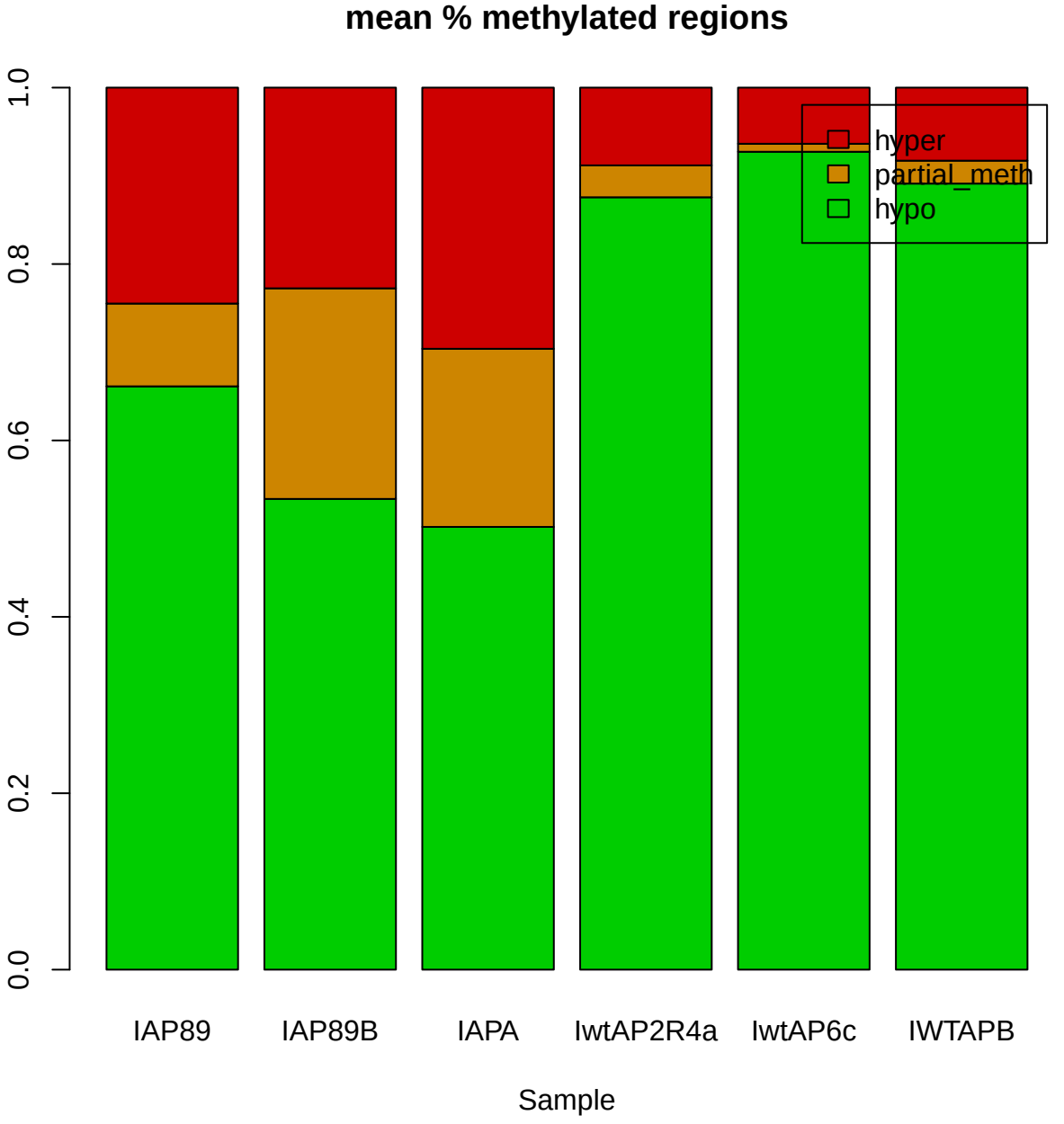
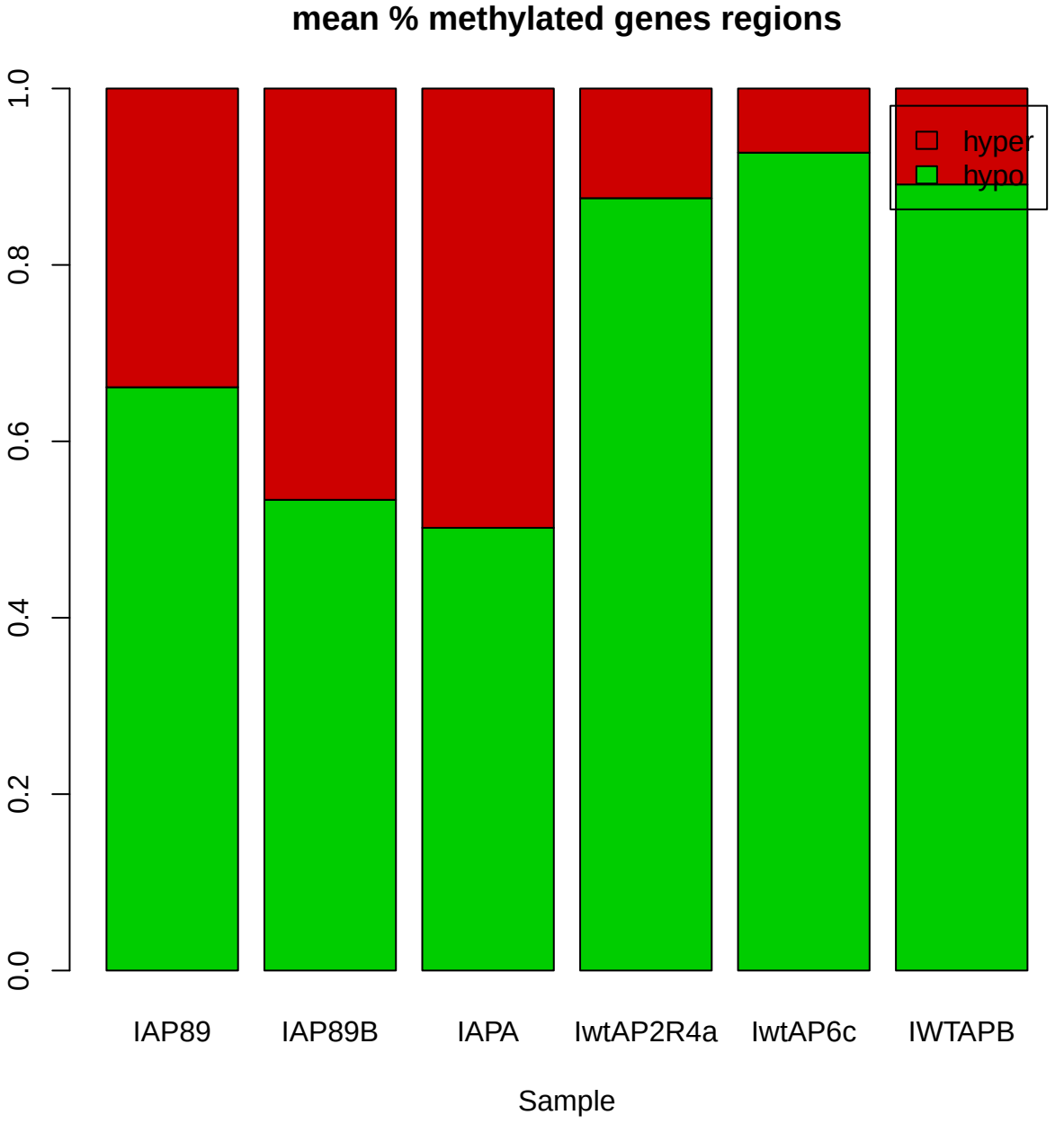
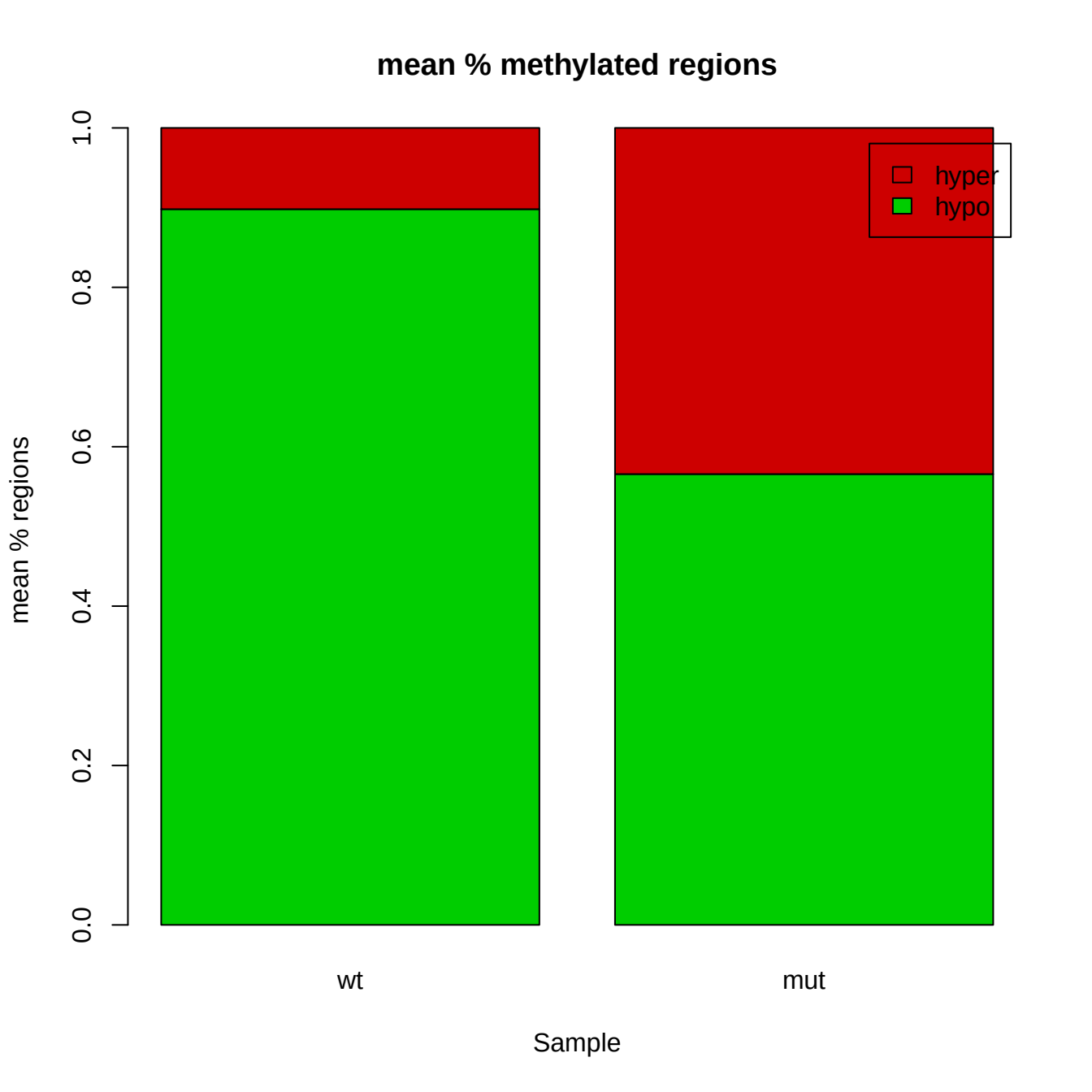

mean % methylated regions
1.0
0.8
0.6
0.4
0.2
0.0
hyper
partial_meth
hypo
IAP89
IAP89B
IAPA
IwtAP2R4a
IwtAP6c
IWTAPB
Sample


mean % methylated genes regions
1.0
0.8
0.6
0.4
0.2
0.0
hyper
hypo
IAP89
IAP89B
IAPA
IwtAP2R4a
IwtAP6c
IWTAPB
Sample


mean % methylated regions
mean % regions
0.0
0.2
0.4
0.6
0.8
1.0
hyper
hypo
wt
mut
Sample