

**Table S2:** Primer sequence for quantitative RT-PCR analyses

<b>Gene</b>	<b>Primers</b>	
<i>TWIST1</i>	Forward	GGAGTCCGCAGTCTTACGAG
	Reverse	CCAGCTTGAGGGTCTGAATC
<i>POLR2A</i>	Forward	TCTTCTGCTCCAAACTGCTTGTGG
	Reverse	GTCGTAGACATGTGTGAGC
<i>RPL13A</i>	Forward	CATGAGGCTACGGAAACAGGC
	Reverse	AAGGGCAGGCAACGCATGA
<i>YAP1</i>	Forward	CACAGCATGTTCGAGC
	Reverse	GATGCTGAGCTGTGGGTGTA
<i>JUP</i>	Forward	CAGCAGCCCTACACGGATGGTGTGA
	Reverse	CGGTTCATGGGGTCCCCGGGC
<i>VIM</i>	Forward	ATCCAAGTTTGCTGACCTCTCTGAG
	Reverse	AGGGACTGCACCTGTCTCCGGT
<i>CDH1</i>	Forward	GGAACTATGAAAAGTGGGCTTG
	Reverse	AAATTGCCAGGCTCAATGAC
<i>CDH2</i>	Forward	CTTGTCAGGATCAGGTCT
	Reverse	GAAGATACCAGTTGGAGGCT
<i>SNAIL</i>	Forward	CTTCCAGCAGCCCTACGAC
	Reverse	CGGTGGGGTTGAGGATCT
<i>ZEB1</i>	Forward	ACTGCTGGGAGGATGACAGA
	Reverse	ATCCTGCTTCATCTGCCTGA
<i>18S</i>	Forward	TCCCCCAACTTCTTAGAGG
	Reverse	CTTATGACCCGCACTTACTG