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Clinical Validation of HPV ctDNA for Early Detection of Residual Disease Following Chemoradiation in Cervical Cancer

Presented by: Kathy Han, MD, MSc Princess Margaret Cancer Center

Disclosure & Study Team

- Employer: Princess Margaret Cancer Centre, University of Toronto
- Advisory Board: AstraZeneca
- Patent: HPV ctDNA (HPV seq) pending
- This study was supported by funding from Cancer Research Society, Ontario Institute for Cancer Research and the Princess Margaret Cancer Foundation.

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Background

- Tumours continually shed their DNA into the circulation
- Majority (>90%) of cervical cancer caused by human papillomavirus (HPV)
- Non-invasive liquid biopsy tests can detect HPV circulating tumor DNA (ctDNA) in the blood as a measure of disease burden
- Our pilot study showed detectable HPV ctDNA at the end of chemoradiation is associated with worse progression-free survival



Han et al, JCO Prec Oncol, 2018; 2:1-8 Leung*, Han*, et al. Clin Cancer Res 2021; 5857-68

Newer HPV ctDNA detection technologies

(A) Droplet-based digital polymerase chain reaction (dPCR)



Sample partitioned into thousands of nanolitersized droplets, each with independent amplification events



dPCR probes target <u>E6 & E7</u> regions (B) Next generation sequencing



Rostami, Bratman, Han. Clin Cancer Res 2021; 5158-60

Method

- Prospective multicenter validation study
- 70 patients with stage IB-IVA HPV+ cervical cancer treated with definitive (chemo)radiation between 2017-2022
- HPV genotype determined using baseline plasma sample with HPV-seq
- HPV genotype-specific plasma DNA levels quantified using dPCR & HPV-seq



Results: dPCR

End of chemoradiation 4 – 6 weeks post CRT 3 months post CRT

(CRT) Progression-Free Survival 1.00 1.00 1.00 0.75 0.75-0.75 p<0.001 p<0.001 HR=8.58 HR=6.29 0.50-0.50 0.50 95% CI (3.56,20.71) 95% CI (2.71,14.59) p=0.036 0.25 0.25 dPCR 4-6w postCRT dPCR 3m postCRT dPCR end CRT HR=2.53 Negative 95% CI (1.06,6.03) Negative Negative Positive Positive - Positive 0.00 21 28 35 42 <u>1</u>9 56 63 56 28 35 56 35 63 21 63 Time (months) Time (months) Time (months) 33 26 43 0 13 3 2 0 0 ctDNA-positive 5 12 7 5 2 Δ 19 16 11 ctDNA-positive 2 25 Numbers at risk Numbers at risk Numbers at risk

— Undetectable HPV ctDNA — Detectable HPV ctDNA

Results: HPV-seq

End of chemoradiation 4 – 6 weeks post CRT 3 months post CRT



— Undetectable HPV ctDNA — Detectable HPV ctDNA

Conclusions

- HPV-seq enables determination of HPV type (genotyping) directly from plasma
- Persistent HPV ctDNA following chemoradiation is independently associated with inferior progression-free survival
- HPV ctDNA testing can be used to identify, as early as at the end of chemoradiation, patients at high risk of recurrence

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