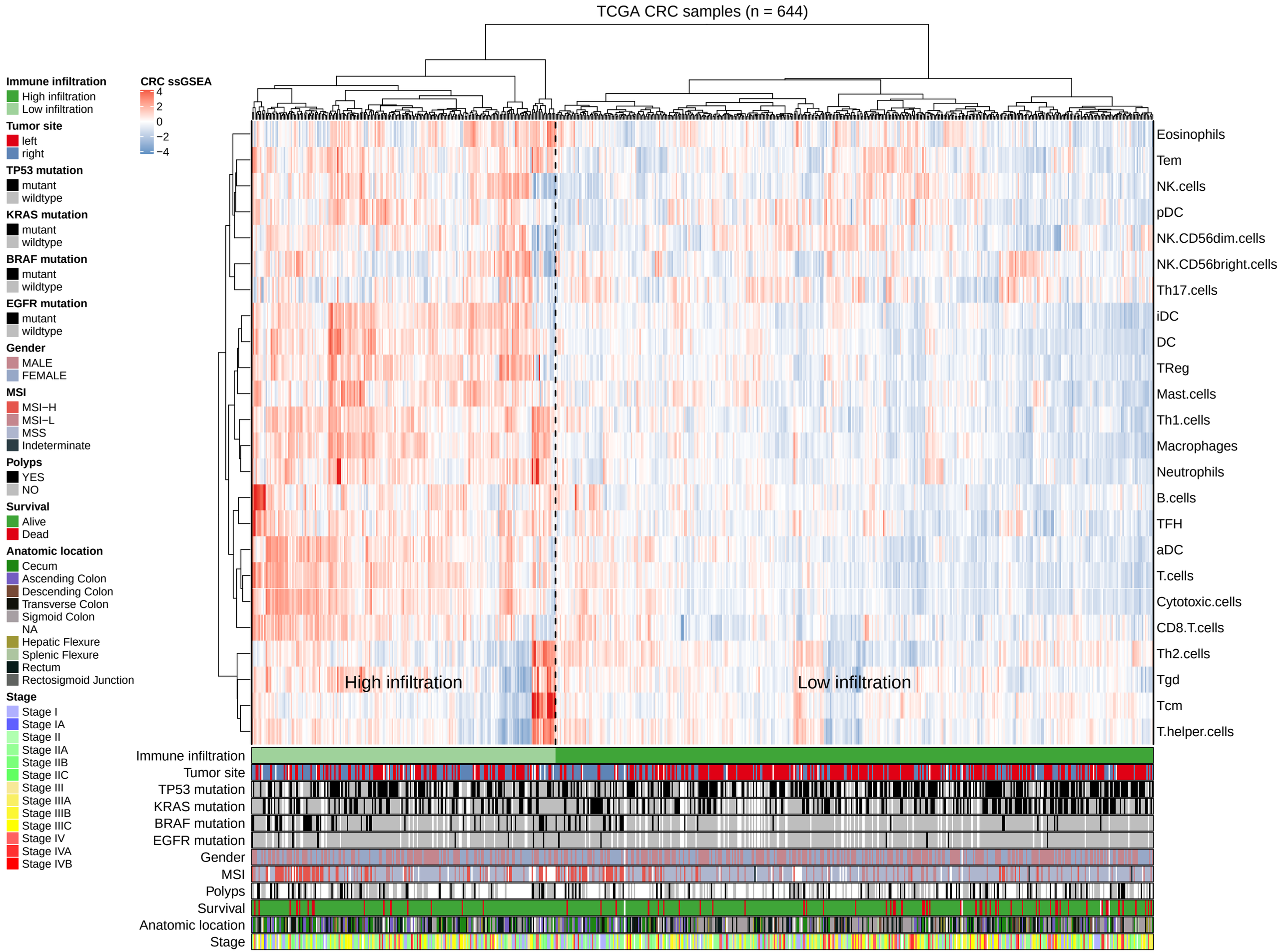
TCGA CRC samples (n = 644)
Immune infiltration
High infiltration
Low infiltration
Tumor site
left
right
TP53 mutation
mutant
wildtype
KRAS mutation
mutant
wildtype
BRAF mutation
mutant
wildtype
EGFR mutation
mutant
wildtype
Gender
MALE
FEMALE
MSI
MSI−H
MSI−L
MSS
Indeterminate
Polyps
YES
NO
Survival
Alive
Dead
Anatomic location
Cecum
Ascending Colon
Descending Colon
Transverse Colon
Sigmoid Colon
NA
Hepatic Flexure
Splenic Flexure
Rectum
Rectosigmoid Junction
Stage
Stage I
Stage IA
Stage II
Stage IIA
Stage IIB
Stage IIC
Stage III
Stage IIIA
Stage IIIB
Stage IIIC
Stage IV
Stage IVA
Stage IVB
CRC ssGSEA
4
2
0
−2
−4
Eosinophils
Tem
NK.cells
pDC
NK.CD56dim.cells
NK.CD56bright.cells
Th17.cells
iDC
DC
TReg
Mast.cells
Th1.cells
Macrophages
Neutrophils
B.cells
TFH
aDC
T.cells
Cytotoxic.cells
CD8.T.cells
Th2.cells
Tgd
Tcm
T.helper.cells
High infiltration
Low infiltration
Immune infiltration
Tumor site
TP53 mutation
KRAS mutation
BRAF mutation
EGFR mutation
Gender
MSI
Polyps
Survival
Anatomic location
Stage