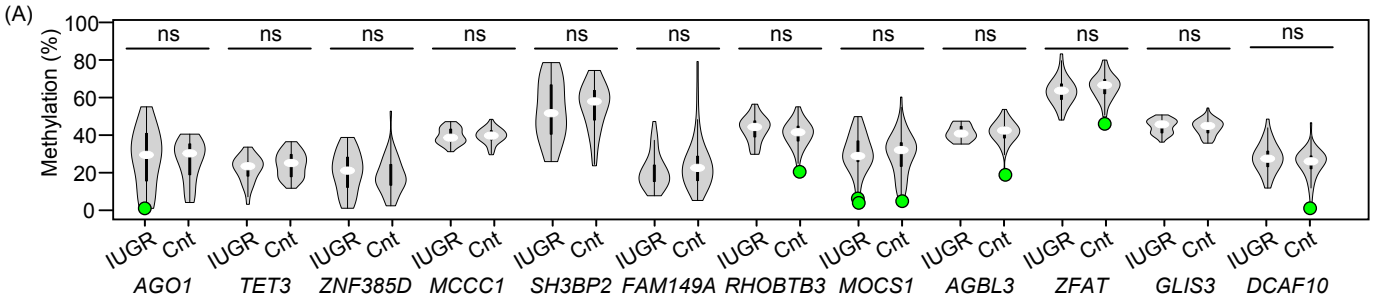
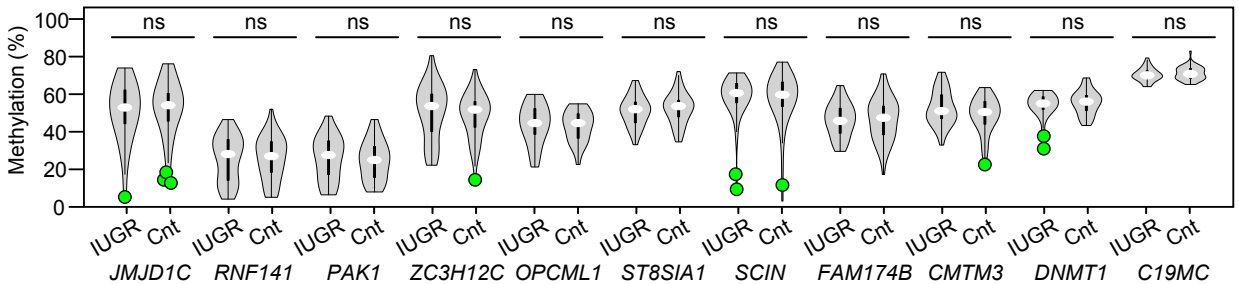
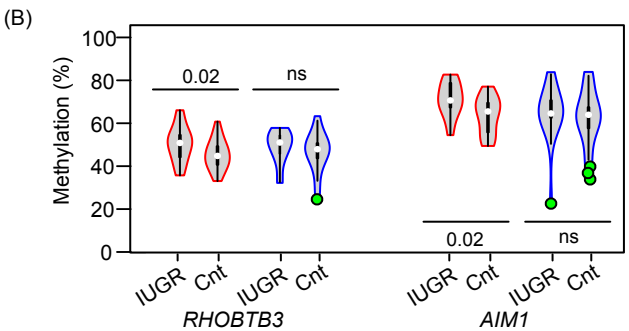
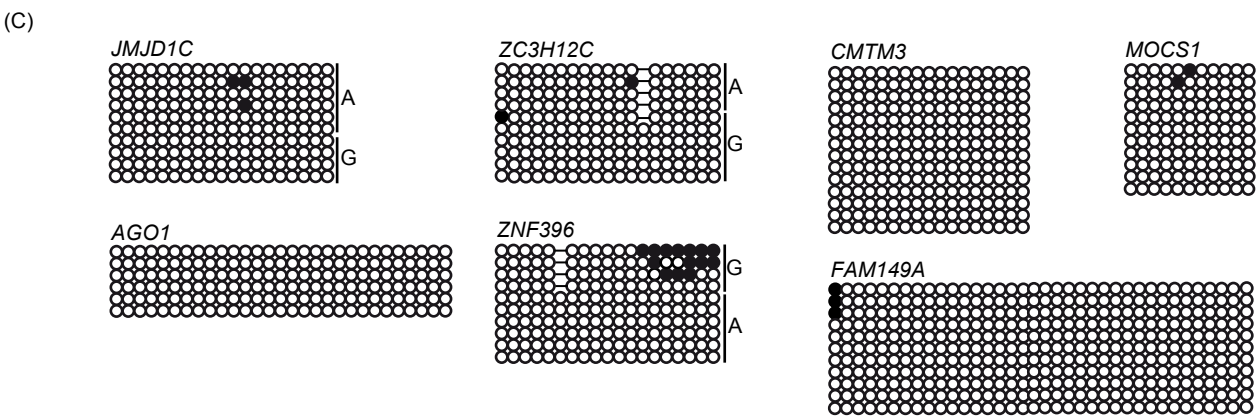

(A)
Methylation (%)
100
80
60
40
20
0
ns
IUGR
Cnt
AGO1
TET3
ZNF385D
MCCC1
SH3BP2
FAM149A
RHOBTB3
MOCS1
AGBL3
ZFAT
GLIS3
DCAF10


Methylation (%)
100
80
60
40
20
0
ns
IUGR
Cnt
JMJD1C
RNF141
PAK1
ZC3H12C
OPCML1
ST8SIA1
SCIN
FAM174B
CMTM3
DNMT1
C19MC


(B)
Methylation (%)
100
80
60
40
20
0
0.02
ns
IUGR
Cnt
RHOBTB3
AIM1


(C)
JMJD1C
A
G
ZC3H12C
A
G
CMTM3
MOCS1
AGO1
ZNF396
G
A
FAM149A