## Electronic Supplementary Information Metal-Organic Gel Coupled Entropy-Driven Circuit for Fluorescence Detection of miR-155

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Name	Sequence (5'-3')			
SS	ACCCCTATCACGATTAGCATTAAAGGGCCGTAAGAGAGCT			
	GTAGATTGGATCG			
OS	FAM-CCCTTTAATGCTAATCGTGAT			
AS	CGATCCAATCTACAGCTCTCTTACGG			
fuel	CGATCCAATCTACAGCTCTCTTACGGCCCTTTAATGCTAAT			
	CGTGAT			
miR-155	UUAAUGCUAAUCGUGAUAGGGGU			
miD-155	TTAATGCTAATCGTGATAGGGGT			
Mis-1	UUAAU <u>C</u> CUAAUCGUGAUAGGGGU			
Mis-2	UUAAU <u>C</u> CUAAUCGUGAUAG <u>C</u> GGU			
Mis-3	UUAAU <u>C</u> CUAAU <u>A</u> GUGAUAG <u>C</u> GGU			
Mis-4	UUAAU <u>C</u> CUAAU <u>A</u> GUG <u>U</u> UAG <u>C</u> GGU			
miR-21	UAGCUUAUCAGACUGAUGUUGA			
let-7a	UGAGGUAGUAGGUUGUAUAGUU			

 Table S1. Oligonucleotide sequences used in this work



**Fig. S1.** The quenching constant of TD (A) and OS (B). The concentration of TD and OS were 100 nM and 200 nM, respectively.



**Fig. S2.** The examination of Cu-MOG hydrostability. (A) The XRD spectra of Cu-MOG in the aqueous solutions with different pH values and inorganic ions. (B) The SEM images of Cu-MOG in the aqueous solutions with different pH values and inorganic ions.



**Fig. S3**. The fluorescence curve of Cu-MOG. (A) The excitation and emission spectra of Cu-MOG. (B) The fluorescence curve of Cu-MOG in the presence of miR-155 under excitation wavelength of 338 nm.



**Fig. S4.** Optimization of experimental conditions. (A) Concentration optimization of Cu-MOG; (B) EDC reaction temperature optimization. (C) EDC reaction time optimization. (D) Optimization of incubation time of Cu-MOG. The experimental conditions were as follows: the concentrations of TD and fuel were 100 nM, and the concentration of miR-155 was 3 nM. The ribonuclease inhibitor concentration was 25 U/mL.



Fig. S5. The adsorption kinetics of OS and TD on Cu-MOG.



**Fig. S6.** The anti-interference ability of this method. The concentration of miR-155 was 3 nM, and the concentrations of  $K^+$  and  $Na^+$  were 5 mM and 140 mM, respectively. In addition, the low concentrations of these proteins were all 50 nM and the high concentrations of these proteins were all 100 nM. The red dashed line showed the value of the miR-155 response fluctuating by 10%.

Sample	Added (nM)	Found (nM)	Recovery(%)	RSD (%)
		Mean <sup>b</sup> ±SD <sup>c</sup>		
1	1.00	$1.06 \pm 0.01$	106.33±1.25	1.17
2	3.00	3.12±0.10	104.00±3.31	3.18
3	5.00	$4.88 \pm 0.07$	97.67±1.41	1.44

Table S2. Recovery tests of miR-155 detection in 10000-fold diluted human serum samples  $(n=3)^{a}$ .

<sup>*a*</sup> The human serum samples were acquired from the Southwest Hospital, Chongqing, China. <sup>*b*</sup> The mean of three determinations. <sup>*c*</sup> SD = standard deviation.



Fig. S7. The SEM images of Cu-MOG before (A) and after (B) sensing.



Fig. S8. The cyclic voltammetry curves of OS. Inset: amplification of delineating potential.



**Fig. S9.** The cyclic voltammetry curves of Cu-MOG. The figure above showed the whole and the figure below showed the amplification of delineating potential.



**Fig. S10.** The absorption of OS (A) and Cu-MOG (B). Inset: Optical bandgap determined from Taut plot  $(Ahv)^2 vs.$  (hv) of indirect semiconductor and  $(Ahv)^{1/2} vs.$  (hv) of direct semiconductor, where A is absorbance.