



## INTRODUCTION

- Intra-tumor heterogeneity and the tumor sampling strategies for profiling greatly influence the consensus molecular subtyping [1,2].
- Molecular subtypes correlate with tumor morphology [3].
- Morphological features can predict the CMSs [4,5]

### AIM

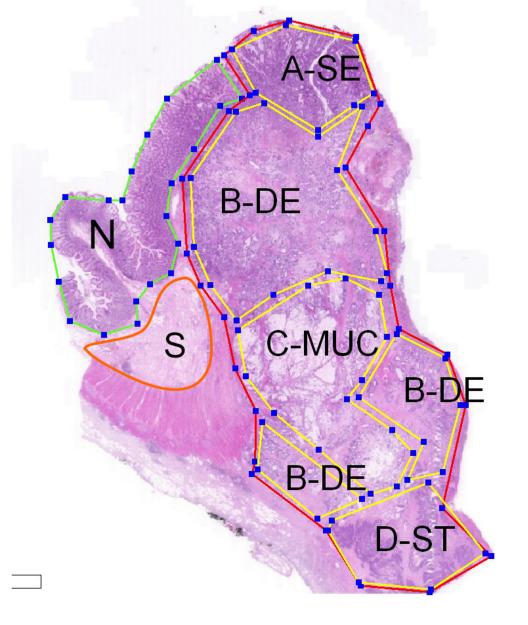
#### Use morphological regions to anchor the profiling and study the CMS mixture

- Study intra-tumor heterogeneity from CMS perspective
- Improve stability of the gene signatures
- Trade-off between whole-tumor and TME profiling

## METHOD

#### Morphology-guided transcriptomics

- Use whole tumor profile for baseline
- CMScaller for classifying resulting profiles
- GSEA analysis





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## **CONSENSUS MOLECULAR SUBTYPES OF MORPHOLOGICAL REGIONS ON COLORECTAL TUMORS**

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## RESULTS

- 100 different cases led to 152 regional RNA profiles + 100 whole-tumor
- No subtype was assigned to 22% (whole-tumor) and 23% (regional) profiles, respectively

plex tubular	СТ	34	Desmoplastic	DE	14
inous	MU	17	Normal	NR	17
lary	PP	6	Polyp	ΡΥ	2
ated	SE	48	Stroma	ST	8
trabecular	ТВ	4			

#### Whole-tumor CMS classification vs regional classification:

- CMS1,2,3 classification had good concordance between whole-tumor and at least one regional profile
- CMS4 less stable: for whole tumor CMS4, 44% of the regions were CMS4, and 21% CMS2

- (32%);

MTORC1 Signaling Protein Secretion Interferon Alpha Response Angiogenesis Extracellular Matrix mCRC E2F Targets G2M Checkpoint Crypt:Proliferation Fatty acids:react Liver TGFB down CRC MSS-MSI up WNT repressed Crypt:Late TA Gastro-Intestina Stromal estimate HNF1A up

Epithelial Mesenchymal Transition

dn 10 0 10 up -log<sub>10</sub>(p)

## CONCLUSIONS

- morphological regions may have a different molecular subtype than the whole tumor gene expression classifiers are sensitive to tumor sampling protocol need for clear specification of region(s) used for RNA profiling multi-region sampling may lead to CMS refinement CMS4 EMT characteristics are most probably due to the desmoplastic reaction.

• Complex tubular (CT): mostly labeled as CMS1 (41%) and CMS2

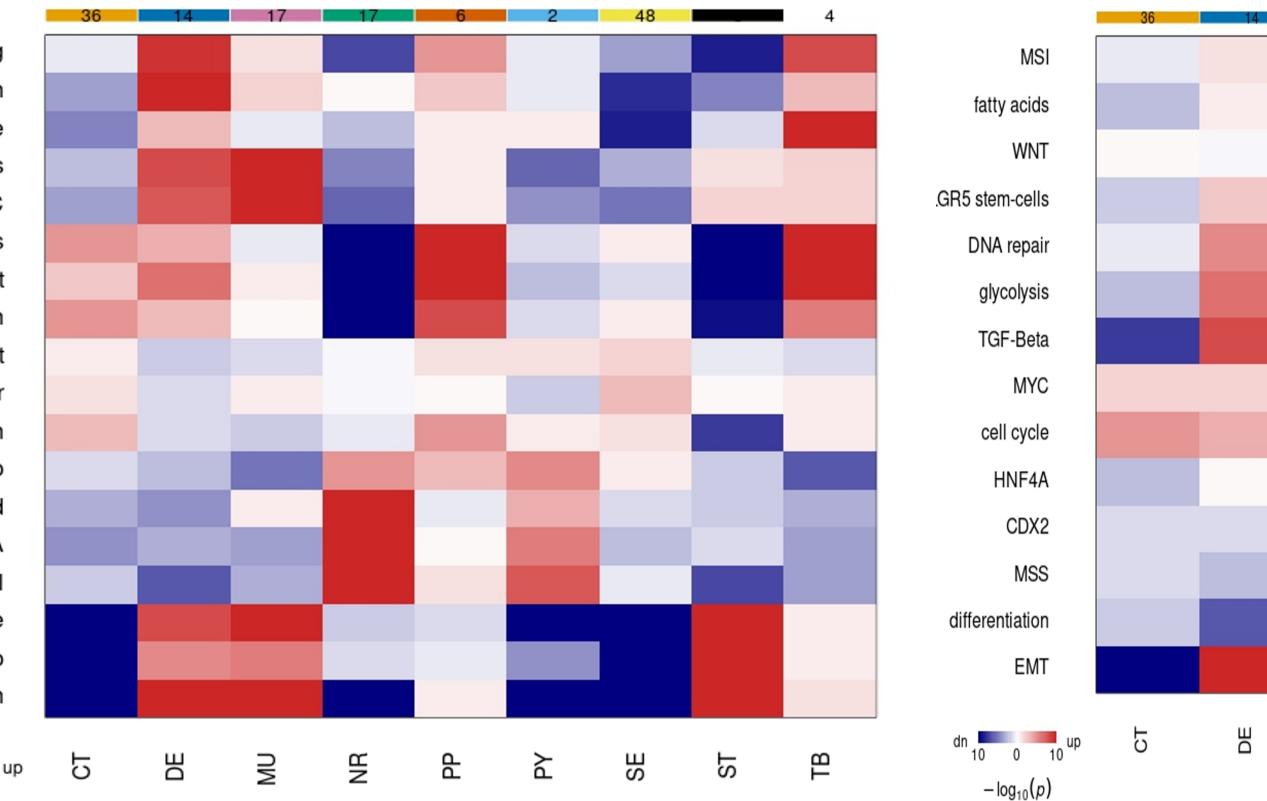
Desmoplastic (DE): CMS4 (54%) and CMS1 (31%)

• The subtype of DE, when in combination with any other region types, determined the whole tumor subtype

• Serrated (SE): mostly labeled as CMS2 (42%) and CMS3 (32%);

All tumor-adjacent stroma (ST) was labeled as CMS4.

• Mucinous (MU): CMS4 (60%).



## REFERENCES

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3. Budinska et al, Gene expression patterns unveil a new level of molecular heterogeneity in colorectal cancer. J. Pathol. 231(1), 2013

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Decion	<b>F</b>
Region	Enri
СТ	MYC
SE	DNA
DE	EMT
MU	EMT

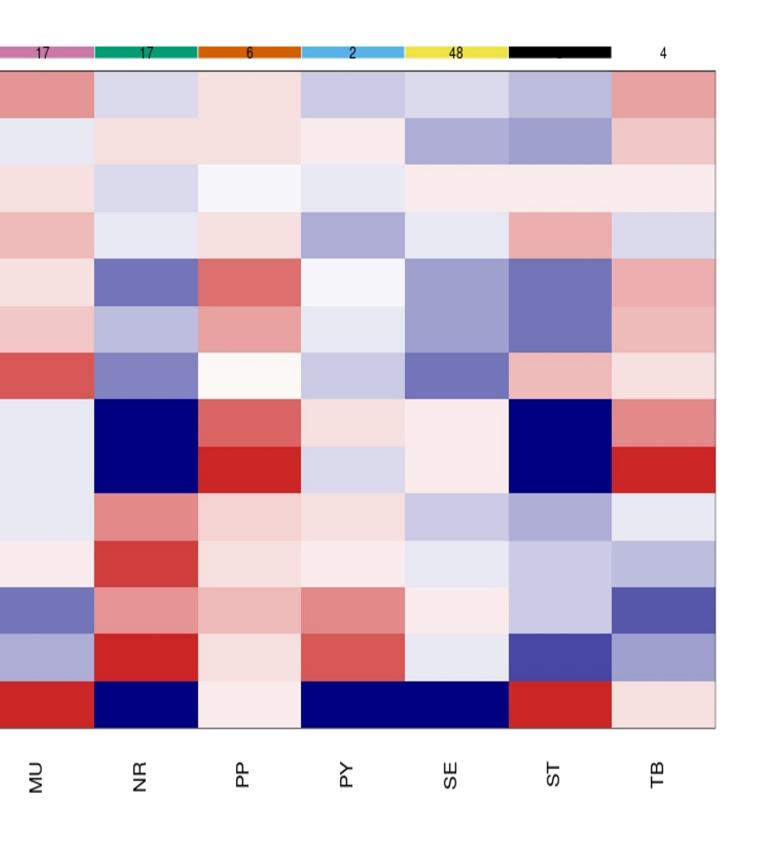
GSEA analysis – hallmark signatures



# MUNI RECETOX SCI

ichment vs rest

- C-targets, DNA repair, MTORC1, unfolded protein response
- A replication, regulation of apoptosis, MYC targets, Wnt/B-catenin
- , TGFb, unfolded protein response, apoptosis
- , TGFb, inflammatory response, KRAS signaling



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