505$ nm) solution. The data obtained from fluorescence titration in Fig.2



Fig S13 Changes in $I_{505 nm} / I_{404 nm}$ of A3 (10 μ M, $\lambda_{ex} = 360$ nm) as a function of externally added Lys in HEPES buffered (0.1 M; MeOH: H₂O, 4:1, v/v; pH 7.4) solution



Fig S14 Linear portion of the $I_{505 nm} / I_{404 nm}$ of A3 (10 μ M, $\lambda_{ex} = 360$ nm) as a function of externally added Lys in HEPES buffered (0.1 M; MeOH: H₂O, 4:1, v/v; pH 7.4)



Fig S15 Normalized excitation spectra of A3 (10 μ M) (left) and A3 (10 μ M) + 2 equiv. Lys (right).



Fig S15a Normalized excitation spectra of A3 (10 μ M) + 5 equiv. Lys.



Fig S16 The ratio I_{505nm} / I_{404nm} for [A3 + Lys] system in presence of other amino acids



Fig S17a Changes in the emission spectra of A3 (10 μ M, CH₃OH) with varying water percentage



Fig S17b Changes in the absorption spectra of A3 (10 μ M, CH₃OH) with varying water percentage



Fig S18a Changes in the absorption spectra of A3 (30 μ M) after addition of Lys (1800 μ M) in MeOH: H₂O, 4:1, v/v (inset show the visible colors of the respective solutions)



Fig S18b Changes in the emission spectra of A3 (30 μ M) after addition of Lys (1800 μ M) in MeOH: H₂O, 4:1, v/v (inset shows the UV light exposed colors of the respective solutions)



Fig S19 DLS of A3 (10 μ M) with increasing Lys content (3.0 equiv., 5.0 equiv. and 10.0 equiv.)



Fig S20 ¹H NMR spectral changes of A3 (CD₃OD) upon gradual addition of Lys (D₂O)



Fig S20a Aromatic region of the 1 H NMR spectra of A3 (CD₃OD) upon addition of 2.0 equiv. other amino acids (D₂O

| Empirical formula | C ₂₆ H ₂₁ NO |
|---|--|
| Formula weight | 363.44g/mol |
| Temperature | 296(2) K |
| Wavelength | 0.71073 Å |
| Crystal system | Monoclinic |
| Space group | Plcl |
| Unit cell dimensions | a = 28.0546(17) Å; b = 9.6810(5) Å; c = 15.0716(8) Å; |
| | $\alpha = 90^{\circ}, \beta = 109.781(6)^{\circ}, \gamma = 90^{\circ}$ |
| Volume | 3851.9(4) Å ³ |
| Ζ | 8 |
| Density (calculated) | 1.253 g/cm^3 |
| Absorption coefficient | 0.076 mm ⁻¹ |
| F(000) | 1536 |
| Theta range for data collection | 2.52 to 28.48° |
| Index ranges | -37<=h<=37, -12<=k<=12, -20<=l<=19 |
| Reflections collected | 35240 |
| Independent reflections | 4832 [R(int) = 0.0775] |
| Completeness to theta = 22.21° | 99.4% |
| Refinement method | Full-matrix least-squares on F ² |
| Data / restraints / parameters | 4832 / 0 / 255 |
| Goodness-of-fit on F ² | 1.006 |
| Final R indices [I>2sigma(I)] | R1 = 0.0489, wR2 = 0.1161 |
| R indices (all data) | R1 = 0.1117, wR2 = 0.1454 |
| Largest diff. peak and hole | 0.185 and -0.198 eÅ ⁻³ |
| R.M.S. deviation from mean | 0.046 eÅ ⁻³ |

Table S1.Crystal data and structure refinement for A3 (CCDC No. 1007420)

References

1. H. A. Benesi and J. H. Hildebrand, J. Am. Chem. Soc., 1949, 71, 2703-2707.