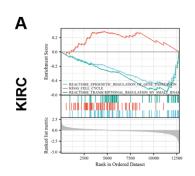
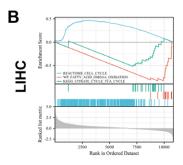
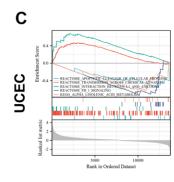
Figure S4. Gene Set Enrichment Analysis. The roles of CDKN2A in KIRC (A), LIHC (B), UCEC (C) by performing Gene Set Enrichment Analysis (GSEA).



ID	ES	NES	p.adjust	FDR
REACTOME_EPIGENET0.482		-1.656	0.067	0.056
REACTOME_TRANS	CRI0.560	-1.869	0.067	0.056
KEGG_CELL_CYCLE	0.287	1.342	0.182	0.151



ID	ES	NES	p.adjust	FDR	
REACTOME_CELL_C	YCLE 0.472	1.783	0.074	0.061	
WP_FATTY_ACID_O	ME0.831	-2.619	0.089	0.073	
KEGG_CITRATE_CYC	CL0.519	-1.968	0.089	0.074	



ID	ES	NES	p.adjust	FDR
REACTOME_TRANSMIS 0.469		1.844	0.119	0.112
REACTOME	INTERACT 0.679	1.964	0.119	0.112
REACTOME	PD_1_SIG0.710	-2.243	0.178	0.167
REACTOME	_APOPTOTI0.562	-1.937	0.205	0.193
KEGG_ALPH	A_LINOLE0.625	-1.956	0.225	0.211