Catalyst-Free Facile Synthesis of Polycyclic Indole/Pyrrole Substituted-1,2,3-triazoles

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ESI/MS experiments:

ESI/MS experiments were performed to gain evidence for the possible intermediates in the proposed mechanism. The experiment was conducted under the optimal reaction condition and the reaction was monitored after 1.5 h. The ESI/MS analyses showed a peak of intermediates, which was identified as a triazoline species. We tried to isolate and characterize the intermediate. We could not isolate dihydrotirazole, only we were able to isolate 1u' under optimized reaction condition. We characterized and included NMR spectrum of compound 1u'.





Figure S1: ESI/MS spectra



Determination of rate limiting step

The reaction of **1u** (0.1 mmol) and sodium azide (0.13 mmol) was carried out in NMR tube using DMSO- d_6 as a solvent. The reactions were conducted for the specified time (such as 15 min, 30 min, 1 h, 2 h, 3 h, 6 h, 10 h and 20 h), then the progress of the reaction was monitored by ¹H NMR spectroscopy. Conversion of **1u** to **1u'** (replacement of bromo to azide) which is giving the corresponding cyclized product (**2u**) was estimated by the relative integration of aliphatic proton(**1u**: 4.82,3.82, **1u'**: 4.59, 3.68 and **2u**: 5.04, 4.69) and α , β -unsaturated ketone protons. The results were listed in figure S2. We observed that the intramolecular azide-alkene cycloaddition is slower than bromo to azide formation (**1u** to **1u'**). Based on the result of figure S2, it may be concluded that the intramolecular azide-alkene cycloaddition step is the rate limiting step.



Figure S2: Progress of the reaction (1u-2u) monitored by ¹H NMR (500 MHz, DMSO- d_6).

Table S1: Screening of solvents^a



^{*a*}Reaction conditions: **1a** (0.5 mmol) and NaN₃ (0.6 mmol), solvent (5 mL) and the reaction was performed at room temp for 20 h under air atmosphere. ^{*b*}Yield of the products. Reaction was performed at room temp for 36 h under air atmosphere

Materials and Method:

Cell Lines

Multiple human cancer cell lines including HT-29 (colorectal adenocarcinoma), A549 (lung carcinoma), PC-3 (prostate adenocarcinoma), MIA PaCa-2 (Pancreatic Carcinoma) and MCF-7 (breast ductal carcinoma) were obtained from ATCC, USA. Early passage cells were resuscitated from liquid nitrogen vapour stocks as needed and cultured according to the manufacturer's instructions. Cells were routinely inspected microscopically for stable phenotype, and all experiments were performed within early passages (within 10) of individual cells.

Evaluation of *in-vitro* anticancer activity of synthetic compounds by Sulphorhodamine B (SRB) calorimetric assay

Efficacy of the compounds as anticancer agents on various cancer cell lines was assessed using standard Sulphorhodamine B (SRB) calorimetric assay for cytotoxicity screening as described by before.^{1,2} Briefly, appropriate number of cells in 5% FBS containing medium were seeded in 96 well plates and incubated at 37 °C in a humidified incubator with 5% CO₂ until the cells attached completely. The cells were treated with a single dose of compounds and were incubated for further 48 hours. After an incubation period, cell monolayers were fixed with 10% (wt/vol) trichloroacetic acid followed by washing under slow running tap water. The cells were stained with SRB dye for 30 minutes, after which the excess dye was removed by washing 3-4 times with 1% (vol/vol) acetic acid. The protein-bound dye is dissolved in 10 mM Tris base solution for OD determination at 510 nm using a microplate reader (Epoch Microplate Reader, Biotek, USA). Percentage of cell growth inhibition was calculated using the formula [100-(Absorbance of compound treated cells/ Absorbance of control cells)] X 100.²

References

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Table S2: Percentage Growth inhibition of the tested compounds against human cancer cell line at 10 μ M.

	HT-29	MCF-7	A549 (Lung)	MIA PaCa-2	PC-3
	(Colon)	(Breast)		(Pancreatic)	(Prostate)
Compound	Average \pm SE	Average \pm SE	Average ± SE	Average ± SE	Average \pm SE
2a	9.95± 6.86	21.2±3.06	8.55±5.79	10.81±0.16	9.47±0.00
2b	15.39±8.9	25.43±0.95	17.22±6.83	14.96±0.03	16.62±2.10
2c	26.91±7.07	24.89±1.33	19.97±5.99	24.84±2.33	13.27±1.38
2d	22.75±4.62	30.39±0.60	14.61±7.15	16.16±0.86	16.78±3.80
2e	7.64±3.87	20.47±0.93	9.69±8.72	0.39±2.43	8.70±0.92
2f	40.1±5.45	33.29±1.11	22.79±10.67	30.28±2.17	16.05±4.25
2g	5.71±6.59	19±5.68	8.41±10.22	20.71±12.23	18.89±4.31
2h	15.38±6.20	20.29±4.73	10.24±10.35	17.33±1.53	14.95±3.05
2i	-3.18±8.45	33.43±5.47	5.3±1.38	5.27±0.60	2.27±3.05
2j	8.1±3.24	26.31±7.16	10.86±3.18	19.87±1.39	18.53±1.47
2k	17.13±1.07	32.05±6.15	18.04±2.44	21.09±0.75	7.0 <u>5±0.91</u>
2m	7.75±1.18	23.07±0.92	13.12±3.60	7.91±1.00	3.07±0.71
2n	38.51±4.75	47.25±0.40	40.78±3.74	36.42±1.87	37.45±1.00

20	0.57±3.50	27.79±2.00	8.54±5.04	7.17±0.08	8.36±3.97
2p	3.19±4.41	27.85±6.22	16.68±10.04	7.90±4.42	11.17±0.33
2q	10.5±4.67	34.58±2.81	22.32±3.98	19.90±6.80	29.27±5.74
2r	1.59±2.70	27.4±0.65	13.27±10.46	19.97±10.22	19.67±3.82
2s	19.67±0.56	23.81±2.53	14.41±11.12	18.73±1.95	17.34±2.75
2t	-3.9±3.48	45.48±6.01	36.54±1.62	37.80±1.83	39.64±1.00
2u	-9.42±5.33	33.07±5.13	14.11±2.52	15.41±1.19	6.30±1.92
2v	-8.57±0.31	23.68±7.18	10.57±0.34	5.10±4.53	-3.20±0.51
2x	34.57±14.5	40.85±6.76	38.79±5.19	27.89±17.75	29.99±1.54
4a	21.25±0.59	39.36±7.39	29.74±2.51	14.62±4.52	20.19±0.47
4b	-13.93±3.28	18.81±2.03	12.68±1.92	26.99±13.70	-8.01±0.21
4c	NT	NT	NT	NT	
4d	-10.4±5.49	33.95±6.31	3.02±1.52	11.92±8.50	-7.34±3.90
4f	8.3 <u>2</u> ±7.39	35. <u>8</u> ±4.17	18.8±6.40	21.66±13.92	6.31±0.05
4g	-1.76±3.06	28.37±6.48	13.18±1.24	5.35±0.70	3.37±3.20
4h	-9.07±4.34	18.91±5.65	10.37±1.95	16.71±3.20	-1.61±2.43

NT = Not Tested

Antibiotic susceptibility testing against ESKAP pathogen panel

Antibiotic susceptibility testing was carried out on the newly synthesized compounds by determining the Minimum Inhibitory Concentration (MIC) according to the standard CLSI guidelines.^{3,4} MIC is defined as the minimum concentration of compound at which visible bacterial growth is inhibited. Bacterial cultures were grown in Mueller-Hinton cation supplemented broth (CA-MHB). Optical density (OD₆₀₀) of the cultures was measured, followed by dilution for ~10⁶ CFU/mL. This inoculum was added into a series of test wells in a microtitre plate that contained various concentrations of compound under test ranging from 64-0.03 μ g/mL. Controls i.e., cells alone and media alone (without compound+cells) and

Levofloxacin used as a reference standard. Plates were incubated at 37 °C for 16-18 h followed by observations of MIC values by the absence or presence of visible growth. For each compound, MIC determinations were performed independently thrice using duplicate samples each time.

References

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	Gram +ve	Gram -ve			
Compound	S. aureus ATCC 29213	E. coli ATCC 25922	K. pneumoni ae BAA- 1705	<i>A. baumannii</i> BAA 1605	P. aeruginosa ATCC 27853
2a	>64	>64	>64	>64	>64
2b	64	>64	>64	>64	>64
2c	>64	>64	>64	>64	>64
2d	>64	>64	>64	>64	>64

Table S3: MIC (μ g/mL) of indolo- and pyrrolo[1,4]diazepines/pyrazines against ESKAP pathogen panel

2e	>64	>64	>64	>64	>64
2f	>64	>64	>64	>64	>64
2g	16	>64	>64	>64	>64
2h	>64	>64	>64	>64	>64
2i	>64	>64	>64	>64	>64
2j	>64	>64	>64	>64	>64
2k	8	>64	>64	>64	>64
2m	>64	>64	>64	>64	>64
2n	8	>64	>64	>64	>64
20	>64	>64	>64	>64	>64
2p	>64	>64	>64	>64	>64
2q	>64	>64	>64	>64	>64
2r	>64	>64	>64	>64	>64
2s	32	>64	>64	>64	>64
2t	>64	>64	>64	>64	>64
2u	>64	>64	>64	>64	>64
2 v	>64	>64	>64	>64	>64
2x	NT	NT	NT	NT	NT
4a	>64	>64	>64	>64	>64
4 b	NT	NT	NT	NT	NT
4 c	>64	>64	>64	>64	>64
4d	>64	>64	>64	>64	>64
4 e	NT	NT	NT	NT	NT
4f	>64	>64	>64	>64	>64
4g	>64	>64	>64	>64	>64
4h	>64	>64	>64	>64	>64
Levofloxacin	<0.5	<0.5	64	8	1

NT = Not Tested

X-ray crystallography information and data



Figure S3. The molecular structure of KA555 (compound **2j**) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radius. **CCDC 1887445** contains the supplementary crystallographic data.

Sample preparation: Compound **2j** was dissolved in ethanol (100 mg in 3 mL ethanol) and allowed for crystallization at room temperature for 7-8 days.

Table S4: Crystal data and structure refinement.

Identification code	KA555
Chemical formula	$C_{22}H_{19}N_5O_2$
Molecular weight	385.42
Temperature	293(2)K
Wavelength	0.71073Å
Crystal system ; space group	monoclinic; P 21/c
Unit cell dimensions	$A = 14.4652(12)\text{\AA}$ $b = 15.665(3)\text{\AA}$ $c = 8.435(2)\text{\AA}$ $\alpha = 90^{\circ}$ $\beta = 91.931(7)^{\circ}$ $\gamma = 90^{\circ}$
Volume	1910.2(6) Å ³
Z, Calculated density	4, 1.340 g/cm ³
Absorption coefficient	0.090 1/mm
F(000)	808
Theta range for data collection	2.600° to 27.500°
Limiting indices	$-18 \le h \le 18$; $-20 \le k \le 20$; $-10 \le l \le 10$

Reflection collected / unique 34927 / 4384 [R(int) = 0.0710]99.9 % Completeness to theta max Refinement method Full-matrix least-square on F² 4384 / 72 / 296 Data / restraints / parameters Goodness of fit on F² 1.172 Final R indices [I>2sigma(I)] R1 = 0.0533; wR2 = 0.1497R1 = 0.0772; wR2 = 0.1703Final R indices [all data] 0.376 and -0.251 e/Å³ Largest diff peak and hole

Data collection and structure solution of KA555 (2j): Single crystal X-ray data for two compounds were collected at room temperature on a Bruker D8 QUEST equipped with a fourcircle kappa diffractometer and Photon 100 detector. An Ius microfocus Mo source (l=0.71073Å) supplied the multi-mirror monochromated incident beam. A combination of Phi and Omega scans were used to collect the necessary data. Unit cell dimensions were determined using 9898 reflections. Integration and scaling of intensity data were accomplished using SAINT program.¹ The structures were solved by Direct Methods using SHELXS97² and refinement was carried out by full-matrix least-squares technique using SHELXL-2014/7.²⁻³ Anisotropic displacement parameters were included for all non-hydrogen atoms. All H atoms were positioned geometrically and treated as riding on their parent C atoms with C-H distances of 0.93--0.97 Å, and with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}$ for methyl atoms. CCDC 1887445 contains the supplementary crystallographic data for this paper which can be obtained free of https://summary.ccdc.cam.ac.uk/structure-summary-form charge or from the at Cambridge Crystallographic Data Centre (CCDC), 12 Union Road, Cambridge CB2 1EZ, UK; fax: +44(0) 1223 336 033; email: deposit@ccdc.cam.ac.uk.

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Scheme S1. Gram-scale reaction.



¹³C NMR (125 MHz, DMSO- d_6) spectrum of compound A











¹³C NMR (125 MHz, DMSO- d_6) spectrum of compound **D**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1a**









¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1d**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 1e



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1f**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1g**







¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1i**



¹³C NMR (125 MHz, DMSO- d_6) spectrum of compound **1**j



 ^{13}C NMR (125 MHz, DMSO- d_6) spectrum of compound 1k



¹³C NMR (125 MHz, DMSO- d_6) spectrum of compound **1m**



 13 C NMR (125 MHz, DMSO- d_6) spectrum of compound **1n**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **10**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1p**



 13 C NMR (125 MHz, DMSO- d_6) spectrum of compound 1q



 13 C NMR (125 MHz, DMSO- d_6) spectrum of compound 1r



¹³C NMR (125 MHz, DMSO-d₆) spectrum of compound 1s


 13 C NMR (125 MHz, DMSO- d_6) spectrum of compound 1t



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1u**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1v**



 13 C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1w**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **1**x



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **3a**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **3b**



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **3c**





¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **3e**



 13 C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **3f**



 ^{13}C NMR (125 MHz, DMSO- d_6) spectrum of compound 3g



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **3h**











¹³C NMR (125 MHz, CDCl₃) spectrum of compound 2c



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **2d**









 ^{13}C NMR (125 MHz, CDCl₃) spectrum of compound 2g







¹³C NMR (125 MHz, CDCl₃) spectrum of compound **2i**





¹³C NMR (125 MHz, CDCl₃) spectrum of compound 2k











¹³C NMR (125 MHz, CDCl₃) spectrum of compound 20



¹³C NMR (125 MHz, CDCl₃) spectrum of compound **2p**



¹³C NMR (125 MHz, CDCl₃ + DMSO- d_6)) spectrum of compound **2q**



 ^{13}C NMR (125 MHz, CDCl₃) spectrum of compound 2r



¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **2s**



¹³C NMR (125 MHz, CDCl₃ + DMSO- d_6) spectrum of compound **2t**







¹³C NMR (125 MHz, DMSO- d_6) spectrum of compound 2v





¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound **4a**


¹³C NMR (125 MHz, CDCl₃) spectrum of compound **4b**









¹³C NMR (125 MHz, CDCl₃) spectrum of compound 4e



 ^{13}C NMR (125 MHz, CDCl₃) spectrum of compound **4f**





¹³C NMR (125 MHz, CDCl₃) spectrum of compound **4h**