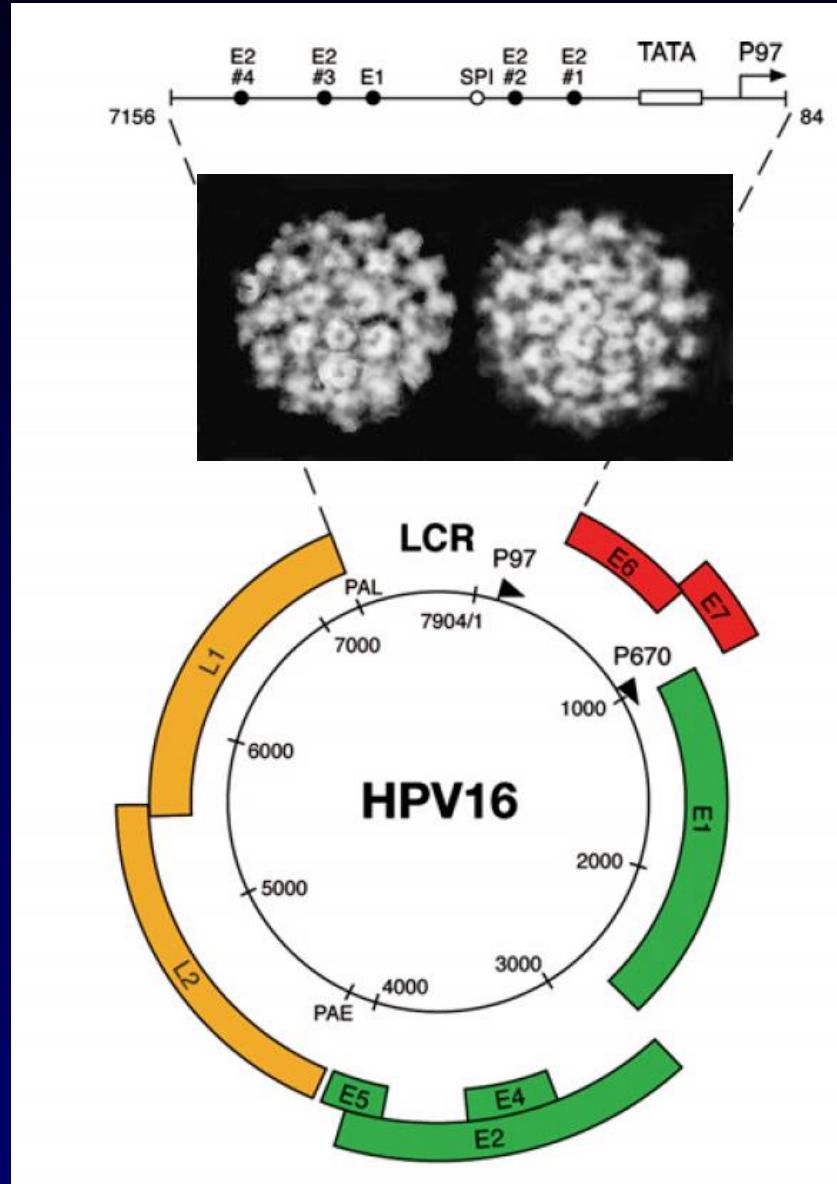


# **HPV and cervical cancer epidemiology**

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# Human papillomaviruses (HPVs)

- Small DNA viruses
  - Circular, double stranded DNA genome (8 kb); 6 early ORFs (E1-E7), 2 late ORFs (L1, L2)
- More than 200 different HPV types identified; strictly epitheliotropic: cutaneous or mucosal
- More than 40 mucosal HPV types: low-risk or high-risk, depending on their association with benign or (pre)malignant lesions



## GP5+/6+ consensus PCR system (L1 region)

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**GP5+** 5'- TTTGTTACTGTGGTAGATACTAC -3'

**GP6+** 3'- CTTATACTAAATGTCAAATAAAAAG -5'

**HPV 16** TTTGTTACTGT**TGTT**GATACTAC (2)

**HPV 18** TTTGTTACTGTGGTAGATA**CCAC** (1)

**HPV 31** TTTGTTACTGTGGTAGATA**CCAC** (1)

**HPV 33** TTTGTTACTGTGGTAGATA**CCAC** (1)

**HPV 35** TTTGTTACTGT**AGTT**GATA**CAAC** (3)

**HPV 39** TTT**CTT**ACTGT**TGTGGAC**ACTAC (4)

**HPV 45** TTTGTTACTGT**AGTGGAC**ACTAC (3)

**HPV 51** TTT**ATTAC**CTGT**GTTG**ATACTAC (6)

**HPV 52** TTTGT**CACAG**TTGTGGATA**CCAC** (5)

**HPV 56** TTTGTTACTGT**AGTAG**ATACTAC (1)

**HPV 58** TTTGTTAC**CGTGGT**TGATA**CCAC** (3)

**HPV 59** TTT**TTAAC**AG**TTGT**AGATACTAC (4)

**HPV 66** TTTGTTACTGT**TGTGG**ATACTAC (2)

**HPV 68** TTT**CTT**ACTGT**TGTGG**ATA**CCAC** (4)

CTTATACTAAATGTCAAATAAAAAG

CTTATACTAAAC**GTCAAATAAAAAG** (1)

CTTAA**ACTAAATGT**TAAATATAAAG (3)

CTTATACTA**GATGTCAAAC**AAAAAG (2)

CTTATACTAAATGTCAAATAAAAAG

**CTCATACTAAATGT**TAAATATAAAG (3)

CTTATACTAAATGTCAAATAAAAAG

**CTCATACTTAAAC**GTCAAATAAAAAG (4)

CTTAA**ACTAAATGT**TAAATAAAAAG (2)

CTTATACT**TAATGT**TAAACAAAAAG (3)

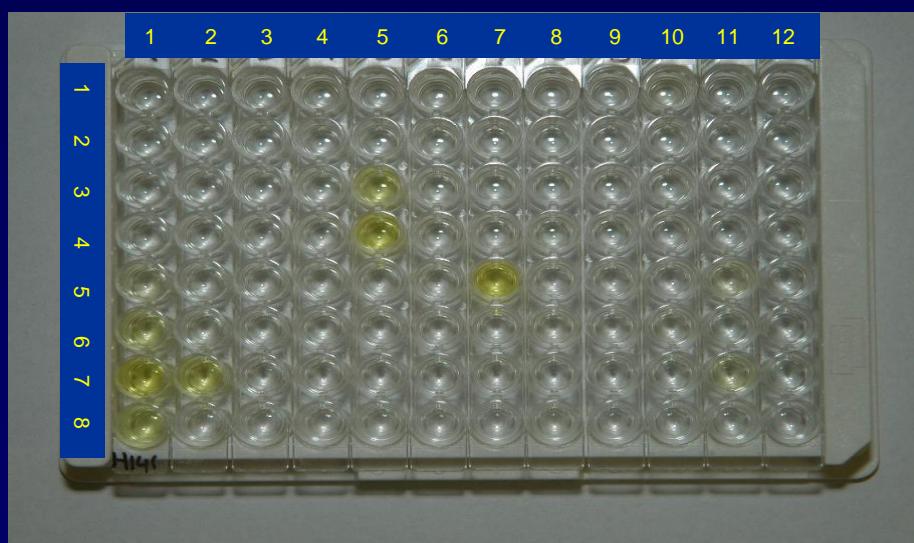
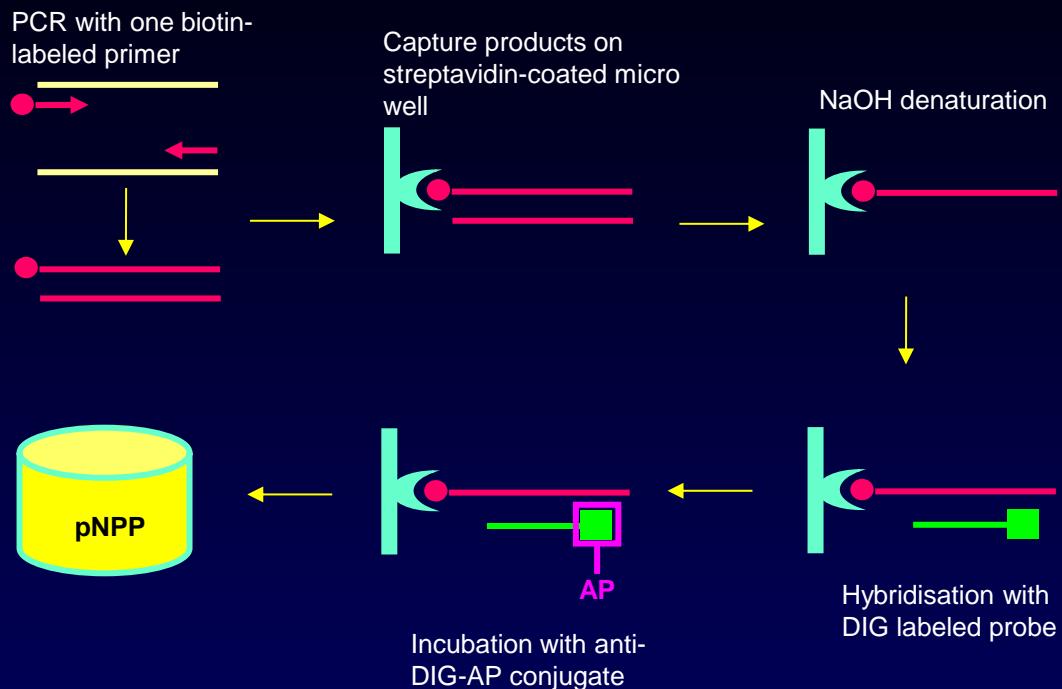
CTTATACT**GAATGTCAAAC**AAAAAG (2)

CTTAA**ACTAAAC**GTCAAATAAAG (3)

CTTATACT**TGATGTCAAAC**ACAAAG (4)

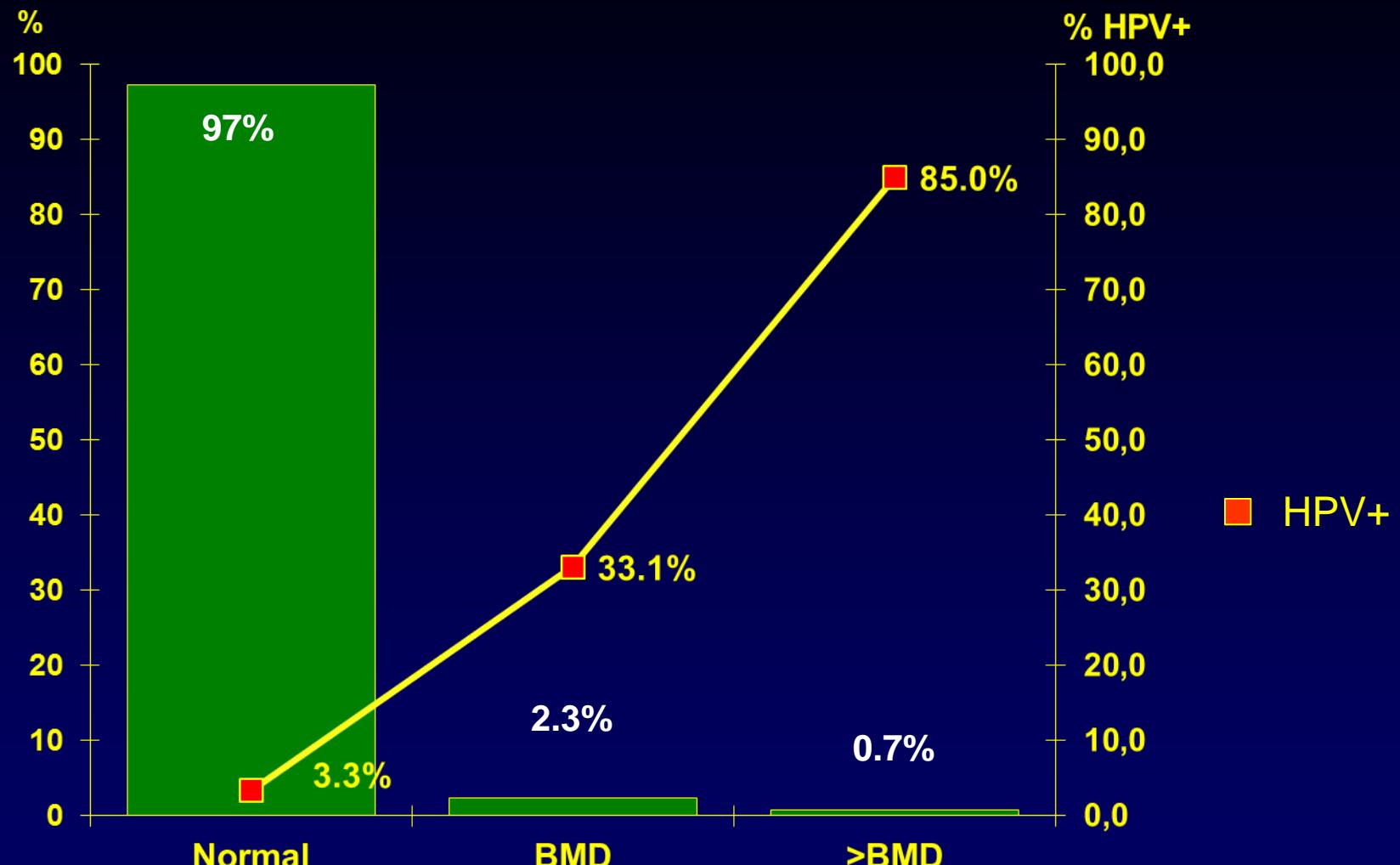
CTTATACTAAAC**GT**TAAATATAAAG (3)

# GP5+/6+-PCR read-out for high-throughput analysis



# **Association with disease**

# High-risk HPV prevalence per cytology class



HPV prevalence increases proportional to disease severity

BMD= borderline or mild dyskaryosis (ASCUS/LSIL)

# HUMAN PAPILLOMAVIRUS IS A NECESSARY CAUSE OF INVASIVE CERVICAL CANCER WORLDWIDE

JAN M. M. WALBOOMERS<sup>1\*</sup>, MARCEL V. JACOBS<sup>1</sup>, M. MICHELE MANOS<sup>2</sup>, F. XAVIER BOSCH<sup>3</sup>, J. ALAIN KUMMER<sup>1</sup>,  
KEERTI V. SHAH<sup>2</sup>, PETER J. F. SNIJDERS<sup>1</sup>, JULIAN PETO<sup>4</sup>, CHRIS J. L. M. MEIJER<sup>1</sup> AND NUBIA MUÑOZ<sup>5</sup>

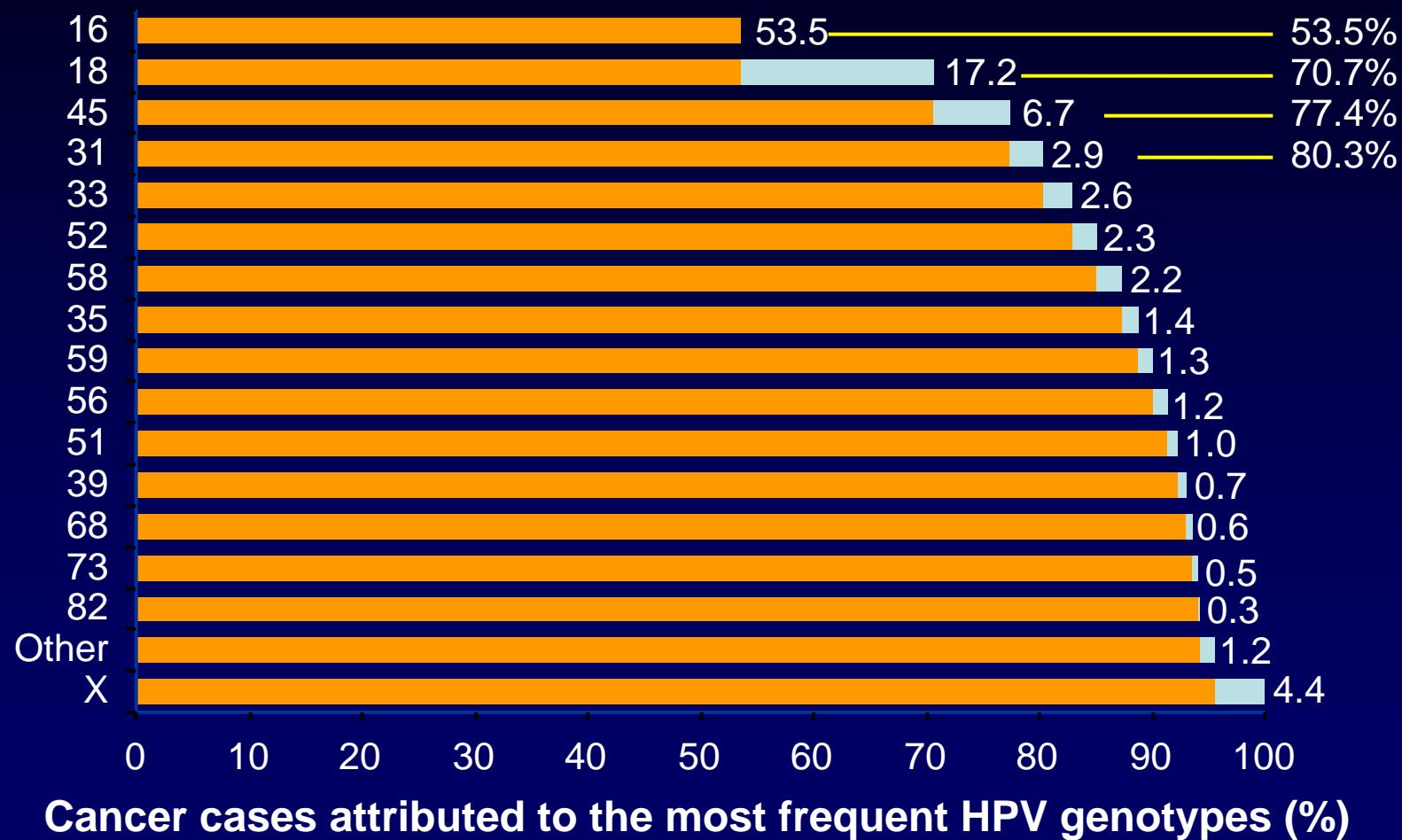
ORIGINAL ARTICLE

# Epidemiologic Classification of Human Papillomavirus Types Associated with Cervical Cancer

Nubia Muñoz, M.D., F. Xavier Bosch, M.D., Silvia de Sanjosé, M.D.,  
Rolando Herrero, M.D., Xavier Castellsagué, M.D., Keerti V. Shah, Ph.D.,  
Peter J.F. Snijders, Ph.D., and Chris J.L.M. Meijer, M.D., for the International  
Agency for Research on Cancer Multicenter Cervical Cancer Study Group\*

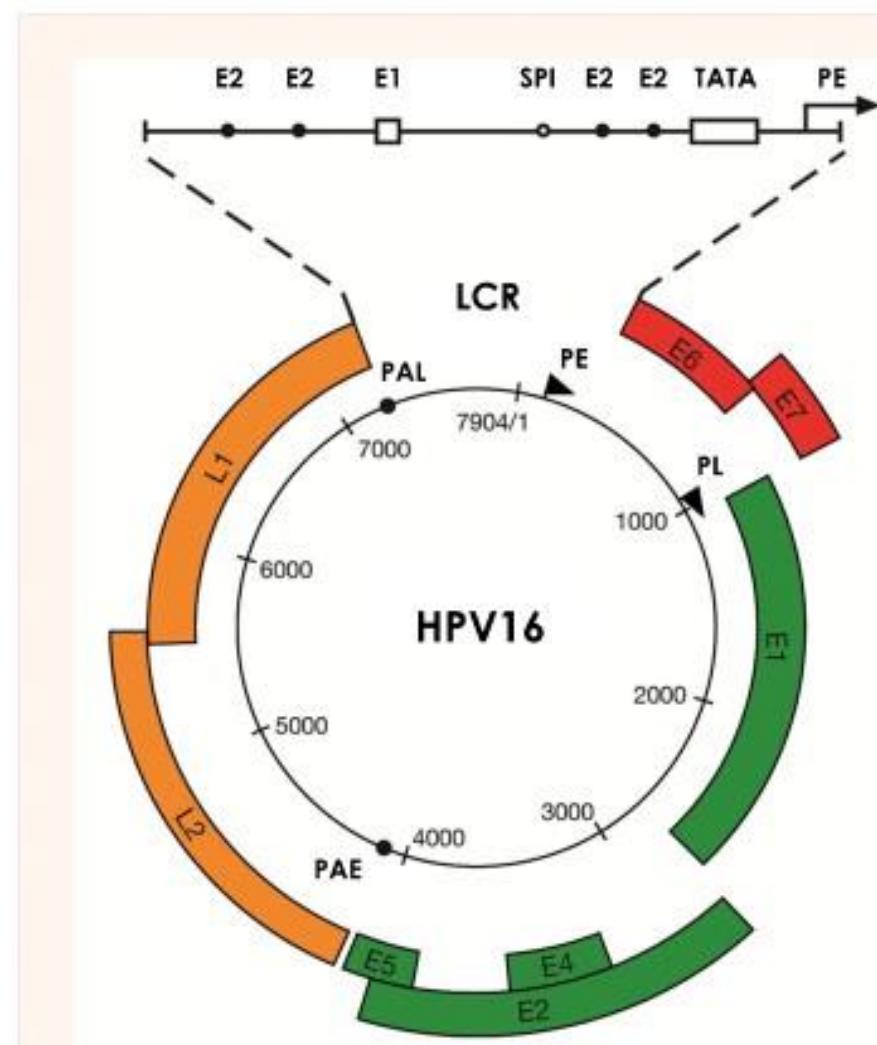
# HPV types in cervical cancer worldwide

## HPV genotype



**A**

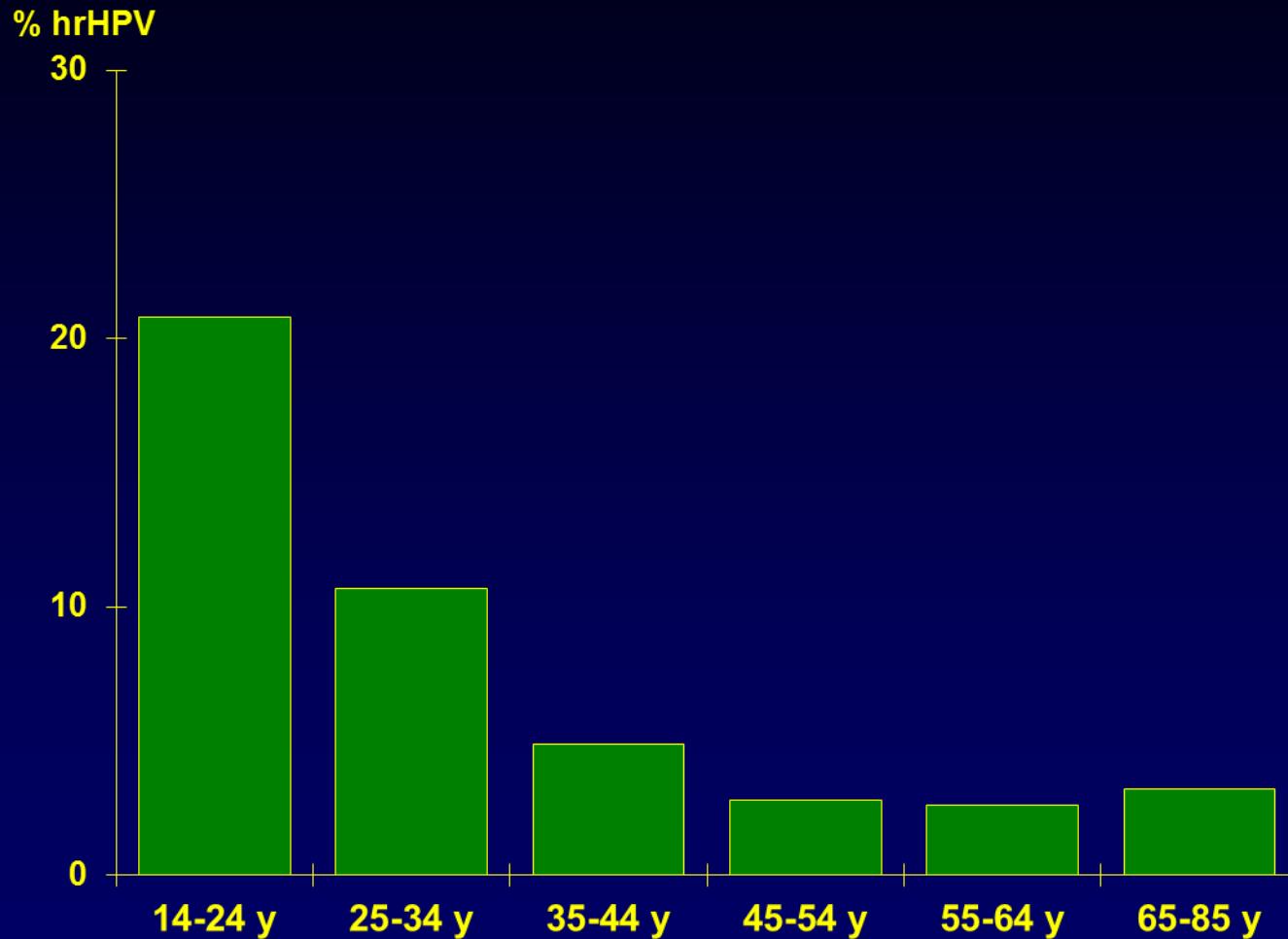
Genus + Species	Type	Invasive Cervical Cancer	IARC Category	Squamous Cell Carcinoma	Adeno Carcinoma	Tropism
Alpha 1	HPV32		3			mucosal
Alpha 2	HPV42					
	HPV3		3			
	HPV10		3			
	HPV28		3			
	HPV29		3			
	HPV77		3			
	HPV94		3			
	HPV117		3			
	HPV125		3			
Alpha 3	HPV61	0.01	3			
	HPV62		3			
	HPV72		3			
	HPV81		3	0.4		
	HPV83		3	0.4		
	HPV84		3			
	HPV86		3			
	HPV87		3			
	HPV89		3			
	HPV102		3			
Alpha 4	HPV2		3			
	HPV27		3			
	HPV57		3			
Alpha 5	HPV26	0.37	2B	0.22		
	HPV51	1.25	1	0.75	0.54	
	HPV69	0.08	2B			
	HPV82	0.07	2B	0.26		
Alpha 6	HPV30	0.37	2B			
	HPV53	0.26	2B	0.04		
	HPV56	0.84	1	1.09		
	HPV86	0.08	2B	0.19		
Alpha 7	HPV18	10.28	1	11.27	37.3	
	HPV39	1.67	1	0.82	0.54	
	HPV45	5.68	1	5.21	5.95	
	HPV59	1.08	1	1.05	2.16	
	HPV68	1.04	2A	0.37		
	HPV70	0.11	2B			
	HPV85		2B			
	HPV97					
Alpha 8	HPV7		3			
	HPV40		3			
	HPV43		3			
	HPV91	0.01	3			
Alpha 9	HPV16	61.35	1	54.38		
	HPV31	3.35	1	3.82	0.54	
	HPV33	3.83	1	2.06		
	HPV35	1.94	1	1.27		
	HPV52	2.71	1	2.25		
	HPV58	2.22	1	1.72		
Alpha 10	HPV67	0.31	2B	41.62		
	HPV6	0.11	3	1.08		
	HPV11	0.02	3	0.54		
	HPV13		3			
	HPV44	0.01	3			
	HPV74	0.01	3			
Alpha 11	HPV34	0.07	3			
	HPV73	0.52		0.49		
Alpha 12	HPV73					
Alpha 13	HPV54					
Alpha 14	HPV71					
	HPV90					
	HPV106		3			

**B**

IARC category 1=carcinogenic, 2A=possibly carcinogenic, 2B= possibly carcinogenic because of phylogenetic relationship with 1 or 2A, 3=non-carcinogenic

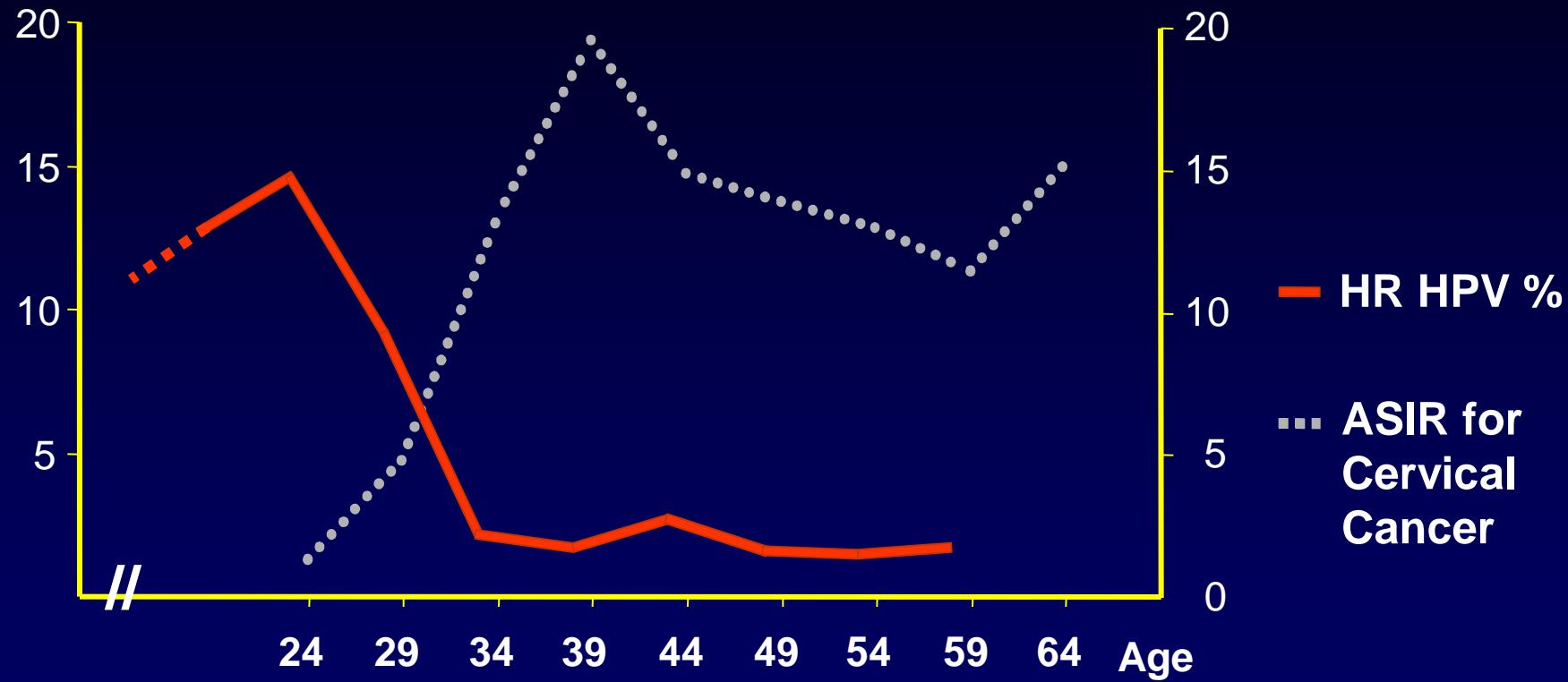
**Temporality: does hrHPV  
infection precede lesion  
development?**

# HPV prevalence according to age



HPV prevalence age-dependent: peak prevalence below 20 years of age

# Age specific prevalence of hrHPV-DNA and ASIR cervical cancer in the Netherlands



Peak HPV prevalence precedes that of ASIR cervical cancer

# Incident CIN3 is preceded by hrHPV infection

Case	Age (years)	Routine screening (Pap)	NNS (Pap)	High risk HPV	Follow up time (years)	CIN
1	34	1	1	Yes	2.1	3
2	35	2	3a2	Yes	2.6	3
3	35	1	1	Yes	3.4	3
4	35	2	2	Yes	2.6	3
5	35	1	1	Yes	5.7	3
6	36	1	1	Yes	2.9	3
7	38	1	3a2	Yes	3.0	3
8	38	1	2	Yes	1.5	3
9	38	1	1	Yes	7.9	3
10	42	1	1	Yes	5.1	3
11	42	1	2	Yes	8.9	3
12	53	2	3a1	Yes	0.9	3
13	41	2	1	No*	4.7	3

# 13 Incident CIN3 from 2250 with normal/ very mildly abnormal Pap enrolled

\* Acquired hrHPV two years following enrolment

**Is hrHPV persistence  
related to disease  
progression?**

# Follow-up of women with abnormal cytology in relation to HPV status: progression/regression

- 353 women referred because of abnormal smear between 1990-1992
  - Monitoring: every 3-4 months by cytology, colposcopy and hrHPV testing
  - Primary endpoint: clinical progression
    - Colposcopical impression of CIN3 over  $\geq 3$  quadrants and/or
    - Cytology suspected of invasive carcinoma (Pap5)
  - Secondary endpoint: histology CIN3 at the end of the study or when reaching primary endpoint.
  - Cytological regression: abnormal cytology returned to normal cytology for at least two consecutive cervical smears
  - HPV clearance: none of the hrHPV types from the previous visit were detected at the next visit
  - Mean age: 32 years (range 18-75)
  - Median follow up time: 33 months (3-74)

# hrHPV status during study and risk for end histology of CIN3

HPV status during study	Number of women	Clinical progression (n=33)		No clinical progression (n=320)				OR for CIN3 histology (95% CI)
		CIN3 (n=32)	CIN2 (n=1)	CIN3 (n=71)	CIN2 (n=29)	CIN1 (n=64)	Normal (n=156)	
HPV persistence	122	32	1	66	7	6	10	327 (42-2468)
HPV clearance or acquisition	150	-	-	4#	15	35	96	2.9 (0.2-20)
Negative	81	-	-	1	7	23	50	1.0

# Two women cleared their infection and acquired another infection persisting for 24 and 32 months, respectively

Two women tested negative at baseline but acquired an infection that persisted for 32 months

# Large Dutch screening studies

POBASCAM: 44,102 women (RCT)

Setting : regular screening program (5 year interval)

Randomised: HPV + cytology vs Cytology (HPV blinded)

Follow-up 5+ years (incl. subsequent screening round)

HPV test: GP5+/6+ PCR EIA

Primary endpoint: nr of CIN3+ in each arm

VUSA-SCREEN: 25,871 women (cohort study)

Setting : regular screening program (5 year interval)

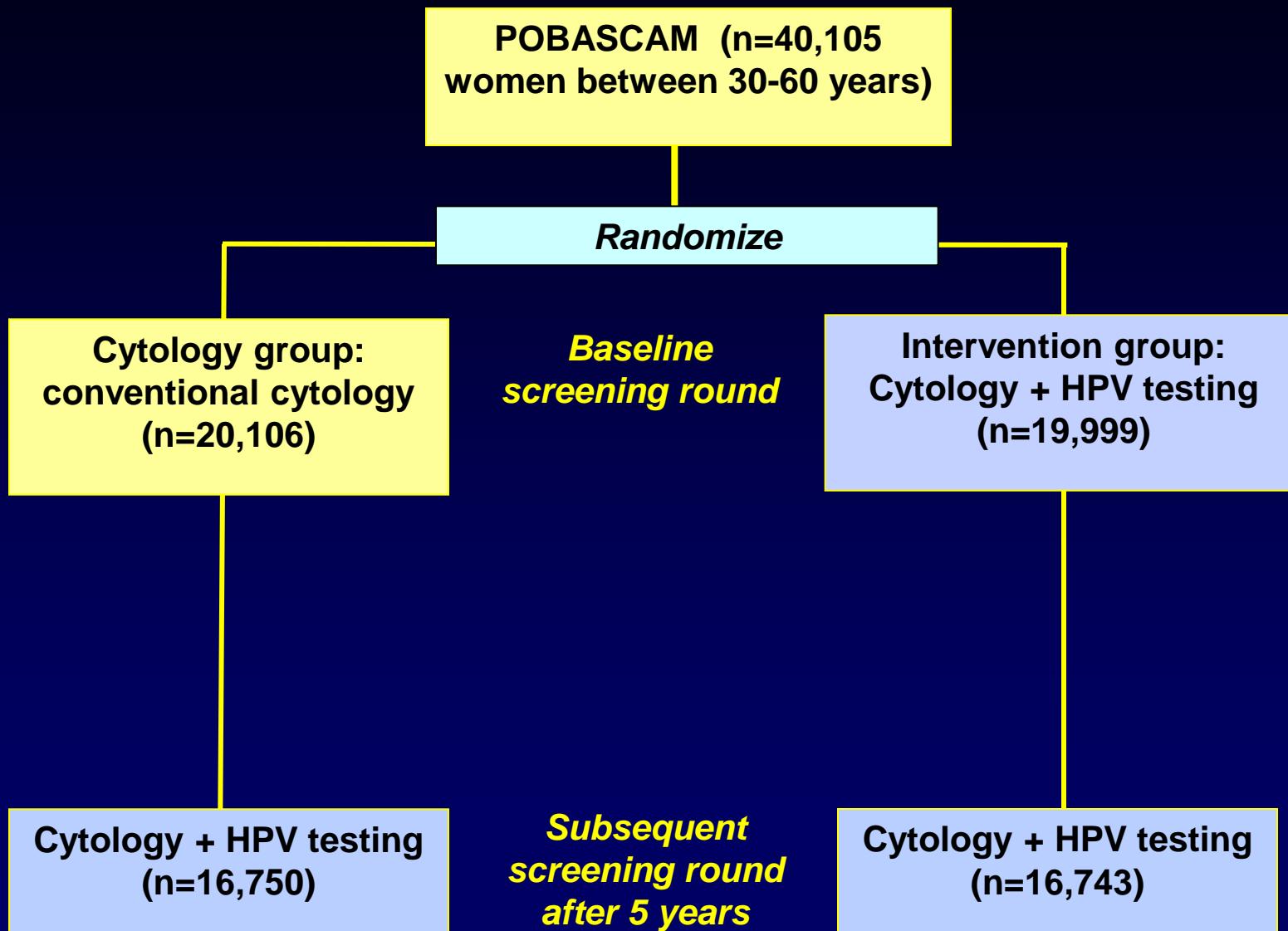
HPV + cytology

Follow-up 2 years (2004-2007)

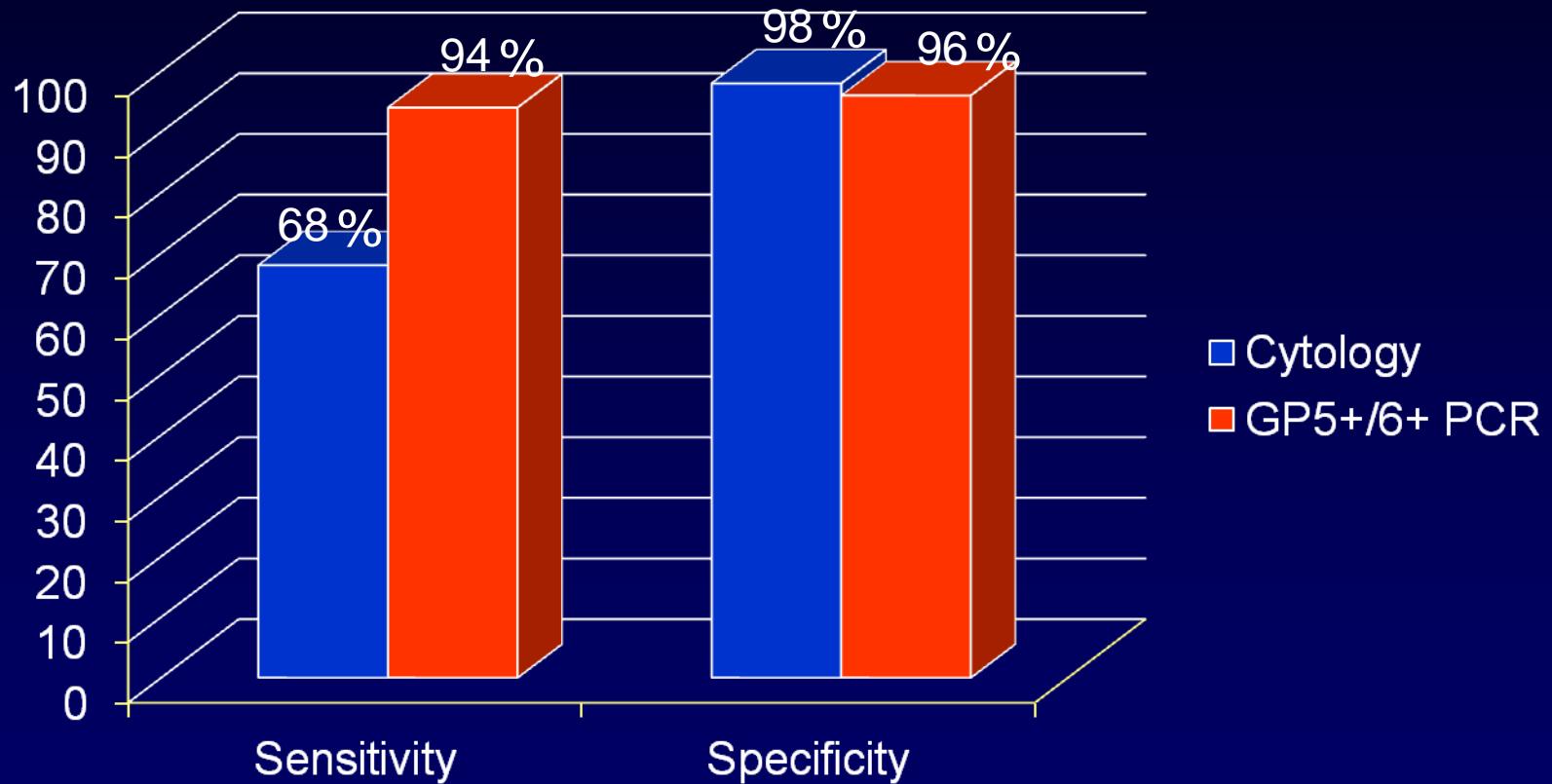
HPV test: Hybrid capture 2 (hc2)

Primary endpoint: CIN 3+

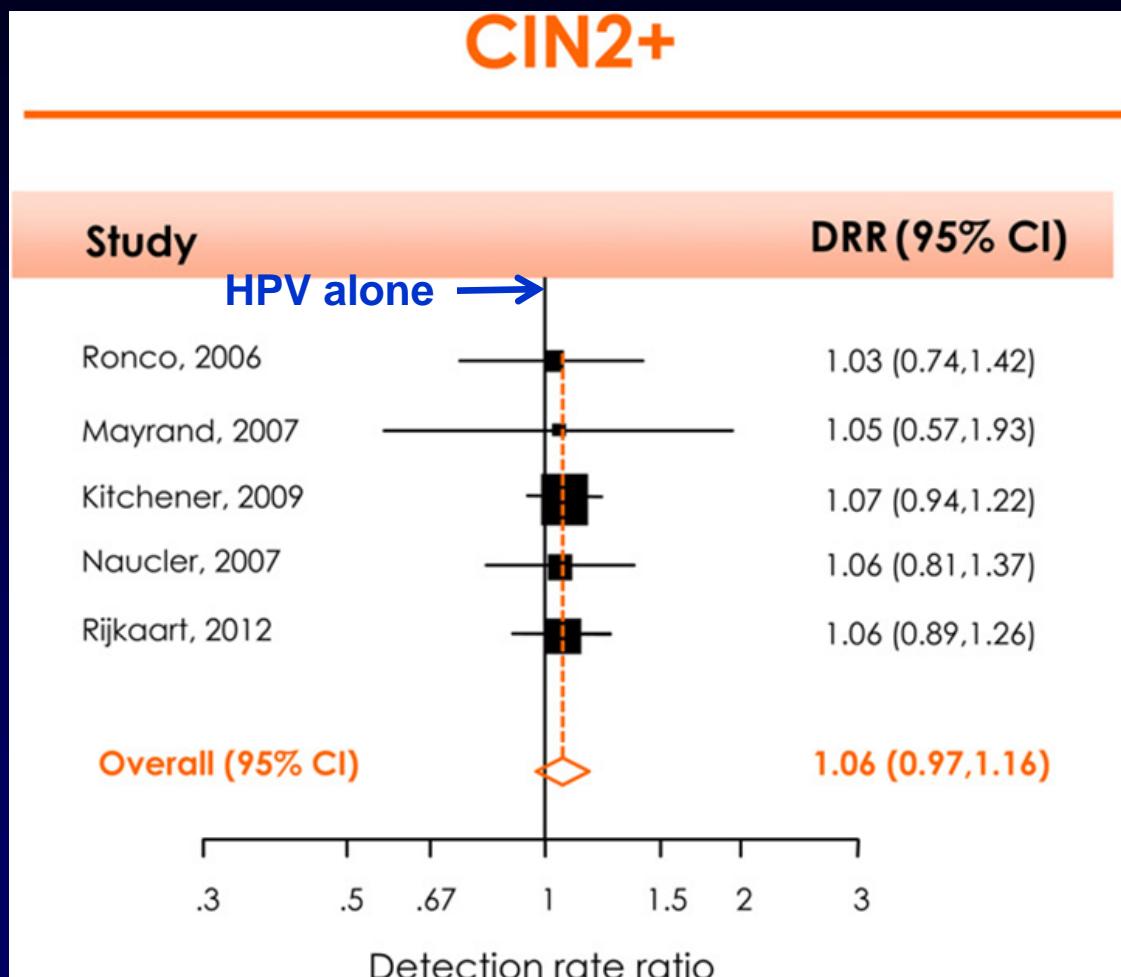
# POBASCAM screening study



# Cross-sectional sensitivity/specificity for CIN2+ of HPV test versus cytology

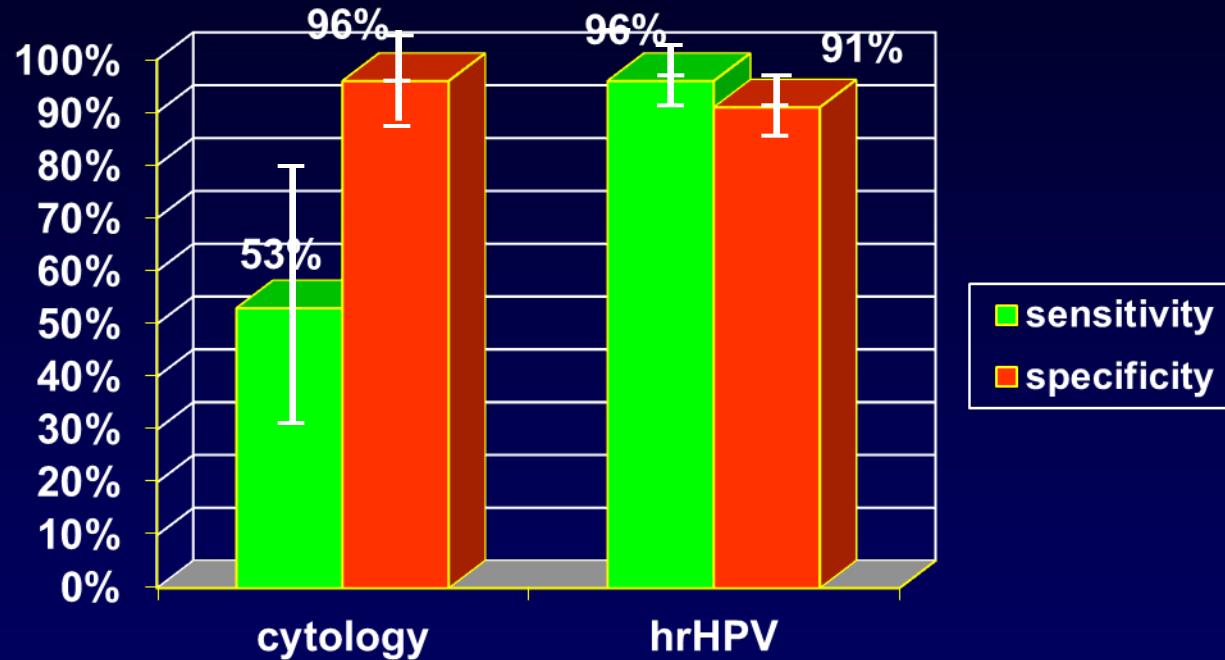


# Performance HPV & Pap (combo) vs HPV test alone



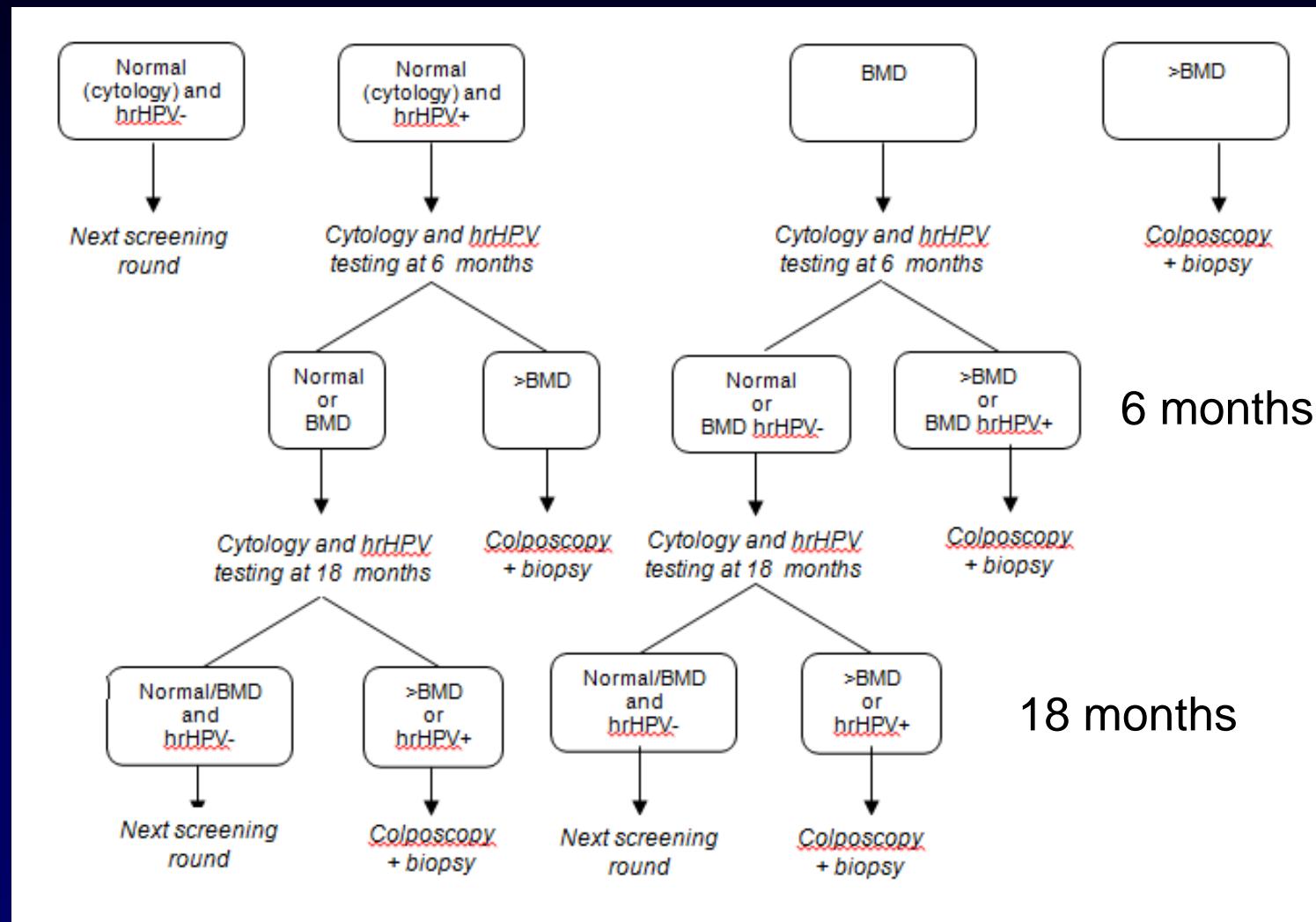
Sensitivity of HPV&Pap is not higher than solely HPV testing

# Sensitivity and specificity for CIN2+: HPV testing vs Cytology in cross-sectional studies



Overview of European and North American studies (n=~ 60,000)

# Management of women in the intervention group at the baseline and subsequent round and the control group at the subsequent round



# POBASCAM trial: cumulative 18-month risk CIN3+

- Overall hrHPV+ (95%CI)
  - Normal 6% (4-9)
  - BMD 20% (16-25)
- HPV16+
  - Normal 14% (9-21); p<.0001
  - BMD 37% (28-48); p<.0001

After excl. HPV16:

- HPV18+
  - Normal 9% (2-23); p=.031
  - BMD 11% (3-27); p=.883
- HPV31+
  - Normal 7% (2-16); p=.049
  - BMD 27% (14-46); p=.016
- HPV33+
  - Normal 10% (0-33); p=.087
  - BMD 22% (9-44); p=.025

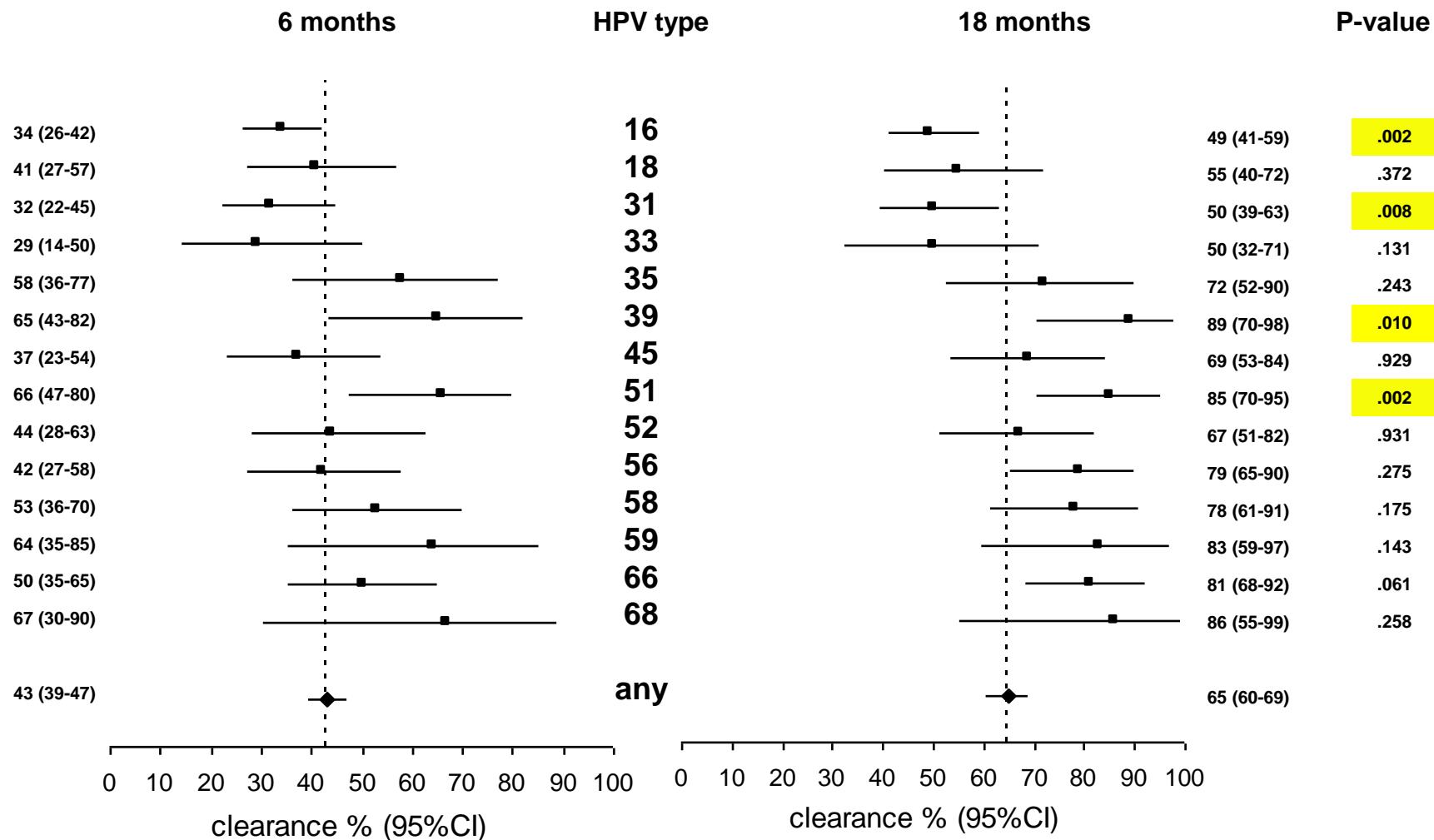
## Summary 18-months risk CIN2+/CIN3+

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- HPV 16+ confers an increased risk of CIN2+/CIN3+ both in women with **normal cytology** and **BMD**
- In the subset of women without HPV 16:
  - HPV 18, 31 and/or 33: increased risk in women with **normal cytology** and/or **BMD**
- After repetitive normal smears:
  - Only HPV 16 and HPV 18 were associated with an increased risk of high-grade CIN ( $p=0.028$ )
    - HPV 16/18+ : 9% (4-48)
    - Non-HPV 16/18: 2% (0-6)
- Opens possibilities for algorithms with HPV16/18 genotyping incorporated to triage women with **normal cytology**

# hrHPV type-specific clearance rates: women with normal cytology

## Clearance Rate % (95%CI)

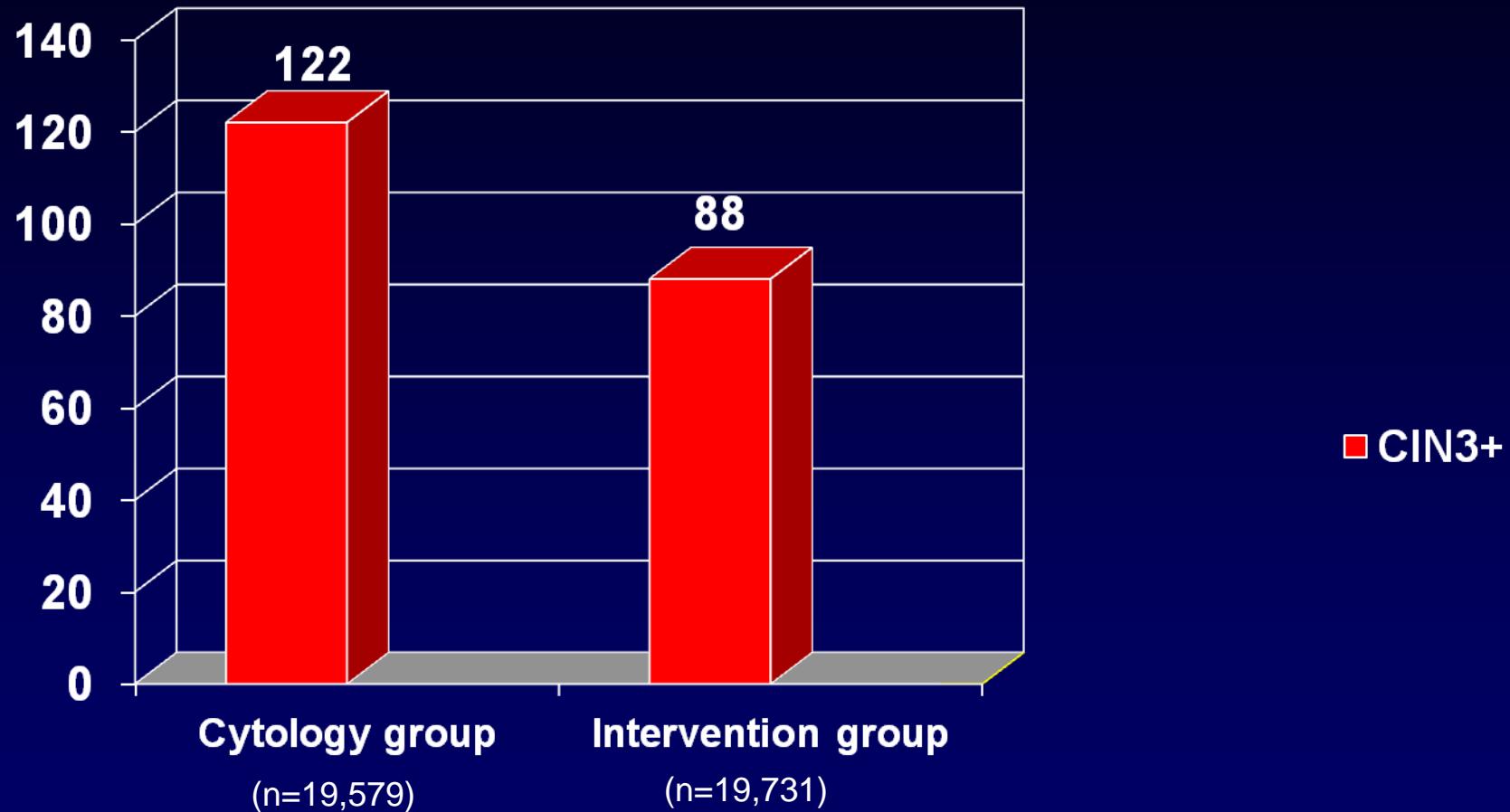


Differential risks at least in part attributable to differential clearance rates

Bulkmans et al Br J Cancer 2007

# **POBASCAM results subsequent screening round**

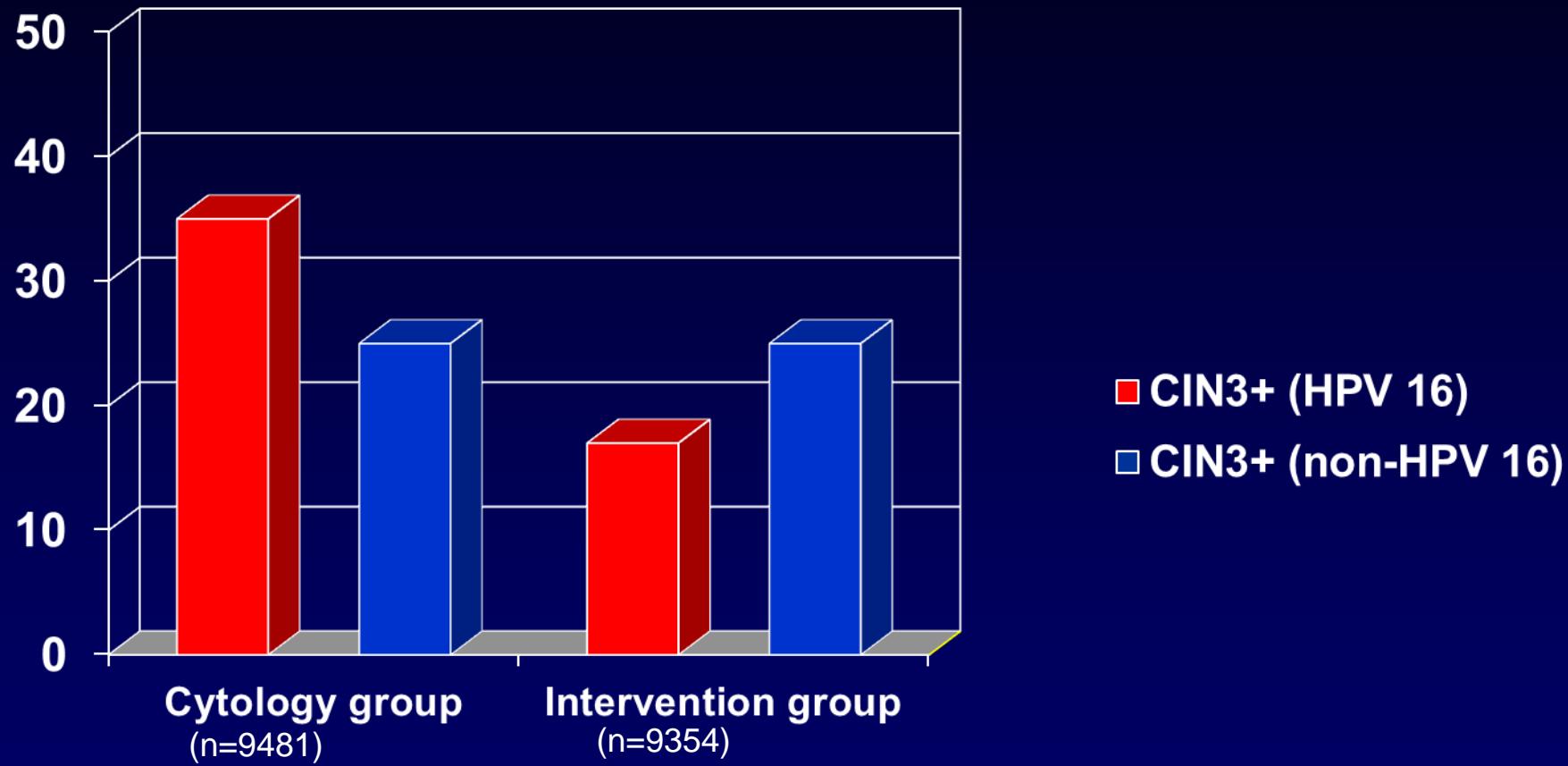
# Final results POBASCAM at subsequent screening round (after 5 years)



27% less CIN3+ in HPV group ( $p=.02$ )

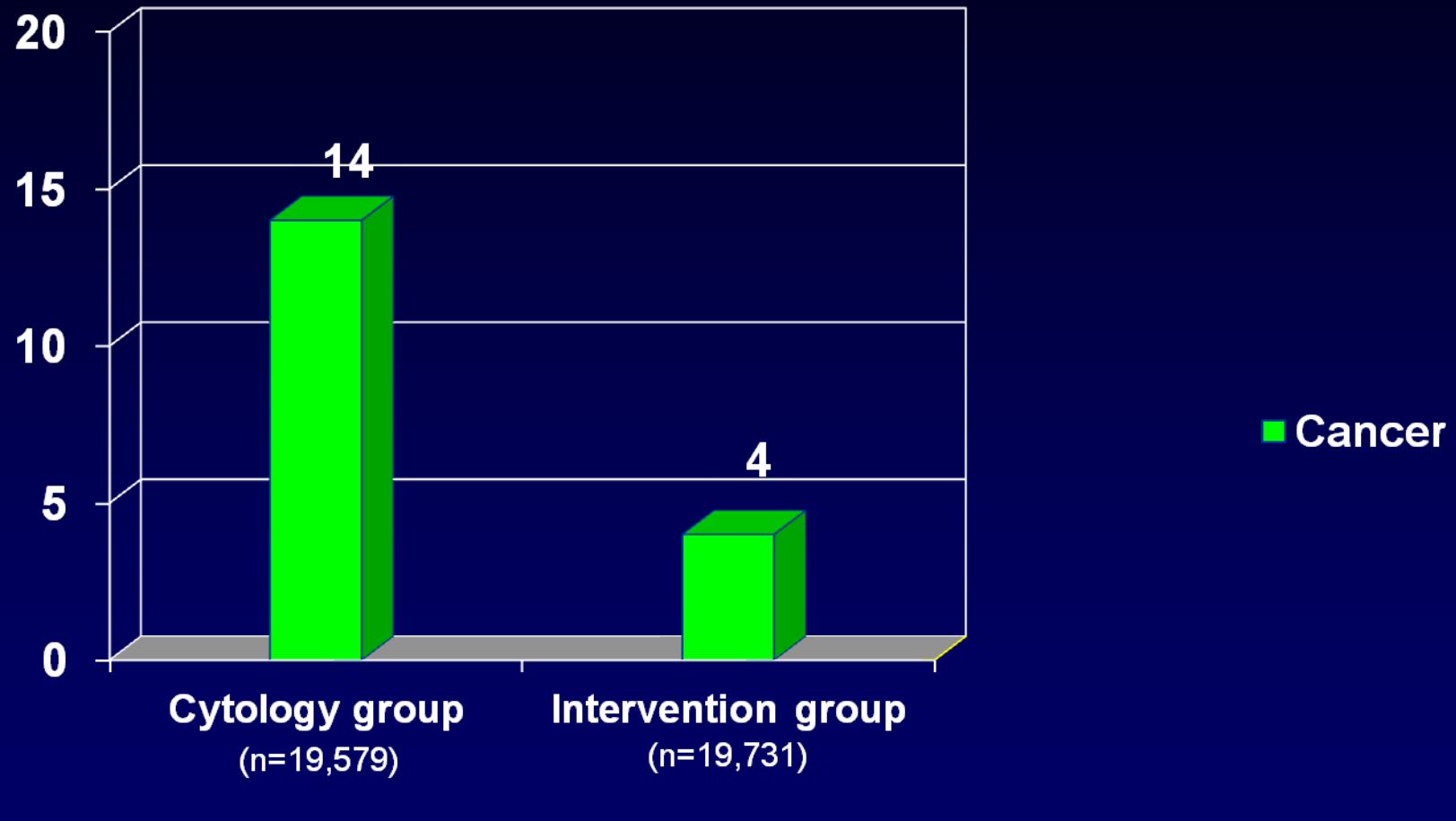
*Rijkaart et al. Lancet Oncol. 2012*

# Final results POBASCAM at subsequent screening round (after 5 years)



**Protective effect largely attributable to HPV16**

# Final results POBASCAM at subsequent screening round (after 5 years)



71% less cancer in HPV group ( $p=.03$ )

# Conclusion

- hrHPV is necessary for development of premalignant cervical disease and progression towards carcinoma

Implications for screening:

- HPV testing allows *earlier* detection of clinically relevant CIN2+ lesions and consequently better prevents invasive cervical cancer than cytology
- This permits extension of the screening interval without increasing interval risk
- HPV test valuable as primary screening test (cytology as triage test for HPV positive women)

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