### Triple Zirconocene/Brønsted Acid/CuO Cooperative and Relay Catalysis System for Tandem Mannich Addition/C-C Formative Cyclization/Oxidation

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### **1.** General procedures

<sup>1</sup>H and <sup>13</sup>C NMR spectra were recorded on a Bruker EQUINX55 (400 MHz for <sup>1</sup>H; 101 MHz for <sup>13</sup>C) spectrometer in CDCl<sub>3</sub>. For <sup>1</sup>H NMR, tetramethylsilane (TMS) served as internal standard ( $\delta = 0$ ) and <sup>1</sup>H NMR chemical shifts are reported in ppm downfield of tetramethylsilane and referenced to residual solvent peak (CDCl<sub>3</sub> at 7.26 ppm) unless otherwise noted. The data are reported as follows: chemical shift, integration, multiplicity (s = singlet, d = doublet, t = triplet, q = quartet and m = multiplet), and coupling constant in Hz. For <sup>13</sup>C NMR, CDCl<sub>3</sub> was used as internal standard ( $\delta = 77.0$ ) and spectra were obtained with complete proton decoupling. HRMS (ESI) analysis was performed and (HRMS) data were reported with sodium mass/charge (m/z) ratios as values in atomic mass units. Column chromatography was performed on silica gel (230-400 mesh) and analytical thin layer chromatography was carried out using 250 µm commercial silica gel plates. Visualization of the developed chromatogram was performed by UV absorbance and stained with an iodine vapor.

### 2. Typical procedures for synthesis of substituted quinolines

A 10 mL test tube, equipped with a magnetic stirrer and a septum, was charged with *i*-PrOH (0.375mL), H<sub>2</sub>O (0.125mL), aldehyde (1.0 mmol), amine (1.1 mmol) and ketone (1.5 mmol), in one portion.  $Cp_2ZrCl_2$  (0.05mmol), trimellitic acid (0.05mmol), and CuO (0.05mmol) were added at 60 °C and stirred until the reaction was completed as indicated by TLC. Upon completion of the reaction, the reaction mixture was quenched with distilled water (5.0 mL). The aqueous phase was extracted with ether (3×5 mL), dried over Na<sub>2</sub>SO<sub>4</sub> and concentrated in vacuo to give desired products. The corresponding solid products were obtained through column chromatography by using 100–200 mesh silica gels.

# **3.** Optimization of reaction conditions for synthesis of substituted quinolines

To further investigate the effect of solvents, next we examined our reaction in different solvent systems. The catalytic activity of zirconocene dichloride, trimellitic acid and CuO in the reaction of anilines, aldehydes and ketones was sligthly influenced by solvents as show in Table S1. At first, non-polar solvent *n*-hexane was evaluated, we can found the catalyst almost inert in CH<sub>2</sub>Cl<sub>2</sub>, *n*-hexane only obtained 20% and 5% yield (entries 1 and 2). DMSO and THF can slightly accelerated Zr-Cu catalyst in this reaction obtained 20% and 32% yields respectively (entries 3, 4). More polar solvent such as EtOH was obviously accelerated Zr-Cu catalyst in this reaction obtained 48% yields (entry 5). So that, several alcohols were examined in this reaction, the result demonstrated that *i*-PrOH was best solvent in this coupling reaction, afforded 54% yield in this reaction (entries 6-12). At the same time, water as a solvent also sreened, only abtained 10%

substituted quinolines, but no byprduct was produced in this reaction. Concentrating this three components coupling reaction process fast in *i*-PrOH, so we combined water and *i*-PrOH, the best result obtained in *i*-PrOH: $H_2O=3:1$ .



Entry	Solvent	Yield (%) <sup>b</sup>		
1	$CH_2CI_2$	29		
2	<i>n</i> -hexane	5		
3	DMSO	20		
4	THF	32		
5	EtOH	48		
6	MeOH	43		
7	PrOH	47		
8	<i>i</i> -PrOH	54		
9	BuOH	50		
10	t-BuOH	44		
11	C <sub>6</sub> H <sub>13</sub> OH	40		
12	C <sub>6</sub> H <sub>15</sub> OH	37		
13	H <sub>2</sub> O	10		
14	<i>i</i> -PrOH/H <sub>2</sub> O=1:1	30		
15	<i>i</i> -PrOH/H <sub>2</sub> O=2:1	47		
16	<i>i</i> -PrOH/H <sub>2</sub> O=3:1	58		
17	<i>i</i> -PrOH/H <sub>2</sub> O=4:1	51		

Table S1 Solvent screening of three-component coupling sequence reaction<sup>a</sup>

 $^{a}$ All reactions were conducted using the aniline (1 mmol),benzaldehyde (1 mmol), methyl pyruvate (1.0 mmol), 50  $^{\circ}$ C, 1h.  $^{b}$ Isolated yields

Table S2 shows the effect of the catalyst amount with increasing of the molar ratio of catalyst trimellitic acid,  $Cp_2ZrCl_2$  and CuO. When the loading of catalyst increased from 1 mol% to 5 mol%, the yield of product increased sharply from 15% to 58%. At a catalyst loading of 5 mol%, the best result can be obtained. Further increasing the amount of catalyst, the yield increase slightly,

therefore 5mol% was selected as the best loading of catalyst. The ration trimellitic acid and  $Cp_2ZrCl_2$  also screened, result demonstrated that 5 mol% trimellitic acid,  $Cp_2ZrCl_2$  and CuO were best. Subsequently, we checked the effect of temperature on the progress of the reaction. According to Table S2, we can see that as the temperature increases the yield decreased, because byproducts increased with temperature increasing, so the optimum temperature is 60°C. Reaction time was also screened, the optical time was 2h.



Entry	Ratio of Cat.	Time	Temp(℃)	Yield(%) <sup>b</sup>
1	1%:1%:1%	1	50	15
2	2%:2%:2%	1	50	30
3	3%:3%:3%	1	50	43
4	5%:5%:5%	1	50	58
5	10%:10%:10%	1	50	60
6	5%:6%:5%	1	50	62
7	5%:7%:5%	1	50	63
8	5%:8%:5%	1	50	64
9	5%:9%:5%	1	50	65
10	5%:10%:5%	1	50	65
11	5%:5%:5%	1	60	70
12	5%:5%:5%	1	70	59
13	5%:5%:5%	1	80	57
14	5%:5%:5%	1	90	55
15	5%:5%:5%	1	100	53
17	5%:5%:5%	1.5	60	85
18	5%:5%:5%	2	60	87
19	5%:5%:5%	2.5	60	88
20	5%:5%:5%	3	60	89
21	5%:5%:5%	3.5	60	89
<sup>a</sup> All reactions were conducted using the aniline (1 mmol), benzaldehyde (1 mmol), methyl pyruvate (1.0 mmol),				

Table S2 Ratio of catalyst and time screening of three-component coupling sequence reaction<sup>a</sup>

ratio of catalyst Cp2ZrCl2: trimellitic acid: CuO, 50  $^\circ \! \mathbb{C}$  , 1 h.  $^b$ Isolated yields based on 2.



Entry	1/2/3	Yield(%) <sup>b</sup>
1	1:1:1	85
2	2:1:1	85
3	3:1:1	85
4	4:1:1	86
5	1:1.1:1	90
6	1:1.2:1	90
7	1:1.3:1	90
8	1:1.4:1	91
9	1:1.5:1	91
10	1:1.1:1.1	73
11	1:1.1:1.2	76
12	1:1.1:1.3	80
13	1:1.1:1.4	84
14	1:1.1:1.5	91
15	1:1.1:1.6	91
16	1:1.1:1.7	91
17	1:1.1:1.8	91
18	1:1.1:2	92

Table S3 Ratio of substrates screening of three-component coupling sequence reaction<sup>a</sup>

<sup>a</sup>All reactions were conducted using aldehyde (1.0 mmol); Cp<sub>2</sub>ZrCl<sub>2</sub> (0.05 mmol, 5 mol%); CuO (0.05 mmol, 5 mol%); trimellitic acid (0.05 mmol, 5 mol%); i-PrOH : H<sub>2</sub>O (3:1, 0.5 mL); All reactions were carried out at 60 °C for 2 h. <sup>b</sup>Isolated yield.

The ratio of substrates is show in Table S3. From this table, we can see the optical ratio of benzaldehyde, aniline and ketone are 1:1.1:1.5.

#### 4. NMR experiments

The interplay of  $Cp_2ZrCl_2$  and trimellitic acid were investigated by <sup>1</sup>H NMR in mechanistic scenario. A general procedure was as follows:  $Cp_2ZrCl_2$  (10 µmol) and trimellitic acid (10 µmol) was placed in  $D_2O$  (0.5mL) and the solution was detected immediately as observed by <sup>1</sup>H NMR

spectroscopy. The mixture was allowed to stand for 1 h and was conducted by <sup>1</sup>H NMR spectroscopyas confirmed by <sup>1</sup>H NMR spectroscopy. After 1.0 equiv. of trimellitic acid was added in the above solution, one new Cp protons singlet appeared at  $\delta$  6.57 ppm, but did not increase as time go on. After 2.0 equiv. aniline was added, Cp<sub>2</sub>ZrCl<sub>2</sub> (I)was consumed gradually in D<sub>2</sub>O in the presence of base and formed new zirconocene species Cp<sub>2</sub>Zr(OOC)<sub>2</sub>PhCOOH. (II).



**Fig. S1** Partial 400 MHz <sup>1</sup>H NMR spectra (D<sub>2</sub>O) of a mixture of Cp<sub>2</sub>ZrCl<sub>2</sub> (1.0 equiv) and trimellitic acid (1.0 equiv) with PhNH<sub>2</sub>(2.0 equiv). 6.49 ppm I $_{\bullet}$ [Cp<sub>2</sub>ZrCl<sub>2</sub>]; 6.57 ppm II $\star$ Cp<sub>2</sub>Zr(OOC)<sub>2</sub>PhCOOH.



**Fig. S2** Partial 400 MHz <sup>13</sup>C NMR spectra ( $D_2O$ ) of a mixture of  $Cp_2ZrCl_2$  (1.0 equiv) and trimellitic acid (1.0 equiv) with PhNH<sub>2</sub> (2.0 equiv).

### 5. HRMS analysis

Mass spectrometric measurements were performed in aBruker EVOQ tandem mass spectrometer. As a general rule, scan mode was Q1MS, positiveion mode (otherwise indicated), tube lens potential was optimized in each case or for aseries of measurements that required equal conditions, a time span of 1 minute was used to collectspectra and average them. The tube lens potential was adjusted in a way that the most interest ions had almost no attenuation (around 70 V).

For CID experiments, the cations of interest were mass-selected using the first quadrupole (Q1) and interacted with argon in the T-wave collision cell at variable collision energies (Elaboratory= 3-15 eV). The ionic products of fragmentation were analyzed with the time-of-

flight analyzer. The isolation width was 1Da and the most abundant isotopomer was massselected in the first quadrupole analyzer.



Fig. S3 The HRMS of trimellitic acid and  $Cp_2ZrCl_2$  (m/z 100-1000)



Fig. S4 The HRMS of trimellitic acid and  $Cp_2ZrCl_2$  (m/z 410-470)



Fig. S5 The HRMS of trimellitic acid and  $Cp_2ZrCl_2$  (m/z 428-438)



Fig. S6 The HRMS of trimellitic acid and  $\mbox{Cp}_2\mbox{ZrCl}_2$ 

## 6. <sup>1</sup>H NMR and <sup>13</sup>C NMR Spectra for All Compounds:

4aa <sup>1</sup>HNMR spectrum





210 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 -10 f1(ppm) 4ac <sup>1</sup>HNMR spectrum



210 200 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 f1(ppm)

4ad <sup>1</sup>HNMR spectrum



210 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 -10 f1(ppm)











4ag <sup>13</sup>C NMR spectrum

- 1672 - 1672 - 1552 - 1490 - 1490 - 1490 - 1490 - 1259 - 1259 - 1289 - 1289 - 1289 - 1188 - 1188 √ 5541 ^ 5261 ~ 3529 ~ 3722





4ai <sup>1</sup>HNMR spectrum



210 200 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 f1(ppm)











4aj <sup>13</sup>C NMR spectrum





4ak <sup>1</sup>HNMR spectrum







4am <sup>1</sup>HNMR spectrum



210 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 -10 f1(ppm)

4an <sup>1</sup>HNMR spectrum









4ap <sup>1</sup>HNMR spectrum







4aq <sup>1</sup>HNMR spectrum





4ar <sup>1</sup>HNMR spectrum







4at <sup>1</sup>HNMR spectrum





4at <sup>13</sup>C NMR spectrum





4bb <sup>1</sup>HNMR spectrum







4bd <sup>1</sup>HNMR spectrum



210 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 -10 f1(ppm)





4bf <sup>1</sup>HNMR spectrum













4bi <sup>1</sup>HNMR spectrum



4bj <sup>1</sup>HNMR spectrum



210 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 -10 f1(ppm)







4bm <sup>1</sup>HNMR spectrum





