

Supporting Information

Endogenous Conjugation of Biomimetic Dinitrosyl Iron Complex with Protein Vehicles for Oral Delivery of Nitric Oxide to Brain and Activation of Hippocampal Neurogenesis

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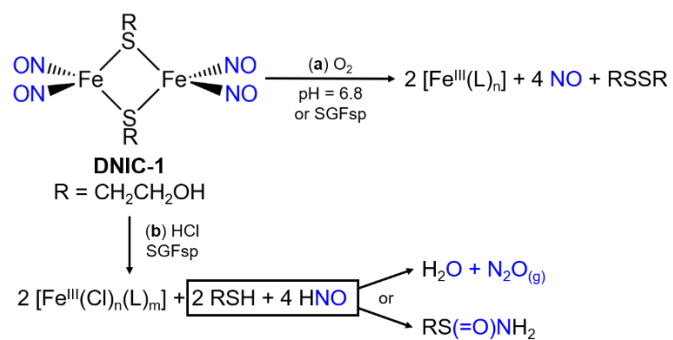
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Scheme S1. Mechanism for degradation of **DNIC-1** (a) at pH 6.8 and (b) in SGFsp, respectively, under aerobic condition.



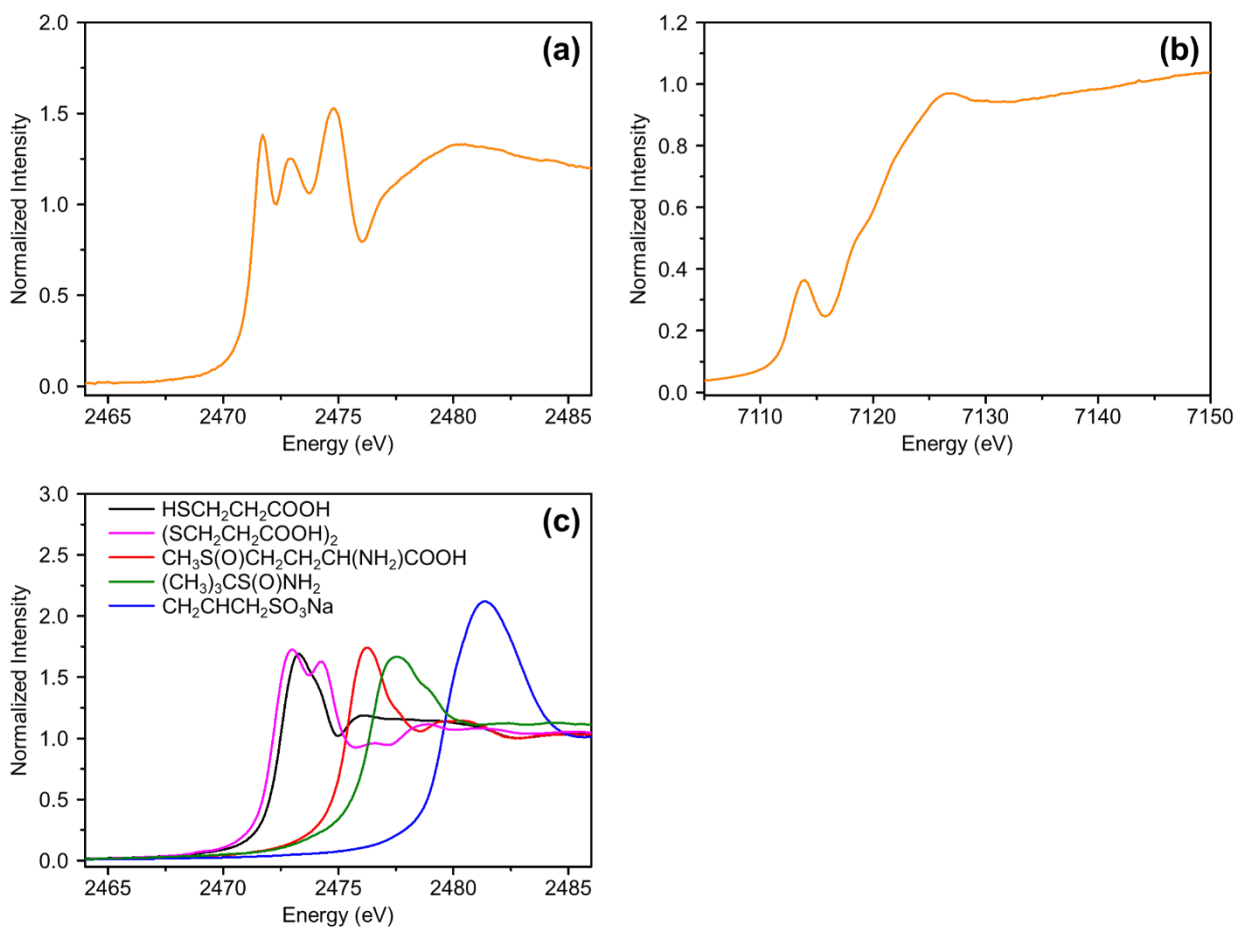


Figure S1. (a) S and (b) Fe K-edge X-ray absorption spectra for **DNIC-1**. (c) S K-edge X-ray absorption spectra for the HSCH₂CH₂COOH (black), (SCH₂CH₂COOH)₂ (magenta), CH₃S(O)CH₂CH₂CH(NH₂)COOH (red), (CH₃)₃CS(O)NH₂ (olive), and CH₂CHCH₂SO₃Na (blue).

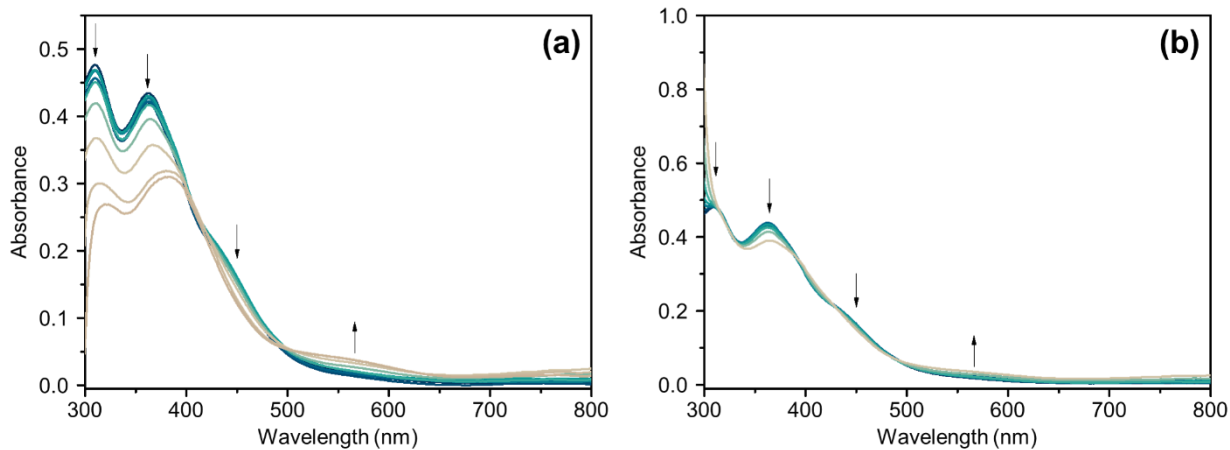


Figure S2. (a) UV-vis spectra for 50 μM of **DNIC-1** after addition of 0.25, 0.5, 0.75, 1.0, 1.25, 2.5, 5, 10, 25, and 50 mg/mL of bovine serum albumin (BSA) in PBS (pH 7.4). Change of UV-vis spectra during the titration was indicated by the arrow. Each UV-vis spectrum for this titration experiment was subtracted with the UV-vis spectrum of BSA at the corresponding concentration. (b) UV-vis spectra for 50 μM of **DNIC-1** after addition of 0.25, 0.5, 0.75, 1.0, 1.25, 2.5, 5, 10, 25, and 50 mg/mL of bovine serum albumin (BSA) in PBS (pH 7.4).

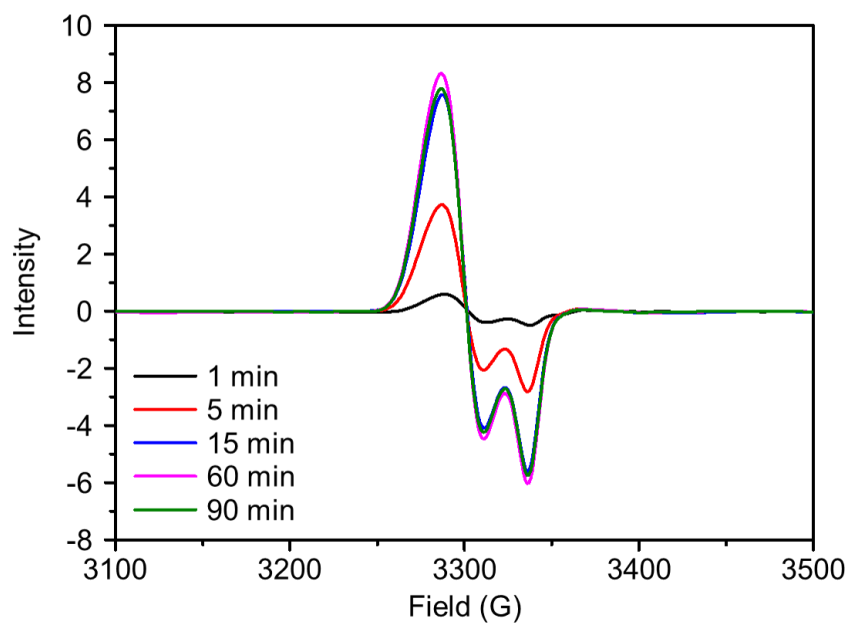


Figure S3. Reaction of **DNIC-1** and BSA monitored by EPR, which is performed under anaerobic condition.

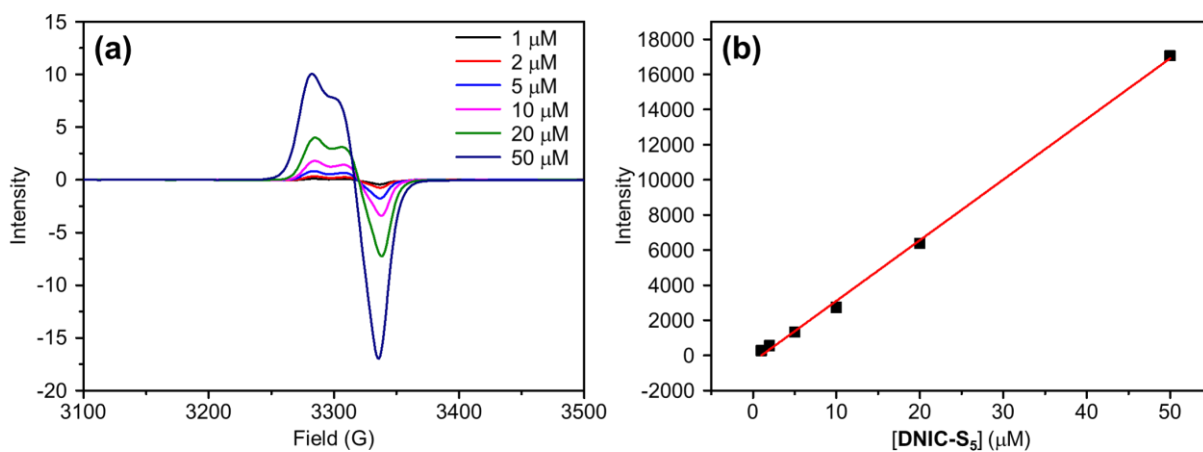


Figure S4. (a) EPR spectra for different concentration of [PPN][(NO)₂Fe(S₅)] (**DNIC-S₅**) in THF. (b) Calibration curve for **DNIC-S₅** in THF.

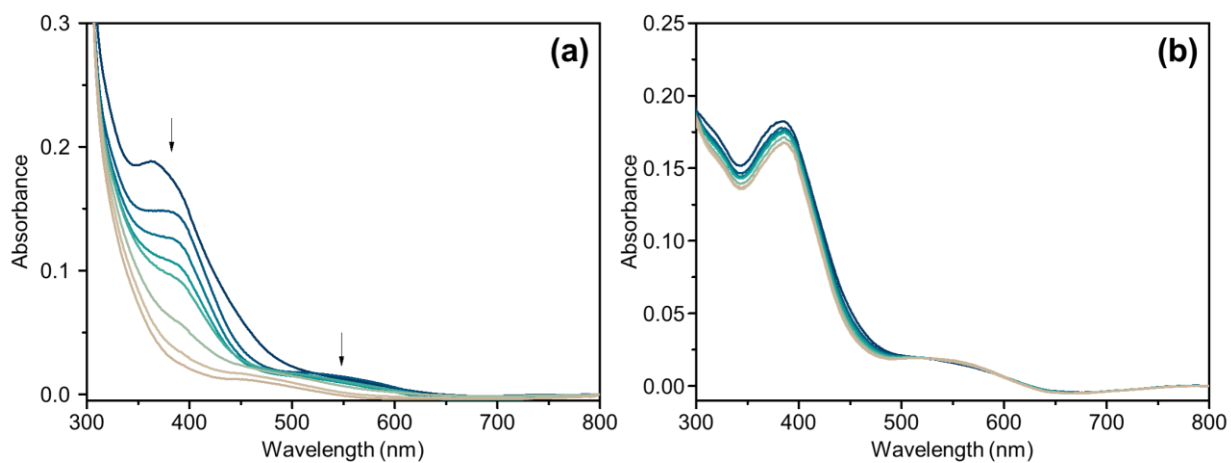


Figure S5. (a) Time-dependent change of UV-vis spectra for **DNIC-1** in the presence of BSA in PBS (pH 7.4) under aerobic condition at 37°C. (b) Time-dependent change of UV-vis spectra for **DNIC-1** in the presence of BSA in PBS (pH 7.4) during its incubation under anaerobic condition at 37°C for 24 h.

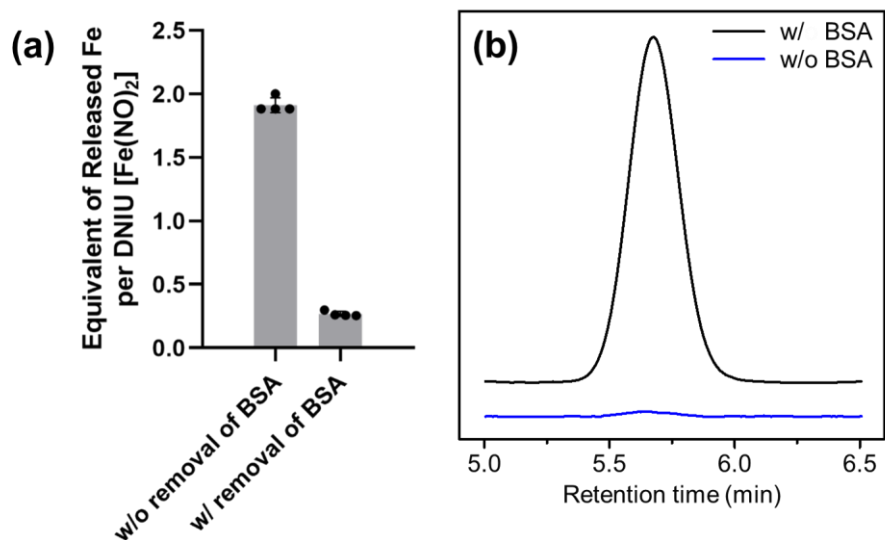


Figure S6. (a) Equivalent of released Fe per DNIU [Fe(NO)₂] after degradation of **DNIC-1** in the presence of BSA. ICP analysis on the supernatant solution was performed with or without removal of BSA adduct(s) by spin column. After removal of BSA adduct(s), significant decrease of equivalent of released Fe indicates the formation of BSA-bound Fe after degradation of **DNIC-1** in the presence of BSA. (b) GC chromatograms for the gaseous byproduct derived from decomposition of **DNIC-1** with (black) or without (blue) the presence of BSA in PBS (pH = 7.4).

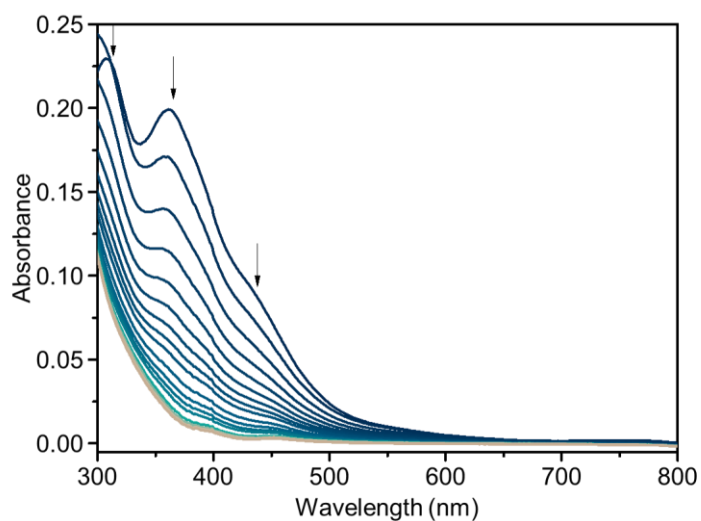


Figure S7. Time-dependent change of UV-vis spectra for 25 μM of **DNIC-1** in the presence of 150 μM of L-cysteine in PBS (pH 7.4) at 37°C under aerobic condition.

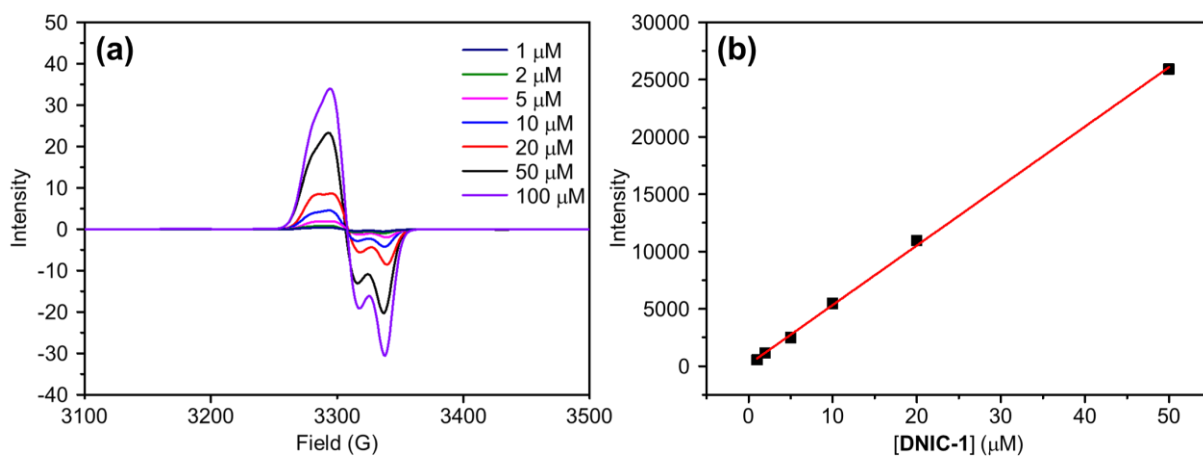


Figure S8. (a) EPR spectra for the isolated plasma after treatment with different concentration of **DNIC-1**. (b) Calibration curve for **DNIC-1** in the isolated plasma.

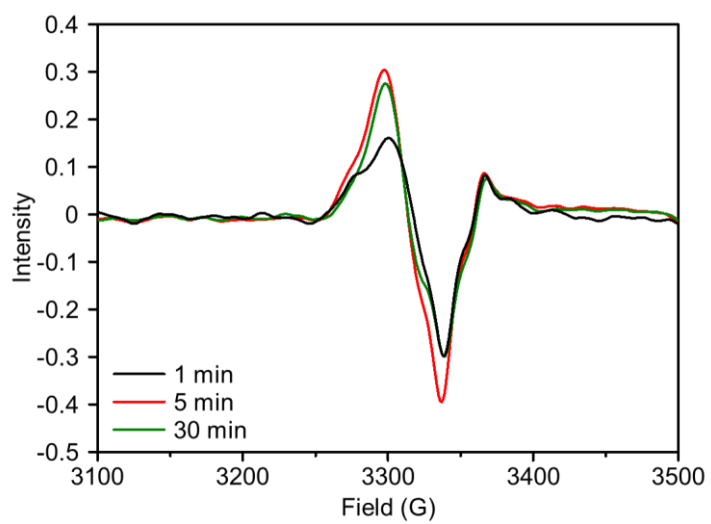


Figure S9. Reaction of **DNIC-1** and porcine stomach mucin in 100 mM phosphate buffer (pH 7.4) monitored by EPR, which is performed under aerobic condition.

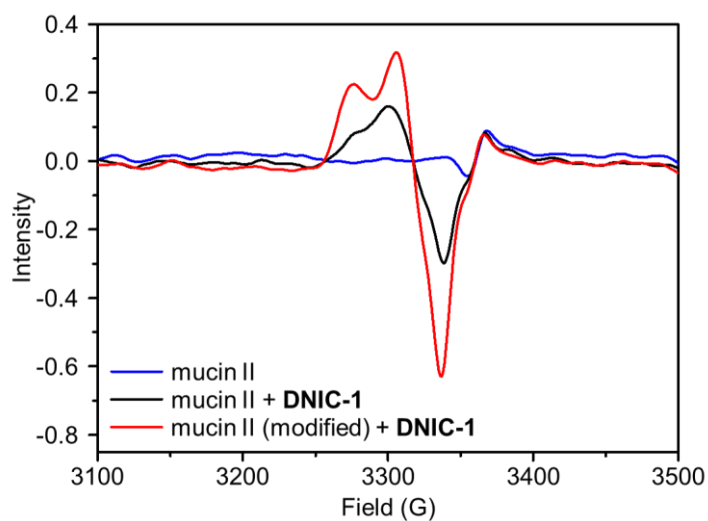


Figure S10. EPR spectra for reaction of **DNIC-1** with native (black) and NEM-modified mucin (red), respectively, in 100 mM phosphate buffer (pH 7.4). EPR spectrum of native mucin is shown in blue. Although the treatment of NEM fails to inhibit the formation of EPR-active and mucin-bound DNIC, the distinctive EPR spectra derived from reaction of **DNIC-1** with native and NEM-modified mucin, respectively, demonstrates the critical role of cysteine residue(s) for the formation of mucin-bound DNIC.

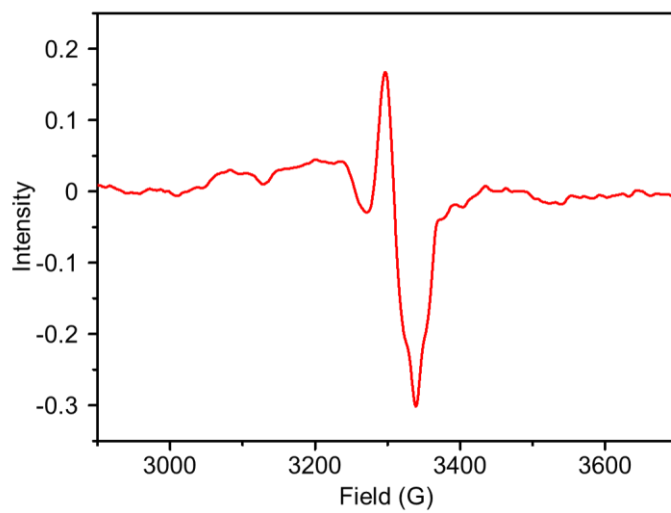


Figure S11. EPR spectrum for the stomach isolated from the mice with the oral administration of **DNIC-1**. This EPR spectrum for the stomach was collected after perfusion and removal of mucus.

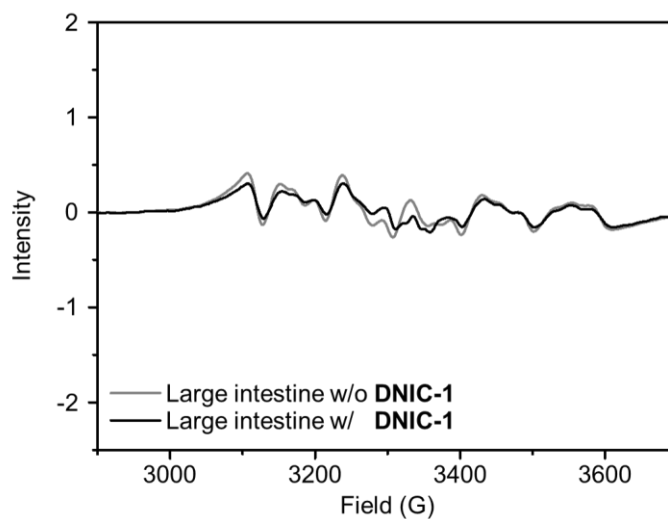


Figure S12. EPR spectra of large intestine isolated from the mice with (black) or without (gray) the oral administration of **DNIC-1**.

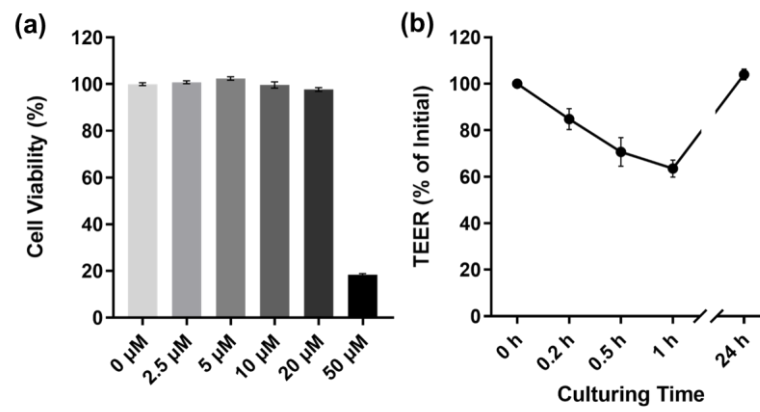


Figure S13. (a) Cell viability assay of the Caco-2 human intestinal epithelial cell treated with different concentration of **DNIC-1** for 24 h. (b) Effect of **DNIC-1** on time-dependent reduction of TEER in Caco-2 cell monolayer.

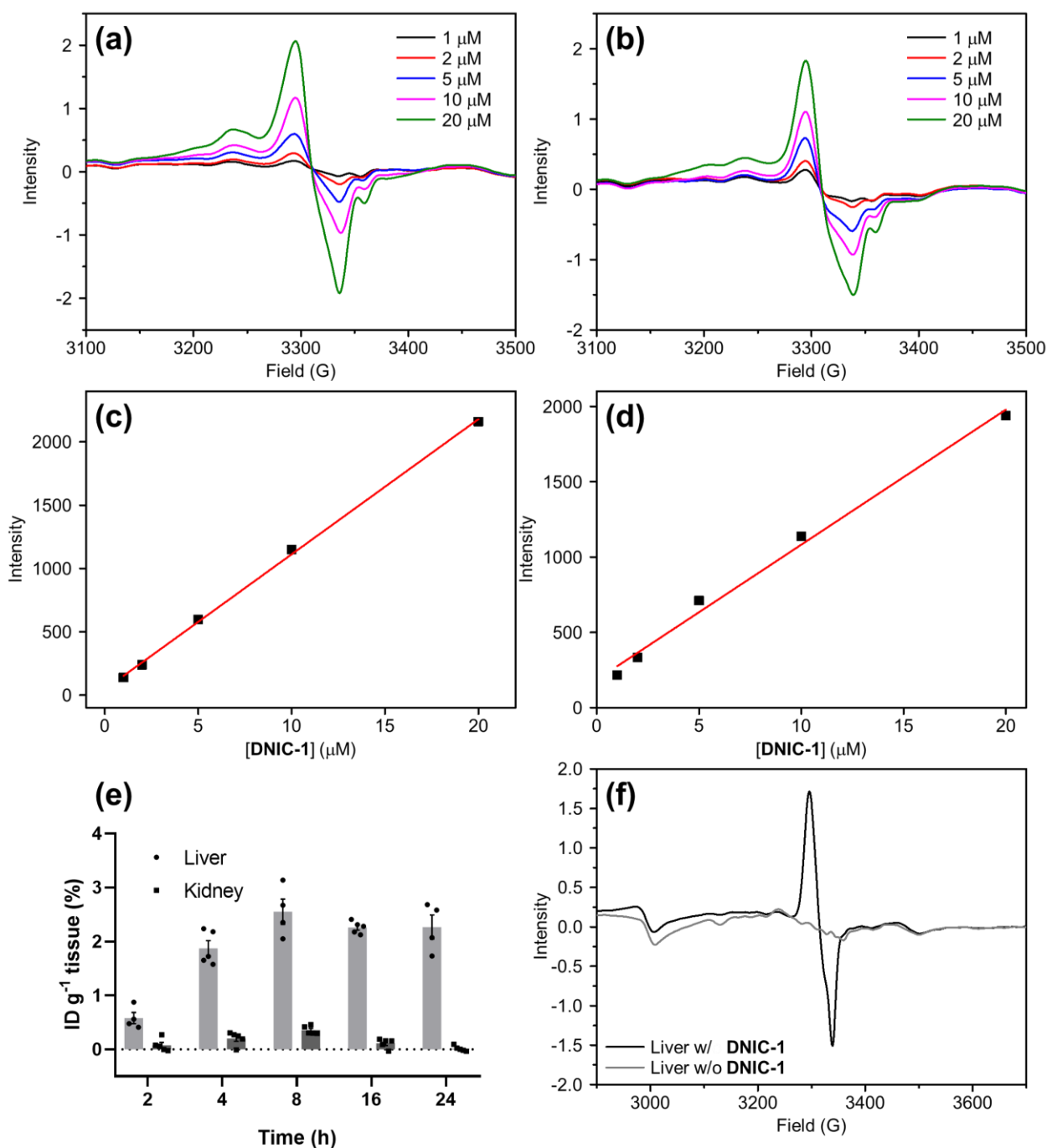


Figure S14. EPR spectra for different concentration of **DNIC-1** in the lysates of (a) liver and (b) kidney. Calibration curve for **DNIC-1** in the lysates of (c) liver and (d) kidney. (e) Time-dependent biodistribution study of **DNIC-1** in the liver (light gray) and kidney (dark gray). Data represent the mean \pm SEM (n=4-5). ID g^{-1} tissue, injected dose per gram of tissue. (f) EPR spectra of liver isolated from the mice with (black) or without (gray) the oral administration of **DNIC-1**. Based on the EPR signal at $g = 2.040, 2.028, 2.014$, formation of protein-bound DNIC in the liver is relevant to the biomimetic DNICs $[(\text{NO})_2\text{Fe}(\text{SR})(\text{L})]^-$ (Table S1, L = thiolate, amid, imidazolate, carboxylate, or phenoxide).

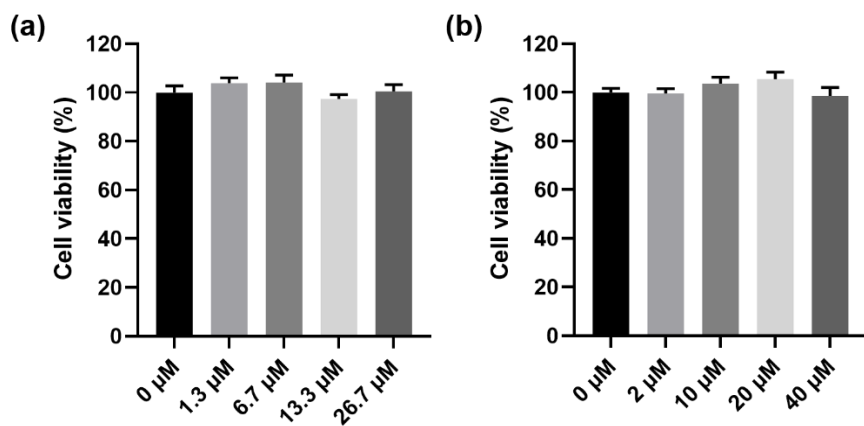


Figure S15. Cell viability assay of the N2A cells treated with different concentration of (a) 2-(*N,N*-diethylamino)-diazonolate 2-oxide (DEA NONOate) and (b) sodium nitroprusside (SNP), respectively, for 24. In this figure, the concentration of released nitric oxide from DEA NONOate (or SNP) is relevant to that from **DNIC-1** shown in Figure 4b.

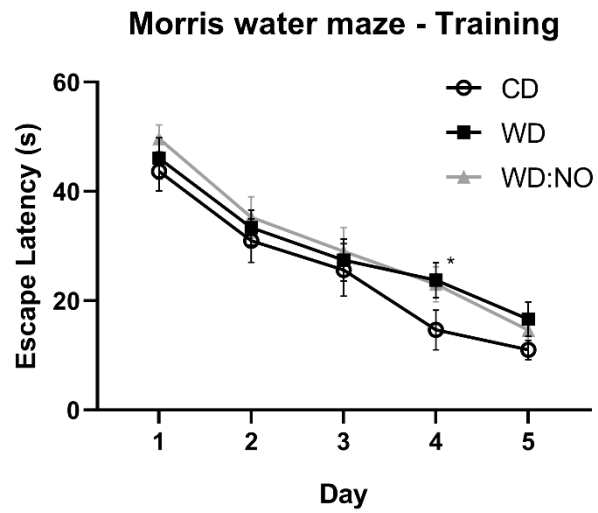


Figure S16. The time spent to reach the hidden platform (escape latency) during the acquisition phase of training (days 1-5 during week 17) in the Morris water maze task. *P < 0.05 compared with the CD.

Table S1. EPR Parameters for Biomimetic, Peptide-bound, and Protein-bound {Fe(NO)₂}⁹ DNICs.^a

Biomimetic {Fe(NO) ₂ } ⁹ DNICs Featuring Tetrahedral Geometry ^b			
DNIC	Ligation Mode	EPR Parameters ^c (g ₁ , g ₂ , g ₃)	Reference
[(NO) ₂ Fe(S(CH ₂) ₃ S)] ⁻	(S,S)	(2.048, 2.033, 2.015)	1
[(NO) ₂ Fe(S ₅)] ⁻	(S,S)	(2.049, 2.027, 2.015)	2
[(NO) ₂ Fe(SEt) ₂] ⁻	(S,S)	(2.042, 2.027, 2.014)	3
[(NO) ₂ Fe(S ^t Bu) ₂] ⁻	(S,S)	(2.039, 2.027, 2.013)	4
[(NO) ₂ Fe(SC ₆ H ₄ - <i>o</i> -NCOPh)] ⁻	(S,N)	(2.052, 2.033, 2.011)	5
[(NO) ₂ Fe(SEt)(Im)] ⁻	(S,N)	(2.037, 2.028, 2.014)	6
[(NO) ₂ Fe(S ^t Bu)(Im)] ⁻	(S,N)	(2.037, 2.027, 2.014)	6
[(NO) ₂ Fe(SC ₆ H ₄ - <i>o</i> -COO)] ⁻	(S,O)	(2.054, 2.036, 2.012)	5
[(NO) ₂ Fe(SC ₄ H ₃ S)(OPh)] ⁻	(S,O)	(2.038, 2.027, 2.013)	7
[(NO) ₂ Fe(Im) ₂] ⁻	(N,N)	(2.040, 2.022, 2.013)	6
[(NO) ₂ Fe(N ₃) ₂] ⁻	(N,N)	(2.043, 2.023, 2.012)	5
[(NO) ₂ Fe(C ₁₂ H ₈ N) ₂] ⁻	(N,N)	(2.029, 2.018, 2.011)	8
[(NO) ₂ Fe(N(Mes)(TMS)) ₂] ⁻	(N,N)	(2.022, 2.013, 2.005)	9
[(NO) ₂ Fe(OPh)(Im)] ⁻	(O,N)	(2.046, 2.021, 2.013)	7
[(NO) ₂ Fe(OPh) ₂] ⁻	(O,O)	(2.041, 2.022, 2.013)	7
[(NO) ₂ Fe(OPh-F) ₂] ⁻	(O,O)	(2.041, 2.024, 2.013)	7
Peptide-/Protein-bound {Fe(NO) ₂ } ⁹ DNICs			
Peptide/Protein	Ligation Mode	EPR Parameters	Ref.
L-cysteine	(S,S)	(2.04, 2.01) ^f	10
Glutathione	(S,S)	(2.04, 2.01) ^f	11
KCAACK	(S,S) ^d	(2.040, 2.032, 2.014)	12
KC(A) ₂ CK	(S,S) ^d	(2.039, 2.029, 2.014)	12
KC(A) ₃ CK	(S,S) ^d	(2.040, 2.030, 2.014)	12
KC(A) ₄ CK	(S,S) ^d	(2.040, 2.029, 2.014)	12
Aconitase	(S,S)	(2.037, 2.031, 2.012)	13
SoxR	(S,S) ^d	(2.03, 2.012) ^f	14
FNR	(S,S)	(2.038, 2.034, 2.016)	15
Rieske-type [2Fe-2S] cluster	(S,S)	(2.032, 2.014) ^f	16
Endonuclease III	(S,S)	(2.042, 2.036, 2.015)	17
Hybrid cluster protein	(S,S)	(2.042, 2.028, 2.014)	18
Bovine serum albumin	(S,S)	(2.044, 2.034, 2.014)	This work
	(S,N)	(2.05, 2.04, 2.01)	10
	(N,N)	(2.055, 2.035, 2.01)	10
Porcine stomach mucin	(S,S)	(2.038, 2.028, 2.014)	This work
	(S,N)/(S,O)	(2.052, 2.026, 2.015)	This work
Mammalian ferritin	(S, S)	(2.033, 2.014) ^f	19
	(N,N)	(2.055, 2.033, 2.015)	19
Mammalian ferrochelatase	(S, S)	(2.033, 2.014) ^f	20
Human glutathione transferase (P1-1)	(S,O) ^e	(2.038, 2.028, 2.015)	21

^aThe EPR parameter for DNIC [(NO)₂Fe(S₅)]⁻ was recorded at 4.2 K, whereas the EPR parameters for all the other DNICs collected in this Table were recorded at 77 K. ^bFive-coordinate and six-coordinate {Fe(NO)₂}⁹ DNICs featuring EPR signal at g_{av} = 2.012-2.018 are not included in this Table. ^cBiomimetic DNICs [(NO)₂Fe(SR)(L)]⁻ (L = thiolate, amido, imidazolate, carboxylate, or phenoxide) feature a g₂ value of 2.036-2.027, whereas biomimetic DNICs [(NO)₂Fe(L)₂]⁻ (L = imidazolate, azide, carbazolate, amide, and phenoxide) exhibit a smaller g₂ value of 2.024-2.013. ^dThe ligation mode is verified by EXAFS. ^eThe ligation mode is verified by single-crystal X-ray diffraction. ^fAxial EPR signal.

Table S2. Pharmacokinetic parameters for oral administration of DNIC-1 in C57BL/6JNarl mice.

	t_{1/2} (h)	CL (L/h)	VD (L)	Bioavailability
DNIC-1	0.94±0.19	0.29	0.32	6.5%

Table S3. Body Weight and Biochemical Parameters of Aging Mice under Control Diet (CD), Western Diet (WD), and Western Diet in Combination with Daily Treatment of DNIC-1 (WD:NO) for 16 Weeks.^a

Group	CD	WD	WD:NO
Body Weight (g)	32.26±4.11	43.37±8.45 ^{***}	40.51±6.32 ^{***}
Plasma glucose (mg/dL)	132.00±41.17	223.92±47.34 ^{***}	152.75±45.75 ^{##}
Serum TG (mg/dL) ^b	72.33±26.17	101.00±41.38	73.03±13.47
Serum TC (mg/dL) ^b	101.95±40.48	189.46±55.10 ^{***}	165.95±39.80 ^{**}
CRE (mg/dL) ^b	0.36±0.05	0.26±0.05 [*]	0.32±0.04
UA (mg/dL) ^b	2.38±0.27	2.70±0.25	3.62±0.38
GOT (U/dL) ^b	73.20±66.19	133.80±98.25	76.00±57.65
GPT (U/dL) ^b	30.80±14.57	36.20±9.55	33.60±10.36 ^{***, ##}

^aEach value represents the mean±SD (n = 11-12 for body weight in each group; n = 7-12 for plasma glucose, serum TG, and serum TC in each group; n = 5 for CRE, UA, GOT, and GPT in each group).

^{*}P < 0.05, ^{**}P < 0.01, and ^{***}P < 0.001 compared with the CD; ^{##}P < 0.01 compared with the WD.

^bTG = triacylglycerol, TC = total cholesterol, CRE = creatinine, UA = uric acid, GOT = glutamate oxaloacetate transaminase, GPT = glutamate pyruvate transaminase

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