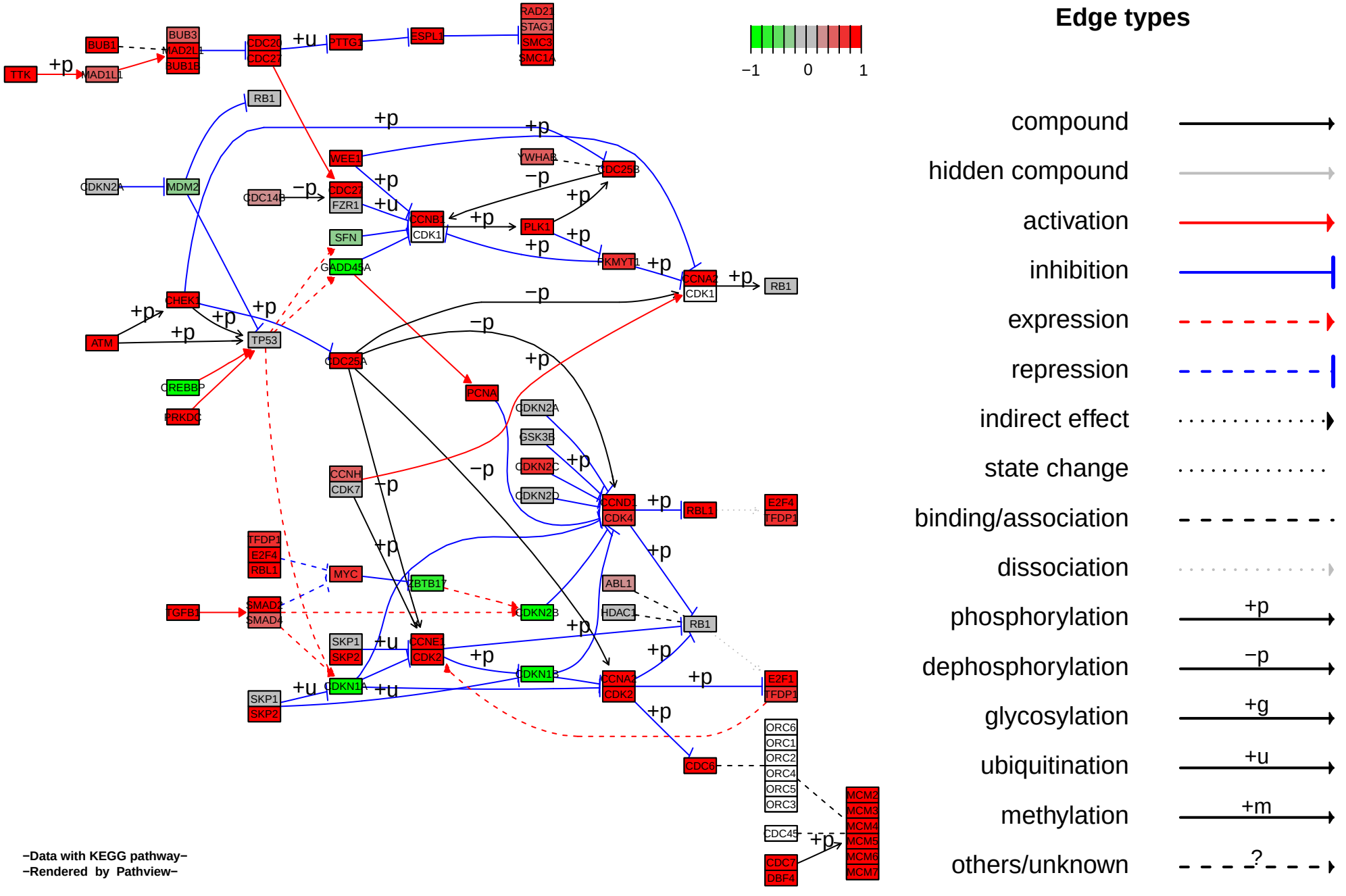
Edge types
−1
0
1
compound
hidden compound
activation
inhibition
expression
repression
indirect effect
state change
binding/association
dissociation
phosphorylation
dephosphorylation
glycosylation
ubiquitination
methylation
others/unknown
+p
−p
+g
+u
+m
?
TTK
BUB1
MAD1L1
BUB3
MAD2L1
BUB1B
CDC20
CDC27
PTTG1
ESPL1
RAD21
STAG1
SMC3
SMC1A
RB1
CDKN2A
MDM2
CDC14B
WEE1
YWHAB
CDC25B
CDC27
FZR1
CCNB1
CDK1
PLK1
SFN
GADD45A
PKMYT1
CCNA2
CDK1
CHEK1
ATM
TP53
CDC25A
CREBBP
PRKDC
PCNA
CDKN2A
GSK3B
CDKN2C
CDKN2D
CCNH
CDK7
CCND1
CDK4
RBL1
E2F4
TFDP1
TFDP1
E2F4
RBL1
MYC
ZBTB17
ABL1
TGFB1
SMAD2
SMAD4
CDKN2B
HDAC1
RB1
SKP1
SKP2
CCNE1
CDK2
CDKN1B
CCNA2
CDK2
E2F1
TFDP1
SKP1
SKP2
CDKN1A
CDC6
ORC6
ORC1
ORC2
ORC4
ORC5
ORC3
CDC45
CDC7
DBF4
MCM2
MCM3
MCM4
MCM5
MCM6
MCM7
−Data with KEGG pathway−
−Rendered by Pathview−