

SUPPLEMENTARY TABLE S3. ΔC_T VALUES FOR ALL THE 10 MARKERS TESTED
IN QUANTITATIVE REVERSE TRANSCRIPTION REAL TIME POLYMERASE CHAIN REACTION

<i>Gene name</i>	<i>Day 8</i>	<i>Day 21</i>	<i>Day 43</i>	<i>Normal tissue</i>	<i>Tumor tissue</i>	<i>BCPAP</i>	<i>KTC1</i>
<i>PTCSC2-spliced</i>	13.51±0.92	16.57±1.14	16.43±0.32	10.10±0.04	11.69±0.10	Undetectable	15.95±1.01
<i>PTCSC2-unspliced</i>	8.36±1.44	10.79±0.36	11.00±0.36	2.29±0.27	3.38±0.14	15.39±0.67	10.05±0.29
<i>PTCSC3</i>	13.62±1.36	15.60±1.47	10.96±0.04	5.93±0.10	8.48±0.43	15.59±0.33	20.09±0.99
<i>TSHR</i>	8.45±0.27	13.31±1.48	11.13±0.25	2.49±0.18	2.64±0.32	17.18±0.31	Undetectable
<i>TPO</i>	6.76±1.14	11.13±2.36	9.94±0.17	1.56±0.20	5.12±0.32	Undetectable	19.38±2.85
<i>TG</i>	6.97±0.76	11.56±0.94	12.13±0.57	-1.65±0.33	-0.15±0.21	19.70±0.27	13.19±0.29
<i>CDH1</i>	4.36±1.63	4.69±0.34	8.79±2.07	-0.18±0.55	0.98±0.61	18.46±0.44	19.80±0.94
<i>SI00A4</i>	5.04±0.84	6.54±0.79	6.83±0.51	2.72±0.95	1.34±0.72	7.11±0.21	5.79±0.20
<i>FNI</i>	2.75±1.49	1.15±0.33	-0.41±0.77	4.85±2.63	-1.68±1.45	1.12±0.09	-2.15±0.11
<i>COL1A1</i>	5.05±2.15	-0.33±1.34	-1.05±1.28	3.34±0.90	-1.20±1.54	7.30±0.25	-0.28±0.32

Data were shown as mean ± standard variation, and all the measurements were made from three to five repeated experiments.