

Supplementary materials

Comprehensive characterization of genes associated with the TP53 signal transduction pathway in various tumors

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Supplementary Tables

Supplementary Table 1

Histopathological characteristics of patients.

Supplementary Table 2

The number of tumor-specific non-synonymous somatic mutations found in 19 tumor types identified using whole exome sequencing (n = 907).

Supplementary Table 3

Association between patient characteristics and TP53-related mutations in many type of tumors

Supplementary Table 1 Histopathological characteristics of patients

Tumor types		Cases	Age	Tumor types		Cases	Age
		n	mean± SD			n	mean± SD
Stomach		116	71.1± 9.3	Breast		60	57.3±13.0
	Moderately differentiated tubular adenocarcinoma	41			Ductal carcinoma	41	
	Poorly differentiated adenocarcinoma	43			Lobular carcinoma	7	
	Well differentiated tubular adenocarcinoma	11			Metaplastic carcinoma	1	
	Signet-ring cell carcinoma	4			Metaplastic carcinoma and invasive ductal carcinoma	1	
	Mucinous adenocarcinoma	3			Metaplastic carcinoma and squamous cell carcinoma	2	
	Papillary adenocarcinoma	3			Mixed invasive cribriform carcinoma and ductal carcinoma	1	
	Neruoendocrine carcinoma	2			Mixed mucinous and invasive ductal carcinoma	1	
	Adenosquamous carcinoma	3			Mucinous carcinoma	1	
	Mixed adenocarcioma and neuroendocrine carcinoma	3			Noninvasive ductal carcinoma	3	
	Small cell neuroendocrine carcinoma	1			Spindle cell carcinoma	1	
	AFP-producing (hepatoid) adenocarcinoma	1			Micropapillary carcinoma	1	
	Endocrine carcinoma	1					
Lung		176	68.6±9.9	Liver		61	70.3± 8.7
	Adenocarcinoma	122			Hepatocellular carcinoma	53	
	Squamous cell carcinoma	33			Intrahepatic cholangiocarcinoma	3	
	Adenoid cystic carcinoma	2			Intrahepatic peripheral cholangiocarcinoma	1	
	Adenosquamous carcinoma	1			Combined hepatocellular and cholangiocarcinoma	3	
	Combined small cell carcinoma and adenocarcinoma	1			Cholangiocarcinoma	1	
	Diffuse malignant mesothelioma	1					
	Enteric adenocarcinoma	1		Pancreas		18	67.3±15.5
	Large cell neuroendocrine carcinoma (LCNEC)	7			Adenocarcinoma	12	
	Leiomyomatous hamartoma	1			Adenosquamous carcinoma	3	
	Schwannoma	2			Endocrine tumor	1	
	Mucinous AIS-like bronchiolar cell hyperplasia	1			Neuroendocrine tumor	1	
	Pleomorphic carcinoma	2			Solid-pseudopapillary neoplasm	1	
	Small cell carcinoma	1					
	Solitary fibrous tumor	1		Kidney		13	65.3±13.5
Colon		311	66.0±11.3		Clear cell renal cell carcinoma	6	
	Cecum	21			Chromophobe renal cell carcinoma	2	
	Well differentiated tubular adenocarcinoma	14			Renal cell clear cell carcinoma	2	
	Moderately differentiated tubular adenocarcinoma	6			Carcinoid tumor	1	
	Mucinous adenocarcinoma	1			Urothelial carcinoma	1	
	Ascending colon	47			Papillary renal cell carcinoma	1	
	Moderately differentiated tubular adenocarcinoma	19		Esophagus		18	66.5±8.9
	Well differentiated tubular adenocarcinoma	23			Esophageal cancer	8	
	Poorly differentiated adenocarcinoma	1			Moderately differentiated squamous cell carcinoma	1	
	Papillary adenocarcinoma	1			Squamous cell carcinoma	7	
	Mucinous adenocarcinoma	3			Esophagogastric junction cancer	10	
	Anus	1			Well differentiated tubular adenocarcinoma	1	
	Well differentiated tubular adenocarcinoma	1			Moderately differentiated adenocarcinoma	4	
					Pooly differentiated adenocarcnioma	3	

Descending colon	13		Papillary adenocarcinoma	1	
Moderately differentiated tubular adenocarcinoma	8		AFP-producing hepatoid adenocarcinoma	1	
Well differentiated tubular adenocarcinoma	5				
Sigmoid colon	55		Uterus	12	55.3±12.8
Moderately differentiated tubular adenocarcinoma	24		Uterine body	8	
Well differentiated tubular adenocarcinoma	25		Endometrioid adenocarcinoma	7	
Mucinous adenocarcinoma	4		Clear cell adenocarcinoma	1	
Poorly differentiated adenocarcinoma	1		Uterine cervix	4	
Papillary adenocarcinoma	1		Adenocarcinoma	1	
Transverse colon	22		Adenosquamous carcinoma	2	
Moderately differentiated tubular adenocarcinoma	12		Squamous cell carcinoma	1	
Mucinous adenocarcinoma	3				
Well differentiated tubular adenocarcinoma	6		Sarcoma	16	45.4±20.4
Poorly differentiated adenocarcinoma	1		Spindle cell sarcoma	2	
Duodenum	2		Osteosarcoma	2	
Adenocarcinoma	2		Synovial sarcoma	1	
Jejunum	1		Leiomyosarcoma	2	
Adenocarcinoma	1		Dedifferentiated liposarcoma	1	
Appendix	1		Dermatofibrosarcoma protuberans	2	
Well differentiated tubular adenocarcinoma	1		Myxofibrosarcoma	2	
Rectum	148		Rhabdomyosarcoma	1	
Moderately differentiated tubular adenocarcinoma	70		Round cell liposarcoma	1	
Well differentiated tubular adenocarcinoma	75		Chondrosarcoma	1	
Poorly differentiated adenocarcinoma	3		Carcinosarcoma	1	
Head and Neck	73	62.5±14.8	GIST (gastrointestinal stromal tumor)	9	70.4±11.0
Arytenoid fold	1				
Squamous cell carcinoma	1		Melanoma	5	61.6±16.9
External ear canal	1				
Squamous cell carcinoma	1		Thymus	6	63.0±9.0
Gingiva	6		Thymic carcinoma	1	
Squamous cell carcinoma	6		Thymoma	5	
Gum	2				
Squamous cell carcinoma	2		Ovary	4	57.5±14.6
Larynx	5		Clear cell adenocarcinoma	3	
Squamous cell carcinoma	5		Sex-cord stromal tumor	1	
Oropharynx	1				
Squamous cell carcinoma	1		Skin	3	52.0±14.8
Thyroid gland	4		Multiple neurofibroma	1	
Adenomatous goiter	1		Squamous cell carcinoma	1	
Papillary carcinoma	3		Eccrine poroma	1	
Buccal mucosa	1				

Squamous cell carcinoma	1
Cheek	3
Squamous cell carcinoma	3
Hard palate	1
Acinic cell carcinoma	1
Hypopharynx	6
Squamous cell carcinoma	6
Lingual root	1
Squamous cell carcinoma	1
Lip	1
Squamous cell carcinoma	1
Mandible	2
Squamous cell carcinoma	2
Maxillary sinus	1
Adenoid cystic carcinoma	1
Mouth	4
Intermediate malignant tumor	1
Squamous cell carcinoma	3
Palate	1
Pleomorphic adenoma with pseudoepitheliomatous hyperplasia	1
Parotid gland	6
Acinic cell carcinoma	1
Carcinoma ex basal cell adenoma	1
Carcinoma ex pleomorphic adenoma (Salivary duct carcinoma)	2
Residual salivary duct carcinoma	1
Squamous cell carcinoma	1
Retromolar region	1
Squamous cell carcinoma	1
Submandibular region	1
Squamous cell carcinoma	1
Tongue	21
Squamous cell carcinoma	21
Tonsil	3
Spindle cell carcinoma	1
Squamous cell carcinoma	2

Brain	3	58.3±8.1
Glioblastoma	1	
Low grade glioma	1	
Oligodendroglioma	1	
Bile duct	2	72.0±8.5
Adenocarcinoma	2	
Gallbladder	1	71
Adenocarcinoma	1	
Total		907 65.5±13.0

Supplementary Table 2 The number of tumor-specific non-synonymous somatic mutations found in 19 tumor types identified using whole exome sequencing (n=907)

Gene	Esophagus	Colorectum	Stomach	Lung	Head&Neck	Pancreas	Sarcoma	Liver	Breast	Kidney	Melanoma	Thymus	GIST	Uterus	Others**	Ovary	Bile duct	Skin	Gallbladder
No. of cases	18	311	116	176	73	18	16	61	60	13	5	6	9	12	3	4	2	3	1
ABL1	2	10	6	2	2		1		1										
ACVR1B*		3	3	1										1					
AKT1		3	3					2	4										
ALK	1	15	6	8	2									1					
AMER1		19	2	3	1		2	1						3					
APC		348	8	4	1									7			1		
AR	1	13	4	1					1			1		2					
ARID1A	4	18	17	10	3			10	4	1				12		1			
ARID1B*		2	2					2	2					3					
ARID2	1	20	6	6	2			4		2				2					
ASXL1	1	10	1	5				4	1					3		1			
ATM	2	27	7	9	2			11	2					6					
ATRX		11	3	8	1				2					3					
AXIN1*	1	3	3		1			4						1					
B2M*		8	3		1									1					
BAP1	1	3	3	3										1					
BCL2		3	2		1														
BCOR*		6	2	3	1		1												
BRAF		23	6	4	1						1			1					
BRCA1*		6		3				1	1					2					
BRCA2*		7	1	2	2			1	1					7					
CARD11		9	4	4				1		1			2	4	1				
CASP8*		5	5		5			1						1				1	
CBL		6	3	5						1				1					
CCND1		2	2	4					1	1				1					
CDC73		2	2		1									3					
CDH1		4	13	2				1	3					1					
CDKN2A	1	2	3	3	7	2		1						1			1		1
CDKN2C				2												1			
CEBPA								1		1									
CIC		8	9	4			1		2					4	1	1	1		
CREBBP		16	6	8	3	1		2		1				1					
CRLF2*		1																	
CSF1R		6	2	3					1					3					
CTNNB1		12	8	8		1		18						2					
CYLD		2		2	1	1		1											
DAXX		3	3	3										4					
DNMT1*		3	4		2			1	1							1			
DNMT3A	1	11	5	5	1			4	1				1	2					
EGFR		5	6	73	1									2					
EP300	1	9	7	5	2		1	1	2	1				5		1			
ERBB2		11	14	4					2				1	2			1		
EZH2		3	1	1					1										
FBXW7	2	58	4	2	3			1	1					2				1	
FGFR2		12	2	1				1	1										
FGFR3	2	7	4	3	4		1		1							1			
FLT3		9	2	3				1	1					2					
FOXL2		1	1																
FUBP1*		1		1										1					

Gene	Esophagus	Colorectum	Stomach	Lung	Head&Neck	Pancreas	Sarcoma	Liver	Breast	Kidney	Melanoma	Thymus	GIST	Uterus	Brain	Ovary	Bile duct	Skin	Gallbladder
No. of cases	18	311	116	176	73	18	16	61	60	13	5	6	9	12	3	4	2	3	1
GATA1		3	2		2				1										
GATA2		3	2	2															
GATA3		8		4				1	9										
GNA11				1				1								1			
GNAQ		3	1	1										1					
GNAS		22	6	8	2			2	2					2					
H3F3A*		1	1						2										
HIST1H3B*				1	1														
HNF1A		4	2	4				2											
HRAS		1	3		3					1		1							
IDH1		1					1							1	1				
IDH2		1	1		2														
IKZF1		12	4	2				1	1										
JAK1		4	3	1					1										
JAK2	1	3	3	2				1						3					
JAK3	1	4	2	6	2			1	1					1					
KDM5C		5	4	2					1	1				2					
KDM6A		7	8	3	2			2						4					
KIT		7	5	2				1					15	3					
KLF4*																			
KMT2C		26	18	10	6	2		9	6	1				4					
KMT2D	2	50	23	30	12	1		9	7	2			2	4			1		
KRAS		132	12	27		8	1	1						2					
LMO1*																			
MAP2K1		3	3																
MAP2K4		4		3				2	1					3					
MAP3K1*		4	3	2				2	3					2					
MDM2		1		1	1														
MDM4		2		2				1											
MED12*		5	3	2	2		1							5					
MEN1		7	5	3	4		2	3		1				1					
MET		13	5	2				1	1	1				3					
MLH1		5	2		1				1					2					
MPL		4	2	3										2					
MSH2		2	2	3				1						1			1		
MSH6		8	3	5	1									4					
MYC		1			2									1					
MYCL		2	1	1										1					
MYCN																			

Supplementary Table 2 (Continued)

Gene	Esophagus	Colorectum	Stomach	Lung	Head&Neck	Pancreas	Sarcoma	Liver	Breast	Kidney	Melanoma	Thymus	GIST	Uterus	Brain	Ovary	Bile duct	Skin	Gallbladder
No. of cases	18	311	116	176	73	18	16	61	60	13	5	6	9	12	3	4	2	3	1
<i>NRAS</i>		8	1																
<i>PAX5</i>		3	2	5	1			1											
<i>PBRM1</i>		7	7	2	1		1	5		4				3					
<i>PDGFRA</i>		12	3					2						5					
<i>PHF6*</i>																			
<i>PIK3CA</i>	4	49	12	8	9		3	2	21					11					
<i>PIK3R1</i>		14	4	3	1		2	1	2					4					
<i>PPP2R1A</i>		8		1	1			1											
<i>PRDM1</i>		1	1	1															
<i>PTCH1</i>	2	13	5	8	3			2						5					
<i>PTEN</i>		27	7	7	3				7					10					
<i>PTPN11</i>		5		3	1														
<i>RB1</i>		5	4	9			1	2	2					5					
<i>RET</i>	1	14	5	8	3				2					1					
<i>RNF43*</i>	1	10	4	1										1					
<i>RUNX1</i>		6	1						1					1					
<i>SETBP1*</i>		7	2	7	1			2	2					2					
<i>SETD2</i>		4		5				3		2		1		3					
<i>SF3B1</i>		3	7	4	2				2	1									
<i>SKP2*</i>		1		1					1	1									
<i>SMAD2</i>		12	1	2															
<i>SMAD4</i>	1	22	9	2	1														
<i>SMARCA4</i>		21	17	11	2		2	2						2		1			
<i>SMARCB1</i>		5	1	1	2									1					
<i>SMO</i>		3	2	1	1		1												
<i>SOCS1*</i>																			
<i>SOX9*</i>		9	3	1												1			
<i>SPOP*</i>			1	1										2					
<i>SRSF2*</i>			1		1														
<i>STAG2*</i>			1		1				2					4					
<i>STK11</i>		3	2	7				1						1					
<i>TET2</i>		10	4	3	1			2	1			1		4					
<i>TNFAIP3</i>		2		1					1										
<i>TP53</i>	11	224	69	86	42	7	3	12	14					2		1	1	1	2
<i>TRAF7*</i>		3	2		1		1							1					
<i>TSC1</i>		7	3	4				2						1					
<i>TSHR</i>		7	2	1	1									1					
<i>U2AF1*</i>			1	3	1									2					
<i>VHL</i>		1								9									
<i>WT1</i>		5	2	1	1				1	9				1					

Nonsynonymous somatic mutations included missense, nonsense, frame-shift insertion or deletion (indel) and splice-site mutations.

In each case, if multiple mutations were found in a gene, all were counted.

*An asterisk denotes gene that is not involved in CCP genes

Supplementary Table 3 Association between patient characteristics and TP53-related mutations in many type of tumors

Cases (n)		TP53	TP63	TP73	TP53BP1	TP53BP2	TP53BP3	TP53BP11	TP53BP13	TP53BP1P	TP53BP2G5	TP53BP2NP2	TP53BP3K	ATM	ATR	PTEN	RB1	CDKN1A	CDKN2A	MDM2	AKT1	BAX	CCND1	CCNE1	PHLDA3	PUMA (BBC3)	CREBBP	EP300
Total (cases)	113	52	3	6	4	1	0	0	0	2	0	1	0	5	3	3	2	0	2	0	2	0	0	0	0	0	5	4
Female	33	13	1	3	2	1	0	0	0	0	0	0	1	3	2	1	1	0	0	0	0	0	0	0	0	2	0	
Male	80	39	2	3	2	0	0	0	0	2	0	1	0	4	0	1	0	0	2	0	2	0	0	0	0	3	4	
P-value*	0.411	1	0.356	0.579	n.a.	n.a.	n.a.	n.a.	1	n.a.	n.a.	n.a.	0.0233	0.204	0.501	n.a.	1	n.a.	1	n.a.	1	n.a.	n.a.	n.a.	n.a.	0.628	0.320	
>70	25	4	1	0	0	1	0	0	0	2	0	1	0	3	2	1	1	0	0	1	0	0	0	0	0	4	0	
<=70	53	27	1	2	2	0	0	0	0	0	0	0	0	2	1	3	1	0	1	0	0	0	0	0	0	1	0	
P-value	0.350	1	0.683	1	n.a.	n.a.	n.a.	n.a.	0.498	n.a.	n.a.	n.a.	1	1	0.1001	1	n.a.	0.498	n.a.	1	n.a.	n.a.	n.a.	n.a.	n.a.	0.369	0.121	
Non-smoker	32	12	2	2	2	0	0	0	0	1	0	0	0	2	2	2	1	0	0	0	1	0	0	0	0	2	0	
Smoker	81	40	1	4	2	1	0	0	0	1	0	1	0	3	1	1	1	0	2	0	1	0	0	0	0	3	4	
P-value	0.298	0.193	1	0.318	n.a.	n.a.	n.a.	n.a.	0.49	n.a.	n.a.	n.a.	0.621	0.193	0.193	0.488	1	n.a.	0.488	n.a.	0.488	n.a.	n.a.	n.a.	n.a.	0.621	0.576	
Non-diabetes	94	40	1	6	4	1	0	0	1	0	0	0	0	5	3	2	2	0	0	2	0	0	0	0	0	4	0	
Diabetes	19	12	2	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	
P-value	0.131	0.0728	0.587	1	n.a.	n.a.	n.a.	n.a.	0.309	n.a.	n.a.	n.a.	0.587	1	0.428	1	n.a.	1	n.a.	1	n.a.	n.a.	n.a.	n.a.	n.a.	1	0.0006	
Information missing**	3																											
Lung																												
Total (cases)	174	76	1	0	4	0	1	1	0	1	0	0	0	8	2	7	7	1	2	1	0	1	1	1	0	0	7	1
Female	62	21	0	0	1	0	0	0	0	1	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	1	0
Male	112	55	1	0	3	0	1	1	0	0	0	0	0	6	2	5	7	1	2	0	0	1	1	1	0	6	0	
P-value	0.0572	n.a.	n.a.	1	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	0.713	0.539	1	0.0514	n.a.	0.539	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	0.424	n.a.	
>70	84	32	1	0	2	0	0	0	0	1	0	0	0	3	0	5	4	0	2	0	0	1	0	1	0	0	5	1
<=70	90	44	0	0	2	0	1	1	0	0	0	0	0	5	2	2	3	1	0	1	0							

Kidney	13	6	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1	0
Total (cases)	13	6	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1	0
Female	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Male	7	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
P-value	0	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
>70	5	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<=70	8	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
P-value	0	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
Non-smoker	5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Smoker	8	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0
P-value	0	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
Non-diabetes	12	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0
Diabetes	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
P-value	0	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
Esophagus																											
Total (cases)	17	10	0	0	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	1
Female	3	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
Male	14	9	0	0	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
P-value	0	0.537	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	1	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
>70	5	3	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<=70	12	7	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	1	0
P-value	0	1	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	0.515	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.	n.a.
Non-smoker	5	4	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1
Smoker	12	6	0	0																							

Supplementary Table 3 (continued)

Ovary

[illegible]

Skin

[illegible]

Brain

[illegible]

Bile duct

[illegible]

Gallbladder

[illegible]

**P* value by Fisher exact test.

**No information about smoking status or diabetes history.

n.a.; Statistical calculation is not applicable
Data in bold indicates statistically significant

Data in bold indicate statistically significant differences ($P < .05$)