

DNA methylation in 291 HCCs carrying wild-type CTNNB1 and 79 HCCs carrying mutated *CTNNB1* from the TCGA dataset

Gene symbol	logFC	AveExpr	<i>t</i>	P.Value	adj.P.Val	B	abs
<b><i>TMPRSS2</i></b>	1,030306228	-1,302330529	5,423771614	9,78E-08	3,27E-06	5,922974484	1,030306228
<b><i>ACE2</i></b>	-0,602063465	0,636097685	-4,558113165	6,74E-06	9,21E-05	1,834523098	0,602063465
<b><i>DPP4</i></b>	0,472521117	2,05063653	4,344378674	1,75E-05	0,000195669	0,923653403	0,472521117