Evidence for clinical utility of extended HPV genotyping in persistence tracking and follow-up after abnormal results and colposcopy and test-of-cure

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Declaration of Conflict of Interest: Worldwide Medical Director, Women's Health & Cancer, BD Life Sciences.

Natural history of HPV infections Average clearance, persistence, & progression Persistence Schiffman M, Castle PE, Jeronimo J, Rodriguez AC, Wacholder S. Human papillomavirus and

Systematic Review Results

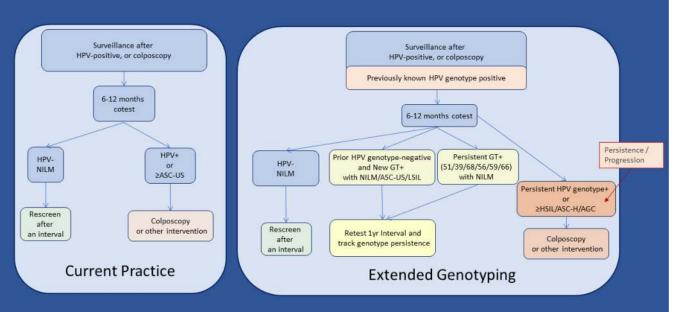
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- · Follow-up after abnormal screening result
 - among those with repeated HPV-positive results, same genotype persistence was associated with substantially elevated risk for high-grade lesions compared to the pooled risk of positive for any type
 - · HPV 16 had the greatest tendency to persist and the highest probability for progression when persistent
 - · HPV 18, 31, 33 were the high-risk HPV types other than HPV 16 associated with high absolute risks for progression
 - · Women with low-grade cytology and positive HPV, those testing negative for the highest-risk types of HPV (16/18/31/33/45/52/58) may not need immediate colposcopy and biopsy, if clinical decisions are risk-based

Systematic Review Results

- Test-of-cure
 - same genotype persistence was associated with substantially elevated risk for high-grade lesions compared to the pooled risk of positive for any type
 - · genotyping had significantly higher sensitivity and specificity to predict residual/recurrent high-grade CIN
 - · almost all disease found at ToC visit was associated with type-persistent hrHPV – 100% in most studies
 - specificity (same genotype clearance) was 100% in most studies
 - proportion of HPV-positive women at ToC with same-genotype persistence (versus new HPV) varied by population from 30% to 80%
 - HPV 16 persistence conferred highest risk

Persistence tracking after screening/colposcopy, with extended genotyping (apply equal management for equal risk)



Citations Follow-Up Abnormal and Colposcopy

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Follow-up guidelines and standards of care for surveillance

- HPV-positive, non-16/18, NILM cytology
 - · Cotest in 12 months
- Colposcopy for lesser abnormalities, impression negative or low-grade, biopsy ≤CIN1 Cotest in 6-12 months
- Excisional treatment of high-grade CIN
 - Cotest in 6-12 months

Follow-up cotest results

- HPV-negative and cytology NILM
- HPV-negative and cytology ASC-US/LSIL
- cytology HSIL/ASC-H/AGC and any HPV result
- HPV-positive and cytology NILM
 - Clearance of prior genotype and new infection OR same-genotype persistence?
- HPV-positive and cytology ASC-US/LSIL
 - Clearance of prior genotype and new infection OR same-genotype persistence?

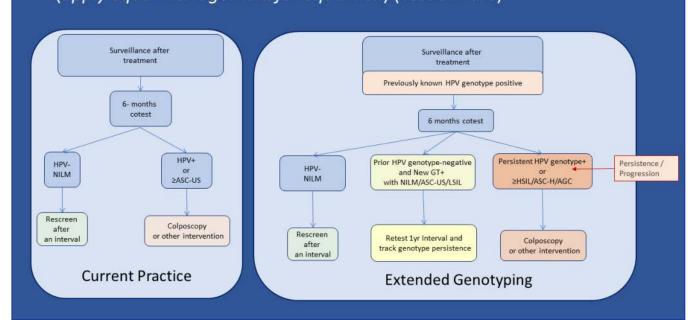
Systematic Review Results

- Post-colposcopy follow-up
 - among those with repeated HPV-positive results, same genotype persistence was associated with substantially elevated risk for high-grade lesions compared to the pooled risk of positive for any type
 - An invited editorial accompanying one of the articles stated, "The future of screening and management will rely on the identification of patients who have the highest risk for developing CIN 2+ and, when appropriate, aggressively treating those at highest risk. There are many known risk factors for the development of CIN such as HPV status, persistence of HPV, known history of CIN, smoking, and HPV vaccine status, and it has become evident that HPV persistence is perhaps the most important risk factor." and "Although present treatment guidelines do not specifically stratify treatment and follow-up based on the presence of certain genotypes, it is perhaps time to seriously consider this: it is time to graduate to greater utilization of typespecific testing."[Huh 2017 AJOG]

Systematic Review Conclusions

- Women with follow-up cotesting may have type-specific persistence, or clearance, or a new HPV infection
 - the risk posed by a new genotype infection is significantly lower than the risk posed by a type-specific persistent infection
 - Risk-based management should differ between these
- Genotype results could be grouped into 3-4 risk tiers (stratification) for NILM results with same genotype persistence
 - genotypes will be grouped into tiers according to similar risk values, based on local thresholds for management actions
- need more studies, meta-analysis
- need clinically validated assays to be tested

Persistence tracking after treatment of CIN, with extended genotyping (apply equal management for equal risk) (Test of Cure)



Citations Test-of-Cure

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