

**Figure S1**. mRNA expression changes of ENaC subunits (-α, -β, -γ) in HT-29/B6-GR/MR cells 48 hours after *Camplyobacter concisus* infection. (A) mRNA expression of α-ENaC gene (*SCNN1A*) in controls, DBA-stimulated controls and *C. concisus*-infected cell monolayers (n = 4 each, ns = not significant, *p*>0.05). (B) mRNA expression of β-ENaC gene (*SCNN1B*) in controls, DBA-stimulated controls and *C. concisus*-infected cell monolayers (n = 4 each, \*\*\**p* < 0.01, \*\**p* < 0.01). (C) mRNA expression of γ-ENaC gene (*SCNN1G*) in controls, DBA-stimulated controls and *C. concisus*-infected cell monolayers (n = 4 each, \*\**p* < 0.01, \**p* < 0.05). For statistical analysis, one-way ANOVA with Bonferroni’s Multiple Comparison Test was used and *p* < 0.05 was considered statistically significant. DBA = Dexamethasone, Butyrate, and Aldosterone.