## Circulating tumor cell (CTC) enumeration in patients (pts) with metastatic genitourinary (mGU) tumors treated in a phase I study of cabozantinib and nivolumab (CaboNivo) +/- ipilimumab (CaboNivolpi)

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## BACKGROUND

- Circulating Tumor Cells (CTCs) and circulating T-Cells may serve as biomarkers for clinical outcomes in GU tumor patients.
- Cabozantinib may have immunomodulatory properties that counteract tumor-induced immunosuppression, providing a rationale for combining cabozantinib with checkpoint inhibitors.
- We examined the association between CTCs and T-Cell populations at baseline and post-treatment at cycle 2 and 3, with progression free survival and overall survival (OS) and response to therapy with combination cabozantinib and nivolumab or cabozantinib, nivolumab, and ipilimumab.

## METHODS

Blood samples from mGU cancer patients undergoing CaboNivo or CaboNivolpi therapy were collected at baseline and on-therapy and sent to Epic Sciences for processing. Slides were stained with pan-CK/CD45/PD-L1/DAPI for CTC detection or CD4/CD8/Ki-67/DAPI for T-Cell analysis. Approximately 3 million cells per slide were imaged through advanced digital pathology pipelines to detect and quantify changes in immune cell populations and to assess circulating tumor cells burden.



Example Slide with 3x10<sup>6</sup> Cells Plated

The Epic Functional Cell Profiling (FCP) platform enables characterization of CTC and immune cell populations from a single blood sample. software performs cloud-based image Epic's analysis to characterize 3 million cells per slide. Representative slide image (above) are shown for healthy donor samples that were stained with CD4, CD8, Ki-67, and DAPI. The histograms were used to determine mean fluorescent intensity (MFI) thresholds for each marker and to quantify leukocyte subpopulations.





detect and characterize CTC and T-cell at the single cell level with a sensitivity of 10<sup>-6</sup> - 10<sup>-7</sup> from a single tube of blood

High CTC counts at Cycle 2, and low %CD4 in GU-cancer patients is associated with lower response to therapy and shorter survival Ongoing efforts include morphology analysis of T-cell populations and single cell sequencing of CTC subtypes associations with response