

# The Human Biofluid RNA Atlas

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AACR – Advances in Liquid Biopsies January 13-16, 2020 Miami, Florida

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\*CRIG illumina







no conflicts of interest to declare

#### expanding the liquid biopsy field beyond the blood stream



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RNA sequencing allows detailed analysis of the transcriptome



#### spike-in RNAs as processing controls and normalization tool



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## highly variable RNA concentrations amongst different biofluids



Preprint on BioRxiv

## highly variable RNA concentrations amongst different biofluids



#### assessment of the tissues of origin in the different biofluids



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#### different cell types contribute to the RNA content of pancreatic cyst fluid



RNA seq of pancreatic cyst fluid

Hulstaert et al, in review Preprint on BioRxiv

### different cell types contribute to the RNA content of pancreatic cyst fluid



differentially expressed mRNAs in urine of bladder cancer patients vs healthy controls

6 KRT5 PLAC4 4 RGS18 47 downregulated 867 upregulated MNDA TGM4 **mRNAs** mRNAs MMP1 MDH1B PLBD1-AS1 2 NEFHP1 ATP5MDP1 0 Hulstaert et al, in review -6 -2 2 6 0 4 -4 -1 Preprint on BioRxiv

volcanoplot (bladder cancer versus control)

log2 fold change

#### circRNAs are enriched in biofluids compared to tissues







circRNA

#### circRNAs are enriched in biofluids compared to tissues



### Take-home messages and future perspectives

- Dare to collect and investigate "alternative" fluids
- RNA is cool!
- Our findings enable a more informed selection of the most relevant biofluid to monitor individual cancer types
- Broader sample collections in different cancer types are ongoing
- We are open to new collaborations!



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