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Supporting information

3Dgraphene/hydroxypropyl- β -cyclodextrin nanocomposite as electrochemical chiral sensor for the recognition of tryptophan enantiomers

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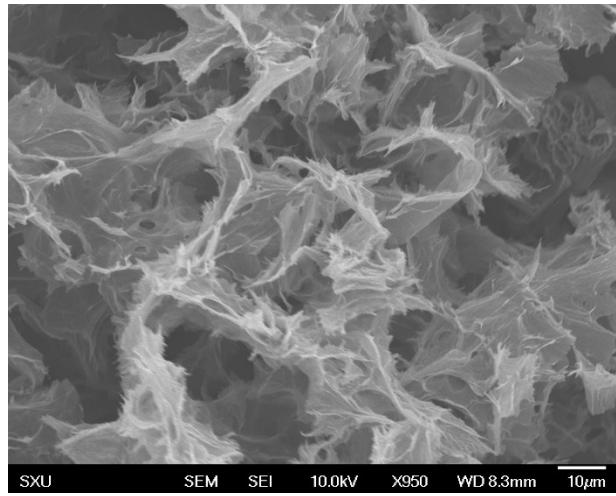


Fig. S1. SEM image of 3D-G

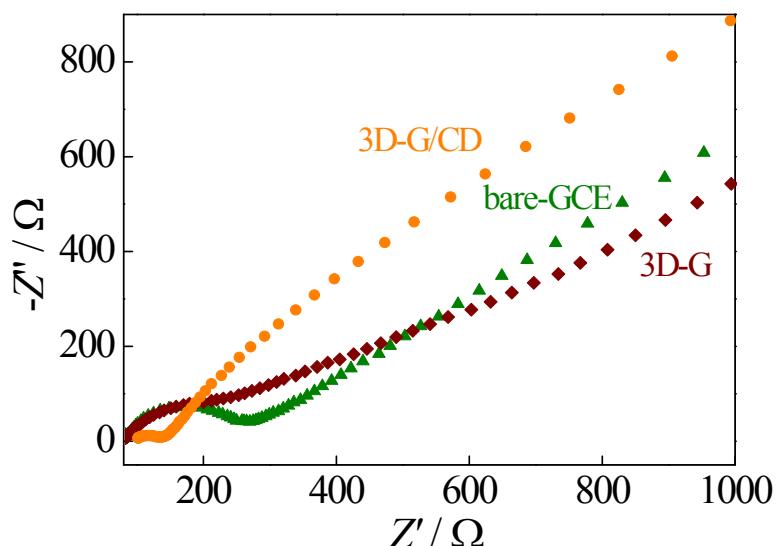


Fig. S2. Nyquist plots of bare GCE, 3D-Gand 3D-G/HP- β -CD modified GCE in 0.1 M KCl containing 10 mM $[\text{Fe}(\text{CN})_6]^{3-4-}$ (pH 7.0) at 25 °C.

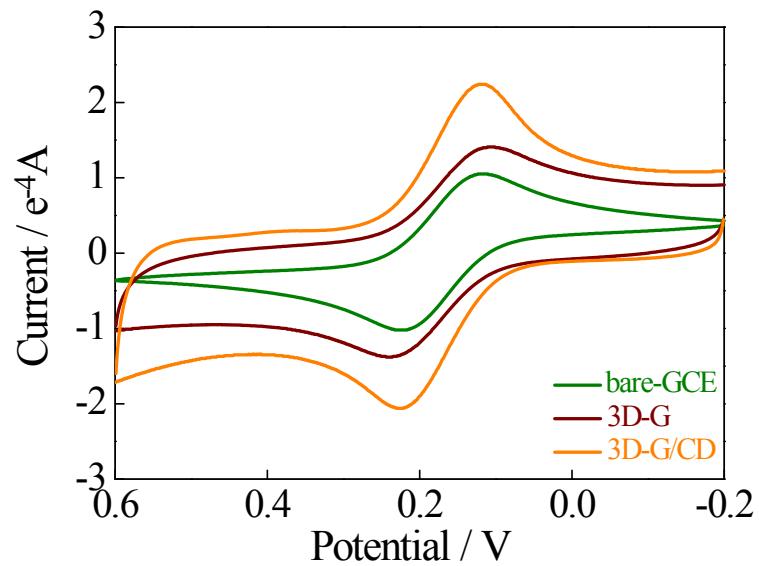


Fig. S3.Cyclic voltammograms of bare GCE, 3D-G and 3D-G/HP- β -CD modified GCE in 0.1 M KCl containing 10 mM $[\text{Fe}(\text{CN})_6]^{3-/-4-}$ (PBS of pH 7.0) at 25 °C.

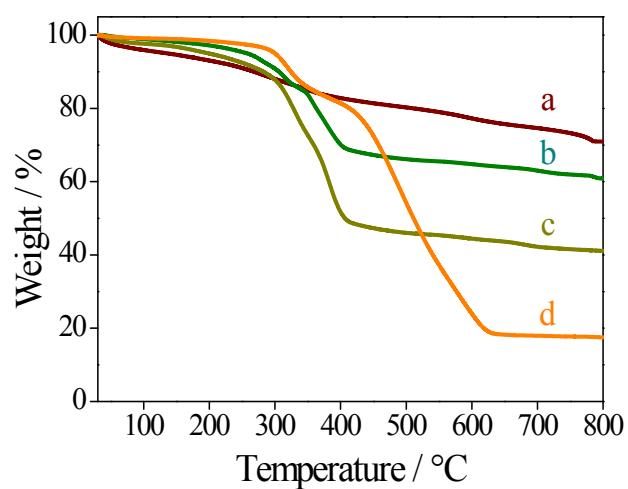


Fig. S4. TGA of 3D-G (a) and 3D-G/HP- β -CD containing different amounts of HP- β -CD (b-d).

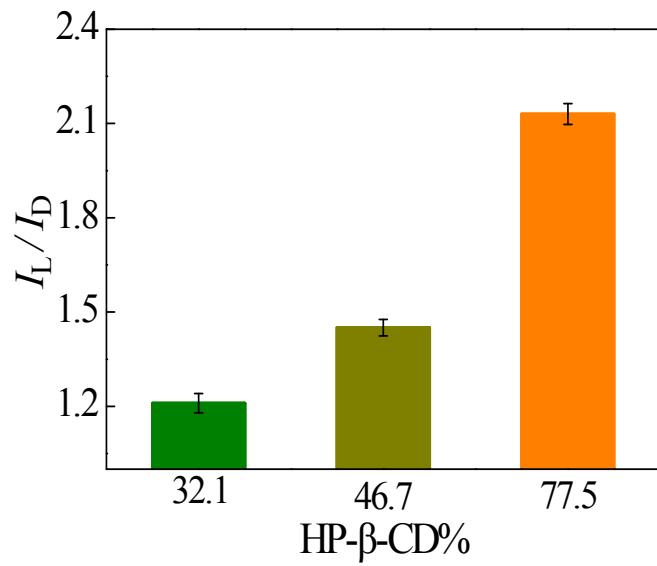


Fig. S5. The HP- β -CD amounts of 3D-G/HP- β -CD (32.1%, 46.7% and 77.5%) influence on the enantioresognition efficiency (I_L/I_D) of 3D-G/HP- β -CD/GCE toward Trp isomers. Errors bars represent the standard deviation for three independent measurements.

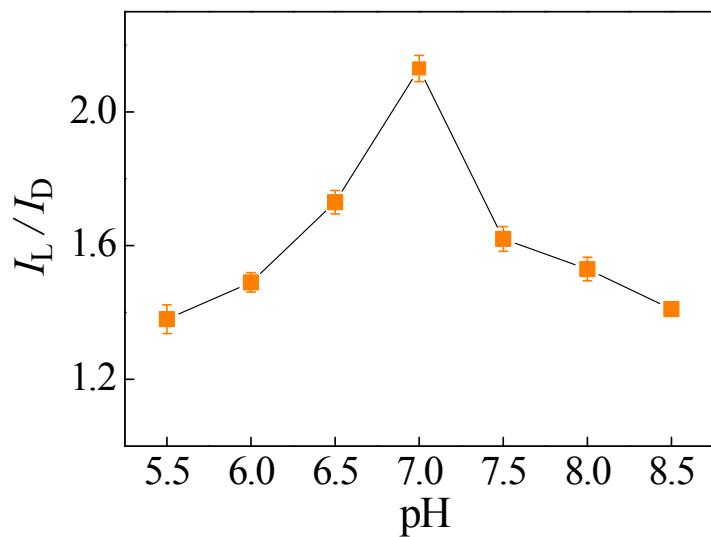


Fig. S6. Influence of pH on the enantioresognition efficiency (I_L/I_D) of 3D-G/HP- β -CD/GCE toward Trp isomers at 25 °C. Errors bars represent the standard deviation for three independent measurements.

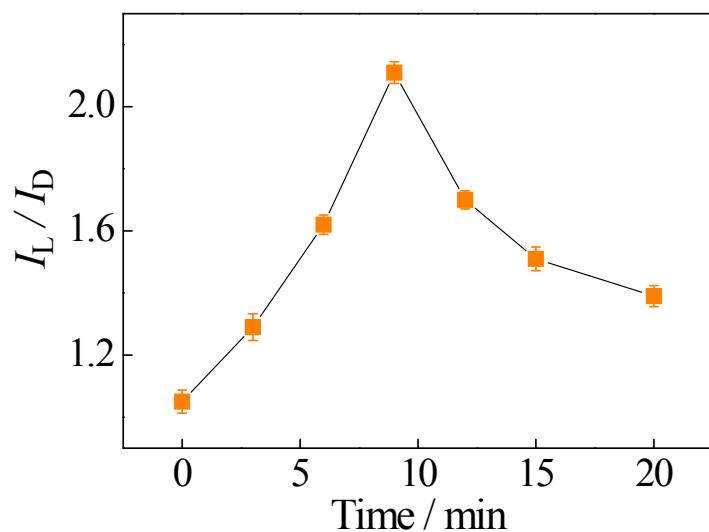


Fig. S7. Influence of the accumulation time on the enantioresognition efficiency (I_L/I_D) of 3D-G/HP- β -CD/GCE toward Trp isomers at 25 °C. Errors bars represent the standard deviation for three independent measurements.

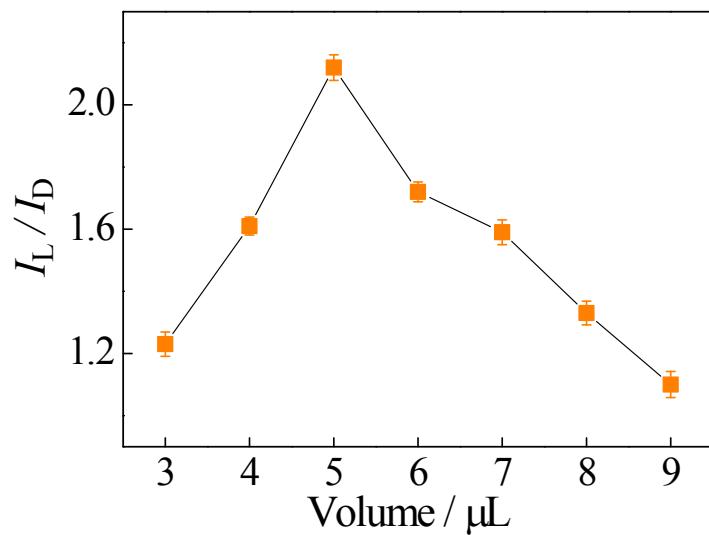


Fig. S8. Influence of the 3D-G/HP- β -CD amount on the enantioresognition efficiency (I_L/I_D) of 3D-G/HP- β -CD/GCE toward Trp isomers at 25 °C. Errors bars represent the standard deviation for three independent measurements.

Table S1. Comparison with other electrochemical methods for the recognition of tryptophan enantiomers

Modified electrode	Peak current ratio (L-Trp to D-Trp)	Linear range (mol L ⁻¹)	Detection limit (mol L ⁻¹)	Ref.
PTCA-CS/GCE	2.60	—	—	40
P-L-Glu/β-CD/GCE	2.30	—	—	9
GQDs/β-CD/GCE	2.10	—	—	39
GQDs-CS/GCE	2.06	—	—	10
Pd-Cu@Cu ₂ O/N-RGO/GCE	—	1.0 × 10 ⁻⁸ – 4.0 × 10 ⁻⁵	1.9 × 10 ⁻⁹ (L/D)	41
β-CD-MNPs/GCE	—	8.0 × 10 ⁻⁷ – 3.0 × 10 ⁻⁴	5.0 × 10 ⁻⁷ (L/D)	42
ds-DNA/Thi-GR/GCE	—	5.0 × 10 ⁻⁷ – 2.5 × 10 ⁻³	1.7 × 10 ⁻⁷ (L/D)	43
NH ₂ -GQDs/β-CD/GCE	—	1.0 × 10 ⁻⁶ – 3.0 × 10 ⁻⁵	6.5 × 10 ⁻⁷ (L) 1.2 × 10 ⁻⁷ (D)	44
NH ₂ -β-CD/Au@Pt/PEI/ MWCNTs/GCE	—	1.0 × 10 ⁻⁵ – 5.0 × 10 ⁻³	4.3 × 10 ⁻⁶ (L) 5.6 × 10 ⁻⁶ (D)	45
PLC/MWCNTs/GCE	—	1.0 × 10 ⁻⁴ – 1.0 × 10 ⁻³	3.3 × 10 ⁻⁵ (L/D)	46
mNBFs/mNC-GPEs	1.40	1.0 × 10 ⁻⁷ – 1.0 × 10 ⁻⁵	—	47
β-CD-PtNPs/GNs/GCE	1.30	5.0 × 10 ⁻⁵ – 5.0 × 10 ⁻³	1.7 × 10 ⁻⁵ (L) 2.1 × 10 ⁻⁵ (D)	36
3D-G/HP-β-CD/GCE	2.13	5.0 × 10 ⁻⁷ – 1.75 × 10 ⁻⁴	9.6 × 10 ⁻⁹ (L) 3.8 × 10 ⁻⁸ (D)	This work

Table S2. Inclusion constant (K) for the 1:1 inclusion complexation of L-, D-phenylalanine, L, D-tryptophan and L-, D-tyrosine with HP- β -CD.

Chiral amino acids	K (M $^{-1}$) of L-isomer with HP- β -CD	K (M $^{-1}$) of D-isomer with HP- β -CD
Phenylalanine (Phe)	23.75	19.73
Tyrosine (Tyr)	32.02	25.31
Tryptophan (Trp)	235.3	176.2