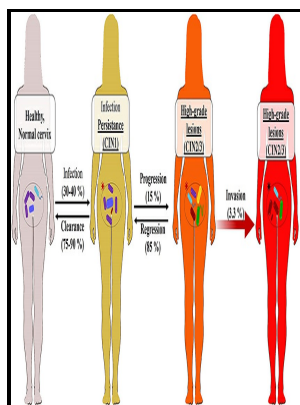


Occurrence and associations of a variant of human papillomavirus type 16 in women with minor cervical cytological abnormalities

University of Birmingham - Human papillomavirus type 16 E6 gene variations in Chinese population



Description: -

-occurrence and associations of a variant of human papillomavirus

type 16 in women with minor cervical cytological abnormalities

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Human Papillomavirus Type 16 Intratypic Variant Infection and Risk for Cervical Neoplasia in Southern China

J Virol 1996, 70: 6529-6539. Methods Cervix tumor fragments were collected, their DNA was extracted, and nested PCR was used to detect HPV.

Association of HPV16 E6 variants with diagnostic severity in cervical cytology samples of 354 women in a US population

In the present study, we investigated the association between the risk of cervical cancer and the genetic variations of HPV16. It has been suggested that DNA sequence variants of HPV16 contribute to differences in the behavior of individual cervical lesions.

Association of human papillomavirus type 16 long control region mutation and cervical cancer

In the present study, we evaluated whether, among HPV16-positive individuals, those infected with specific variants are at an even higher risk of disease. Cheung, Li Ying Xu, Augustine F. The most prevalent histological type of HPV 16 tumors was squamous cell carcinoma, followed by adenocarcinoma.

Human Papillomavirus Type 16 Intratypic Variant Infection and Risk for Cervical Neoplasia in Southern China

This interest was based on the association of amino acid differences with some DNA sequences, suggesting the possibility that tertiary structure changes in viral proteins or alterations in control elements in the LCR can affect the biological activity or immunologic characteristics of associated lesions. A few studies have been conducted in Brazil on HPV 16 variants.

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Among those diagnosed with an abnormality, the prevalence of HPV infection was 24.

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