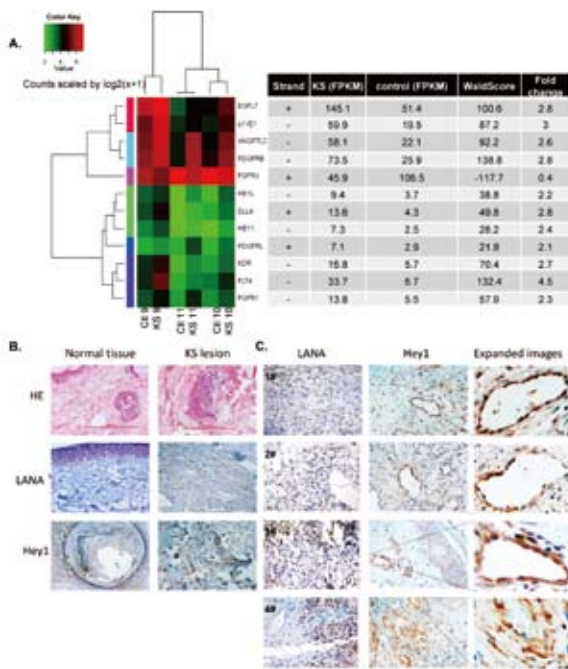


# Novel Mechanism for Kaposi's Sarcoma-Associated Herpesvirus Mediated Oncogenesis

On February 12, 2014, the international academic journal *Cancer Research* published an article online entitled "Latency Associated Nuclear Antigen of Kaposi's Sarcoma Associated Herpesvirus Promotes Angiogenesis through Targeting Notch Signaling Effector Hey1", as a result from the research conducted by Prof. LAN Ke's lab at the Institut Pasteur of Shanghai, Chinese Academy of Sciences.

Notch signaling is implicated in the pathogenesis of Kaposi's sarcoma (KS), which is an angioproliferative neoplasm that originates from Kaposi's sarcoma-associated herpesvirus (KSHV) infection. In their previous research, the team revealed that the KSHV LANA protein can stabilize intracellular Notch in KSHV-infected tumor cells and promote cell proliferation. However, Notch signaling functions in pathological angiogenesis of KS remained largely unknown. Hey1, an essential downstream effector of the Notch signaling pathway, has been demonstrated in the literature to play a fundamental role in vascular development. In the present study, researchers perform whole transcriptome, paired-end sequencing on three patient-matched clinical KS specimens and their corresponding adjacent stroma samples, with an average depth of 42 million reads per sample, and reveal that *Dll4*, *Hey1* and *HeyL* display significant upregulation in KS. Further verification based on immunohistochemistry analysis proves that *Hey1* is highly expressed in KS lesions. Using the matrigel plug assay, down-regulation of *Hey1* and gamma-secretase inhibitor (GSI) treatment causes dramatic reduction in the formation of new blood vessels in mice. Interestingly, LANA is responsible for the elevated level of *Hey1* through inhibition of its degradation. More importantly, *Hey1* stabilized by LANA promotes the neoplastic vasculature. The data suggest that hijacking of the pro-angiogenic property of *Hey1* by LANA is an important strategy utilized by KSHV to achieve pathological angiogenesis.



LANA and Hey1 are highly expressed in pathological vascular endothelial cells in KS lesions. (A) Heat map representation of differentially regulated genes performed on RNA extracted from three KS lesions and their patient-matched normal tissues. The genes were ranked according to absolute values of log2 relative change. (B) LANA and Hey1 were highly specifically distributed in KS tumor rather than its adjacent stromal tissue on IHC analysis. Original magnification,  $\times 5$  in HE and  $\times 40$  in IHC. (C) IHC analysis of LANA and Hey1 co-localization in pathological vessels in KS lesions. Original magnification,  $\times 40$  in all examinations.

The research was supervised by Prof. LAN Ke in collaboration with Prof. WEN Hao from Xinjiang Medical University and was completed by postdoctoral fellow WANG Xing and other group members. The study was supported by the Key Project of Natural Science Foundation of China, the National Basic Research Program (973) and Sanofi-Aventis-SIBS scholarship.

The link for this article: <http://cancerres.aacrjournals.org/content/early/2014/02/12/0008-5472.CAN-13-1467.long>