mutation type
missense mutation
nonsense mutation
frame_shift_ins
frame_shift_del
in_frame_ins
in_frame_del
splice_site
total
TCGA
ICGC
clinical classification
primary solid tumor
recurrent solid tumor
primary blood derived cancer - peripheral blood
recurrent blood derived cancer - bone marrow
additional - new primary
metastatic
additional metastatic
clinical significance
pathogenic
likely pathogenic
uncertain significance
likely benign
benign
somatic mutations
gene copy number
ABL1
ABCA13
ABCB1
ACAN
ACSM1
ACSM2A
ADAM29
ADAM7
ADAMTS12
ADAMTS20
ADAMTSL4
AFAP1L2
AGL
AHNAK
AHNAK2
AIP
AKAP13
AKAP9
ANK1
ANK2
ANKRD36A
ANKRD36C
ANO1
APC
APLNR
APOB
ARID1A
ARID2
ASCL1
ASPM
ASXL1
ATM
ATR
ATRX
AXIN1
AXIN2
B2M
BAP1
BCL9
BCL9L
BIRC6
BMPR1A
BRAF
BRCA1
BRCA2
BRD4
BRIP1
BTK
BUB1B
C10orf90
C2
C7
CACNA1D
CACNA1E
CACNA1S
CAPN1
CASR
CCDC168
CCSCTB
CDH10
CDH11
CDH17
CDH23
CDH7
CDH8
CDH9
CDKN2A
CELSR2
CENPE
CNTNAP1
CNTNAP2
CNTNAP5
COL18A1
COL1A1
COL2A1
COL3A1
COL4A4
COL5A1
CPS1
CR1
CR1L
CSMD1
CSMD2
CSMD3
CTNNB1
CUX2
CYP11B1
CYP2C8
CYP2D6
DCHS1
DCHS2
DDX11
DDX3X
DDX60
DDX60L
DGKB
DHX15
DHX34
DHX57
DHX9
DNAH1
DNAH10
DNAH11
DNAH14
DNAH17
DNAH2
DNAH5
DNAH7
DNAH8
DNAH9
DNAI1
DNAI2
DNAJC13
DNAJC6
DNAJC7
DNAJC8
DNAJC9
DNAL1
DNALI1
DNAH12
DNAH6
DNAH3
DNAH13
DNAH4
DNAH10L
DNAH5L
DNAH7L
DNAH8L
DNAH9L
DNAI3
DNAI4
DNAI7
DNAJA4
DNAJB13
DNAJB14
DNAJB4
DNAJB5
DNAJB6
DNAJB7
DNAJB8
DNAJB9
DNAJC1
DNAJC10
DNAJC11
DNAJC12
DNAJC14
DNAJC15
DNAJC16
DNAJC17
DNAJC18
DNAJC19
DNAJC2
DNAJC21
DNAJC22
DNAJC24
DNAJC25
DNAJC27
DNAJC28
DNAJC3
DNAJC30
DNAJC4
DNAJC5
DNAJC5B
DNAJC5G
DNAJC6
DNAJC7
DNAJC8
DNAJC9
DNAL4
DNAL1
DNAL4
DNAAF1
DNAAF2
DNAAF3
DNAAF4
DNAAF5
DNAAF6
DNAAF8
DNAAF9
DNAH1
DNAH10
DNAH11
DNAH12
DNAH14
DNAH17
DNAH2
DNAH3
DNAH4
DNAH5
DNAH6
DNAH7
DNAH8
DNAH9
DNAI1
DNAI2
DNAJC13
DNAJC6
DNAJC7
DNAJC8
DNAJC9