HPV and pathogenesis of SIL

Behrooz Shokouhi, MD ACP

cancer of the cervix

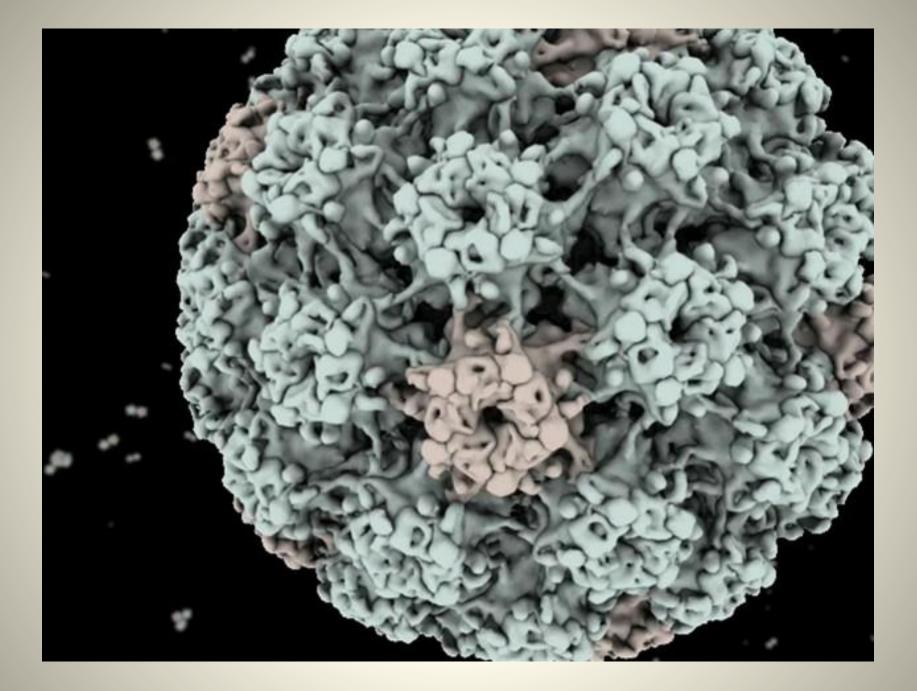
- a burden of human suffering and mortality
 is disproportionate to its size
- due to the susceptibility of the epithelium of the cervical transformation zone to infection by oncogenic HPV
- cervical screening programs
- Vaccination against HPV
- remains one of the most common cancers

HPV and the Lower Female Genital Tract

a rapidly advancing field

The role of HPV in the pathogenesis

- of both condyloma and carcinoma
- Relatively recently was unsuspected
- greater than 99% of cervical carcinomas are associated with oncogenic HPV
- the molecular mechanisms of virus-mediated oncogenesis are understood in considerable detail



HPV, a DNA virus

- is classified on the viral genome into different genotypes
- HPV-2 is the most common cause of verruca vulgaris
- the genital HPV genotypes, transmitted by sexual contact
 - associated with condyloma, the low-risk HPV genotypes, 6 & 11
 - associated with cervical carcinoma, the high-risk genotypes (oncogenic), the risk varies, 16 & 18
 - -possibly oncogenic, of uncertain oncogenic potential

Table 1 Papillomavirus types in genital lesions			
Type of genital lesion	HPV type Less prevalent	More prevalent	
Condylomata acuminata	42,44,51,53,83	6,11	
Intraepithelial neoplasias	6,11,18,26,30,31,33,34,35,39,40,42,43, 45,51,52,53,54,55,56,57,58,59,61,62,64, 66,67,68,69,70,71,73,74,79,81,82,83,84	16	
Cervical and other anogenital cancers	(6,11),18,31,33,35,39,45,51,52,54,56, 58,59,66,68,69	16	
Human papillomavirus (HPV) types in brackets indicate extremely rare prevalence.			

Human papillomavirus (HPV) types in brackets indicate extremely rare prevalence.

The genital HPVs are epitheliotropic

- the high-risk HPV types, in particular, have a tropism for the metaplastic squamous cells at the squamocolumnar junction of cervix
- This has very important consequences for cytologic screening of cervix
- Also for native squamous epithelium of other sites, in contrast to most cervical cases

cytologic screening of cervix

- -The most precursor lesions and carcinomas arise at SCJ
- is amenable to sampling with a brush or spatula
- if the lesions of HPV-associated neoplasia of the genital tract were randomly distributed within the squamous mucosa, screening would have failed
 - as it has in the oral cavity, pharynx, esophagus, larynx, skin, vulva, vagina and ectocervix

Potentially infectious virus particles

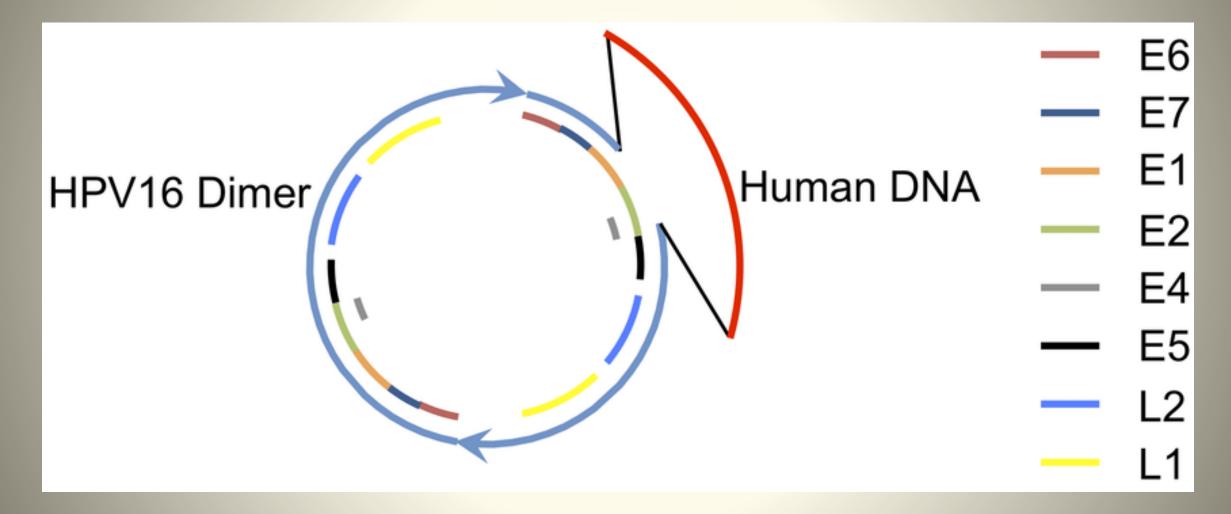
- must reach the proliferating basal cells for productive infection to take place
- a micro-injury to the squamous epithelium is required
- In the case of glandular epithelium
 - the stem cell compartment should be more accessible
 - infection of cervical glandular epithelial cells is less likely

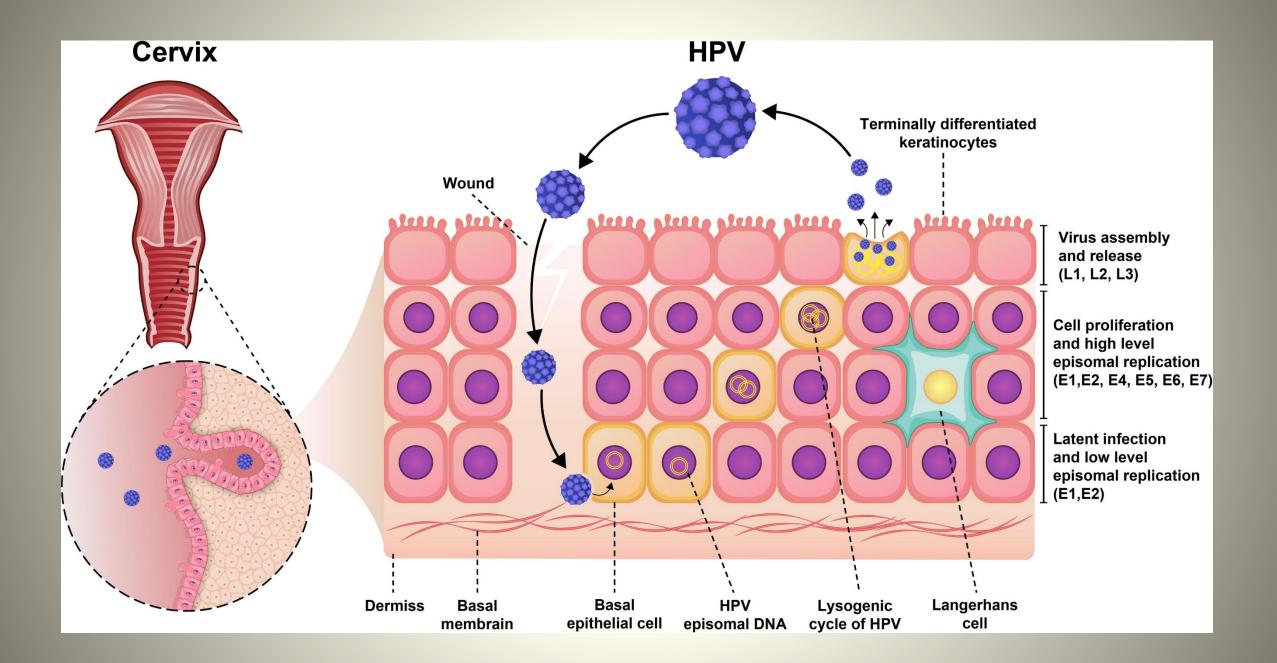
silent infection productive infection

silent infection

- Once the virus particles are taken up by the basal epithelial cells
- methylation of the viral genome
- Episomic viral DNA remains in the cell
- but is not transcribed or translated

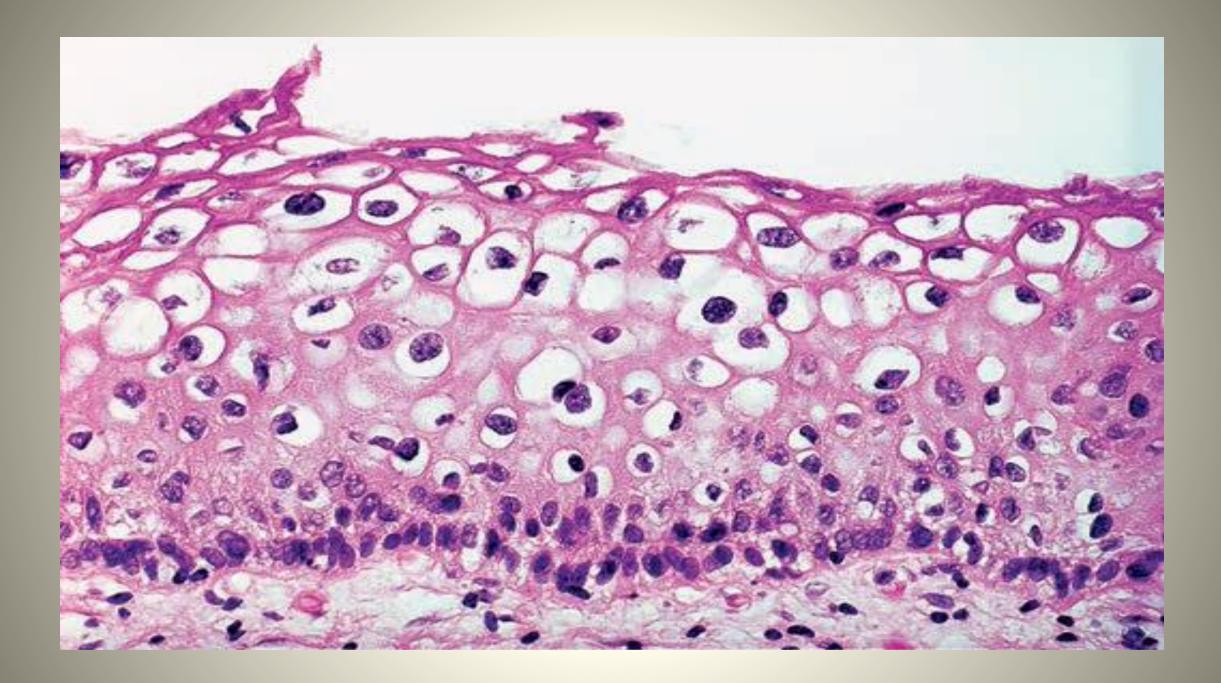
the HPV Dimer-Human DNA Hybrid Episome





productive infection may result

- may result in orderly expression of viral genes
- the host squamous cells mature
- assembly and release of infectious virus particles at the epithelial surface
- This is typically associated with **koilocytic change**
- a viral cytopathic effect: the E4 protein encoded by the viral genome causes disruption in the cytoplasmic keratin matrix

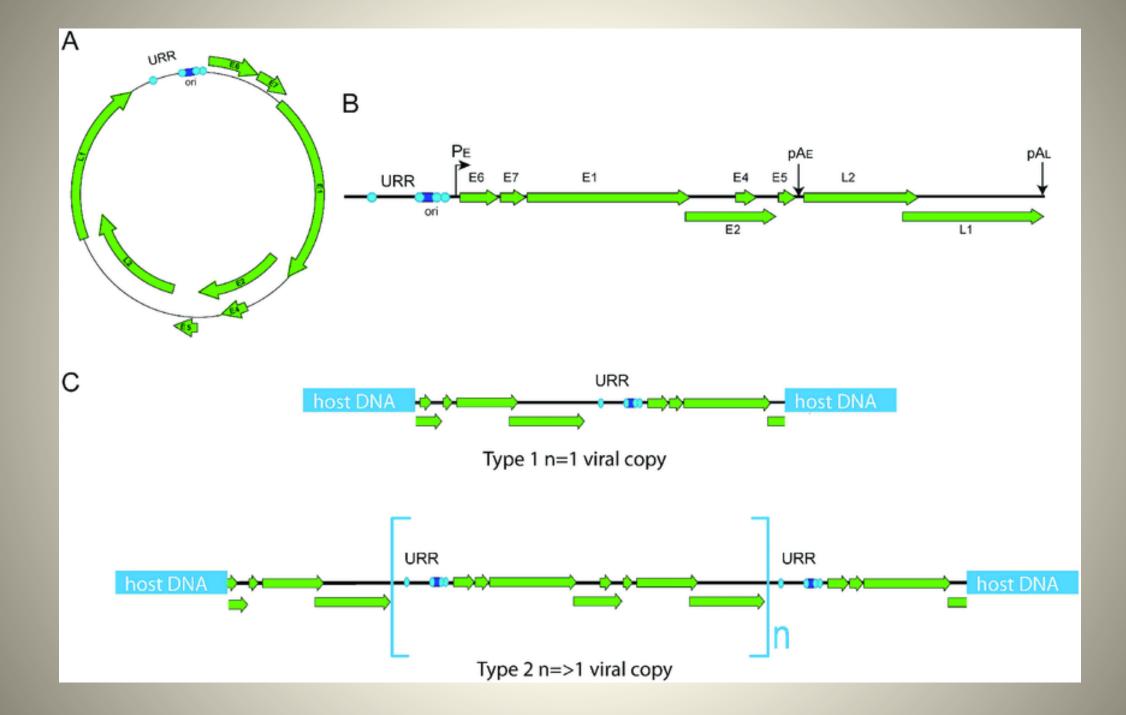


LSIL

- either a diploid or polyploidy nuclear DNA distribution
- orderly expression of the viral genome
- the squamous cells mature and move to the surface

high-risk/oncogenic HPV infection

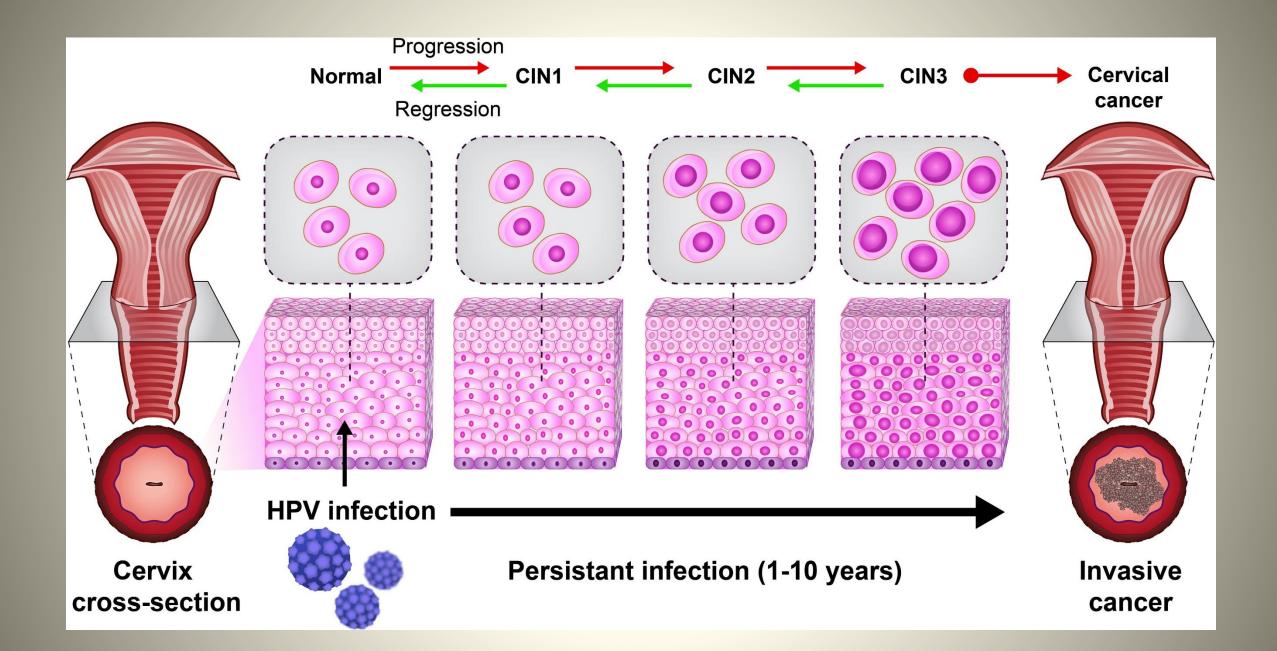
- deregulated expression of the viral genome
- The E6 and E7 proteins encoded by the genomes
- bind and inactivate the proteins encoded by the TP53 and RB anti-oncogenes
- unfettered proliferation of the host cell
- decreased E4 expression and virion production
- decreased or absent koilocytic change



increased likelihood of:

- integration of the viral genome into the host genome
- acquisition of other genetic abnormalities in the host cell

malignant transformation



Not all viral infections lead to transformation

most are cleared by the immune system of the host through cell-mediated immunity

productive infection due to

• By low-risk HPV, 6 or 11

-LSIL/condyloma with viral cytopathic effects

- By high-risk HPV, most commonly 16 or 18

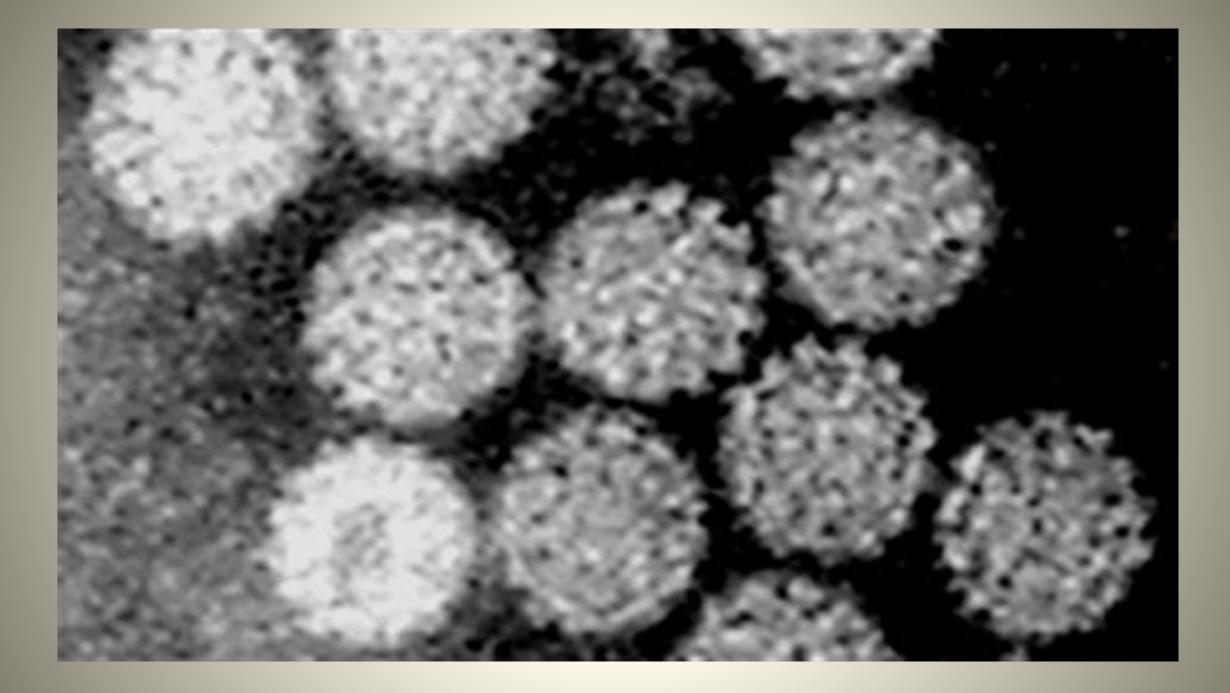
 –LSIL
 - -progress to HSIL
 - -invasive squamous cell carcinoma

HPV can be detected

electron microscopy:

intranuclear crystalline occasionally filamentous inclusions

- specific identification: nucleic acid hybridization
 - -with or without amplification
 - -from DNA or RNA
 - -either a liquid-based or in situ analysis



The p16 antioncogene

- in cells infected by high-risk HPV
- inactivation of the RB protein by the viral E7 protein
- phosphorylated RB protein
- absence of the normal downregulation of p16 expression
- high-level expression of p16
- strong nuclear and cytoplasmic immunoreactivity

Immunostaining for p16 (p16INK4a)

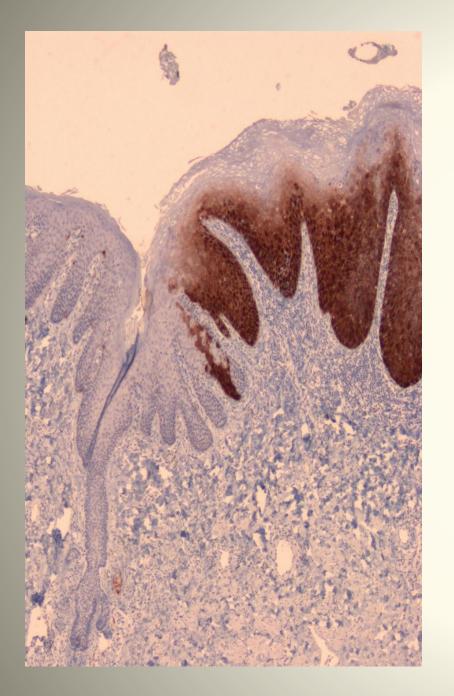
- a sensitive and specific surrogate for detection of HPV-associated versus HPV-independent vulvar squamous neoplasia
 - sensitivity of 100%
 - specificity of 98%–99%
- performed on intraepithelial lesions or early invasive carcinomas
 - p16 expression can be lost during tumor progression

Correct interpretation of p16 immunostaining

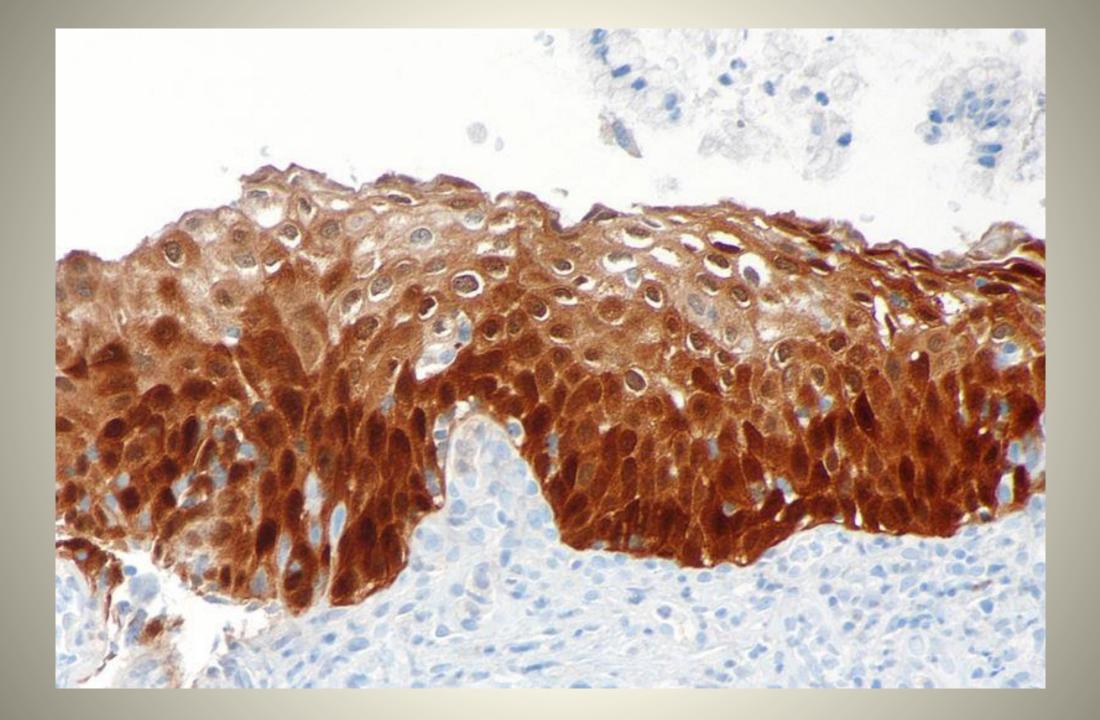
- the staining protocol has been properly validated
- appropriate controls run
- a number of antibody clones specific for p16
- it is incumbent on each lab to optimize the staining for the clone they have chosen
- a responsibility that is sometimes neglected

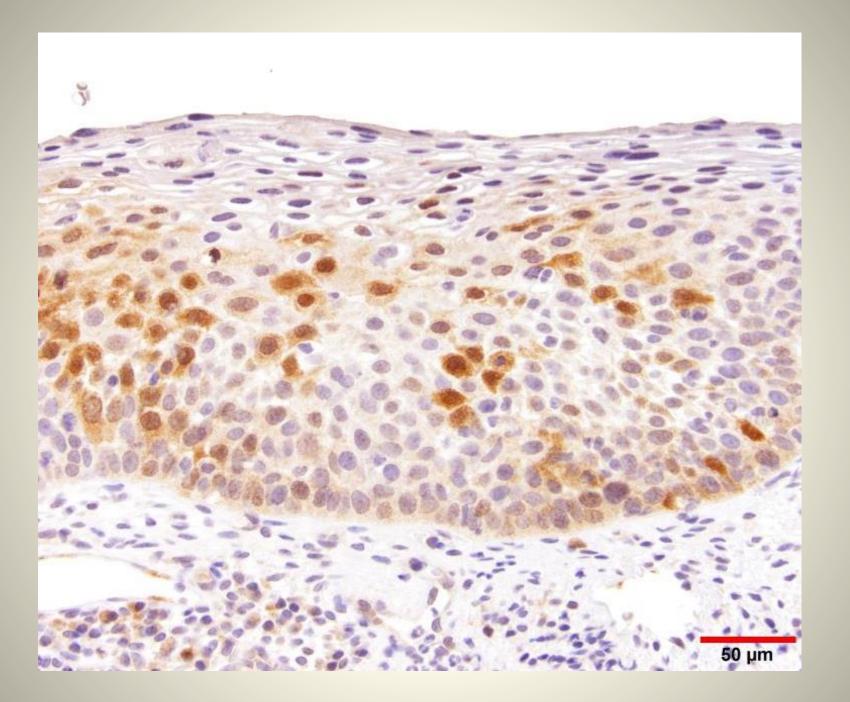
block positivity for p16

- indicative of high-risk HPV infection
- strong nuclear and cytoplasmic staining of every cell in the basal third of the epithelium
- typically extending into the middle and upper thirds



"Block" immunoreactivity in HSIL a test for the presence of oncogenic HPV





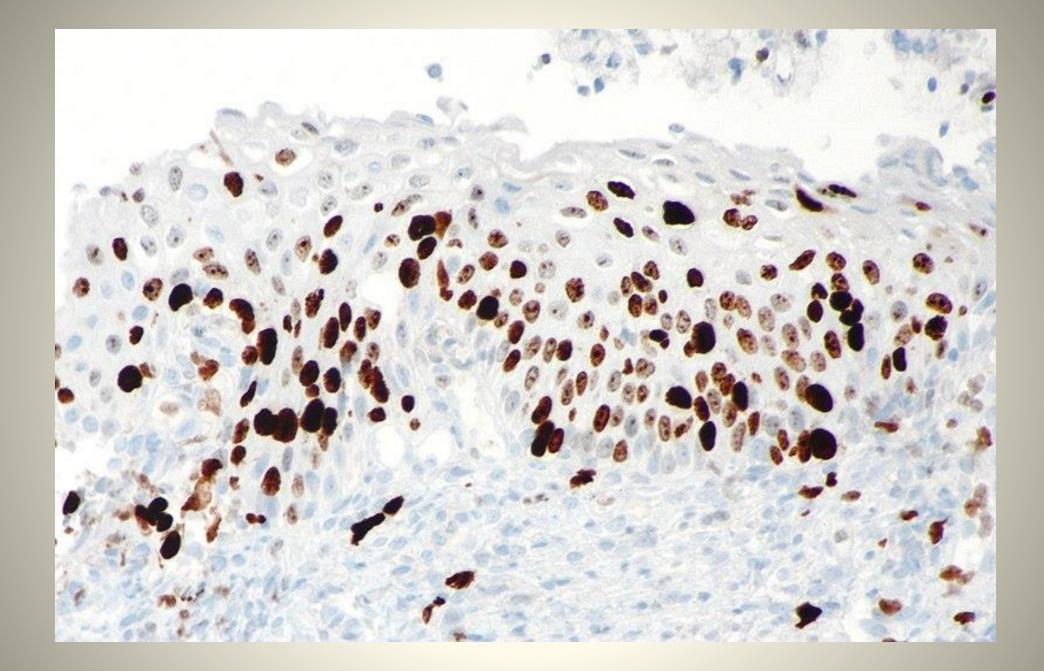
p16 immunostaining		HSIL n (%)	LSIL n (%)
Percentage of positive cells (%)	0(<5%)	5 (41.7%)	11(57.9%)
	1(5-49%)	4(33.3%)	6(31.6%)
	2 (50-80%)	3 (25.0%)	2(10.5%)
Intensity of reaction	0 (No reaction)	1 (8.3%)	3 (15.8%)
	1(Weak)	3 (25.0%)	10 (52.6%)
	2 (Variable)	4 (33.3%)	4 (21.1%)
	3 (Strong)	4 (33.3%)	2 (10.5%)
Cellular reaction pattern	0 (No reaction)	1 (8.3%)	3 (15.8%)
	1 (Focal)	10 (83.3%)	16 (84.2%)
	2 (Diffuse)	1 (8.3%)	0 (0%)
p16 Negative (0–3)	0	1 (8.3%)	3 (15.8%)
	2	3 (25.0%)	7 (36.8%)
	3	1 (8.3%)	4 (21.1%)
Positive (4–8)	4	4(33.3%)	2 (10.5%)
	5	0 (0%)	2 (10.5%)
	6	2 (16.7%)	1(5.3%)
	7	1 (8.3%)	0 (0%)

Immunoreactivity for p16

- is not specific for HPV
 - other tumor types, unrelated to HPV, dysregulation of RB:
 - high-grade serous carcinoma of tubo-ovarian or endometrial origin
- can be lost through mutation, as cervical cancer progresses
 - rare in the HSIL or in situ adenocarcinoma

Combined staining for Ki-67 and p16

✓ a higher proliferative index
✓ The strong, diffuse, block positivity for p16



le 3. Ki-67 grading in HSIL and LSIL

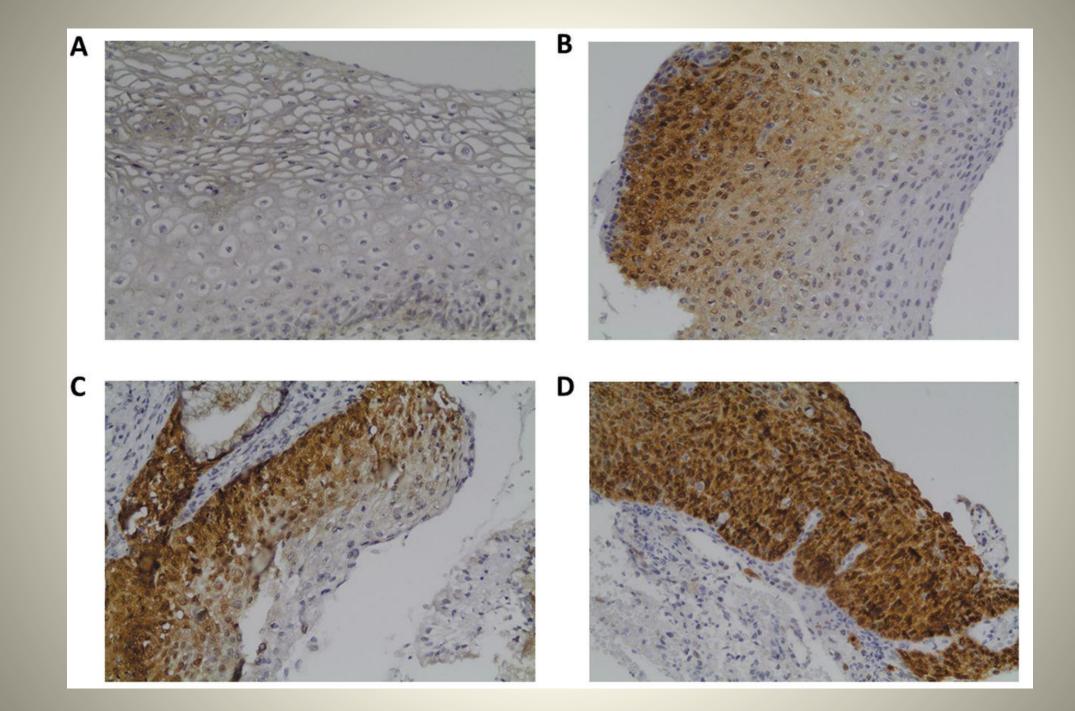
Ki67	HSIL n (%)	LSIL n (%)	Chronic Cx
0	0(0%)	0(0%)	16(80%)
1 (<5%)	6 (50.0%)	15 (78.9%)	4(20%)
2 (5-30%)	4 (33.3%)	4 (21.1%)	0(0%)
3 (>30%)	2 (16.7%)	0 (0%)	0(0%)

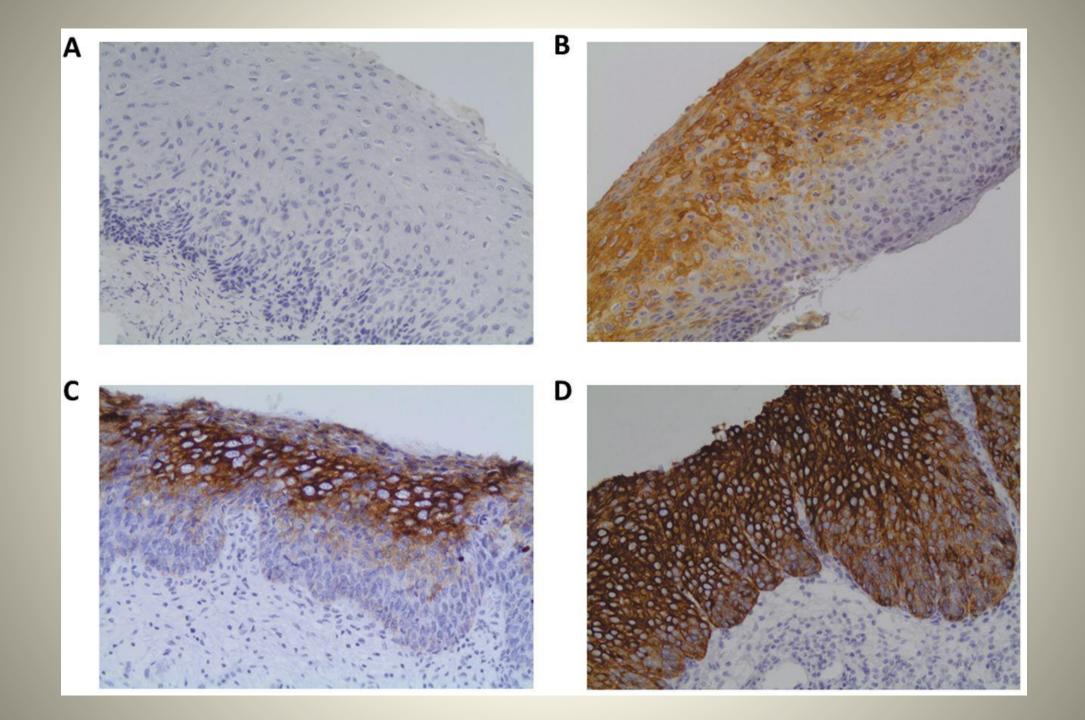
Risk factors for progression

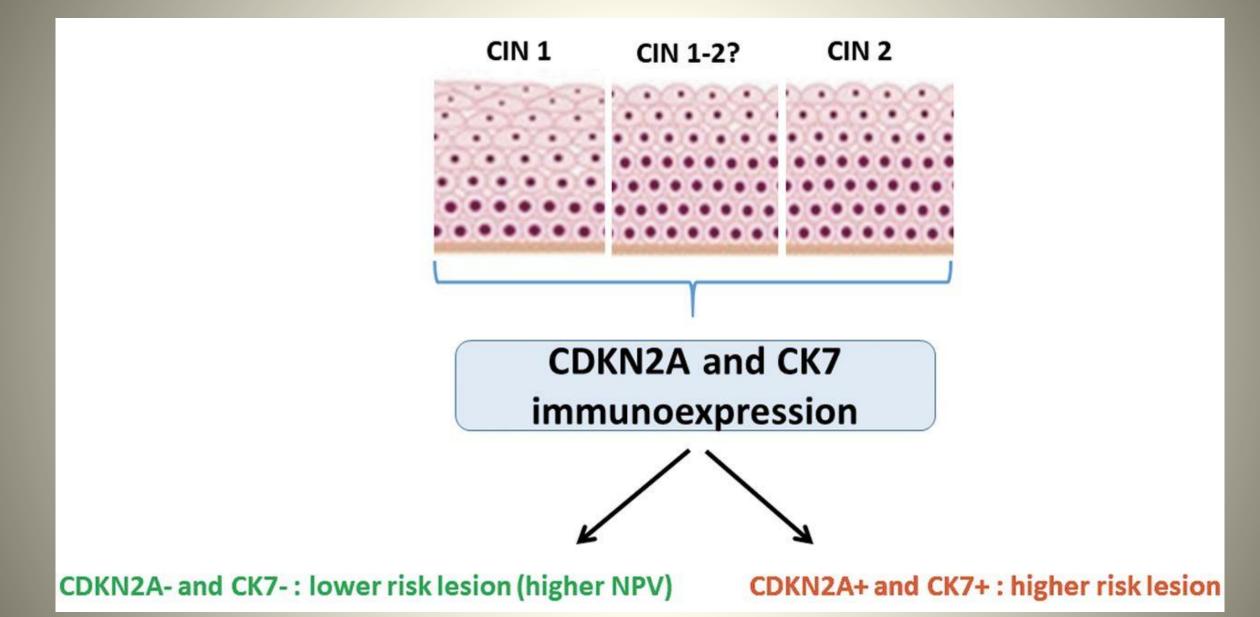
- the presence of high-risk HPV
- ✓ high-risk HPV are present in most LSIL

• Cytokeratin 7

- ✓ a marker of metaplastic cells of the transformation zone
- ✓ a marker of LSIL that is more likely to progress
- ✓ LSIL/condyloma can arise on the ectocervix or elsewhere in the lower female genital tract: little risk of progression to HSIL
- ✓ further validation is required before it enters routine practice







Other tests

 In addition to p16 immunostaining:
 can be valuable in establishing HPV status
 for cases where the clinical, histomorphologic, and p16 immunostaining data are not concordant

Other tests

- PCR for HPV DNA
 - -viral genotyping
- The Hybrid Capture HPV test
- In situ hybridization for HPV
- viral E6/E7 oncoprotein mRNA

PCR for HPV DNA

- allows for viral genotyping
- is highly sensitive
- lacks specificity
- may be present as a bystander if there is no transcription of the viral oncogenes

presence of HPV DNA may reflect

(1) virus particles on the surface of the epithelium secondary to **recent exposure**, without infection

(2) productive infection that **will be cleared completely** by the host

(3) a **latent** infection

(4) **oncogenic** HPV DNA that is integrated into the host genome

identification of the specific genotype of HPV

- is of import
- but does not mean premalignant change or malignancy
- Especially in the young, HPV may be cleared by the host

The Hybrid Capture HPV test

- a signal-amplified hybridization microplate-based array
- detect multiple HPV genotypes
- a liquid based nucleic acid detection technique
- more sensitivity than ISH techniques

In situ hybridization for HPV

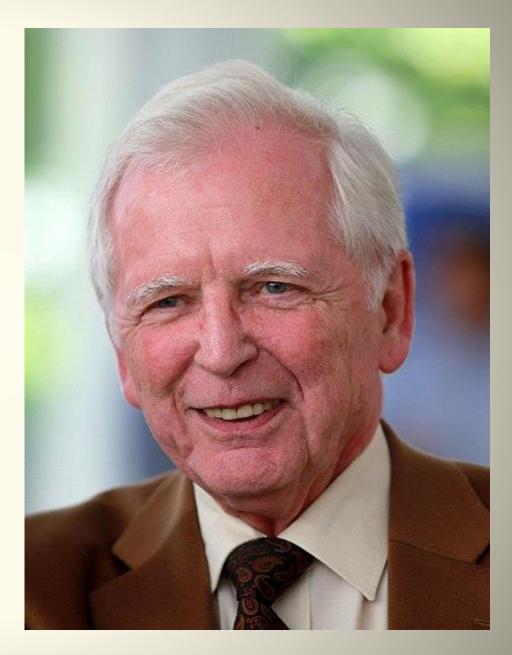
- is highly specific
- is more expensive than immunostaining
- has a longer turnaround time

Detection of mRNA of HPV oncogene E6/E7

- Expression of the oncoprotein mRNA
- the virus genome is being transcribed
- a better marker of HPV
- establish an etiologic role: a causative role in oncogenesis
- is currently **not widely available** in clinical laboratories

Harald zur Hausen هار الد تسور هاوزن

 deservedly won the Nobel prize in 2008 for his work establishing the link between HPV and cervical cancer



role of HPV in cervical cancer

- researchers started to postulate and analyse a possible role of HPV in cervical cancer, 1974
- the appearance of Koilocytes in cervical smears indicates the presence of a papillomavirus infection, 1976
- The demonstration of **heterogeneity** within the papillomavirus family
- HPV16 and HPV18 were cloned in 1983 and 1984

role of HPV in cervical cancer

- expression of specific viral genes (such as E6 and E7) was shown in cervical cancer biopsies, 1985
- viral ring molecule was shown for **integrated** genome copies
- the **immortalization** property of viral DNA, 1987
- viral oncogene expression for the maintenance of the malignant phenotype, 1992
- E6-p53 and E7-pRB interactions, 1994

1994-2008

- a more detailed knowledge of the natural history of HPV infection
- epidemiological studies have also been performed
- high-risk HPV as the primary risk factor
- **persisting** HPV infections were the most significant risk factor



HPV pathogenesis

The papillomavirus life cycle

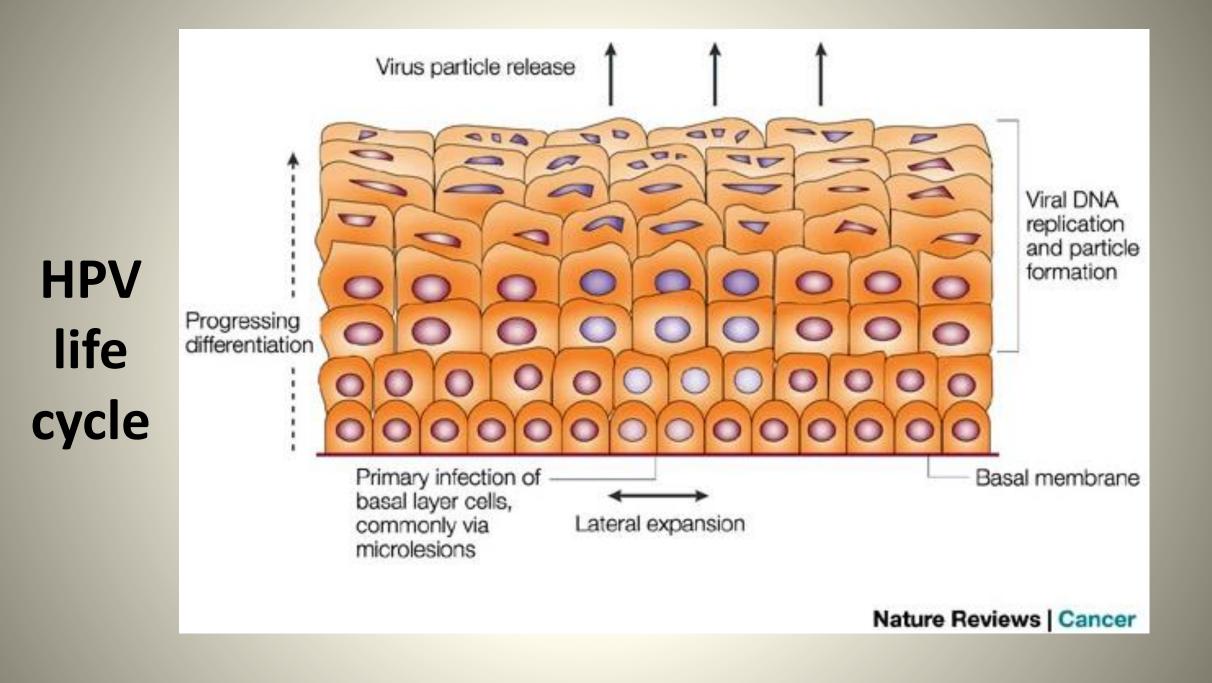
- differs from all other virus families
- the availability of epidermis or mucosal epithelium that are still able to proliferate (basal cells)
- microlesions of skin or mucosa

The papillomavirus life cycle

- viral gene expression is largely suppressed
- But limited expression of specific 'early' viral genes (E5, E6 and E7) results in enhanced proliferation and their lateral expansion
- The infected cell divides
- the population spreads laterally
- Some migrate into the upper layers

the upper differentiating cells

- 'late' viral gene expression is initiated
- the circular viral genome is then replicated
- structural capsid proteins are formed
- Viral particle formation ensues: complete viral particles are assembled
- Particles are released at the surface and might then infect additional tissues



HPV genome

- 6800-8000 base pairs
- eight open reading frames E6, E7, E1, E2, E4, E5, L2 and L1 coding for 'early' (E) or 'late' (L) functions
- Three genes possess proliferation-stimulating activity
- **√E5**

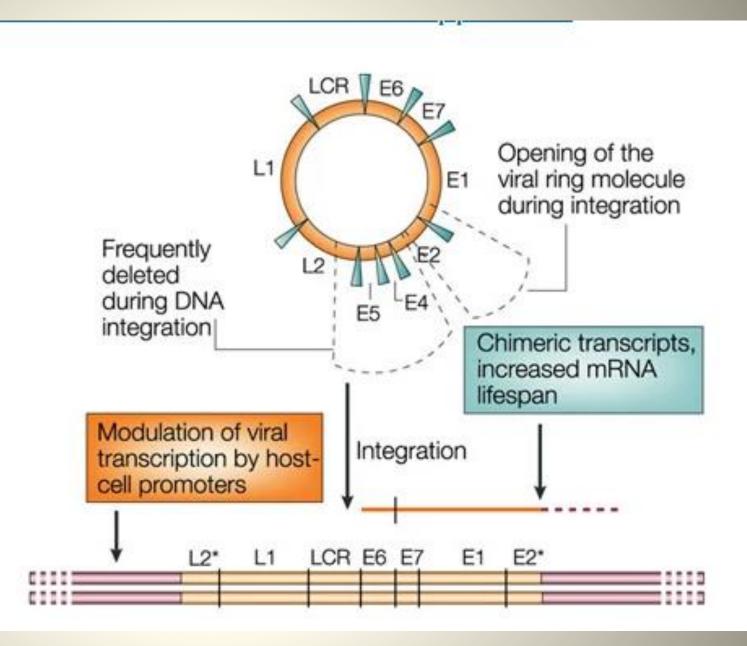
√E6

√F7

E5 is important in the early course of infection

- **stimulates cell growth** by forming a complex with the EGFR, PDGF-βR and the CSF-1R
- prevent apoptosis following DNA damage
- as HPV-infected lesions progress to cervical cancer, the episomal viral DNA frequently becomes integrated into hostcell DNA, and a substantial part of the genome, (including the E5 gene) is deleted
- E5 is not obligatory in late events of carcinogenesis

The circular HPV DNA and its integration



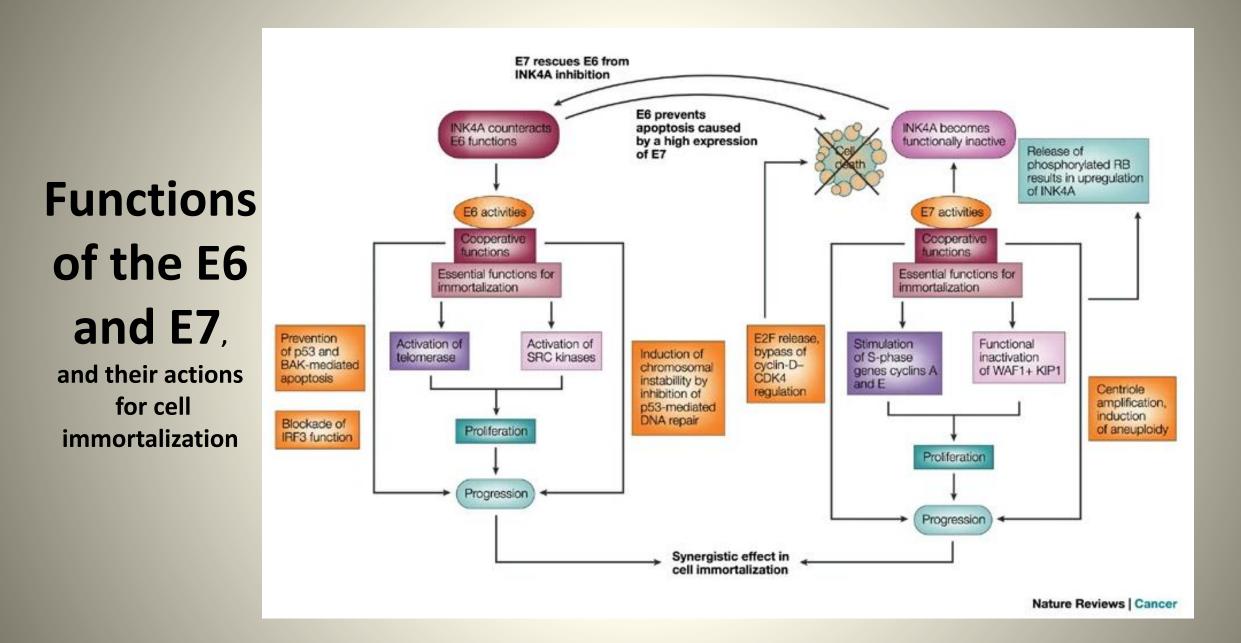
Integration of viral molecule into host-cell DNA

- The ring molecule is opened
 Part of E2 and L2 and whole genes E4, E5 are deleted
- Viral transcripts uniformly span the E6 and E7 region
- are often linked to flanking cellular sequences

 transcription is enhanced by flanking host-cell promoters

E6 and E7 genes (their proteins)

- A more significant role for malignant transformation
- are **consistently expressed** in malignant tissue
 - inhibiting expression blocks the malignant phenotype
- They are independently able to immortalize various human cell types in tissue culture
- efficiency is increased when they are expressed together



functions for E6 and E7

- E6 interacts with p53
- E7 interacts with RB
- block the activity of these tumor suppressors

Immortalization

deregulated cellular growth and genomic instability

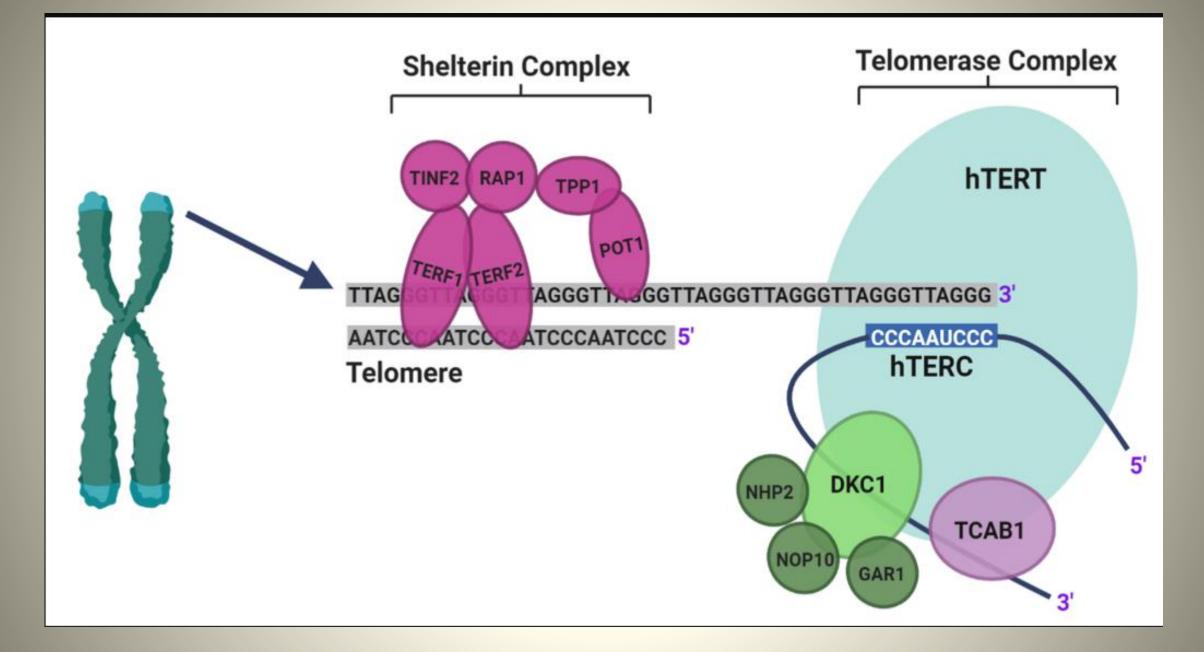
- both of which are characteristics of immortalized cells
- complex with p53
- resulting in the ubiquitination & degradation of p53
- Loss of p53

✓increased chromosomal instability

- Due to resistance to apoptosis
 - interacts with p53
 - -interacts with pro-apoptotic protein **BAK**

✓growth stimulation

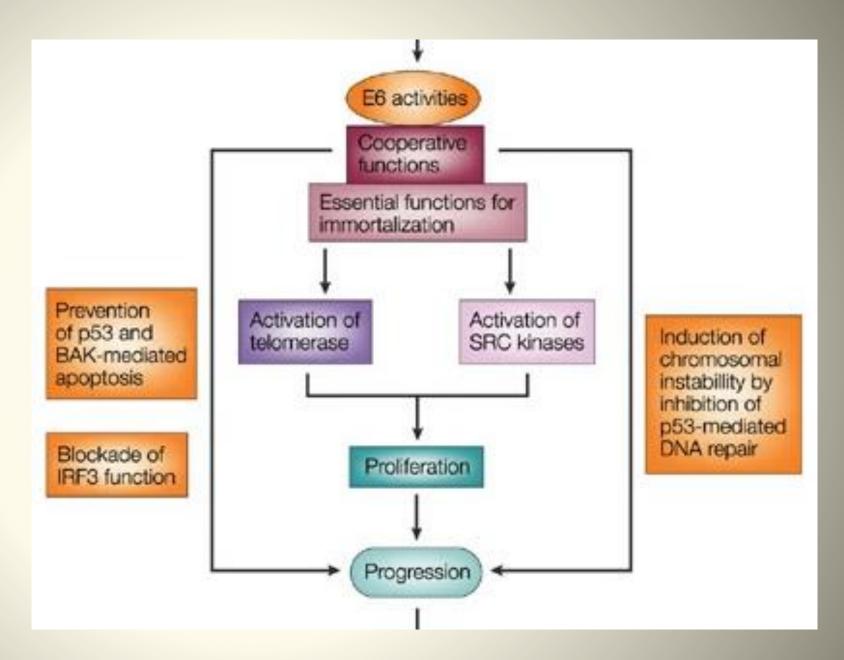
- the activation of telomerase
- inhibition of degradation of activated SRC-family kinases



disrupts antiviral response

- binding to IRF-3 (interferon regulatory factor-3)
- the inhibition of its transcriptional activity
- significantly dampens the induction of IFNβ

Functions of the E6



E7 protein

- interacts with and degrades RB
- releases the transcription factor E2F
 - might lead to apoptosis
 - upregulates cyclin-dependent kinase inhibitor INK4A

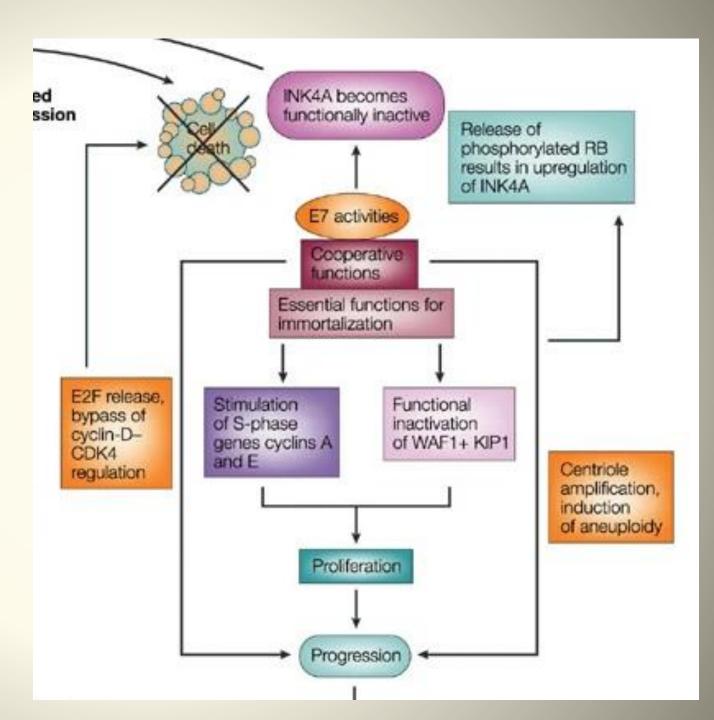
also inactivates INK4A (p16)

• The INK4A (p16) counteract functions of E6 protein

E7 protein

- stimulate cyclins A and E (s phase genes)
- inactivate the cyclin-dependent kinase inhibitors WAF1 (CIP1 and p21) and KIP1 (p27)
- inducing centriole amplification
 - -induces aneuploidy
 - -contributes to tumorigenesis

Functions of the E7

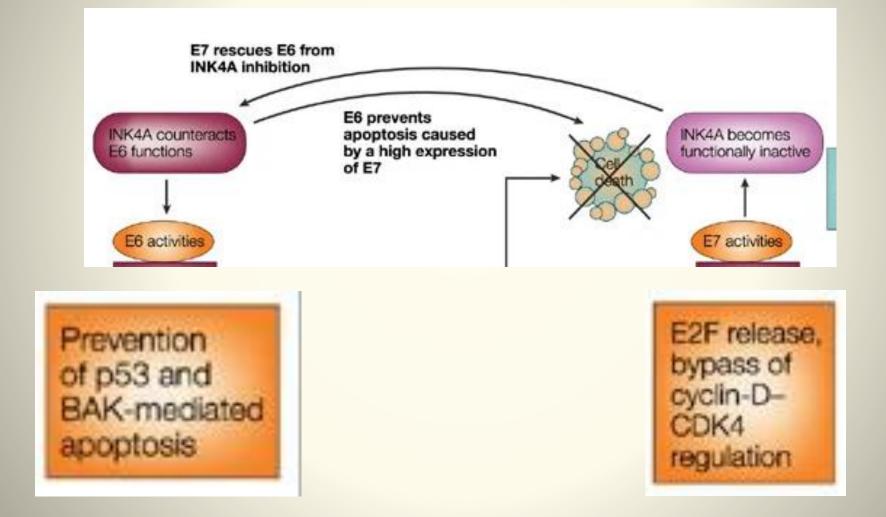


E6 and E7 synergism

- E6 prevents E7-induced apoptosis that is induced by E2F
 By p53 and BAK
- E7 rescues E6 from inhibition by INK4A
 - inactivates INK4A E7
 - directly activating cyclins A and E

their joint function results in a marked increase in immortalization & transformation

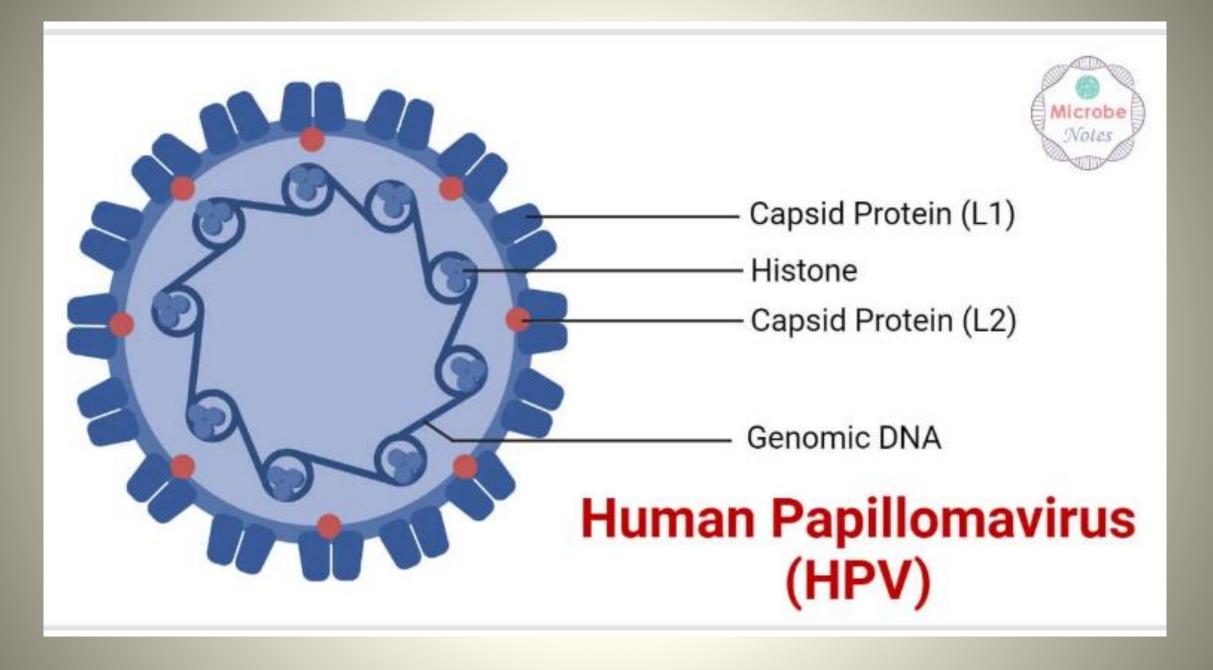
E6 and E7 synergism



two HPV early proteins

structural proteins L1 and L2 are important for vaccine development

	Cervarix	Gardasil, Silgard	Gardasil-9
Valency	2-Valent	4-Valent	9-Valent
Types	HPV16/18	HPV6/11/16/18	HPV6/11/16/18/31/33/45/52/58
Adjuvant	ASO4 (0.5 mg aluminum hydroxide and 50 µg 3-O-desacyl-4"-	0.225 mg aluminum	0.5 mg aluminum
	monophosphoryl lipid A (MPL))	hydroxyphosphate sulfate	hydroxyphosphate sulfate
Expression	Baculovirus-insect cell	Yeast	Yeast
system			



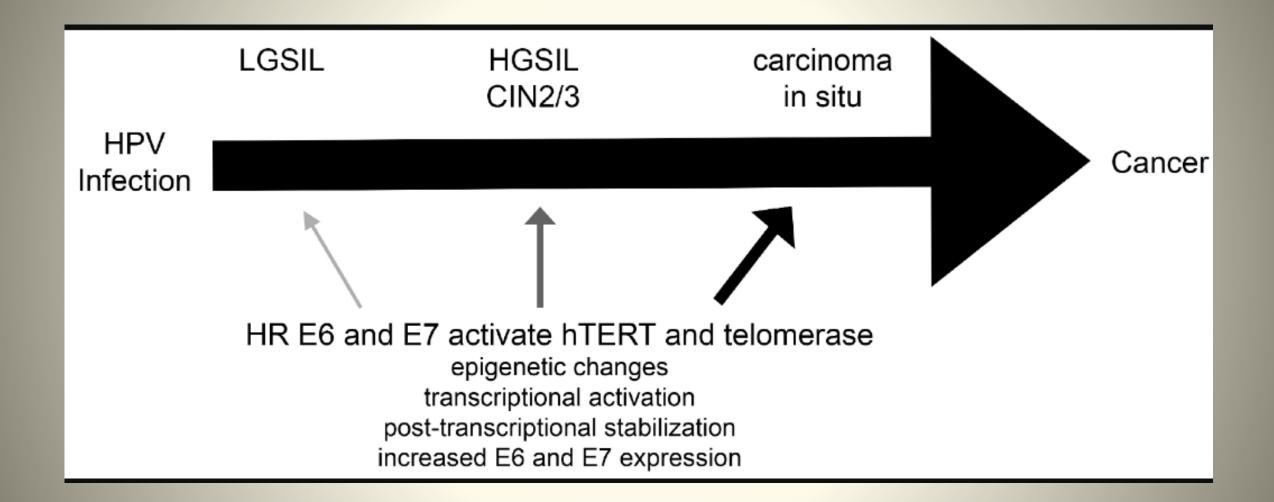
High- and low-risk HPV infections

- 'high-risk' types
 - are found preferentially in cervical and other anogenital cancers
- 'low-risk' types
 - found primarily in genital warts and non-malignant lesions

only the E6 and E7 genes of high-risk types were able to immortalize human cells

High-risk HPV types

- are widespread within all human populations
- Infection is commonly transmitted by sexual contact
- results initially in inconspicuous squamous intraepithelial lesions (SIL) in women
- Most of these lesions are cleared 6–12 months after appearance, probably due to immunological intervention

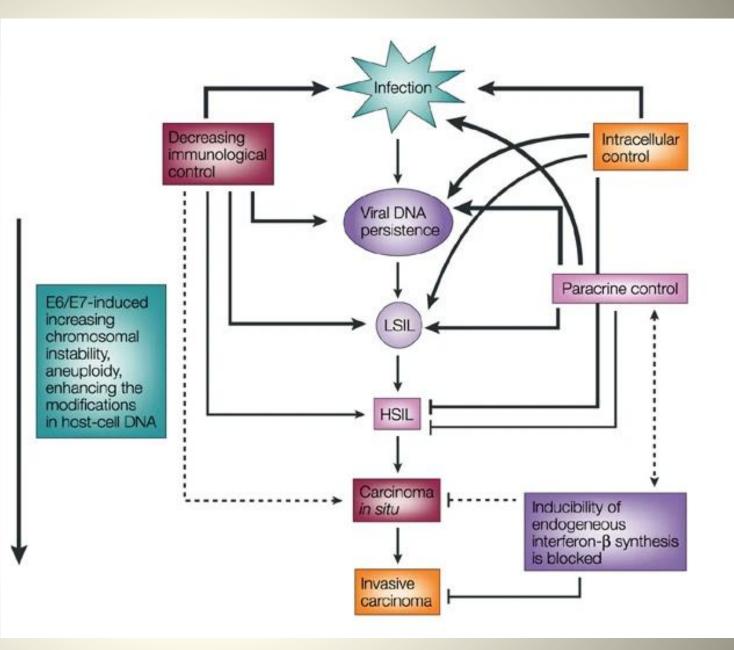


A small percentage

- Persists
- progresses to:
- high-grade SIL
- carcinoma in situ
- Squamous cell carcinoma
- adenocarcinoma

HPV-induced progression:

failing control mechanisms



Host control of HPV infection

- an intracellular control
 - by cyclin-dependent kinase inhibitors
- a paracrine signaling cascade
 - by macrophages and cytokines
 - tumor necrosis factor- α
 - loss of synthesis of interferon- β
- a decreasing immunological control

Host immune response factors

- ✓ Genes in the HLA region of chromosome 6
 - increased susceptibility to the transforming properties of high-risk
 HPV
- ✓ a detectable Humoral and Cellular immune response against HPV antigens during the course of regression
- The escape from immunological control is one important step in the progression of HPV-linked tumors

Impaired immune function

- increased incidence and prolonged persistence of SIL in immunosuppressed women
- ✓ a 5- to 10-fold increased risk
- immunosuppressant therapy in organ transplantation recipients
- HIV infection

High-risk HPV infections progress to cervical cancer

in only a small percentage after a long latency period

Factors that affect HPV malignancy

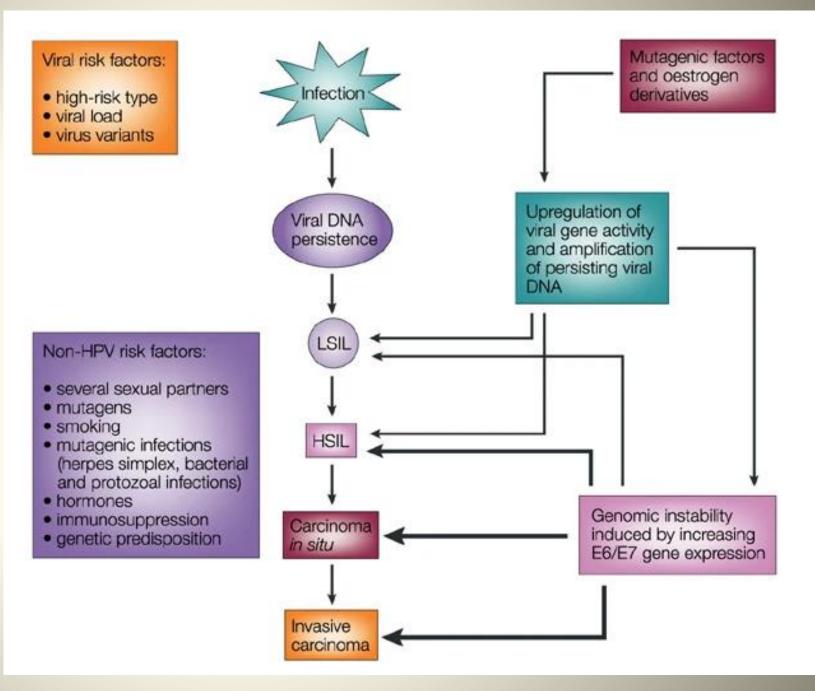
- modifications of cellular genes that influence antigen presentation
- signaling cascades that are engaged in viral oncogene transcription or viral oncoprotein function
- upregulating viral oncogene transcription
- modifying the viral promoter region
- amplifying the persisting viral genomes

HPV and non-HPV factors

that contribute to HPV-induced malignant progression

- Hormones (estrogen) activate the HPV promoter and facilitate immortalization of HPV-infected cells
- Mutagenic agents amplify persisting HPV DNA
- A rising level of E6 and E7 oncogene expression, results in increasing genomic instability and facilitates progression towards invasive growth

HPV and non-HPV factors that contribute to HPV-induced malignant progression



we now see the circle closing

- primary HPV screening to detect premalignant lesions
- HPV immunization to prevent such lesions increasingly being adopted

It is possible to envision

a world where the burden of disease due to HPV is dramatically reduced

SCC and Its Precursors

Squamous Intraepithelial Lesions/ Cervical Intraepithelial Neoplasia

SIL/CIN

- SIL terminology has gained wide acceptance, but there are still holdouts for the CIN terminology
- we will adopt the practice of providing a two-part diagnosis, with SIL first and the equivalent CIN in parenthesis thereafter
- LSIL (CIN1)
- HSIL (CIN2 and CIN3)

CIN2 and CIN3

- the distinction is arbitrary
- not clinically relevant
- Subtle differences in natural history have been reported

• HSIL (CIN2/3) for all high-grade lesions

LSIL (CIN1)

- is inclusive of **condyloma** in the LAST criteria
- The former practice of attempting to determine whether there is or is not dysplasia within condyloma has mercifully been brought to an end.
- condyloma ± CIN1 ???

Condyloma acuminatum

- one or several soft elevated masses of variable size
- is a venereal disease
- in vulva, vagina, cervix, bladder and penis
- is caused by HPV, usually type 6 (low-risk HPV infection)
- should be diagnosed as (VIN1, VaIN, PeIN)
- ✓ we do not recommend adopting LSIL, while we are able to endorse LSIL (CIN1) for cervical condyloma



Large condyloma of vulva

condyloma acuminatum

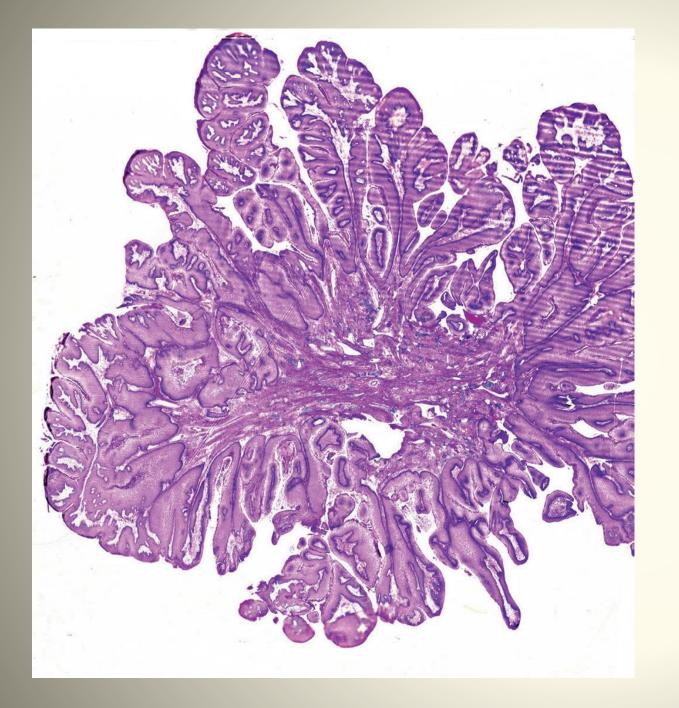
- Exophytic versus flat: grossly as a polypoid lesion
- considerably less common than the latter in cervix
- An undulating appearance of the epithelium on lowpower
- papillomatosis, acanthosis
- expanded or hyperplastic parabasal cell layer

Microscopically

- orderly maturation; a smooth transition to intermediate and superficial cells
- **mitotic activity** confined to the lower third of the epithelium (but few or no abnormal mitoses)
- a variable degree of lymphocytic inflammation in the stroma

Condyloma acuminatum

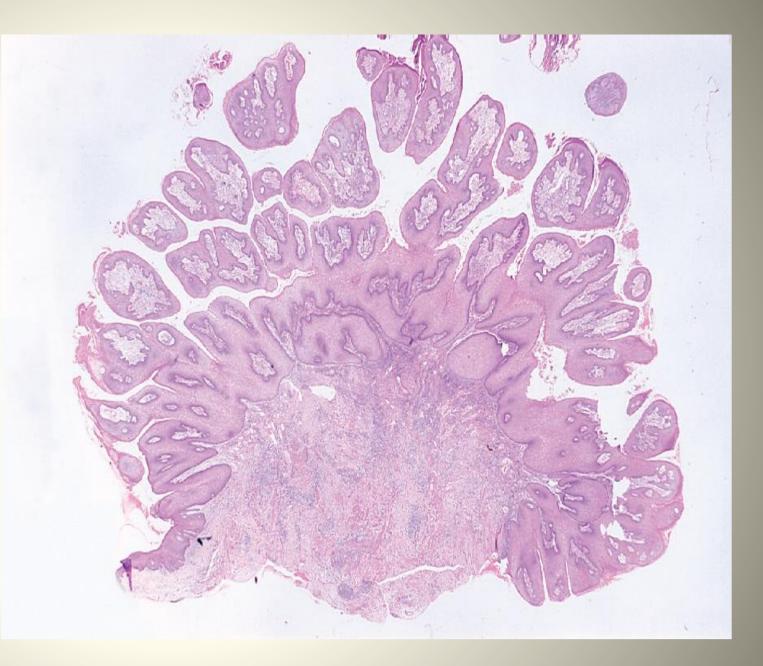
- A mild degree of basal or parabasal atypia is common
- if **more severe**, it should be evaluated and graded as for the flat SIL (is there HSIL [CIN2/3] present?)
- is associated with HPV-6 or HPV-11 in 70%–90% of the cases
- occasionally other types—such as HPV-16: high-grade cytologic atypia may be found

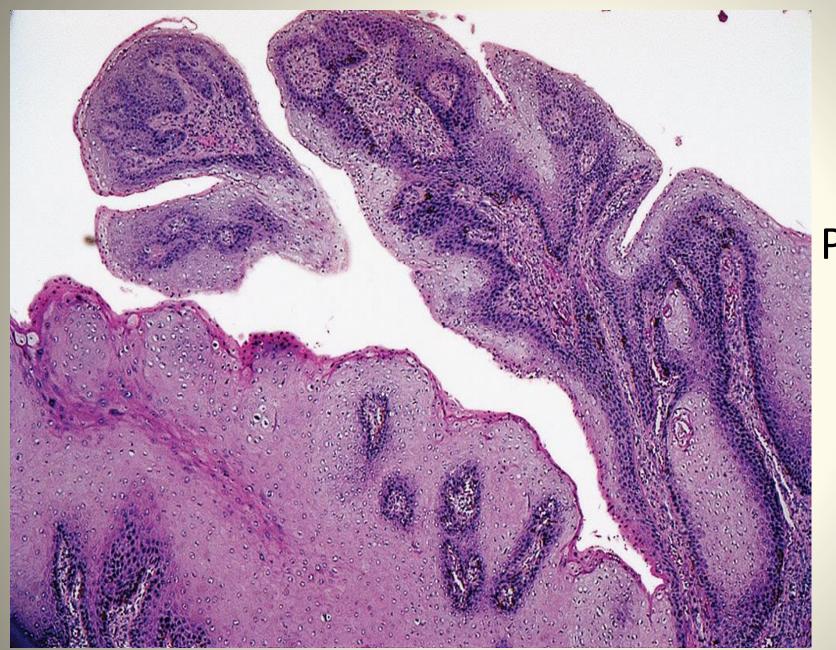


condyloma acuminatum

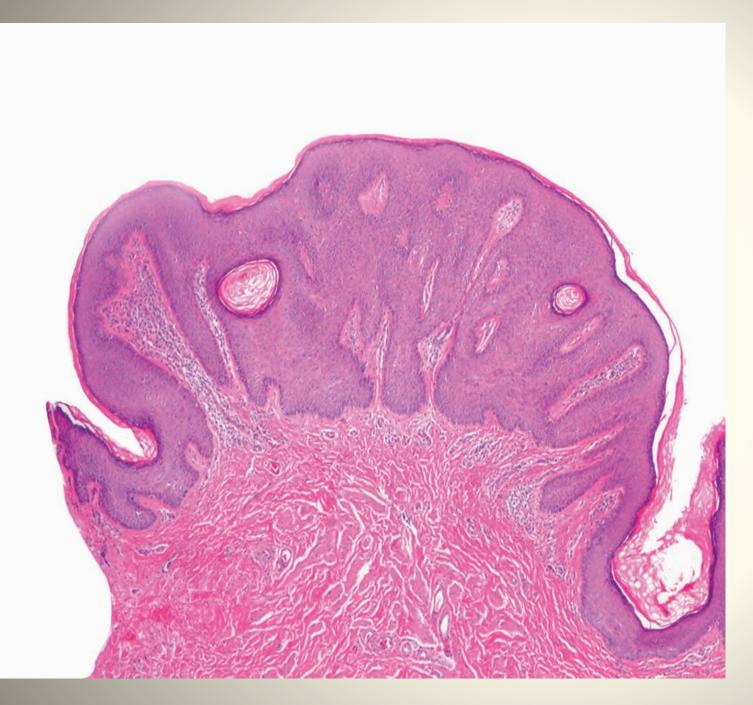
Condyloma Acuminatum

Complex papillary pattern composed of welldifferentiated squamous epithelium





Papillomatous shape of vulvar condyloma



Some condylomas have a low-power appearance simulating seborrheic keratosis

The other form: flat condyloma

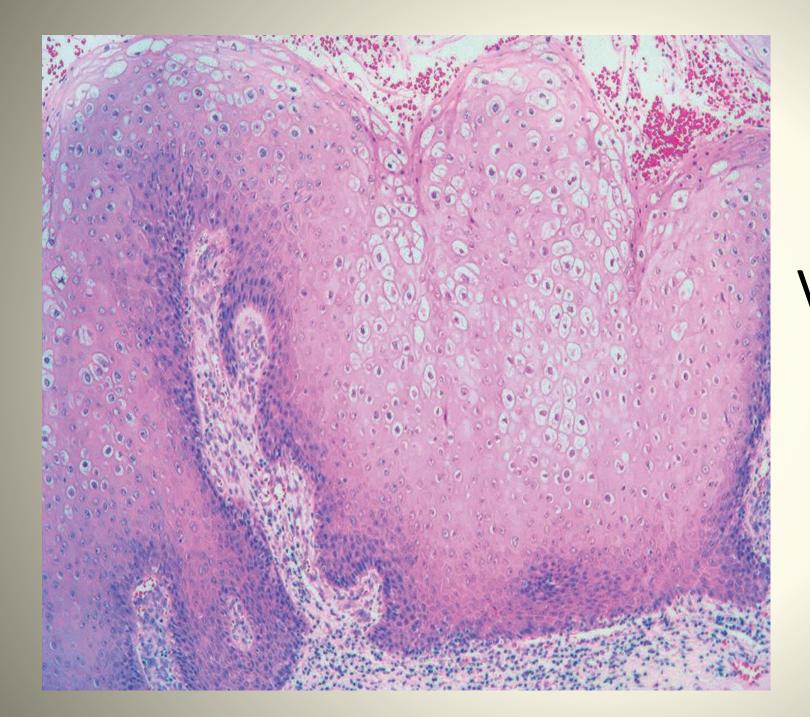
- is more commonly encountered in cervix than condyloma acuminatum
- The cytologic features are similar in both forms
- the classic lesion of LSIL (CIN1)
- It is typically not recognizable grossly

increased proliferative activity

- mitoses may be numerous, but are all typical
- with the Ki-67 stain
 - in contrast to fibroepithelial polyp and squamous papilloma
- The DNA content
 - Diploid
 - Polyploid (tetraploidy and octaploidy)

LSIL (CIN1)

- Koilocytic viral cytopathic effect is the pathognomonic
- it is doubtful whether LSIL (CIN1) can be diagnosed in the absence of koilocytic change
- koilocytic change (HPV effect) with or without dysplasia should not be distinguished

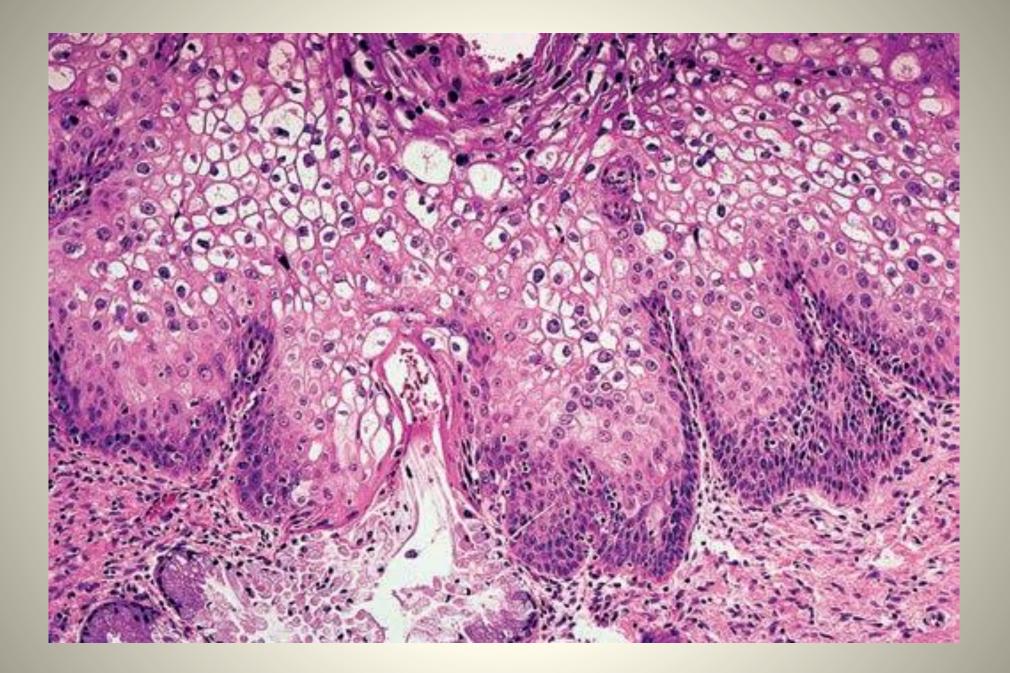


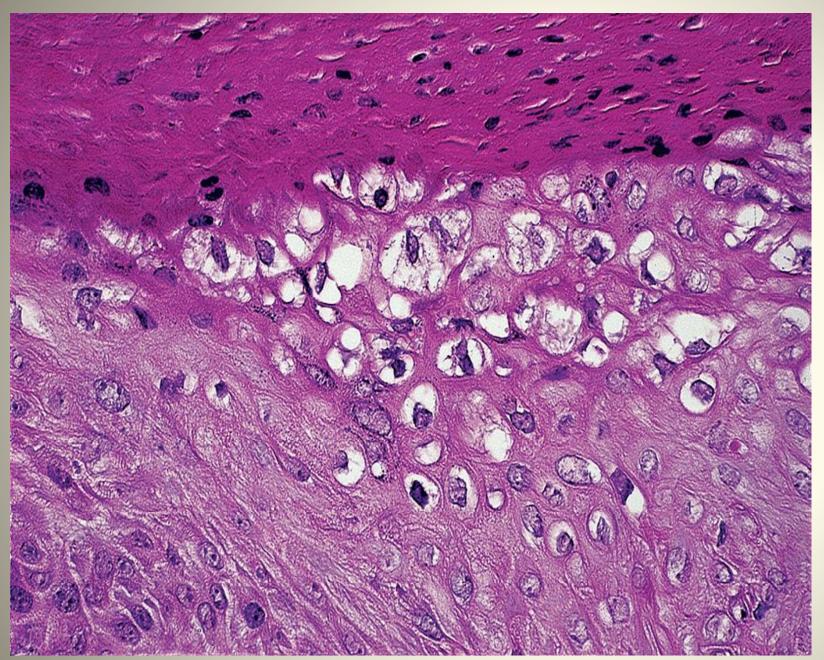
Virus induced cytopathic changes

Koilocytosis of the malpighian epithelium

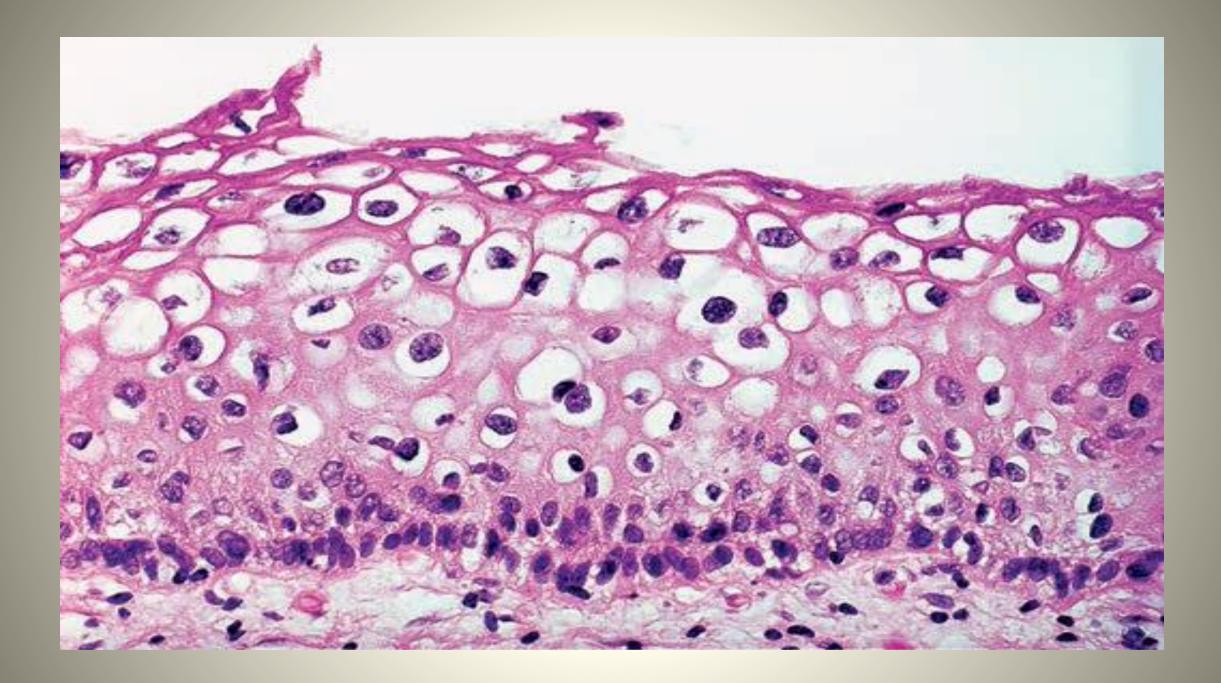
- a superficial or intermediate mature cell
- a sharply outlined perinuclear clear vacuolation
- an enlarged nucleus with an wrinkling or undulating nuclear membrane (prune-like or nuclear "raisins")
- a rope-like chromatin pattern
- dense- and irregular-staining peripheral cytoplasm

- Binucleation and multinucleation
- orderly maturation; a smooth transition to koilocytotic intermediate and superficial cells
- Koilocytosis is not as florid as condylomas of the cervix in other areas





Prominent koilocytotic changes



LSIL

- dysplasia of the lower third of the epithelium
- the distinction between "dysplasia" and reactive expansion of the parabasal layer is highly subjective.

the old chestnut

should be relegated to the bin reserved for medical trivia of historical interest only

- mild, moderate and severe cervical dysplasia (CIN1-3)
- dysplasia in the lower third, lower two thirds, and full thickness of the squamous epithelium

this is still repeated to trainees and has no underlying mechanistic basis

Bethesda classification

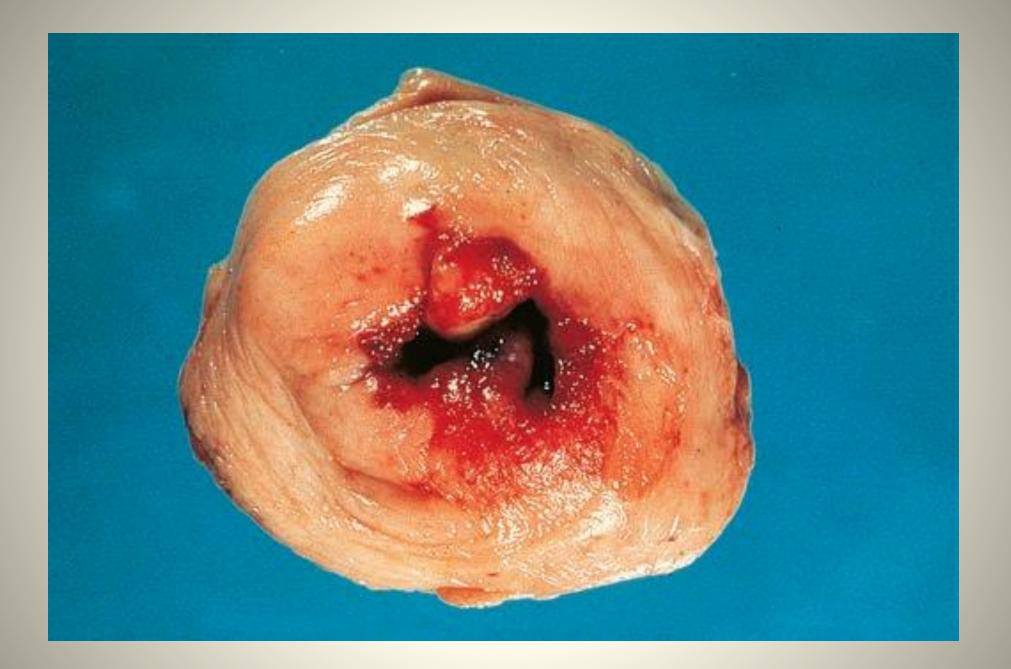
- reflects the biology of in situ squamous neoplasia of the cervix,
- especially as it relates to HPV
- a bipartite LSIL/HSIL system
- The proliferation rate of LSIL (CIN1) is higher than that of inflamed or metaplastic cervical squamous epithelium.

The natural history of LSIL

- most will clear spontaneously
- relatively few will progress to HSIL (CIN2/3)

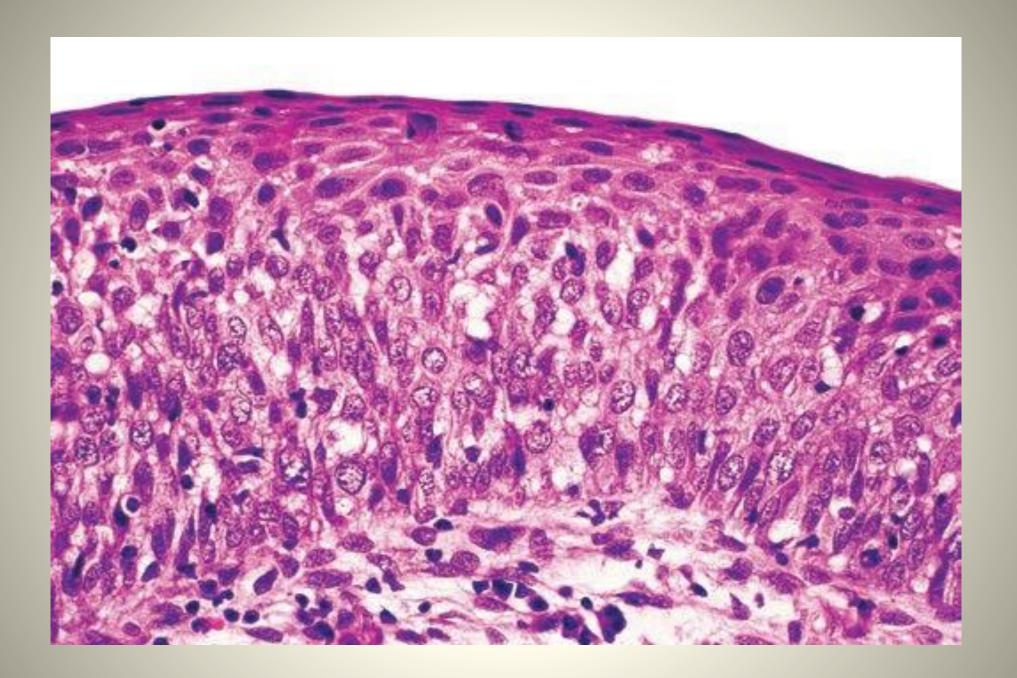
HSIL (CIN2/3)

- is best appreciated colposcopically
- may be seen macroscopically, when extensive



Mioscopically

- high N:C ratio in all layers of the epithelium
- Most importantly by striking nuclear atypia
 nuclear pleomorphism, enlargement, and hyperchromasia
- Prominent mitotic activity, with atypical mitoses
 - in the upper layers of the epithelium



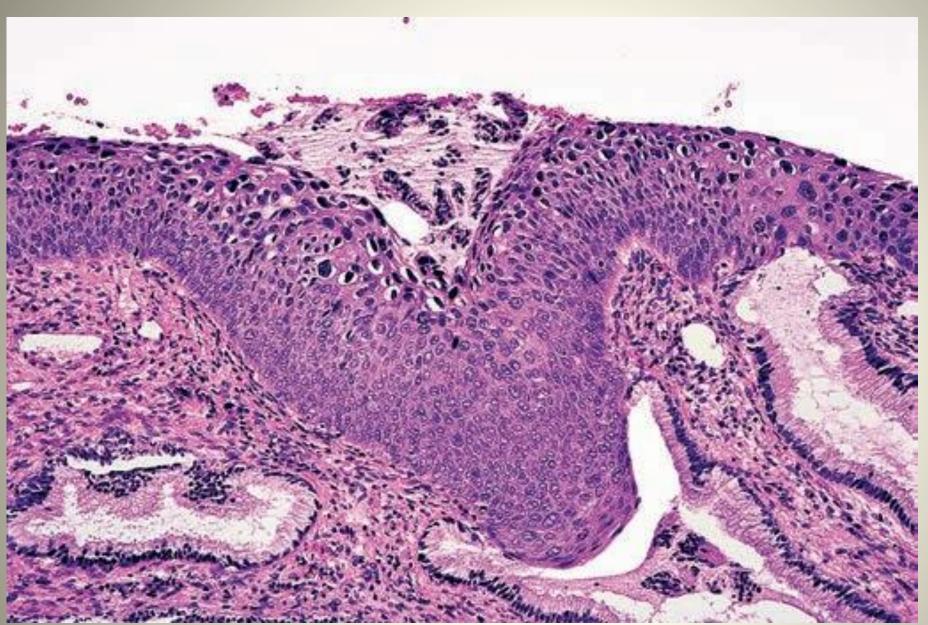
the differential diagnosis between LSIL and HSIL

- is made primarily based on H&E
- Only in selected borderline cases:
- p16 immunostaining strong and diffuse block positivity

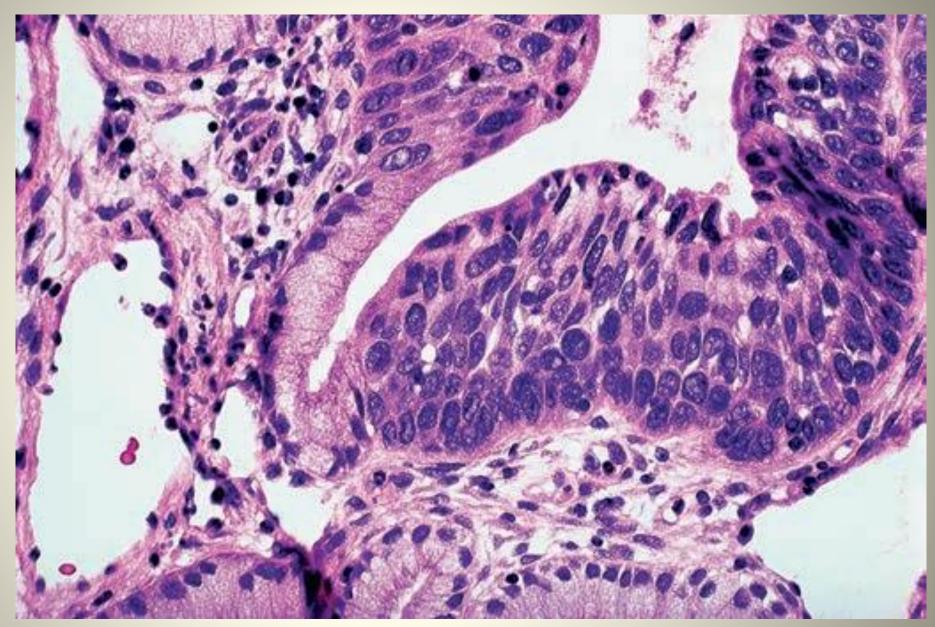
✓ Ki-67 stain: increased proliferative activity

these have to be taken into account

- the surface
- Margins
- the possible depth of involvement
- extension into endocervical glands

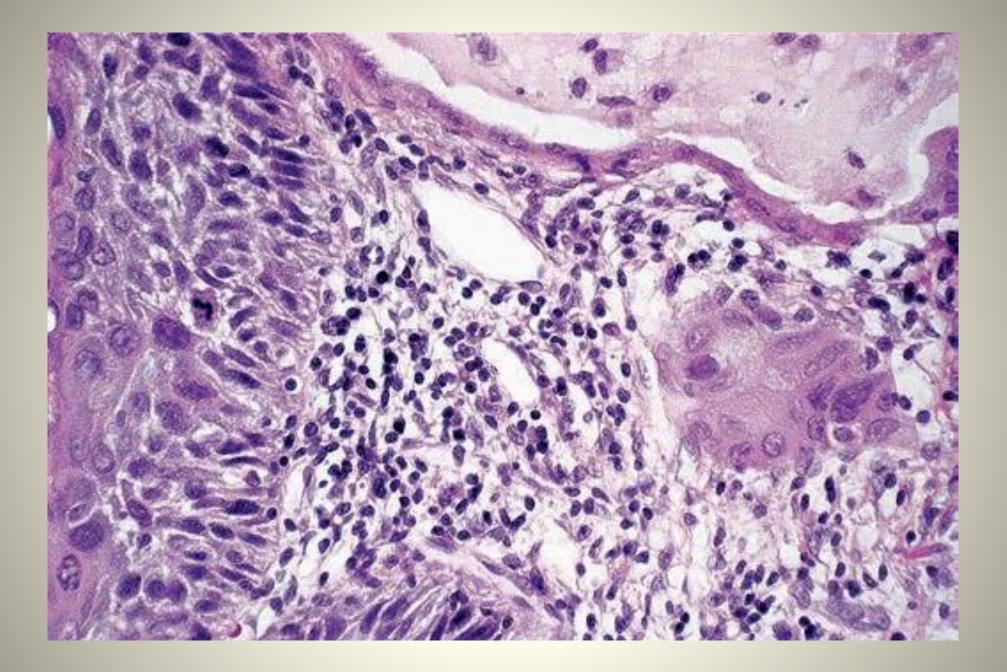


Extensive involvement by HSIL (CIN2/3) of surface epithelium and glands of endocervix Partial replacement of endocervical glandular epithelium by HSIL (CIN2/3)



the presence or absence of invasive carcinoma

- Thorough cervical sampling
- the proper sectioning of tissue
- continuing abnormal Cytology after initial treatment of HSIL (CIN2/3) were found to be 25 times more likely to develop invasive carcinomas than women with normal follow-up cytology



VIN2/3

- abnormal mitoses and nuclear pleomorphism, enlargement, and hyperchromasia in the basal and parabasal cell layers
- "block positivity" on p16 immunostaining,
 - at least the basal third of the epithelium
 - usually extending into the upper half
- The increased proliferative activity with the Ki-67 stain
- The DNA content is aneuploid pattern in most cases
- a significant number of HPV-independent VIN

VaIN2/3

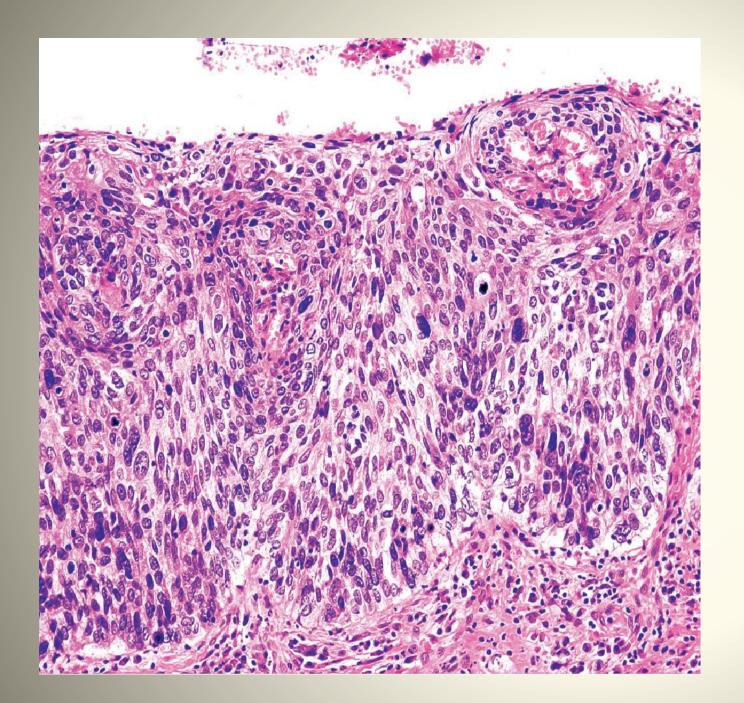
- The upper third of the vagina is the most common site, the vaginal and cervical lesions may be confluent
- arise from **native** squamous epithelium, in contrast to most cervical cases, which originate from *metaplastic* epithelium
- is **multifocal** in about half of cases
- very frequently associated with concomitant, subsequent, or prior (in situ or invasive) neoplasms elsewhere in the lower genital tract, especially the cervix

Penile Intraepithelial Neoplasia (PeIN)

- Human Papilloma Virus Related PelN
 - Basaloid (undifferentiated) PelN
 - Warty (Bowenoid) PelN
 - Warty-basaloid PeIN, shows an admixture
- HPV-unrelated PeIN
 - Differentiated PeIN has been associated with lichen sclerosus or other chronic inflammatory conditions

Basaloid (undifferentiated) PelN

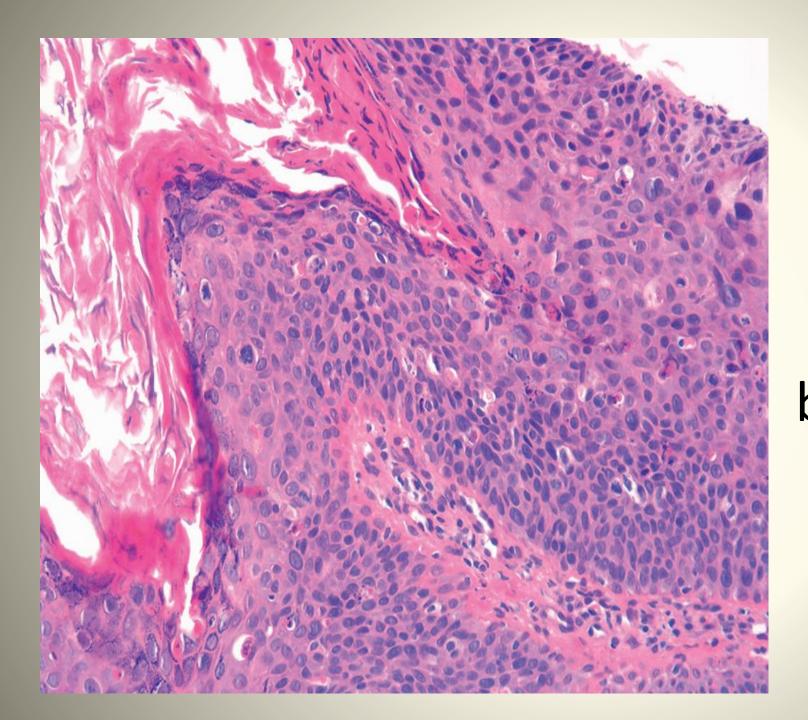
- typically involves the glans penis in young men
- similar to HSIL of the cervix with full-thickness involvement by small monotonous immature cells with high nucleartocytoplasmic ratio
- Mitotic activity and apoptotic bodies may be frequent, but squamous maturation is not characteristic
- show diffuse staining for p16
- a common association with HPV type 16



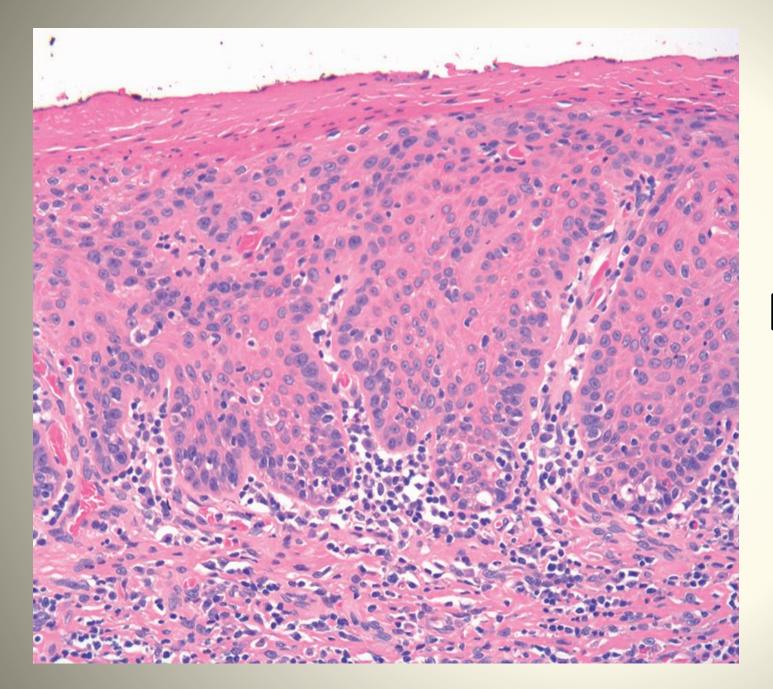
Basaloid PelN

Warty (Bowenoid) PelN

- has more complex architecture than basaloid PeIN
- frequent squamous maturation, a papillomatous surface and more abundant surface keratin, a contrasting point with basaloid PeIN
- more nuclear pleomorphism, and well-developed koilocytotic atypia
- expresses strong p16
- the associated HPV type is more variable



Wartybasaloid PeIN



Differentiated PelN