

**S3 Table. Summarized sequencing results of H/K/Nras cDNAs of SCCs.**

<b>Animal</b>	<b>Tumor</b>	<b>H-ras status</b>	<b>N-ras status</b>	<b>K-ras status</b>	<b>p53 status</b>	<b>Tumor type</b>
#1	T1	wildtype	wildtype	wildtype	mutated	mixed
#2	T1	codon 71 C>T (silent)	wildtype	wildtype	mutated	nKSCC
	T2	wildtype	wildtype	wildtype	mutated	KSCC
#3	T1	wildtype	wildtype	not tested	mutated	KSCC
#4	T1	wildtype	wildtype	wildtype	mutated	nKSCC
#5	T1	wildtype	wildtype	wildtype	mutated	nKSCC
#6	T1	wildtype	wildtype	wildtype	mutated	nKSCC
#7	T1	wildtype	wildtype	wildtype	mutated	KSCC
#8	T1	wildtype	wildtype	wildtype	mutated	nKSCC
#9	T1	wildtype	wildtype	wildtype	wildtype	KSCC
	T2	wildtype	wildtype	wildtype	wildtype	KSCC
	T3	wildtype	not tested	wildtype	wildtype	KSCC
#10	T1	wildtype	wildtype	wildtype	wildtype	nKSCC
		<b>n = 13</b>	<b>n = 10</b>	<b>n = 12</b>		