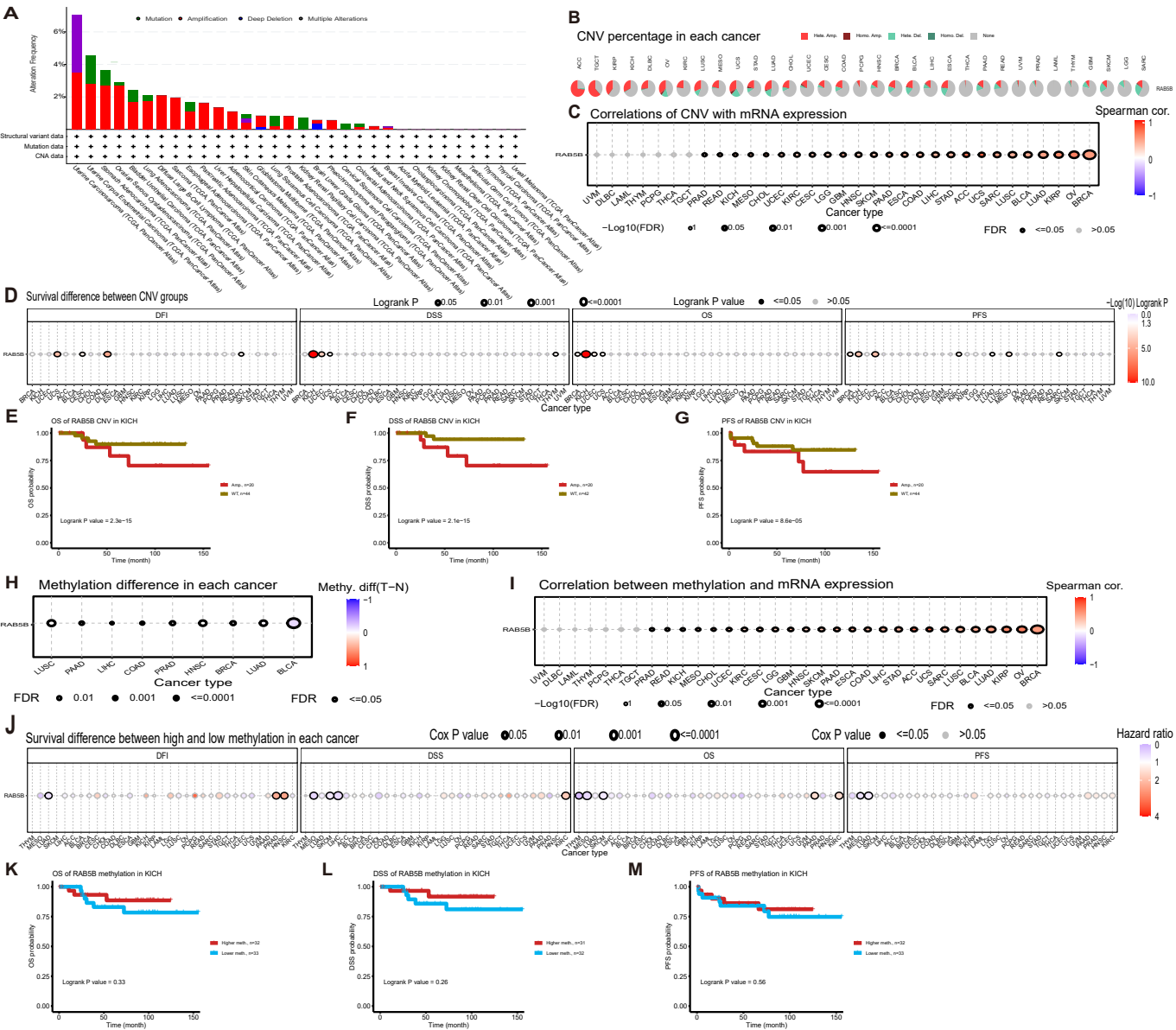
A
Mutation
Amplification
Deep Deletion
Multiple Alterations
Alteration Frequency
Structural variant data
Mutation data
CNA data
B
CNV percentage in each cancer
Hete. Amp.
Homo. Amp.
Hete. Del.
Homo. Del.
None
C
Correlations of CNV with mRNA expression
Spearman cor.
Cancer type
−Log10(FDR)
FDR
D
Survival difference between CNV groups
Logrank P
Logrank P value
DFI
DSS
OS
PFS
−Log(10) Logrank P
Cancer type
E
OS of RAB5B CNV in KICH
OS probability
Logrank P value = 2.3e−15
Time (month)
F
DSS of RAB5B CNV in KICH
DSS probability
Logrank P value = 2.1e−15
G
PFS of RAB5B CNV in KICH
PFS probability
Logrank P value = 8.6e−05
H
Methylation difference in each cancer
Methy. diff(T−N)
Cancer type
I
Correlation between methylation and mRNA expression
Spearman cor.
Cancer type
−Log10(FDR)
FDR
J
Survival difference between high and low methylation in each cancer
Cox P value
Hazard ratio
DFI
DSS
OS
PFS
Cancer type
K
OS of RAB5B methylation in KICH
OS probability
Logrank P value = 0.33
Time (month)
L
DSS of RAB5B methylation in KICH
DSS probability
Logrank P value = 0.26
M
PFS of RAB5B methylation in KICH
PFS probability
Logrank P value = 0.56