## Epstein-Barr Virus LMP-1 Elevates HIF-1 $\alpha$ Gene Transcription in Nasopharyngeal Carcinoma Cells

Wei-Wen Sung<sup>1</sup>, Yi-Chih Chu<sup>2</sup>, Peir-Rong<sup>3</sup> Chen, Ming-Hui Liao<sup>1</sup>, Jeng-Woei Lee<sup>4,\*</sup>

<sup>1</sup>Department of Veterinary Medicine, National Pingtung University of Science and Technology, Pingtung, Taiwan; <sup>2</sup>Department of Molecular Biology and Human Genetics, <sup>4</sup>Department of Life Science, Tzu-Chi University, Hualien, Taiwan; <sup>3</sup>Department of Otolaryngology, Hualien Tzu-Chi Medical Center, Hualien, Taiwan

Hypoxia inducible factor-1 (HIF-1) is a transcriptional factor composed of a constitutive expressed  $\beta$  and an oxygen-dependent  $\alpha$  subunit, which plays a pivotal role in carcinogenesis and metastases. Accumulating evidences reveal elevated expression of HIF-1 $\alpha$  in various tumors, including nasopharyngeal carcinoma (NPC). However, the expression mechanism is still unclear. Here, we showed that transactivation activity of HIF-1 was remarkably induced by Epstein-Barr virus (EBV) oncoprotein, latent membrane protein 1 (LMP1), via up-regulated  $\alpha$  gene expression in NPC cells. Upon LMP1 induction, we also displayed enhanced promoter activity as well as RNA stability of HIF-1 $\alpha$  gene though CTARs (C-terminal of activation regions) of LMP1. Taken together, our data suggest an alternative pathway of HIF-1 $\alpha$  regulation by viral protein under normoxic condition. The detail mechanisms should be further investigated.