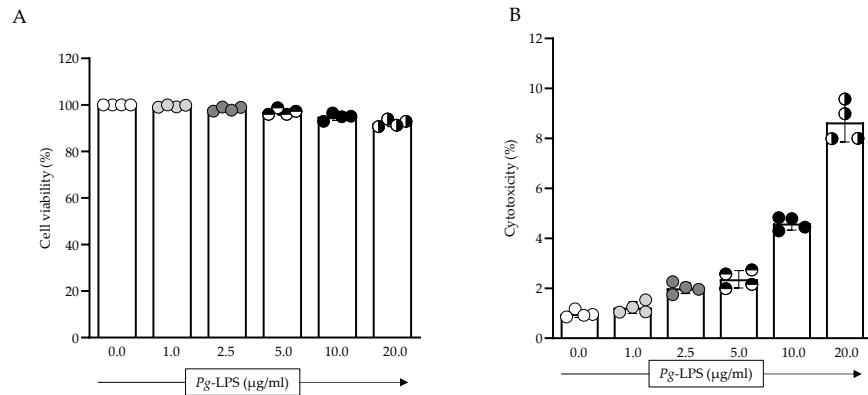
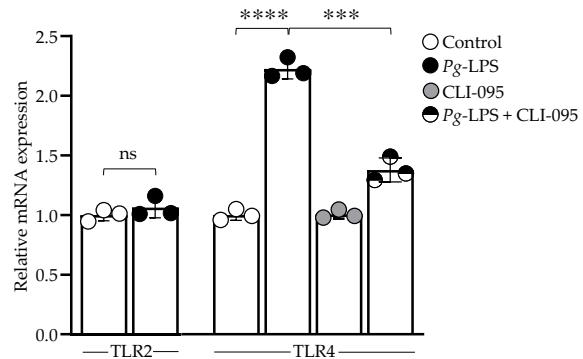


Supplementary Data

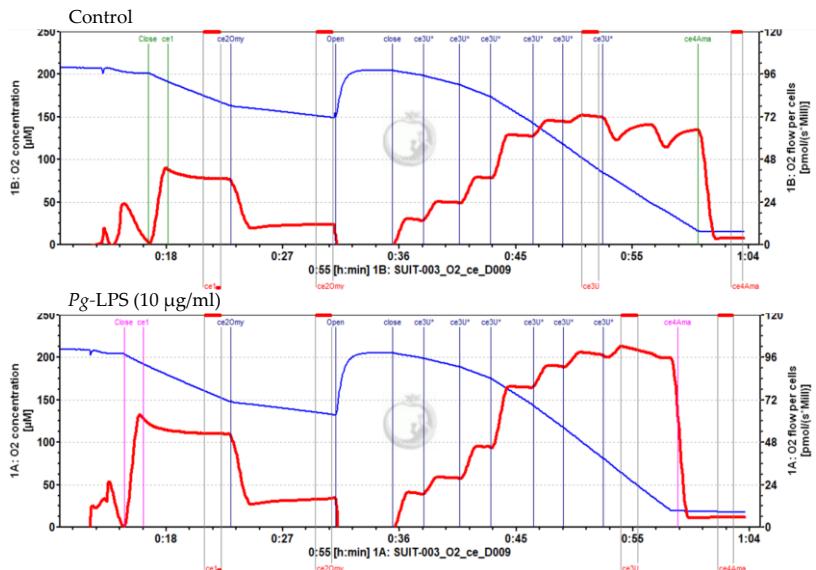


Supplementary Figure S1. Effect of *P. gingivalis*-LPS on cells viability with MTS assay (A) and cytotoxicity with LDH assay (B) in SH-SY5Y cells treated with 0, 1.0, 2.5, 5.0, 10.0 and 20.0 $\mu\text{g/ml}$ concentrations of *P. gingivalis*-LPS for 24 hours and untreated cells were used as control.

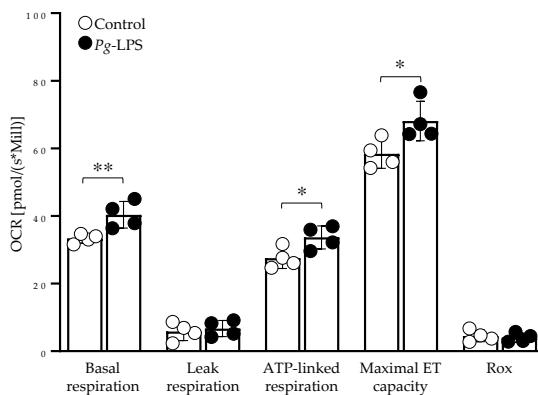


Supplementary Figure S2. Effects of *P. gingivalis*-LPS on mRNA expression of TLR2 and TLR4. RT-qPCR analysis showing the gene expression of TLR2 in untreated control and cells treated with 10 $\mu\text{g/ml}$ concentration of *P. gingivalis*-LPS, and TLR4 gene expression in untreated control, cells treated with 10 $\mu\text{g/ml}$ concentration of *P. gingivalis*-LPS, CLI-095 (1 μM), *P. gingivalis*-LPS + CLI-095 (1 μM) (n=3). ***p < 0.001, ****p < 0.0001, ns: p > 0.05.

A



B



Supplementary Figure S3. High-resolution respiratory analysis of oxidative phosphorylation in Oroboros O2K respirometer. (A) OCR and (B) Basal respiration, leak respiration, ATP-linked respiration, Maximum ET capacity was measured from intact SH-SY5Y cells treated with 10 $\mu\text{g}/\text{ml}$ *P. gingivalis* LPS for 24 hours and untreated cells were used as control ($n=4$). * $p < 0.05$, ** $p < 0.01$, ns: $p > 0.05$.

Supplement Table S1. List of primer sequences used for qPCR analysis.

Primer Name	Primer Sequence (5'-3')
Tau-F	5'-GATTGGGTCCCTGGACAATA-3'
Tau-R	5'-GTGGTCTGTCTTGGCTTG-3'
VEGF-F	5'- TGCAGATTATGCGGATCAAAC C-3'
VEGF-R	5'-TGCATTCACATTGTTGTGCTGTAC-3'
TGF- β -F	5'- GCGTGCTAATGGTGGAAACC-3'

<i>TGF-β</i> -R	5'- CGGAGCTCTGATGTGTTGAAGA-3'
<i>IL-1β</i> -F	5'-ATGGGATAACGAGGCTTATGTG-3'
<i>IL-1β</i> -R	5'-CAAGGCCACAGGTATTTGTC-3'
<i>IL-6</i> F	5'-ACTTGCCTGGTAAAATCAT-3'
<i>IL-6</i> R	5'-CAGGAACCTGGATCAGGACTT-3'
<i>TNF-α</i> -F	5'-TCAGCAAGGACAGCAGAGG-3'
<i>TNF-α</i> -R	5'-CAGTATGTGAGAGGAAGAGAACCC-3'
<i>iNOS</i> -F	5'-GCAGAAATGTGACCACATGG-3'
<i>iNOS</i> -R	5'-ACAACCTTGGTGTGAAGGC-3'
<i>TLR2</i> -F	5'- CTTCACTGCTTCAACTGGTA -3'
<i>TLR2</i> -R	5'-TTGCGGTACAAGACAGAG -3'
<i>TLR4</i> -F	5'-GACTTGCGGGTTCTACATCA-3'
<i>TLR4</i> -R	5'-GAGGTGGCTTAGGCTCTGATA-3'
<i>SRF</i> -F	5'- GGGCCGCGTGAAGATCAA-3'
<i>SRF</i> -R	5'- GTCAGCGTGGACAGCTCATA-3'
<i>p49/STRAP</i> -F	5'- GAAGACACCTGCTGACCCAA-3'
<i>p49/STRAP</i> -R	5'- TCTTCTCCGCCATCACTGTT-3'
<i>PGC-1α</i> -F	5'- ACCCACAGAGAACAGAAACAG-3'
<i>PGC-1α</i> -R	5'- GGGTCAGAGGAAGAGATAAAGTTG-3'
<i>PGC-1β</i> -F	5'- ACTACTCGCTGACACGCAG-3'
<i>PGC-1β</i> -R	5'- CTCTGAGTTCTCTGGGCACC-3'
<i>NRF1</i> -F	5'- GGAAACGGCCTCATGTATTG-3'
<i>NRF1</i> -R	5'- GTTTGGAGGGTGAGATAACAGAG-3'
<i>TFam</i> -F	5'- TTTCTCCGAAGCATGTGGG-3'
<i>TFam</i> -R	5'-GCCAAGACAGATGAAAACCAC -3'
<i>MFN1</i> -F	5'- TGGCATCTGTGGCCGAGTT-3'
<i>MFN1</i> -R	5'-GAAACAGGTTCTGCCATTATGCT -3'
<i>MFN2</i> -F	5'- CGCGCTTATCCACTTCCCTC-3'
<i>MFN2</i> -R	5'- AGAAGAGCAGGGACATTGCG-3'
<i>Fis1</i> -F	5'- TGACATCCGTAAGGCATCG-3'
<i>Fis1</i> -R	5'-CTTCTCGTATTCCCTGAGCCG -3'
<i>Opa1</i> -F	5'- GGAGAACCATATTGTTTGACC-3'
<i>Opa1</i> -R	5'- AGAGCTGTCCTCTTCCCTG-3'
<i>NDUFV1</i> -F	5'- CGGGTATCTGTGCGTTTCAG-3'
<i>NDUFV1</i> -R	5'- GGTCTTCATCCTTCAGCGAG-3'
<i>NDUFV2</i> -F	5'- GGCAAAATCCCCAAAACCAGG-3'
<i>NDUFV2</i> -R	5'- TGCTTGTACACCAAATCGAGG-3'
<i>NDUFAB1</i> -F	5'-TCCAGGACCGTGTCTTACG -3'
<i>NDUFAB1</i> -R	5'- GGTCCAAGACTGTCTAAGCCC-3'
<i>NDUFS1</i> -F	5'- ACTCTGACACCTTATGCACTG-3'
<i>NDUFS1</i> -R	5'- AACATCTGGCTCTCCACAC-3'

F: forward primer; R: reverse primer.