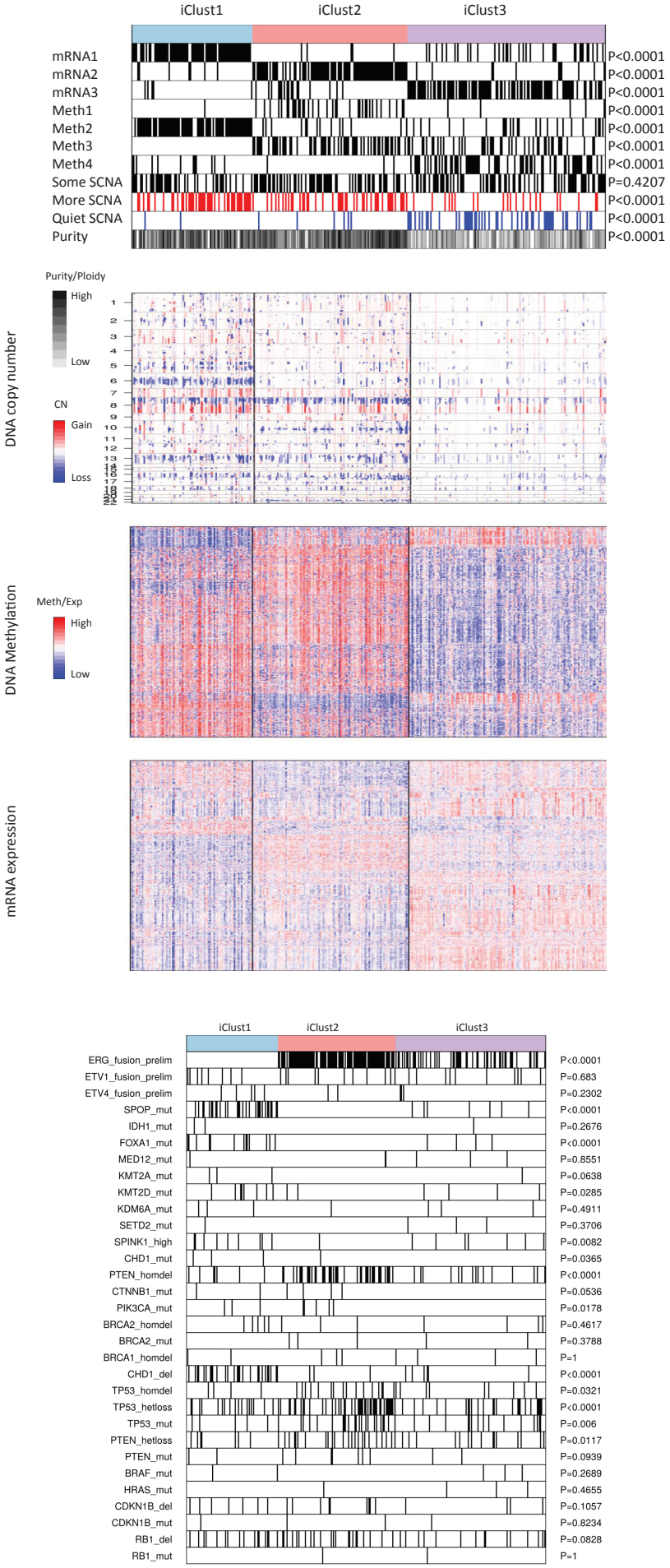
iClust1
iClust2
iClust3
mRNA1 P<0.0001
mRNA2 P<0.0001
mRNA3 P<0.0001
Meth1 P<0.0001
Meth2 P<0.0001
Meth3 P<0.0001
Meth4 P<0.0001
Some SCNA P=0.4207
More SCNA P<0.0001
Quiet SCNA P<0.0001
Purity P<0.0001
Purity/Ploidy
High
Low
CN
Gain
Loss
DNA copy number
Meth/Exp
High
Low
DNA Methylation
mRNA expression
iClust1
iClust2
iClust3
ERG_fusion_prelim P<0.0001
ETV1_fusion_prelim P=0.683
ETV4_fusion_prelim P=0.2302
SPOP_mut P<0.0001
IDH1_mut P=0.2676
FOXA1_mut P<0.0001
MED12_mut P=0.8551
KMT2A_mut P=0.0638
KMT2D_mut P=0.0285
KDM6A_mut P=0.4911
SETD2_mut P=0.3706
SPINK1_high P=0.0082
CHD1_mut P=0.0365
PTEN_homdel P<0.0001
CTNNB1_mut P=0.0536
PIK3CA_mut P=0.0178
BRCA2_homdel P=0.4617
BRCA2_mut P=0.3788
BRCA1_homdel P=1
CHD1_del P<0.0001
TP53_homdel P=0.0321
TP53_hetloss P<0.0001
TP53_mut P=0.006
PTEN_hetloss P=0.0117
PTEN_mut P=0.0939
BRAF_mut P=0.2689
HRAS_mut P=0.4655
CDKN1B_del P=0.1057
CDKN1B_mut P=0.8234
RB1_del P=0.0828
RB1_mut P=1