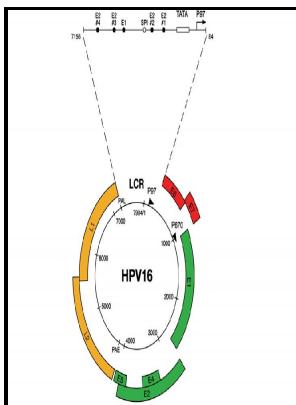


Occurrence and associations of a variant of human papillomavirus type 16 in women with minor cervical cytological abnormalities

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



Description: -

- occurrence and associations of a variant of human papillomavirus type 16 in women with minor cervical cytological abnormalities
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Notes: Thesis (M.D.) - University of Birmingham, Department of Obstetrics and Gynaecology, Faculty of Medicine and Dentistry, 2000.

This edition was published in 1999



Filesize: 33.81 MB

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

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

Martha Hutchinson Womens and Infants' Hospital, Providence, RI for their careful review of cytology specimens, Dr. The substitution s found in variants are indicated by a number for the position; a letter preceding it refers to the substituted nucleotide.

Role of HPV 16 variants among cervical carcinoma samples from Northeastern Brazil

Period and location of study The study was conducted from January 2016 to December 2017, at the High Complexity Care Unit UNACON of the Cancer State Hospital of Maranhão and in the High Complexity Care Center in Oncology CACON of the Aldenora Bello Cancer Hospital.

Role of HPV 16 variants among cervical carcinoma samples from Northeastern Brazil

Suthipintawong C, Siriaunkul S, Tungsirinunkong K, Pientong C, Ekalaksananan T, Karalak A, Kleebkaow P, Vinyuvat S, Triratanachat S, Khunamornpong S, Chongsuwanich T: Human papilloma virus prevalence, genotype distribution, and pattern of infection in thai women. Human papilloma virus HPV prophylactic vaccination: challenges for public health and implications for screening.

Role of HPV 16 variants among cervical carcinoma samples from Northeastern Brazil

This proposal does not negate the aggregate evidence that NE variants, particularly AA, have an increased risk for persistence and progression to cancer.

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

Analysis of HPV16As LCR polymorphisms Forty-seven DNA samples from the HSIL and SCC cases were classified as HPV16As and investigated for LCR polymorphisms.

Prevalence of human papillomavirus infection among women from quilombo communities in northeastern Brazil

J Virol 1997, 71: 2463-2472. Lace MJ, Isacson C, Anson JR, Lorincz AT, Wilczynski SP, Haugen TH, Turek LP: Upstream regulatory region alterations found in human papillomavirus type 16 HPV-16 isolates from cervical carcinomas increase transcription, ori function, and HPV immortalization capacity in culture.

Prevalence of human papillomavirus infection among women from quilombo communities in northeastern Brazil

In the US, Xi et al demonstrated strong evidence of increased association of NE variants, particularly AA, with CIN3 using the ALTS population that was derived from four different centers in the US. Pinto DS, Fuzii HT, Quaresma JAS. The data about the health of the Quilombola population are scarce, especially in relation to prevalence, genotypes, and risk factors for HPV infection.

Prevalence of human papillomavirus type 16 and its variants in abnormal squamous cervical cells in Northeast Thailand

The aim of this study was to understand further the oncogenic potential of HPV16As. According to Akarolo-Anthony et al. Only seven cases 2% showed the As pattern which is included in the EV category.

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