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Electronic Supplementary Information (ESI)

Transition-metal-free Synthesis of 1-Pyrroline Derivatives via

Cyclization of Terminal Alkynes with 2-Azaallyls

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Supplementary Methods:

Preparation of ketimines

Ketimines (2a-2l) were prepared according to literature procedures.¹

Procedure and characterization for the cyclization of ketimines and alkynes Reaction Optimization:

An oven-dried 8 mL reaction vial equipped with a stir bar was charged with ethynylbenzene **1a** (0.2 mmol) and ketimine **2a** (0.1 mmol) under a nitrogen atmosphere in a glove box. A solution of base (0.3 mmol) in 0.5, 1 and 2mL dry solvent was added by a "Titan" brand 1000 μ L pipettor to the reaction vial. The reaction mixture turned to a dark purple color. Then the vial was sealed with a cap, removed from the glove box, and stirred for 12 h at 25, 60, 80 and 100 °C. After the reaction was completed, the reaction mixture was opened to air, quenched with three drops of H₂O, diluted with 3 mL of ethyl acetate, and filtered over a 2 cm pad of MgSO₄ and silica. The pad was rinsed with ethyl acetate (3×2 mL), and the combined solutions were concentrated in vacuo. CDCl₃ and CH₂Br₂ were then added sequentially to determine the yield of the reaction mixture. CH₂Br₂ was used as an internal standard.

Gram scale synthesis of 3ac

An oven-dried 100 mL Schlenk tube equipped with a stir bar was sealed with a rubber septum and degassed, purged with nitrogen (repeated three times). DCM (10 mL) was added under nitrogen via syringe through the rubber septum. (4-methoxyphenyl)methanamine (411.3 mg, 3.0 mmol) and benzophenone imine (543.3 mg, 3.0 mmol) were added under nitrogen via syringe through the rubber septum. The reaction was stirred at 23 °C for 12 h, the solvent was completely removed in *vacuo* and the tube was filled with nitrogen. A solution (prepared in the glove box) of ethynylbenzene **1a** (606.1 mg, 6.0 mmol) in 10 mL anhydrous DME was added to the Schlenk tube via syringe through the rubber septum. Next, a solution of KN(SiMe₃)₂ (1.8 g, 12.0 mmol) in 20 mL anhydrous DME was added by

syringe through the rubber septum. The reaction mixture was stirred for 12 h in total at 100 °C, opened to air, and quenched with 3 mL of H₂O. The layers were separated and the aqueous layer was extracted with DCM (3 X 5 mL). The combined organic layers were concentrated in *vacuo*. The crude product was separated by flash chromatography on deactivated silica gel hexanes to give the product **3ca** in 86% yield (1.04 g).

Mechanistic study

a. Evidence of the mechanism goes through 4aa.



The reaction was performed following the General Procedure with ethynylbenzene **1a** (40.8 mg, 0.4 mmol) and *N*-benzyl-1,1-diphenylmethanimine **2a** (54.2 mg, 0.2 mmol). The crude product was separated by flash chromatography on deactivated silica gel hexanes to give the intermediate **4aa** in 10% yield (7.5 mg).

4aa(major): colorless liquid, $R_f = 0.39$ (hexanes:ethyl acetate = 30:1); ¹H NMR (600 MHz, Chloroform-*d*) δ 7.91 (d, J = 7.6 Hz, 2H), 7.61 – 7.55 (m, 5H), 7.48 (dt, J = 18.9, 7.4 Hz, 7H), 7.43 – 7.29 (m, 6H), 6.70 (dd, J = 16.0, 6.6 Hz, 1H), 6.57 (d, J = 15.9 Hz, 1H), 5.33 (d, J = 6.5 Hz, 1H) ppm; ¹³C{¹H} NMR (150 MHz, Chloroform-*d*) δ 167.7, 143.6, 140.1, 137.4, 137.0, 132.8, 130.3, 129.6, 128.9, 128.9, 128.7, 128.7, 128.6, 128.5, 128.3, 127.9, 127.6, 127.5, 127.1, 126.6, 68.8 ppm; HRMS (ESI⁺) [M+H]⁺ calc'd for C₂₈H₂₄N⁺: 374.1903, found: 374.1906.

b. 4aa is converted to 3aa.



The reaction was performed following the General Procedure with **4aa**. The crude product was separated by flash chromatography on deactivated silica gel hexanes to give the product **3aa** in 99% yield (7.4 mg).

X-ray crystal structures of compound 3ac.

CCDC 2085476 contains the supplementary crystallographic data for compound **3ac**. The data can be obtained free of charge from The Cambridge Crystallographic Data Center via www.ccdc.cam.ac.uk/data_request/cif.



Supplementary References

(1) Matin, J. O'D.; William, D. B.; Wiiiam, A. B.; Wiiiam, N. J.; Keith, K.; Brigitte, L.; Robin, L. P.;

Frederick, G. B.; Susan, R. M., J. Am. Chem. Soc. 1998, 110, 8520-8525

NMR Spectra

FigureS1.¹HNMRspectra(600MHz,Chloroform-d)of2,2,3,5-tetraphenyl-3,4-dihydro-2H-pyrrole (3aa).







FigureS3.¹HNMRspectra(400MHz,Chloroform-d)of2,2,5-triphenyl-3-(p-tolyl)-3,4-dihydro-2H-pyrrole (3ba).









FigureS6.¹³C{¹H}NMRspectra(100MHz,Chloroform-d)of3-(4-(*tert*-butyl)phenyl)-2,2,5-triphenyl-3,4-dihydro-2*H*-pyrrole (3ca).



FigureS7.¹HNMRspectra(600MHz,Chloroform-d)of3-(4-ethoxyphenyl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3da).



FigureS8. $^{13}C{^1H}$ NMRspectra(150MHz,Chloroform-d)of3-(4-ethoxyphenyl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3da).



FigureS9.¹HNMRspectra(400MHz,Chloroform-d)of2,2,5-triphenyl-3-(4-(trifluoromethoxy)phenyl)-3,4-dihydro-2H-pyrrole (3ea).







FigureS11.¹⁹FNMRspectra(376MHz,Chloroform-d)of2,2,5-triphenyl-3-(4-(trifluoromethoxy)phenyl)-3,4-dihydro-2H-pyrrole (3ea).



FigureS13. $^{13}C{^{1}H}$ NMRspectra(100MHz,Chloroform-d)of3-([1,1'-biphenyl]-4-yl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3fa).



FigureS15.¹³C{¹H}NMRspectra(100MHz,Chloroform-d)of3-(4-fluorophenyl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole(3ga).

3-(4-fluorophenyl)-2,2,5-triphenyl-3,4-dihydro-2*H*-pyrrole (3ga).

FigureS17.¹HNMRspectra(600MHz,Chloroform-d)of3-(4-chlorophenyl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3ha).

FigureS19.¹HNMRspectra(600MHz,Chloroform-d)of3-(4-bromophenyl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3ia).

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FigureS21.¹HNMRspectra(600MHz,Chloroform-d)of3-(naphthalen-2-yl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3ja).

FigureS23.¹HNMRspectra(400MHz,Chloroform-d)of3-(4,4-dimethylthiochroman-6-yl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3ka).

FigureS24. $^{13}C{^1H}$ NMRspectra(100MHz,Chloroform-d)of3-(4,4-dimethylthiochroman-6-yl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3ka).

FigureS25.¹HNMRspectra(400MHz,Chloroform-d)of3-(4,4-dimethylthiochroman-6-yl)-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3la).

FigureS27.¹HNMRspectra(600MHz,Chloroform-d)of3-hexyl-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3ma).

FigureS29.¹HNMRspectra(400MHz,Chloroform-d)of3-cyclopentyl-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3na).

FigureS31.¹HNMRspectra(400MHz,Chloroform-d)of3-cyclohexyl-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (30a).

FigureS33.¹HNMRspectra(400MHz,Chloroform-d)of2,2,3-triphenyl-5-(m-tolyl)-3,4-dihydro-2H-pyrrole (3ab).

FigureS34.¹³C{¹H}NMRspectra(100MHz,Chloroform-d)of2,2,3-triphenyl-5-(m-tolyl)-3,4-dihydro-2H-pyrrole (3ab).

FigureS35.¹HNMRspectra(400MHz,Chloroform-d)of5-(4-methoxyphenyl)-2,2,3-triphenyl-3,4-dihydro-2H-pyrrole(3ac).

FigureS37.¹HNMRspectra(400MHz,Chloroform-d)of5-(benzo[d][1,3]dioxol-5-yl)-2,2,3-triphenyl-3,4-dihydro-2H-pyrrole (3ad).

FigureS39.¹HNMRspectra(400MHz,Chloroform-d)of5-([1,1'-biphenyl]-4-yl)-2,2,3-triphenyl-3,4-dihydro-2H-pyrrole (3ae).

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FigureS42.13C{1H}NMRspectra(100MHz,Chloroform-d)of3-cyclohexyl-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3af).

FigureS43.¹⁹FNMRspectra(376MHz,Chloroform-d)of3-cyclohexyl-2,2,5-triphenyl-3,4-dihydro-2H-pyrrole (3af).

Figure S45. $^{13}C{^{1}H}$ NMR spectra (100 MHz, Chloroform-*d*) of 5-(4-chlorophenyl)-2,2,3-triphenyl-3,4-dihydro-2*H*-pyrrole (3ag).

FigureS47.13C{1H}NMRspectra(100MHz,Chloroform-d)of2,2,3-triphenyl-5-(o-tolyl)-3,4-dihydro-2H-pyrrole (3ah).

FigureS49.13C{1H}NMRspectra(100MHz,Chloroform-d)of5-(naphthalen-2-yl)-2,2,3-triphenyl-3,4-dihydro-2H-pyrrole (3ai).

FigureS51 $^{13}C{^1H}$ NMRspectra(100MHz,Chloroform-d)of2-(2,2,3-triphenyl-3,4-dihydro-2H-pyrrol-5-yl)pyridine (3aj).

FigureS53. $^{13}C{^{1}H}$ NMRspectra(100MHz,Chloroform-d)of4-(2,2,3-triphenyl-3,4-dihydro-2H-pyrrol-5-yl)pyridine (3ak).

FigureS55. $^{13}C{^1H}$ NMRspectra(100MHz,Chloroform-d)of5-(furan-2-yl)-2,2,3-triphenyl-3,4-dihydro-2H-pyrrole (3al).

FigureS57. $^{13}C{^{1}H}$ NMRspectra(150MHz,Chloroform-d)of(E)-N-(1,3-diphenylallyl)-1,1-diphenylmethanimine (4aa).

HRMS Spectra

		m/z 🗠	Ion	Formula	Abundance						
•		374.1905	(M+H)+	C28 H24 N	148848.1]					
		Best	Formula (M)	Ion Formula 4	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
	•	V	C28 H23 N	C28 H24 N	97.06		374.1903	-0.55	99.75	91.82	98

MS Formula Results: + Scan (0.7752 min) Sub (ZQX-99-4.d)

	m/z 🗠	lon	Formula	Abundance						
-	388.2065	(M+H)+	C29 H26 N	243512.4						
	Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing
		000,000,00	000 1100 11	00.0		200.200	1.07	00.4	07.70	

3ca

MS Formula Results: + Scan (1.0870 min) Sub (ZQX-99-16.d)

	m/z 🗠	Ion	Formula	Abundance						
-	430.2529	(M+H)+	C32 H32 N	518959.7]					
	Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacir
÷-	×	C32 H31 N	C32 H32 N	99.54		430.2529	-0.01	100	99.4	

60 80 100 120 140 160 180 200 220 240 260 280 300 320 340 360 380 400 420 440 460 480 500 520 540 560 580 600 620 640 660 680 700 720 740 760 780 800 820 Counts vs. Mass-to-Charge (m/z)

		m/z 🗠	lon	Formula	Abundance						
•-		418.2167	(M+H)+	C30 H28 N O	415132.5						
	Γ	Best	Formula (M)	Ion Formula 🛛 🗠	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
	•	V	C30 H27 N O	C30 H28 N O	97.86		418.2165	-0.48	99.79	94.95	97.5

3ea

x10 ⁵	+ESI Scan (0.9968 min) Frag=150.0V ZQX-99-3.d Subtract	
1.1-	458.	1730
1.05-		
1-		
0.95-		
0.9-		
0.85-		
0.8-		
0.75-		
0.7-		
0.65-		
0.6-		
0.55-		
0.5-		
0.45-		
0.4-		
0.35-		
0.3-		
0.25-		
0.2-		
0.15-		
0.1-		
0.05-		
۲0		

50 75 100 125 150 175 200 225 250 275 300 325 350 375 400 425 450 475 500 525 550 575 600 625 650 675 700 725 750 775 800 825 850 875 900 Counts vs. Mass-to-Charge (m/z)

MS Formula Results: + Scan (0.9968 min) Sub (ZQX-99-3.d)

m/z 🗠	lon	Formula	Abundance						
458.173	(M+H)+	C29 H23 F3 N O	107270.9]					
Best	Formula (M)	Ion Formula 🛛 🗠	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spa
 v	C29 H22 F3 N O	C29 H23 F3 N O	95.74		458.1726	-0.92	99.15	90.81	

		m/z 🗠	Ion	Formula	Abundance						
-		450.2216	(M+H)+	C34 H28 N	57708.5						
		Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
	÷.	V	C34 H27 N	C34 H28 N	47.61		450.2216	0.16	99.97	0	0

MS Formula Results: + Scan (0.9127 min) Sub (ZQX-99-1.d)

	m/z 🗠	lon	Formula	Abundance						
	392.1808	(M+H)+	C28 H23 F N	209452.5]					
	Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	S
÷-	V	C28 H22 F N	C28 H23 F N	99.03		392.1809	0.12	99.99	98.11	

60 80 100 120 140 160 180 200 220 240 260 280 300 320 340 360 380 400 420 440 460 480 500 520 540 560 580 600 620 640 660 680 700 720 740 760 780 Counts vs. Mass-to-Charge (m/z)

MS Formula Results: + Scan (0.9565 min) Sub (ZQX-99-6.d)

	m/z 🗠	Ion	Formula	Abundance						
-	408.1518	(M+H)+	C28 H23 CI N	276104.8						
	Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
÷		C28 H22 CI N	C28 H23 CI N	96.12		408.1514	-1.26	98.57	90.5	97.98

	11//2	1011	i officia	Abundance						
	452.1006	(M+H)+	C28 H23 Br N	521714.3]					
	Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
÷[C28 H22 Br N	C28 H23 Br N	98.05		452.1008	0.63	99.61	93.94	99.87

		m/z 🗠	Ion	Formula	Abundance						
-		424.206	(M+H)+	C32 H26 N	568757.1						
	Γ	Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
	÷.	V	C32 H25 N	C32 H26 N	98.42		424.206	-0.02	100	94.51	99.95

MS Formula Results: + Scan (0.8176 min) Sub (ZQX-99-7+.d)

	m/z 🗠	lon	Formula	Abundance						
	474.2248	(M+H)+	C33 H32 N S	66904.7						
Γ	Best	Formula (M)	lon Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing
- d	V	C33 H31 N S	C33 H32 N S	96.55		474.225	0.43	99.81	88.5	

		m/z 🗠	lon	Formula	Abundance						
•		375.1856	(M+H)+	C27 H23 N2	496650.5						
		Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
	÷-	V	C27 H22 N2	C27 H23 N2	97.84		375.1856	-0.19	99.97	92.51	99.96

40 60 80 100 120 140 160 180 200 220 240 260 280 300 320 340 360 380 400 420 440 460 480 500 520 540 560 580 600 620 640 660 680 700 720 Counts vs. Mass-to-Charge (m/z)

MS Formula Results: + Scan (0.8600 min) Sub (ZQX-99-28+.d)

m/z 🗠	lon	Formula	Abundance						
382.2532	(M+H)+	C28 H32 N	533341.3]					
Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Sp

MS Formula Results: + Scan (1.0044 min) Sub (ZQX-99-22.d)

Γ		m/z	lon	Formula 🗠	Abundance						
•		366.2216	(M+H)+	C27 H28 N	1028606.1						
		Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
1	÷	V	C27 H27 N	C27 H28 N	98.74		366.2216	0.16	99.98	97.16	98.15

3oa

MS Formula Results: + Scan (1.1352 min) Sub (ZQX-99-2.d)

		m/z	lon	Formula 🗠	Abundance						
		380.2372	(M+H)+	C28 H30 N	1529291.6						
		Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
G	0-	V	C28 H29 N	C28 H30 N	98.39		380.2373	0.33	99.91	98.96	94.67

		m/z 🗠	lon	Formula	Abundance						
=·[388.2063	(M+H)+	C29 H26 N	159290						
		Best	Formula (M)	Ion Formula	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
	•		C29 H25 N	C29 H26 N	94.25		388.206	-0.91	99.29	81.47	99.53

MS Formula Results: + Scan ((1.3004 min)	Sub (70X-92-8 d)
IND FUITIUIA RESULS. + SCALL	1.3094 11111)	Sub (ZQA-92-0.U)

		m/z 🗠	Ion	Formula	Abundance						
-		404.2008	(M+H)+	C29 H26 N O	172314.1						
	Γ	Best	Formula (M)	lon Formula 🛛 🗠	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
6	•	V	C29 H25 N O	C29 H26 N O	95.83		404.2009	0.17	99.97	86.52	98.7

	m/z 🗠	lon	Formula	Abundance						
	418.1801	(M+H)+	C29 H24 N O2	71435.7]					
	Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Sp
÷	>	C29 H23 N O2	C29 H24 N O2	96.98		418.1802	0.07	100	90.34	

MS Formula Results: + Scan (1.1124 min) Sub (ZQX-92-7.d)

	m/z 🗠	lon	Formula	Abundance						
	450.2217	(M+H)+	C34 H28 N	280195.3]					
ſ	Best	Formula (M)	lon Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Sp
		C34 H27 N	C34 H28 N	97.66		450.2216	-0.15	99.98	92.4	

	m/z 🗠	Ion	Formula	Abundance						
	392.1808	(M+H)+	C28 H23 F N	353691.5						
	Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
÷.	>	C28 H22 F N	C28 H23 F N	97.33		392.1809	0.15	99.98	91.08	99.54

Г

MS Formula Results: + Scan (0.8385 min) Sub (ZQX-92-5.d)

	m/z 🗠	lon	Formula	Abundance						
=-[408.1516	16 (M+H)+ C28 H23 CI N		178678.3						
	Best	Formula (M)	Ion Formula	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
E	>	C28 H22 CI N	C28 H23 CI N	95.93		408.1514	-0.78	99.46	89.55	96.5

MS Formula Results: + Scan (1.2373 min) Sub (ZQX-92-1+.d)

		m/z 🗠	lon	Formula	Abundance						
- ·		424.2061	(M+H)+	C32 H26 N	144377.8						
		Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
d]	▼	C32 H25 N	C32 H26 N	95.33		424.206	-0.4	99.85	86.18	97.28

MS Formula Results: + Scan (0.8300 min) Sub (ZQX-92-3.d)

		m/z 🗠	Ion	Formula	Abundance						
-		375.1859	(M+H)+	C27 H23 N2	62110.9						
	[Best	Formula (M)	Ion Formula	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spacing Match
	•		C27 H22 N2	C27 H23 N2	92.25		375.1856	-0.98	99.2	84.16	88.06

3ak

MS Formula Results: + Scan (0.8980 min) Sub (ZQX-92-15.d)

	m/z 🗠	lon	Formula	Abundance						
	375.1857	(M+H)+	C27 H23 N2	14858]					
Γ	Best	Formula (M)	lon Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spaci
•	V	C27 H22 N2	C27 H23 N2	95.32		375.1856	-0.35	99.9	91.38	

MS Formula Results: + Scan (0.8759 min) Sub (Z	ZQX-92-12.d)
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m/z 🗠	lon	Formula	Abundance						
364.17	(M+H)+	C26 H22 N O	175784.5						
Best	Formula (M)	Ion Formula 🖉	Score	Cross Score	Calc m/z	Diff (ppm)	Mass Match	Abund Match	Spa
	C26 H21 N O	C26 H22 N O	98.79		364.1696	-1.24	98.76	98.59	