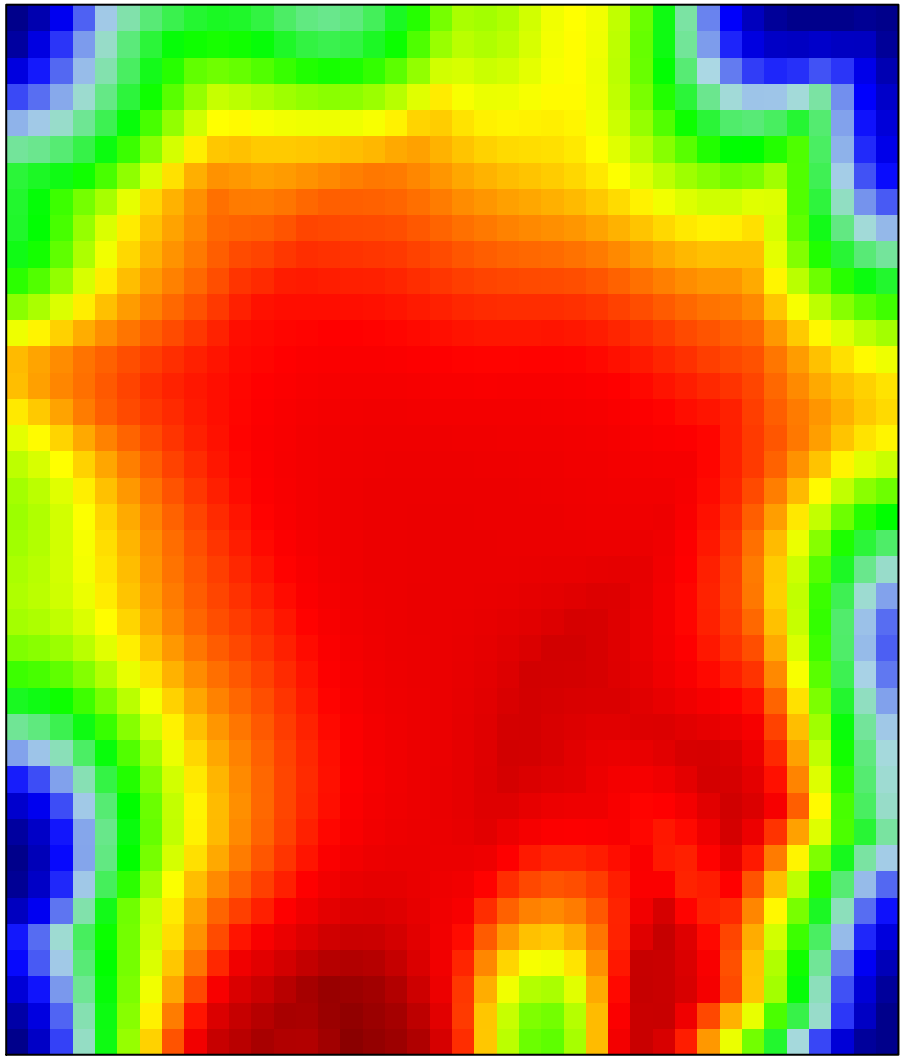
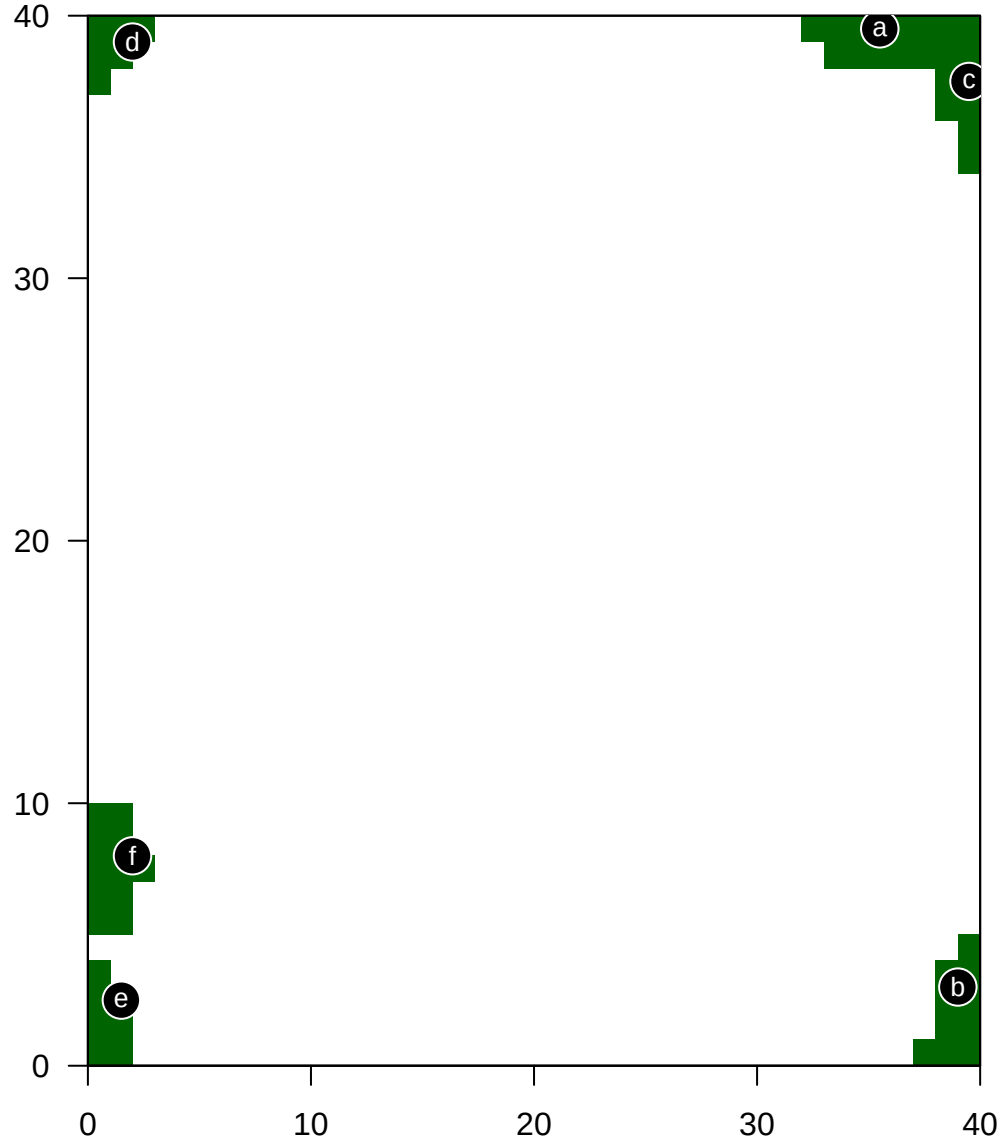
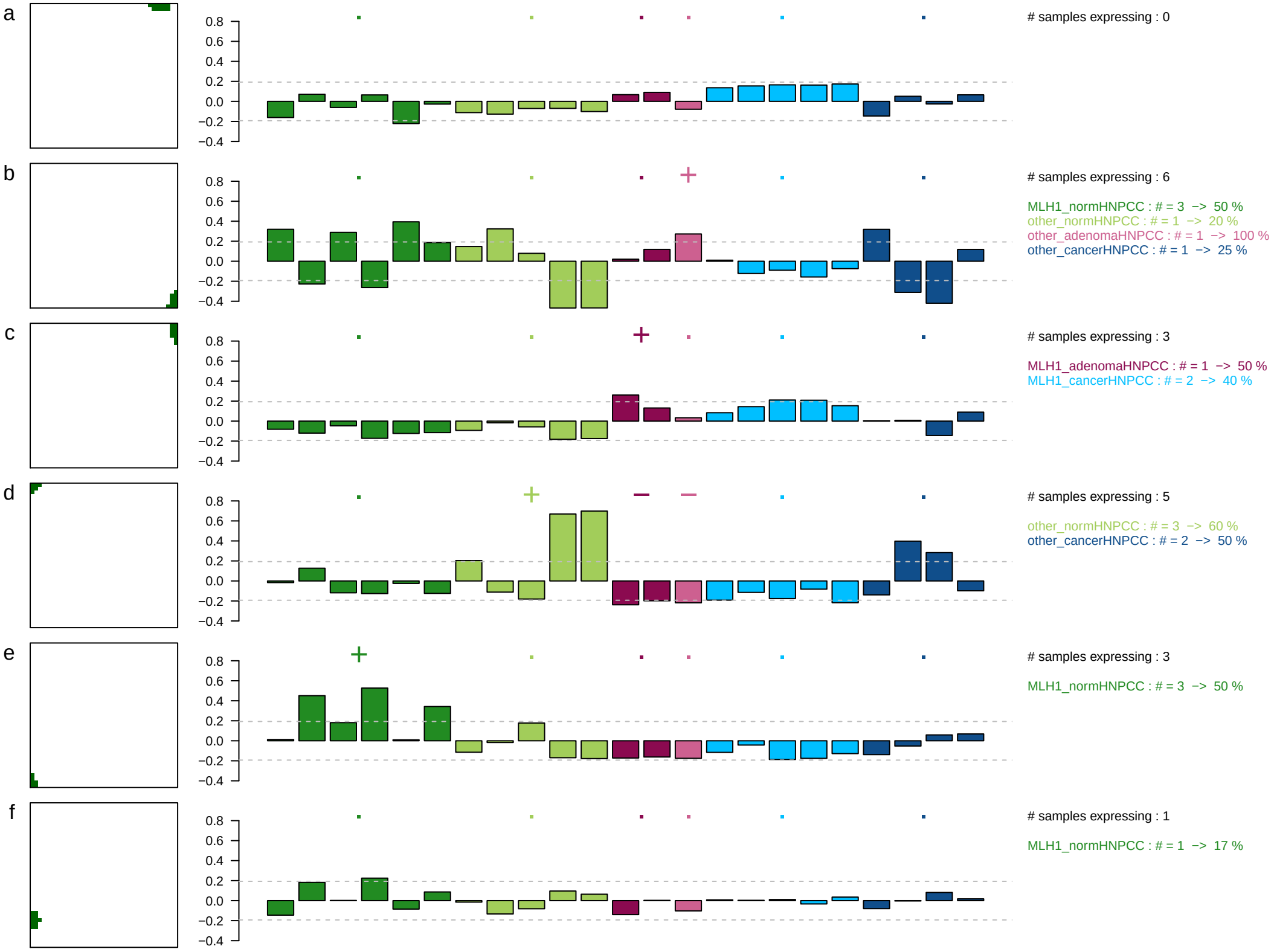
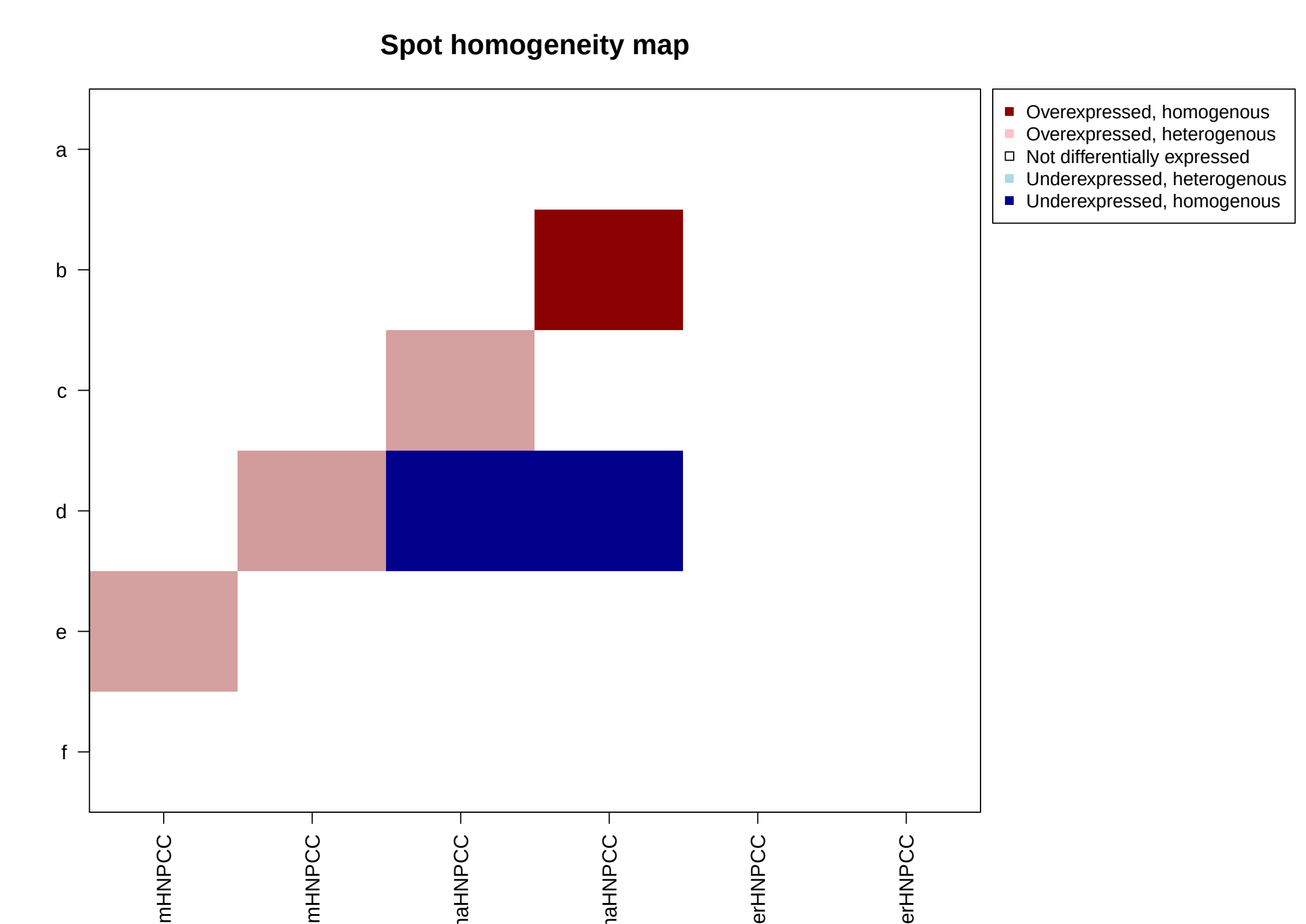
Overview map, Underexpression Spots

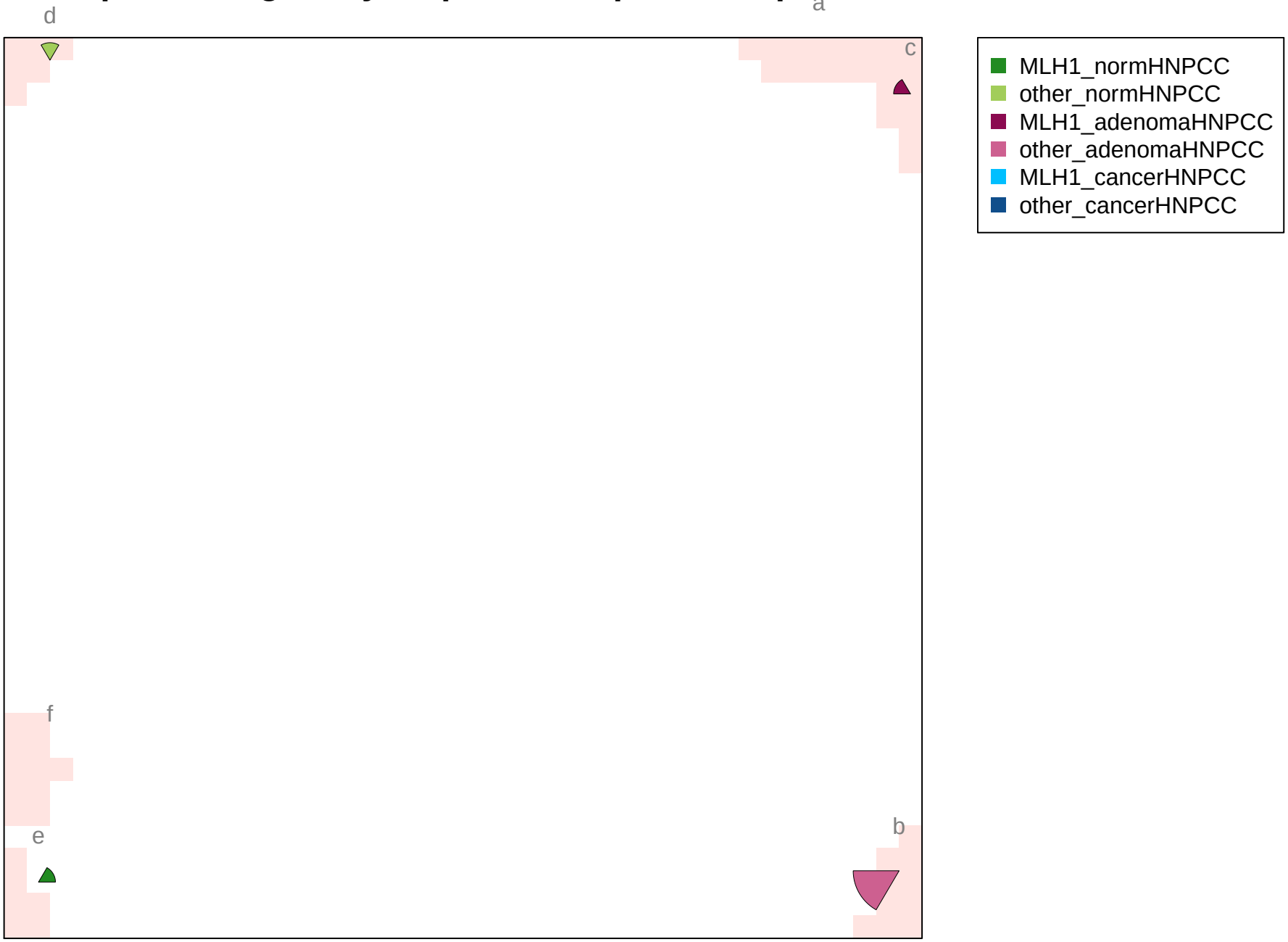


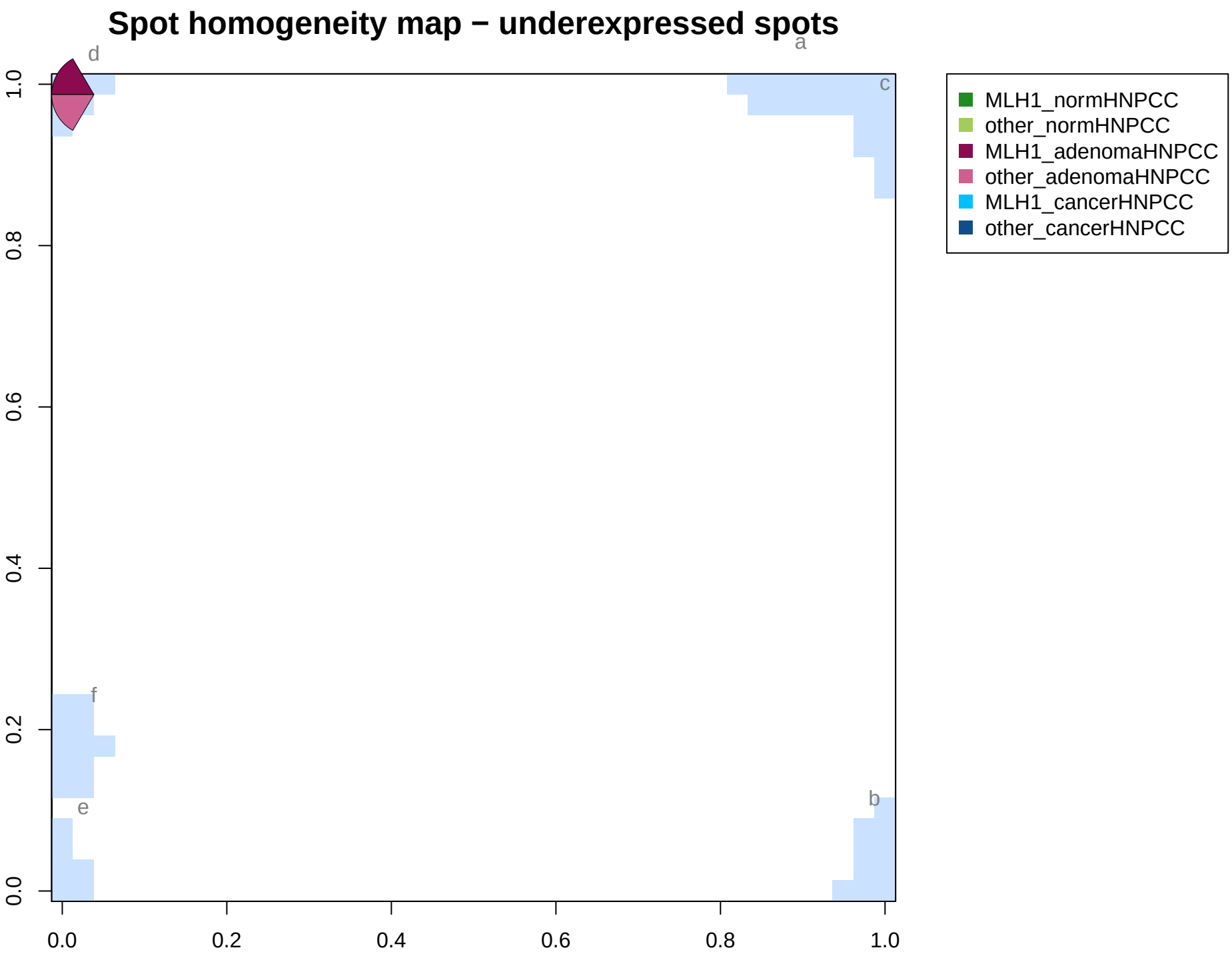


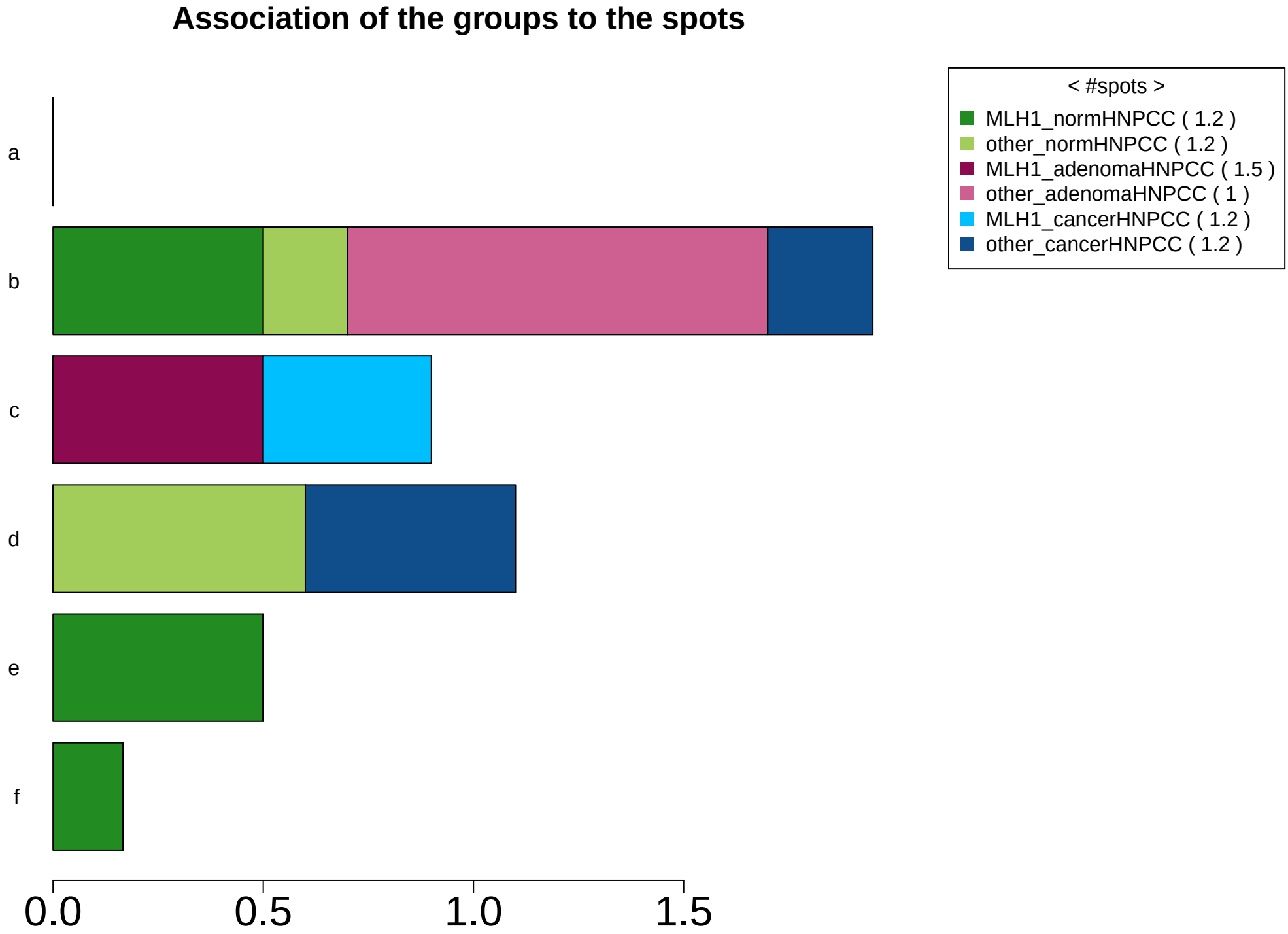
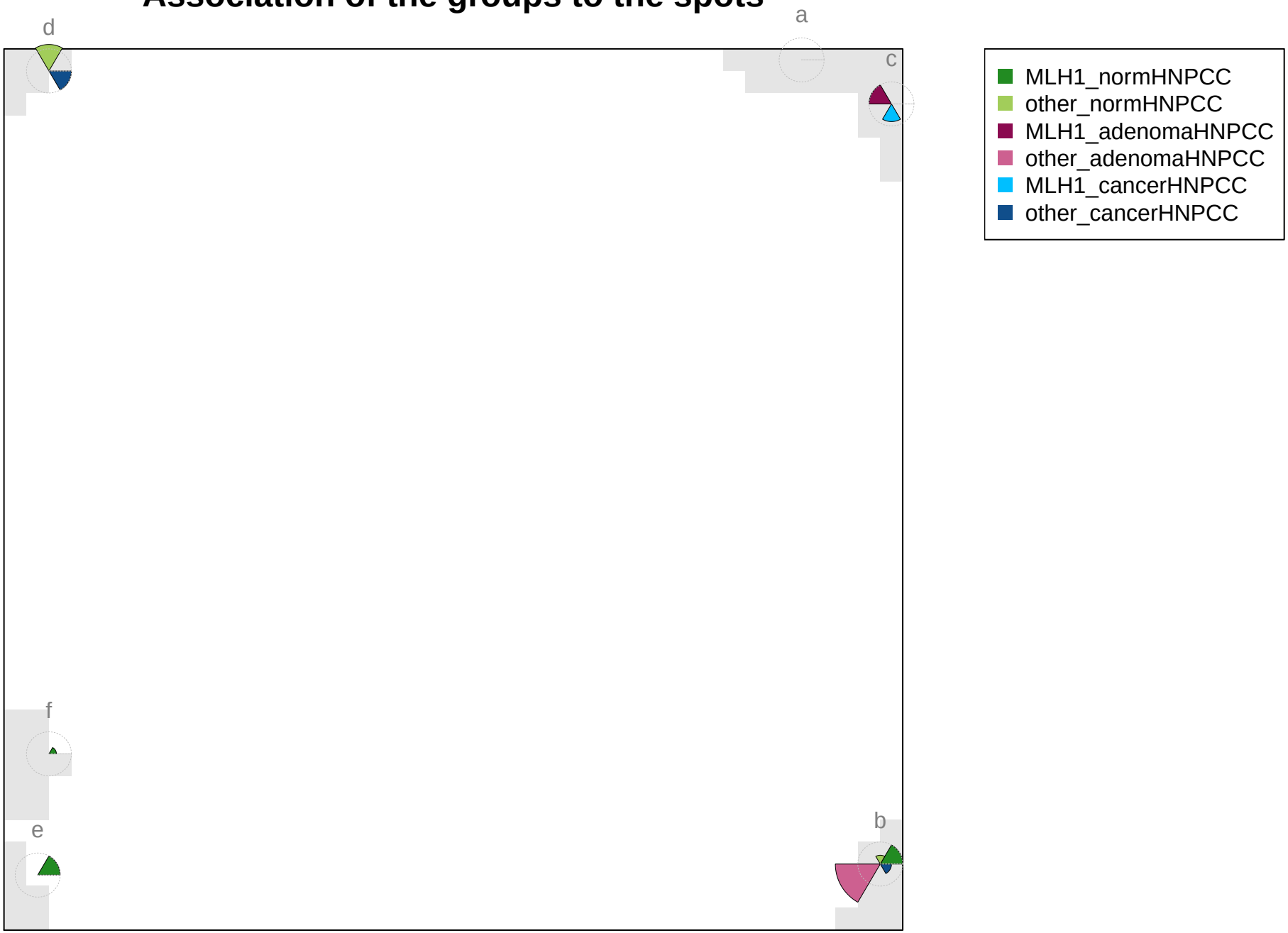

a
b
c
d
e
f
0.8
0.6
0.4
0.2
0.0
−0.2
−0.4
samples expressing : 0
samples expressing : 6
MLH1_normHNPCC : # = 3 −> 50 %
other_normHNPCC : # = 1 −> 20 %
other_adenomaHNPCC : # = 1 −> 100 %
other_cancerHNPCC : # = 1 −> 25 %
samples expressing : 3
MLH1_adenomaHNPCC : # = 1 −> 50 %
MLH1_cancerHNPCC : # = 2 −> 40 %
samples expressing : 5
other_normHNPCC : # = 3 −> 60 %
other_cancerHNPCC : # = 2 −> 50 %
samples expressing : 3
MLH1_normHNPCC : # = 3 −> 50 %
samples expressing : 1
MLH1_normHNPCC : # = 1 −> 17 %


Spot homogeneity map
Overexpressed, homogenous
Overexpressed, heterogenous
Not differentially expressed
Underexpressed, heterogenous
Underexpressed, homogenous
a
b
c
d
e
f
mHNPCC
mHNPCC
naHNPCC
naHNPCC
erHNPCC
erHNPCC

Spot homogeneity map − overexpressed spots




Spot homogeneity map − underexpressed spots
a
b
c
d
e
f
0.0
0.2
0.4
0.6
0.8
1.0
MLH1_normHNPCC
other_normHNPCC
MLH1_adenomaHNPCC
other_adenomaHNPCC
MLH1_cancerHNPCC
other_cancerHNPCC


Association of the groups to the spots
< #spots >
MLH1_normHNPCC (1.2)
other_normHNPCC (1.2)
MLH1_adenomaHNPCC (1.5)
other_adenomaHNPCC (1)
MLH1_cancerHNPCC (1.2)
other_cancerHNPCC (1.2)
a
b
c
d
e
f
0.0
0.5
1.0
1.5

Association of the groups to the spots