Abundance scores, truncated above at 42

SegmentName
- PanCK−pos
- Geometric Segment
- PanCK−neg

ScanName
- AGBT CTA2.0
- CTA 2.0
- AGBT CTA 2.0

Abundance scores

SegmentName
- PanCK−pos
- Geometric Segment
- PanCK−neg

ScanName
- AGBT CTA2.0
- CTA 2.0
- AGBT CTA 2.0

Cell
- fibroblasts
- endothelial.cells
- Treg
- neutrophils
- monocytes.NC.I
- monocytes.C
- mDCs
- pDCs
- NK
- T.CD8.memory
- T.CD8.naive
- T.CD4.memory
- T.CD4.naive
- plasma
- B.memory
- B.naive
- mast
- macrophages

ScanName
SegmentName

Proportion of fitted cells

ScanName
SegmentName

SegmentName
- PanCK−pos
- Geometric Segment
- PanCK−neg

ScanName
- AGBT CTA2.0
- CTA 2.0
- AGBT CTA 2.0

Cell
- fibroblasts
- endothelial.cells
- Treg
- neutrophils
- monocytes.NC.I
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- mDCs
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- NK
- T.CD8.memory
- T.CD8.naive
- T.CD4.memory
- T.CD4.naive
- plasma
- B.memory
- B.naive
- mast
- macrophages