#### **Supplementary Information**

# Direct Inhibition of the First PDZ Domain of ZO-1 by Glycyrrhizin is a Possible Mechanism of Tight Junction Opening of Caco-2 Cells.

### Authors

Emi Hibino<sup>1</sup>, Natsuko Goda<sup>1</sup>, Misaki Hisada<sup>1</sup>, Takeshi Tenno<sup>1,2</sup>, and Hidekazu Hiroaki<sup>1,2,\*</sup>.

<sup>1</sup>Laboratory of Structural Molecular Pharmacology, Graduate School of Pharmaceutical Sciences, Nagoya University, Furocho, Chikusa-ku, Nagoya, Aichi, Japan, 464-8601. <sup>2</sup>BeCellBar LLC., Business Incubation Building, Nagoya University, Furocho, Chikusaku, Nagoya, Aichi, Japan, 464-8601.

**Contents** Figure S1-5

Table S1-2

#### **1** Supplementary Information

2







- 7 chemical formula is shown in the main Figure 2C.
- 8 **B.** The horizontal axis is the mixing time for  ${}^{1}H{}^{-1}H$  NOESY and the vertical axis is the NOE
- 9 peak intensity. H5-H9, shown as a thick red line, was used as the reference.



# 11 Figure S2



- 13 A, B. <sup>1</sup>H-<sup>1</sup>H NOESY spectra of GL at concentrations of 2 mM (A) and 4 mM (B); blue and red
- 14 are the same and opposite sign as the diagonal peak, respectively.



### 16 Figure S3

- 17 The HADDOCK structure with the highest score without distance restrictions.
- 18 A. Only the CSP result from the NMR titration was added as a restriction to the calculations.
- The aglycone part is located away from the PDZ domain. ZO-1(PDZ1) is colored in yellowand GL in cyan.
- 21 B. No restriction was used for calculations. The glucuronate part is away from the PDZ domain.
- 22 C. Overlay of docking structures with (cyan) and without (magenta) restriction.
- 23 D, E. Enlarged view of the box regions is shown in C with ZO-1(1-PDZ1) (yellow) together.
- 24 The dotted lines show hydrogen bonds. In the case of docking without restriction, the
- 25 carbonyl group of GL is oriented in a direction where no hydrogen bond can be formed with
- 26 ZO-1(PDZ1).



# 28 Figure S4

- 29 No binding of ZO-1(PDZ1) to GA or glucuronic acid was detected.
- 30 A. The chemical formula of GA.
- 31 **B**. The chemical formula of glucuronic acid.
- 32 C. HSQC spectra of ZO-1(-PDZ1) with and without two equivalents of GA.
- 33 D. HSQC spectra of ZO-1(-PDZ1) with and without two equivalents of glucuronic acid.



### 35 Figure S5

34

36 TEER changes  $\lim_{n \to \infty} \frac{140}{120}$  daco-2 cells.

37 A. Experimental oprocedures. Pink arrows indicate the administrations of GL or DMSO. Details

- of the timing of TEER measurement compound administration are described in Experimental
  procedures.
- 40 **B**, **C**. Time course of TEER normalized to 100% of TEER before the initial administration. The
- 41 effect of GL administered extracelly arly was not significantly different from DMSO (B) and
- 42 continued administration of DCA weakened cells (C).

atoms	coefficient	distance (Å)
H12-H18	42038.55	2.38
Н5-Н9	39963.17	2.40
H9-H27	28631.79	2.54
H12-H19	11484.73	2.95
H12-H27	10627.50	2.99
H12-H28	10585.28	2.99
H1-H2'	9840.13	3.03
H5-H7	9581.74	3.04
H15-H18	7789.93	3.15
H18-H22	6503.35	3.25
H18-H28	4419.91	3.46

# 44 Table S1

45 The slope of the fitted line and the calculated distance between the hydrogen atoms. This was

46 given as restriction for the calculation in the HADDOCK program.

Restriction	glycyrrhizin	+	-	-
	mZO1-PDZ1	+	+	_
HADDOCI	K Score (a.u.)	-47.8 +/- 1.8	-51.2 +/- 1.1	-38.7 +/- 1.6
Clust	ter Size	5	5	4
RMSD from the o stru	verall lowest-energy icture	0.1 +/- 0.1	0.2 +/- 0.1	0.6 +/- 0.1
Van der W	Vaals energy	-37.2 +/- 2.2	-45.1 +/- 1.2	-21.2 +/- 0.4
Electrost	atic energy	-136.2 +/- 16.7	-108.4 +/- 4.0	-166.8 +/- 3.0
Desolvat	ion energy	3.0 +/- 0.6	4.6 +/- 0.4	3.7 +/- 0.5
Restraints vi	olation energy	0.4 +/- 0.2	1.1 +/- 0.7	0.0 +/- 0.0
Buried Su	ırface Area	784.6 +/- 17.0	866.1 +/- 4.9	518.0 +/- 12.8
Z-9	Score	-1.5	-2.2	-1.6
Fi	gure	3	S3A	S3B

48 Table S2

49 The obtained parameters in the calculation of the HADDOCK simulation.