

S8 Table. Summary of *Trp53* Protein Coding Sequence Mutations

Mutation Class	Location ^a	Alteration	Predicted Protein Change	No Tumours (if >1)
Deletion	Exon 2 – 4	del Exon 2 – 4	Aberrant protein	
Insertion	+726	ins 'GC'	Frameshift	
Point-Mutation	+329	T>G	F110C	
	+404	C>T	A135V	3
	+508	G>A	V170M	
	+578	G>C	R193P	
	+628	C>T	R210C	
	+632	A>G	H211R	
	+638	T>G	V213G	
	+692	A>G	Y231C	
	+722	G>A	G241E	
	+801	T>G	F267L	
	+808	C>T	R270C	

^a Location relative to 'A' of translation start 'ATG' codon.