

ลิขสิทธิ์มหาวิทยาลัยเชียงใหม่

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Appendix A

**Homo sapiens TAp63 alpha mRNA sequence, complete cds (1926 bp),
GenBank accession number AF075430**

1	atgtcccaga	gcacacagac	aatgaattc	ctcagtccag	aggtttcca	gcatactg
61	gattttctgg	aacagcctat	atgttcagtt	cagcccattg	actgaactt	tgtggatgaa
121	ccatcagaag	atggtgagac	aaacaagatt	gagattagca	tggactgtat	ccgcatgcag
181	gactcggacc	tgagtgacc	catgtggcca	cagtacacga	acctggggct	cctgaacagc
241	atggaccagc	agattcagaa	cggtcctcg	tccaccagtc	cctataacac	agaccacgcg
301	cagaacagcg	tcacggcgcc	ctcgccctac	gcacagccca	gctccactt	cgatgtctc
361	tctccatcac	ccgcatccc	ctccaacacc	gactaccag	gcccgcacag	tttcgacgtg
421	tccttcagc	agtcgagcac	cgccaagtgc	gccacctgga	cgtattccac	tgaactgaag
481	aaactctact	gccaattgc	aaagacatgc	cccatccaga	tcaaggtgat	gaccccacct
541	cctcagggag	ctgttatccg	cgccatgctt	gtctacaaa	aagctgagca	cgctacggag
601	gtggtgaagc	ggtgccccaa	ccatgagctg	agccgtgaat	tcaacgaggg	acagattgcc
661	cctcctagtc	atttgattcg	agtagagggg	aacagccatg	cccagtatgt	agaagatccc
721	atcacaggaa	gacagagtgt	gctggtacct	tatgagccac	cccaggttgg	cactgaattc
781	acgacagtct	tgtacaattt	catgtgtaac	agcagttgtg	ttggagggat	gaaccgctgc
841	ccaattttaa	tcattgttac	tctggaacc	agagatgggc	aagtctggg	ccgacgctgc
901	tttgaggccc	ggatctgtgc	ttgccagga	agagacagga	aggcggatga	agatagcatc
961	agaaagcagc	aagtttcgga	cagtacaaag	aacggtgatg	gtacgaagcg	cccgttctgt
1021	cagaacacac	atggtatcca	gatgacatcc	atcaagaaac	gaagatcccc	agatgatgaa
1081	ctgttatact	taccagtgag	gggccgtgag	acttatgaaa	tgtctgtgaa	gatcaagag
1141	tccttgaac	tcatgcagta	ccttctcag	cacacaattg	aaacgtacag	gcaacagcaa
1201	cagcagcagc	accagcactt	acttcagaaa	cagacctcaa	tacagtctcc	atcttcatat
1261	ggtaacagct	ccccacctct	gaacaaaatg	aacagcatga	acaagctgcc	ttctgtgagc
1321	cagcttatca	accctcagca	gcgcaacgcc	ctcactccta	caaccattcc	tgatggcatg
1381	ggagccaaca	ttccatgat	gggcacccac	atgccaatgg	ctggagacat	gaatggactc
1441	agccccacc	aggcactccc	tccccactc	tccatgcat	ccacctccca	ctgcacacc
1501	ccacctcgt	atcccacaga	ttgcagcatt	gtcagttct	tagcgagggt	gggctgttca
1561	tcatgtctgg	actattcac	gaccagggg	ctgaccacca	tctatcagat	tgagcattac
1621	tccatggatg	atctggcaag	tctgaaaatc	cctgagcaat	ttcgacatgc	gatctggaag
1681	ggcatcctgg	accaccggca	gtcccacgaa	ttctctccc	cttctcatct	cctgcggacc
1741	ccaagcagt	cctctacagt	cagtgtgggc	tccagtgaga	ccgggggtga	gcgtgttatt
1801	gatgtctgtc	gattcaccct	ccgccagacc	atctctttcc	caccccgaga	tgagtggaat
1861	gacttcaact	ttgacatgga	tgctcgcgc	aataagcaac	agcgcatcaa	agaggagggg
1921	gagtga					

Appendix B

**Homo sapiens TAp63 alpha amino acid sequence,
coded by GenBank accession number AF075430**

(641 amino acids)

1	atg	tcc	cag	agc	aca	cag	aca	aat	gaa	ttc	ctc	agt	cca	gag
	M	S	Q	S	T	Q	T	N	E	F	L	S	P	E
15	gtt	ttc	cag	cat	atc	tgg	gat	ttt	ctg	gaa	cag	cct	ata	tgt
	V	F	Q	H	I	W	D	F	L	E	Q	P	I	C
29	tca	gtt	cag	ccc	att	gac	ttg	aac	ttt	gtg	gat	gaa	cca	tca
	S	V	Q	P	I	D	L	N	F	V	D	E	P	S
43	gaa	gat	ggt	gcg	aca	aac	aag	att	gag	att	agc	atg	gac	tgt
	E	D	G	A	T	N	K	I	E	I	S	M	D	C
57	atc	cgc	atg	cag	gac	tcg	gac	ctg	agt	gac	ccc	atg	tgg	cca
	I	R	M	Q	D	S	D	L	S	D	P	M	W	P
71	cag	tac	acg	aac	ctg	ggg	ctc	ctg	aac	agc	atg	gac	cag	cag
	Q	Y	T	N	L	G	L	L	N	S	M	D	Q	Q
85	att	cag	aac	ggc	tcc	tcg	tcc	acc	agt	ccc	tat	aac	aca	gac
	I	Q	N	G	S	S	S	T	S	P	Y	N	T	D
99	cac	gcg	cag	aac	agc	gtc	acg	gcg	ccc	tcg	ccc	tac	gca	cag
	H	A	Q	N	S	V	T	A	P	S	P	Y	A	Q
113	ccc	agc	tcc	acc	ttc	gat	gct	ctc	tct	cca	tca	ccc	gcc	atc
	P	S	S	T	F	D	A	L	S	P	S	P	A	I
127	ccc	tcc	aac	acc	gac	tac	cca	ggc	cgg	cac	agt	ttc	gac	gtg
	P	S	N	T	D	Y	P	G	P	H	S	F	D	V
141	tcc	ttc	cag	cag	tcg	agc	acc	gcc	aag	tcg	gcc	acc	tgg	acg
	S	F	Q	Q	S	S	T	A	K	S	A	T	W	T
155	tat	tcc	act	gaa	ctg	aag	aaa	ctc	tac	tgc	caa	att	gca	aag
	Y	S	T	E	L	K	K	L	Y	C	Q	I	A	K
169	aca	tgc	ccc	atc	cag	atc	aag	gtg	atg	acc	cca	cct	cct	cag
	T	C	P	I	Q	I	K	V	M	T	P	P	P	Q

Appendix B (continued)

**Homo sapiens TAp63 alpha amino acid sequence,
coded by GenBank accession number AF075430
(641 amino acids)**

183	gga G	gct A	gtt V	atc I	cgc R	gcc A	atg M	cct P	gtc V	tac Y	aaa K	aaa K	gct A	gag E
197	cac H	gtc V	acg T	gag E	gtg V	gtg V	aag K	cgg R	tgc C	ccc P	aac N	cat H	gag E	ctg L
211	agc S	cgt R	gaa E	ttc F	aac N	gag E	gga G	cag Q	att I	gcc A	cct P	cct P	agt S	cat H
225	ttg L	att I	cga R	gta V	gag E	ggg G	aac N	agc S	cat H	gcc A	cag Q	tat Y	gta V	gaa E
239	gat D	ccc P	atc I	aca T	gga G	aga R	cag Q	agt S	gtg V	ctg L	gta V	cct P	tat Y	gag E
253	cca P	ccc P	cag Q	gtt V	ggc G	act T	gaa E	ttc F	acg T	aca T	gtc V	ttg L	tac Y	aat N
267	ttc F	atg M	tgt C	aac N	agc S	agt S	tgt C	gtt V	gga G	ggg G	atg M	aac N	cgc R	cgt R
281	cca P	att I	tta L	atc I	att I	gtt V	act T	ctg L	gaa E	acc T	aga R	gat D	ggg G	caa Q
295	gtc V	ctg L	ggc G	cga R	cgc R	tgc C	ttt F	gag E	gcc A	cgg R	atc I	tgt C	gct A	tgc C
309	cca P	gga G	aga R	gac D	agg R	aag K	gcg A	gat D	gaa E	gat D	agc S	atc I	aga R	aag K
323	cag Q	caa Q	gtt V	tcg S	gac D	agt S	aca T	aag K	aac N	ggt G	gat D	ggt G	acg T	aag K
337	cgc R	ccg P	ttt F	cgt R	cag Q	aac N	aca T	cat H	ggt G	atc I	cag Q	atg M	aca T	tcc S
351	atc I	aag K	aaa K	cga R	aga R	tcc S	cca P	gat D	gat D	gaa E	ctg L	tta L	tac Y	tta L

Appendix B (continued)

**Homo sapiens TAp63 alpha amino acid sequence,
coded by GenBank accession number AF075430
(641 amino acids)**

365	cca P	gtg V	agg R	ggc G	cgt R	gag E	act T	tat Y	gaa E	atg M	ctg L	ttg L	aag K	atc I
379	aaa K	gag E	tcc S	ctg L	gaa E	ctc L	atg M	cag Q	tac Y	ctt L	cct P	cag Q	cac H	aca T
393	att I	gaa E	acg T	tac Y	agg R	caa Q	cag Q	caa Q	cag Q	cag Q	cag Q	cac H	cag Q	cac H
407	tta L	ctt L	cag Q	aaa K	cag Q	acc T	tca S	ata I	cag Q	tct S	cca P	tct S	tca S	tat Y
421	ggc G	aac N	agc S	tcc S	cca P	cct P	ctg L	aac N	aaa K	atg M	aac N	agc S	atg M	aac N
435	aag K	ctg L	cct P	tct S	gtg V	agc S	cag Q	ctt L	atc I	aac N	cct P	cag Q	cag Q	cgc R
449	aac N	gcc A	ctc L	act T	cct P	aca T	acc T	att I	cct P	gat D	ggc G	atg M	gga G	gcc A
463	aac N	att I	ccc P	atg M	atg M	ggc G	acc T	cac H	atg M	cca P	atg M	gct A	gga G	gac D
477	atg M	aat N	gga G	ctc L	agc S	ccc P	acc T	cag Q	gca A	ctc L	cct P	ccc P	cca P	ctc L
491	tcc S	atg M	cca P	tcc S	acc T	tcc S	cac H	tgc C	aca T	ccc P	cca P	cct P	ccg P	tat Y
505	ccc P	aca T	gat D	tgc C	agc S	att I	gtc V	agt S	ttc F	tta L	gcg A	agg R	ttg L	ggc G
519	tgt C	tca S	tca S	tgt C	ctg L	gac D	tat Y	ttc F	acg T	acc T	cag Q	ggg G	ctg L	acc T
533	acc T	atc I	tat Y	cag Q	att I	gag E	cat H	tac Y	tcc S	atg M	gat D	gat D	ctg L	gca A

Appendix B (continued)

**Homo sapiens TAp63 alpha amino acid sequence,
coded by GenBank accession number AF075430
(641 amino acids)**

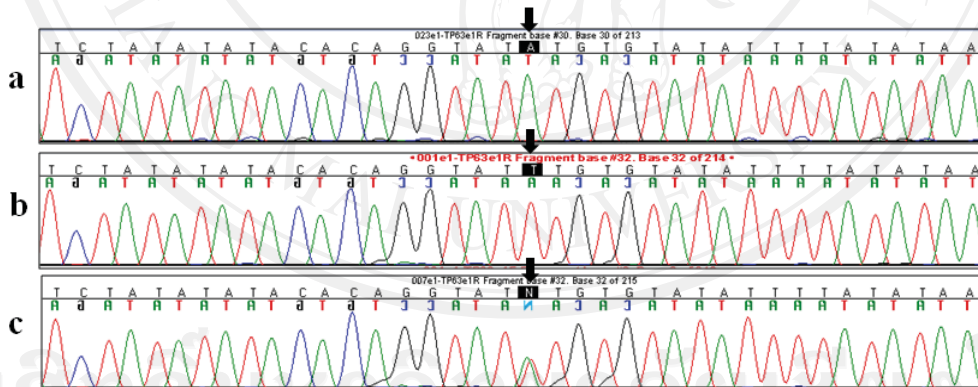
547	agt	ctg	aaa	atc	cct	gag	caa	ttt	cga	cat	gcg	atc	tgg	aag
	S	L	K	I	P	E	Q	F	R	H	A	I	W	K
561	ggc	atc	ctg	gac	cac	cgg	cag	ctc	cac	gaa	ttc	tcc	tcc	cct
	G	I	L	D	H	R	Q	L	H	E	F	S	S	P
575	tct	cat	ctc	ctg	cgg	acc	cca	agc	agt	gcc	tct	aca	gtc	agt
	S	H	L	L	R	T	P	S	S	A	S	T	V	S
589	gtg	ggc	tcc	agt	gag	acc	cgg	ggt	gag	cgt	gtt	att	gat	gct
	V	G	S	S	E	T	R	G	E	R	V	I	D	A
603	gtg	cga	ttc	acc	ctc	cgc	cag	acc	atc	tct	ttc	cca	ccc	cga
	V	R	F	T	L	R	Q	T	I	S	F	P	P	R
617	gat	gag	tgg	aat	gac	ttc	aac	ttt	gac	atg	gat	gct	cgc	cgc
	D	E	W	N	D	F	N	F	D	M	D	A	R	R
631	aat	aag	caa	cag	cgc	atc	aaa	gag	gag	ggg	gag	tga		
	N	K	Q	Q	R	I	K	E	E	G	E	Stop		

Appendix C

The previously reported SNPs of *TP63* found in the study

1. RefSNP ID: rs28673064; dbSNP Database (Appendix C.1)

- a. This SNP was located in 5'UTR of *TP63* in 12 samples, including 5 heterozygous and 7 homozygous states.
- b. HGVS Names:
 - i. NG_007550.1: g.5032A>T
 - ii. NM_003722.4: c.-58A>T



Appendix C.1 Chromatogram of RefSNP ID: rs28673064: (a) Normal sequence.

(b) A heterozygous mutation of A>T within 5'UTR. (c) A homozygous SNP of A>T

within 5'UTR. Arrows indicate the position of the changes. (NM_003722.4: c.-

58A>T) (RefSNP ID: rs28673064; dbSNP Database).

Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

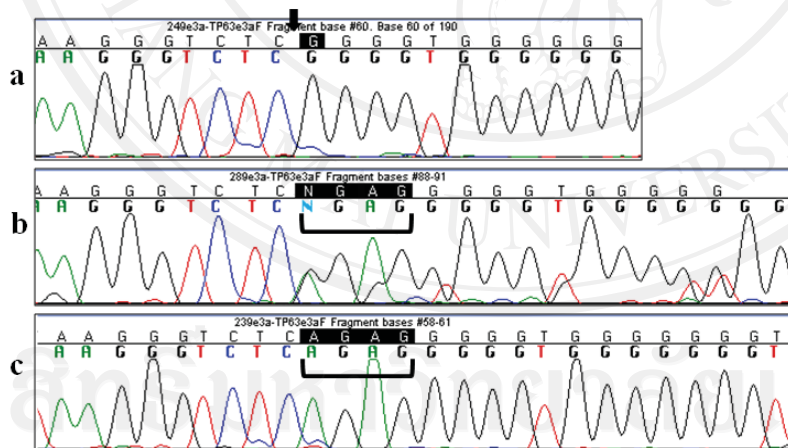
2. RefSNP ID: rs62702062; dbSNP Database (**Appendix C.2**)

a. This SNP was located in Intron 3 of *TP63* in 12 samples, including 8 heterozygous and 4 homozygous states.

b. HGVS Names:

i. NG_007550.1: g.163304_163305ins4

ii. NM_003722.4: c.325-18542_325-18541ins4



Appendix C.2 Chromatogram of RefSNP ID: rs62702062: (a) Normal sequence.

(b) A heterozygous mutation of AGAG insertion within intron 3. (c) A homozygous

SNP of AGAG insertion within intron 3. The arrow indicates the position of

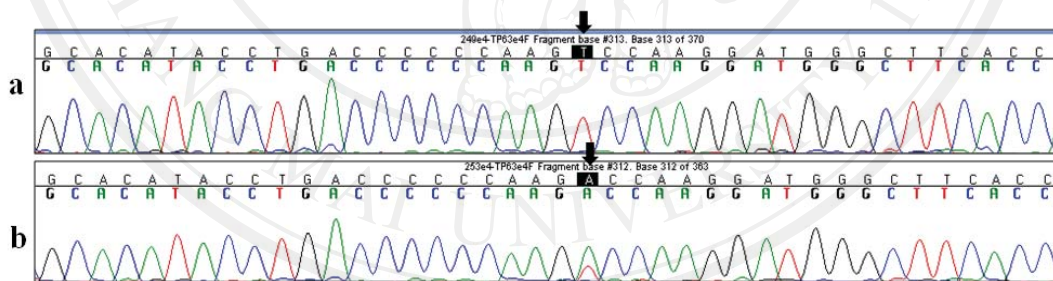
insertion. (NM_003722.4: c.325-18542_325-18541ins4) (RefSNP ID: rs62702062;

dbSNP Database).

Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

3. RefSNP ID: rs34429985; dbSNP Database (**Appendix C.3**)
 - a. This SNP was located in intron 4 of *TP63* in 2 samples, including 2 heterozygous states.
 - b. HGVS Names:
 - i. NG_007550.1: g.182139T>A
 - ii. NM_003722.4: c.579+39G>T



Appendix C.3 Chromatogram of RefSNP ID: rs34429985: (a) Normal sequence.

(b) A heterozygous SNP of T>A within intron 4. Arrows indicate the position of the change. (NM_003722.4: c.579+39G>T) (RefSNP ID: rs34429985; dbSNP Database).

Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

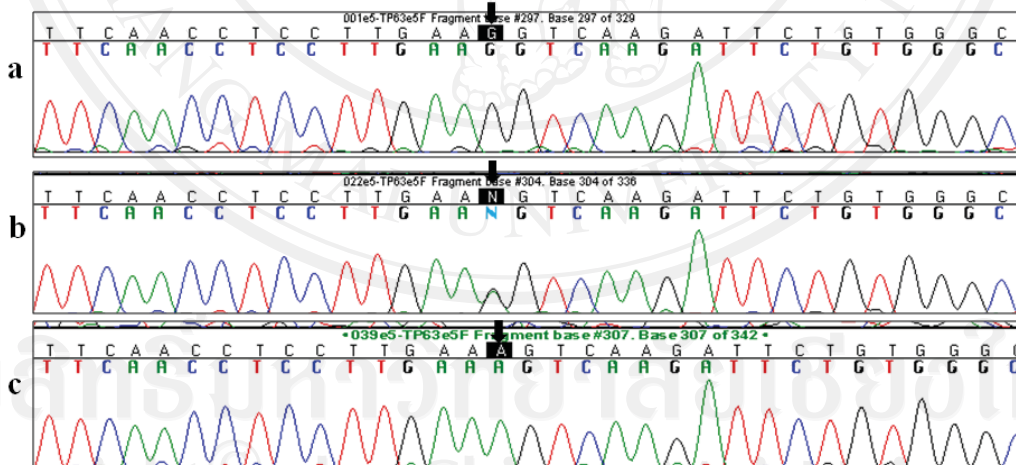
4. RefSNP ID: rs2276792; dbSNP Database (**Appendix C.4**)

a. This SNP was located in intron 5 of *TP63* in 3 samples, including 2 heterozygous and 1 homozygous states.

b. HGVS Names:

i. NG_007550.1: g.238034G>A

ii. NM_003722.4: c.766+42G>A



Appendix C.4 Chromatogram of RefSNP ID: rs2276792: (a) Normal sequence. (b)

A heterozygous SNP of G>A within intron 5. (c) A homozygous SNP of G>A within

intron 5. Arrows indicate the position of the changes. (NM_003722.4:

c.766+42G>A) (RefSNP ID: rs2276792; dbSNP Database).

Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

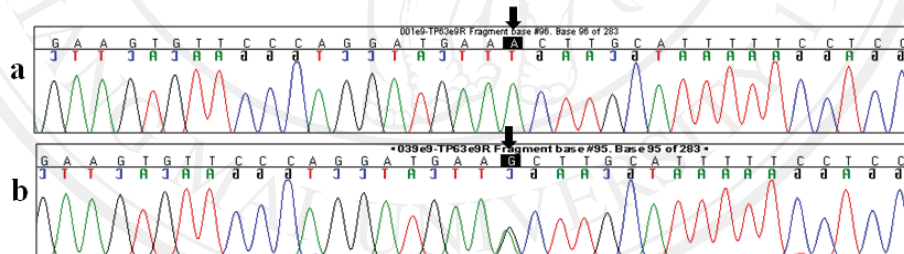
5. RefSNP ID: rs6789961; dbSNP Database (**Appendix C.5**)

a. This SNP was located in intron 8 of *TP63* in 5 samples, including 5 heterozygous states.

b. HGVS Names:

i. NG_007550.1: g.242876A>G

ii. NM_003722.4: c.1130-22C>A



Appendix C.5 Chromatogram of RefSNP ID: rs6789961: (a) Normal sequence. (b)

A heterozygous SNP of A>G within intron 8. Arrows indicate the position of the

change. (NM_003722.4: c.1130-22C>A) (RefSNP ID: rs6789961; dbSNP Database).

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Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

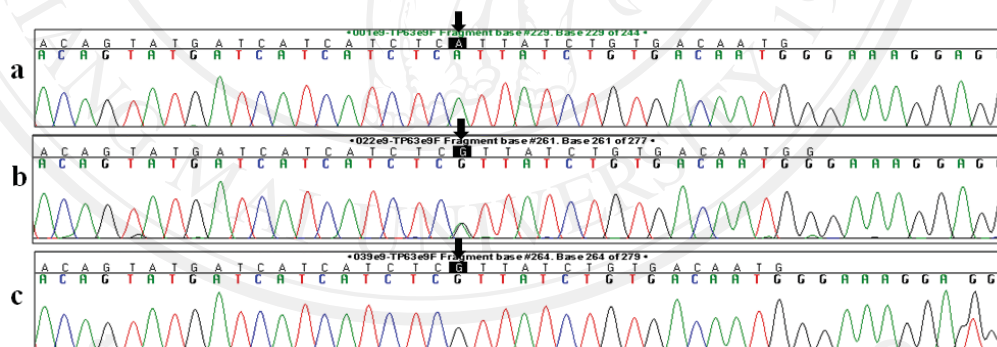
6. RefSNP ID: rs6790167; dbSNP Database (**Appendix C.6**)

a. This SNP was located in intron 9 of *TP63* in 10 samples, including 8 heterozygous and 2 homozygous states.

b. HGVS Names:

i. NG_007550.1: g.243059A>G,

ii. NM_003722.4: c.1212+79A>G



Appendix C.6 Chromatogram of RefSNP ID: rs6790167: (a) Normal sequence. (b)

A heterozygous SNP of A>G within intron 9. (c) A homozygous SNP of A>G within intron 9. Arrows indicate the position of the changes. (NM_003722.4:

c.1212+79A>G) (RefSNP ID: rs6790167; dbSNP Database).

Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

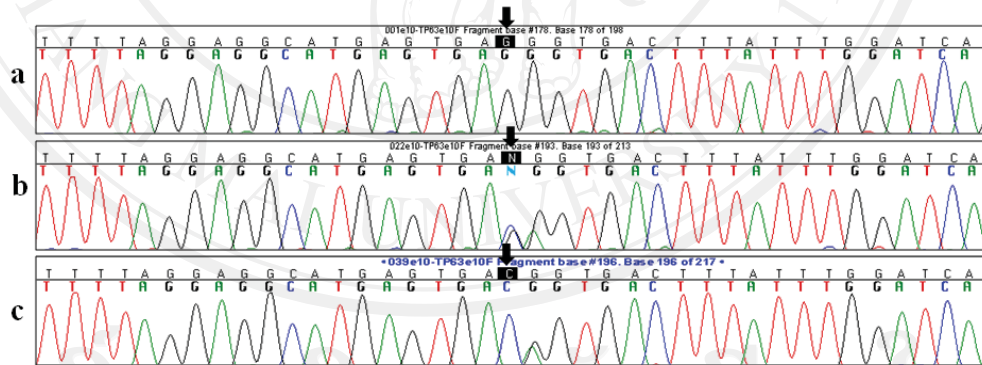
7. RefSNP ID: rs9840359; dbSNP Database (**Appendix C.7**)

a. This SNP was located in intron 10 of *TP63* in 11 samples, including 9 heterozygous and 2 homozygous states.

b. HGVS Names:

i. NG_007550.1: g.246609G>C,

ii. NM_003722.4: c. 1349+40A>G



Appendix C.7 Chromatogram of RefSNP ID: rs9840359: (a) Normal sequence. (b)

A heterozygous SNP of G>C within intron 10. (c) A homozygous SNP of G>C

within intron 10. Arrows indicate the position of the changes. (NM_003722.4:

c.1349+40A>G) (RefSNP ID: rs9840359; dbSNP Database).

Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

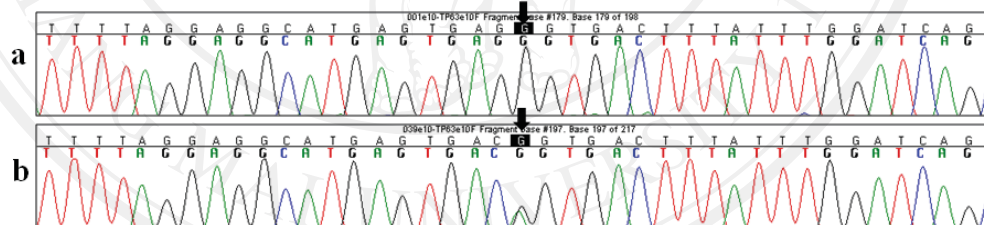
8. RefSNP ID: rs9840360; dbSNP Database (**Appendix C.8**)

a. This SNP was located in intron 10 of *TP63* in 1 sample, a heterozygous state.

b. HGVS Names :

i. NG_007550.1: g.246610G>A

ii. NM_003722.4: c. 1349+41A>G



Appendix C.8 Chromatogram of RefSNP ID: rs9840360: (a) Normal sequence. (b)

A heterozygous SNP of G>A within intron 10. Arrows indicate the position of the

change. (NM_003722.4: c.1349+41A>G) (RefSNP ID: rs9840360; dbSNP

Database).

Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

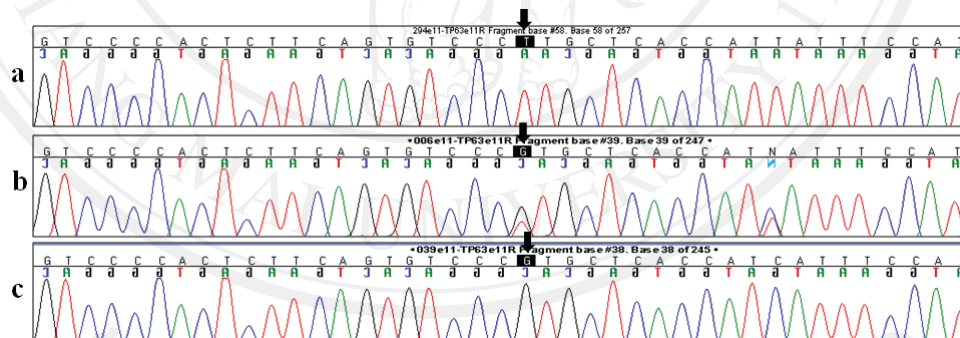
9. RefSNP ID: rs1554131; dbSNP Database (**Appendix C.9**)

a. This SNP was located in intron 10 of *TP63* in 17 samples, including 9 heterozygous and 8 homozygous states.

b. HGVS Names :

i. NG_007550.1: g.259934T>G

ii. NM_003722.4: c. 1350-34G>T



Appendix C.9 Chromatogram of RefSNP ID: rs1554131: (a) Normal sequence. (b)

A heterozygous SNP of T>G within intron 10. (c) A homozygous SNP of T>G

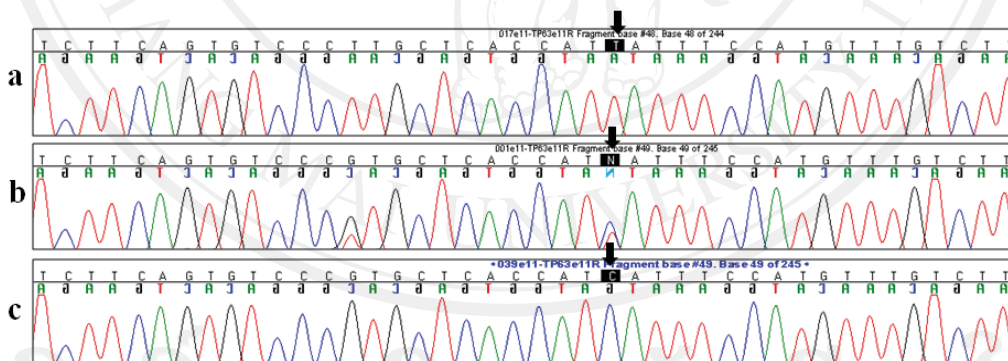
within intron 10. Arrows indicate the position of the changes. (NM_003722.4: 1350-34G>T) (RefSNP ID: rs1554131; dbSNP Database).

Appendix C (continued)

The previously reported SNPs of *TP63* found in the study

10. RefSNP ID: rs1345186; dbSNP Database (**Appendix C.10**)

- a. This SNP was located in intron 10 of *TP63* in 21 samples, including 9 heterozygous and 12 homozygous states.
- b. HGVS Names :
 - i. NG_007550.1: g.259945T>C
 - ii. NM_003722.4: c. 1350-23G>T



Appendix C.10 Chromatogram of RefSNP ID: rs1345186: (a) Normal sequence.

(b) A heterozygous SNP of T>C within intron 10. (c) A homozygous SNP of T>C

within intron 10. Arrows indicate the position of the changes. (NM_003722.4: 1350-

23G>T) (RefSNP ID: rs1345186; dbSNP Database).

Appendix D

List of chemicals and materials were used in the study

Chemicals and materials	Source
Absolute ethanol	Merck KGaA, Darmstadt, Germany
Agarose (Molecular Biology Grade)	Vivantis, Selangor DE, Malaysia
Ammonium bicarbonate (Molecular Biology Grade)	Sigma-Aldrich Quimica S. A. Toluca, Mexico
Ammonium chloride (Molecular Biology Grade)	Sigma-Aldrich Japan K.K. Tokyo, Japan
AmpliTaq Gold [®] Polymerase	Applied Biosystems, USA
Boric acid (Molecular Biology Grade)	Vivantis, Selangor DE, Malaysia
Deoxynucleoside triphosphate (dNTPs) Mix (each 10 mM)	Fermentas UAB, Vilnius, Lithuania
EDTA, Disodium Salt, Dihydrate (Molecular Biology Grade)	Vivantis, Selangor DE, Malaysia
GeneRuler [™] 100 bp Plus DNA Ladder, ready-to-us (Loading dye +Ladder dye) #SM0323	Fermentas UAB, Vilnius, Lithuania
Isopropanol	Merck KGaA, Darmstadt, Germany
MgCl ₂ solution	Applied Biosystems Inc, Foster City, CA, USA
Oligonucleotide primers	1 st BASE Pte Ltd, Singapore
PCR Buffer II (10x)	Applied Biosystems Inc, Foster City, CA, USA
Proteinase K (20 mg/mL)	New England BioLab, Ipswich, MA, USA

Appendix D (continued)**List of chemicals and materials were used in the study**

Chemicals and materials	Source
SYBR [®] Safe DNA gel stain 10,000x concentrate in DMSO	Invitrogen Inc., Carlsbad, CA, USA
Sodium Azide (Molecular Biology Grade)	USB Corporation, Cleveland, OH, USA
Sodium chloride (Molecular Biology Grade)	Sigma-Aldrich Corp., St. Louis, MO, USA
Sodium dodecyl sulfate (SDS) (Molecular Biology Grade)	USB Corporation, Cleveland, OH, USA
Sterile Water (for injection)	General Hospital Products Public, Thailand
Tris base (Molecular Biology Grade)	Vivantis, Selangor ED, Malaysia
Tris-EDTA (TE) Buffer (50x)	USB Corporation, Cleveland, OH, USA

Appendix E

List of instruments were used in the study

Instruments	Source
Adjustable micropipette (Pipet-Lite™)	Rainin Instrument LLC, Oakland, CA, USA
Adjustable Pipette (Rota-Filler 3000™)	Heathrow Scientific®, Vernon Hills, IL, USA
Aluminum Foil	Alcoa, Pittsburgh, PA, USA
BD Vacutainer® (EDTA anticoagulant)	Becton, Dickinson and Company (BD), Franklin Lakes, NJ, USA
Beaker : 100 mL, 250 mL, 600 mL, 1 L	ISOLAB Laborgeräte GmbH, Wertheim, Germany
Beckman Centrifuge	Beckman Coulter, Inc., Brea, CA, USA
Centrifuge tube 15 mL (#430052)	Corning Incorporated, Corning, NY, USA
Gloves	Sempermed®, Thailand
Hettich Centrifuge (MIKRO 200)	Andreas Hettich GmbH & Co.KG, Tuttlingen, Germany
Microcentrifuge tube : 0.2 mL, 1.5 mL	Axygen Scientific Inc, Union City, CA, US
Mini-Centrifuge (Sprout™)	Heathrow Scientific®, Vernon Hills, IL, USA
Molecular Imager Gel Doc XR and System	BIO-RAD, Hercules, CA, USA
Needle	NIPRO, THAILAND
Parafilm	PECHINEY Plastic Packaging Inc, Chicago, IL, USA
Pipette tip : 10 µL, 200 µL, 1,000 µL	Axygen Scientific Inc, Union City, CA, US

Appendix E (continued)**List of instruments were used in the study**

Instruments	Source
Polypropylene 96-Well PCR Plate with flat top and printed grid (0.2 mL)	Labcon North America, Petaluma, CA, USA
PowerPac™ Basic Power Supply	BIO-RAD, Hercules, CA, USA
Power supply model UPS ACURA plus1500	Leonics, Thailand
PYREX® Erlenmeyer flask 125 mL	Corning, Lowell, MA, USA
Reagent bottle : 50 mL, 100 mL, 250 mL, 500 mL, 1 L, 2 L	ISOLAB Laborgeräte GmbH, Wertheim, Germany SCHOTT AG, Mainz, Germany
Refrigerator-Freezers	Mitsubishi, Thailand
Sequencher 4.8	Genecodes, Ann Arbor, MI, USA
Stripette 10 mL Costar® (#4488)	Corning Incorporated, Corning, NY, USA
Sub-Cell® GT Agarose Gel Electrophoresis System	BIO-RAD, Hercules, CA, USA
Syringe	NIPRO, THAILAND
Vortex Mixer	Labnet International Inc., Woodbridge, NJ, USA
Weighing Machine (#PL303)	Mettler-Toledo, Inc., Columbus, OH, USA

Appendix F

List of solutions and buffers were used in the study

1. Red cell lysis buffer (1 L)

1 mM NH_4HCO_3	0.08 g
115 mM NH_4Cl	6.15 g
Add distilled water to	1 L
Sterilize the solution by autoclaving and store at 4°C.	

2. White cell lysis buffer (100 mL)

100 mM Tris-Cl	1.57 g
40 mM EDTA (or 8 μL 0.5 M)	1.5 g
50 mM NaCl	0.3 g
0.2% SDS	0.2 g
0.05% Sodium Azide	0.05 g
Add distilled water to	100 mL

Sterilize the solution by autoclaving and store at room temperature.

3. 6 M NaCl Solution (10 mL)

Sodium chloride	3.5 g
Add distilled water to	10 mL

Appendix F (continued)

List of solutions and buffers were used in the study

4. 1x Tris-EDTA (TE) buffer

50x TE buffer, pH 7.5	1 mL
Add distilled water to	50 mL
Mix the solution and store at room temperature.	

5. 10x Tris-Borate-EDTA (TBE) buffer (1 L)

Tris base	108 g
Boric acid	55 g
Na ₂ EDTA	7.4 g
Add distilled water to	1 L
The pH is 8.3 and requires no adjustment.	
Mix the solution and store at room temperature.	

6. 1% agarose gel (w/v)

Agarose	0.70 g
1x TBE buffer	70 mL

Dissolve by heating in microwave oven and occasional mix until no granules of agarose are visible.

CURRICULUM VITAE

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