a
Myeloid.cells.traffic
Myeloid.cells.traffic
intermediate
subTME−IC1
subTME−IS
subTME−MRM
subTME−PSE
BC
ESCA
b
Epithelial.cells
Epithelial.cells
subTME−IS+MRM
subTME−MRM
subTME−PSE
ESCA
PDAC
c
R = 0.4, p = 4.4e-08
R = 0.68, p < 2.2e-16
R = 0.65, p < 2.2e-16
R = 0.39, p = 1.7e-07
subTME-IS
Angiogenesis
Matrix remodeling
EMT
Tumor proliferation
ESCA
d
R = 0.31, p = 1.9e-05
R = 0.51, p < 2.2e-16
R = 0.41, p = 1.1e-08
R = 0.29, p = 6.7e-05
subTME-IS
Angiogenesis
Matrix remodeling
EMT
Tumor proliferation
PDAC
e
R = − 0.06, p = 0.035
R = 0.23, p = 9.3e-16
R = 0.24, p < 2.2e-16
R = 0.0051, p = 0.86
subTME-IS
Angiogenesis
Matrix remodeling
EMT
Tumor proliferation
BC