



B

Sample	KEAP1 Mutation	CDS Mutation	Tissue	Fold WDR23
TCGA-46-6025-01	p.Q75*	c.223C>T	Lung	-2.51
TCGA-91-6848-01	p.D236N	c.706G>A	Lung	-2.34
TCGA-35-5375-01	p.E593fs*2	c.1777delG	Lung	CNV 4.21
TCGA-49-4488-01	p.Q284L	c.851A>T	Lung	2.06
TCGA-4B-A93V-01	p.L199F	c.597G>T	Lung	4.48
TCGA-55-A4DF-01	p.?	c.1708+1G>T	Lung	3.01
TCGA-73-4670-01	p.V123L	c.367G>T	Lung	2.08
TCGA-73-4677-01	p.K287*	c.859A>T	Lung	3.63
TCGA-86-8358-01	p.G558E	c.1673G>A	Lung	3.58
TCGA-86-A4D0-01	p.G480W	c.1438G>T	Lung	2.53
TCGA-43-6143-01	p.I506V	c.1516A>G	Lung	3.02
TCGA-44-7667-01	p.?	c.1709-1G>T	Lung	4.84

C

Sample	WDR23 Mutation	CDS Mutation	Tissue	Fold KEAP1
TCGA-78-7220-01	p.R145L	c.434G>T	Lung	2.07
TCGA-BR-8361-01	p.R341W	c.1021C>T	Stomach	2.43
TCGA-BR-8372-01	p.Q70R	c.209A>G	Stomach	2.15
TCGA-CG-5721-01	p.R265H	c.794G>A	Stomach	3.41
TCGA-BR-8361-01	p.R341W	c.1021C>T	Stomach	2.43

