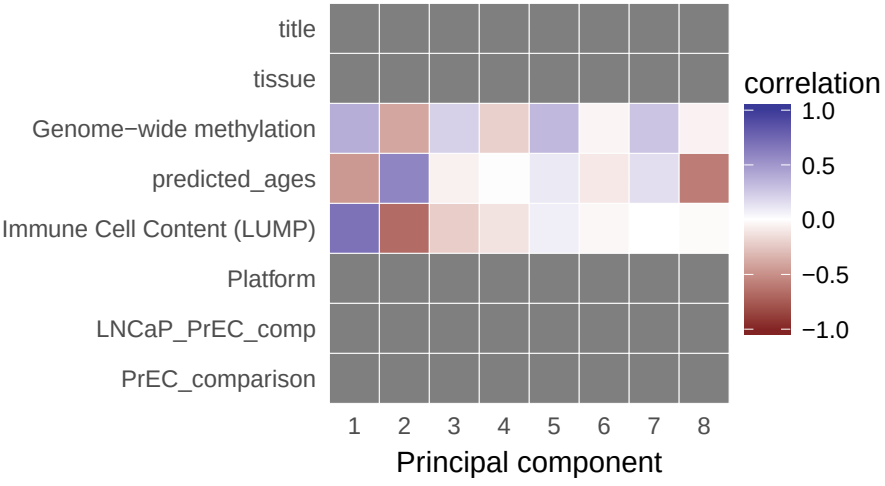
title
tissue
Genome−wide methylation
predicted_ages
Immune Cell Content (LUMP)
Platform
LNCaP_PrEC_comp
PrEC_comparison
1
2
3
4
5
6
7
8
Principal component
correlation
1.0
0.5
0.0
−0.5
−1.0