

Supplementary Information

Table S1 Composition of the experimental diets (%)

| Ingredients (%) | Normal | Model | DHA-PL* | EPA-PL* | DHA-EE* | EPA-EE* |
|---------------------|--------|-------|---------|---------|---------|---------|
| Corn Starch | 46.57 | 46.57 | 46.57 | 46.57 | 46.57 | 46.57 |
| Dextrin | 15.50 | 15.50 | 15.50 | 15.50 | 15.50 | 15.50 |
| Casein-Vitamin Free | 14.00 | 14.00 | 14.00 | 14.00 | 14.00 | 14.00 |
| Sucrose | 10.00 | 10.00 | 10.00 | 10.00 | 10.00 | 10.00 |
| Powdered Cellulose | 5.00 | 5.00 | 5.00 | 5.00 | 5.00 | 5.00 |
| Soybean Oil | 4.00 | 4.00 | 2.00 | 2.00 | 3.39 | 3.35 |
| AIN 93M mineral Mix | 3.50 | 3.50 | 3.50 | 3.50 | 3.50 | 3.50 |
| AIN 93 Vitamin Mix | 1.00 | 1.00 | 1.00 | 1.00 | 1.00 | 1.00 |
| Choline Bitartrate | 0.25 | 0.25 | 0.25 | 0.25 | 0.25 | 0.25 |
| L-Cystine | 0.18 | 0.18 | 0.18 | 0.18 | 0.18 | 0.18 |
| DHA-PL | - | - | 2.00 | - | - | - |
| EPA-PL | - | - | - | 2.00 | - | - |
| DHA-EE | - | - | - | - | 0.61 | - |
| EPA-EE | - | - | - | - | - | 0.65 |

* The mass concentration of DHA or EPA in DHA-PL, EPA-PL, DHA-EE, EPA-EE were 269 mg/g, 255 mg/g, 889 mg/g, 789 mg/g, respectively.

Table S2 Primer sequences for RT-PCR amplification

| Gene | Forward | Reverse |
|---------------|-------------------------------|-------------------------------|
| IFN- γ | CACTGCATCTTGGCTTTGCA | GCTGATGGCCTGATTGTCTTTC |
| TNF- α | GGCAGGTCTACTTTGGAGTCATTG C | ACATTCGAGGCTCCAGTGAATTCG G |
| IL-1 β | ACCTGCTGGTGTGTGACGTT | TCGTTGCTTGGTTCTCCTTG |
| IL-6 | AGCCAGAGTCCTTCAGAGAGAT | GCACTAGGTTTGCCGAGTAGAT |
| FFAR 2 | CCGCCCAAGGCTAAGTCAATA | CGCCAGGGTCAGATTAAGCA |
| Muc-2 | CACACAGCGGCCTTTCTCAT | ACCCTCCTCCTACCACATTG |
| IL-10 | GGCGCTGTCATCGATT | AGCTCTGTCTAGGTCC |

Table S3 Primer sequences for fecal microbiota analysis

| Primer | Sequence |
|---------------------------------|--------------------------|
| PRBA338f | ACTCCTACGGGAGGCAGCAG |
| PRUN518r | ATTACCGCGGCTGCTGG |
| <i>Bifidobacteria</i> F | GGGATGCTGGTGTGGAAGAGA |
| <i>Bifidobacteria</i> R | TGCTCGCGTCCACTATCCAGT |
| <i>Lactobacilli</i> F | AGCAGTAGGGAATCTTCCA |
| <i>Lactobacilli</i> R | CACCGCTACACATGGAG |
| <i>Clostridium coccooides</i> F | CGGTACCTGACTAAGAAGC |
| <i>Clostridium coccooides</i> R | AGTTT(C/T)ATTCTTGCGAACG |
| <i>Roseburia spp.</i> F | GCGGTRCGGCAAGTCTGA |
| <i>Roseburia spp.</i> R | CCTCCGACACTCTAGTMCGAC |
| <i>Enterococcus spp.</i> F | CCCTTATTGTTAGTTGCCATCATT |
| <i>Enterococcus spp.</i> R | ACTCGTTGTACTTCCCATTGT |
| <i>Prevotella spp.</i> F | GGTGTCCGGCTTAAGTGCCAT |
| <i>Prevotella spp.</i> R | CGGA(C/T)GTAAGGGCCGTGC |
| <i>Bacteroides fragilis</i> F | TGATTCCGCATGGTTTCATT |
| <i>Bacteroides fragilis</i> R | CGACCCATAGAGCCTTCATC |
| <i>Enterobacteriaceae</i> F | GTGCCAGCMGCCGCGGTAA |
| <i>Enterobacteriaceae</i> R | GCCTCAAGGGCACAACCTCCAAG |

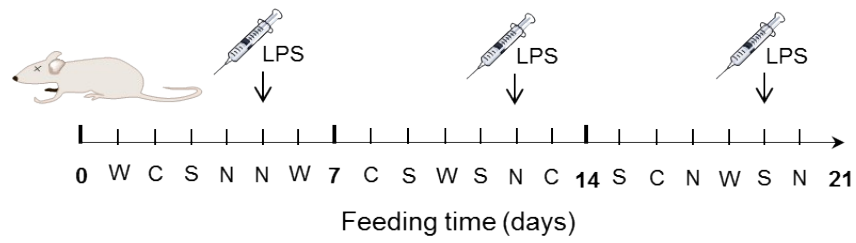


Fig. S1 Schematic illustration of the induction of chronic stress model. W-wet bedding; C-chronic immobilization for 2 h; S-swimming in 24°C water for 25 min; N- noises for 2 h.

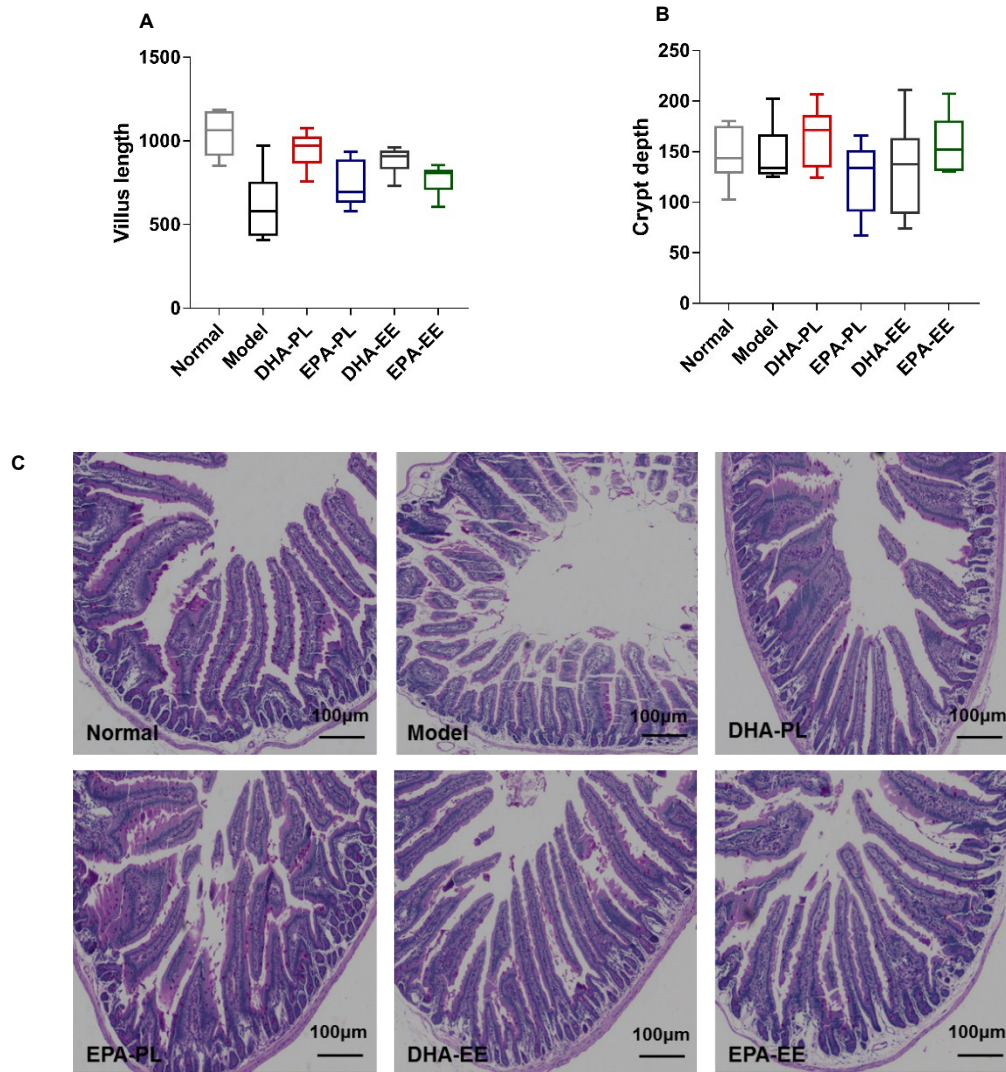


Fig. S2 DHA-PL and EPA-PL improved the histological morphology of small intestine in mice under chronic exercise. (A) Villus length, (B) Crypt depth, (C) Alcian Blue Staining of mucins. Data were mean \pm SEM (n = 10).