

Suppl 5. Correlations between *CLDN8* mRNA expression and clinicopathological parameters of kidney renal clear cell carcinoma patients (* $p<0.05$).

Clinicopathological features		N	<i>CLDN8</i> expression		
			Mean ± SD	p	F/t
Gender	Male	35 2	0.73±1.90	0.25	1.16
	Female	18 6	0.56±1.43		
Age	<50 years	10 7	0.74±1.93	0.64	0.46
	≥50 years	43 1	0.65±1.70		
Ethnicity	Hispanic or Latino	26	0.25±0.92		
	Not Hispanic or Latino	36 1	0.84±1.99	<0.01	11.7
	Not evaluated or not available	15 1	0.33±1.05	*	1
Tumor status	T ₁	27 7	0.80±1.77		
	T ₂	71 17	1.01±2.84	0.88	0.66
	T ₃	9	0.34±0.92		
	T ₄	11	0.66±1.83		
Node status	N ₀	24 1	0.62±1.70	0.51	0.67
	N ₁	16	0.28±0.48		
	N _x	28 1	0.73±1.84		
Metastasis status	M ₀	44 8	0.73±1.86	0.77	0.53
	M ₁	80	0.37±1.02		
	M _x	10	0.18±0.22		
Tumor grade	G ₁	14	0.05±0.08		
	G ₂	23 5	0.83±1.93		
	G ₃	20 6	0.47±1.31	0.09	9.59
	G ₄	75	0.39±1.04		
	G _x	5	6.14±5.64		
Tumor stage	Not available	3	2.20±3.81		
	Stage I	27 1	0.81±1.78		
	Stage II	59	1.18±3.09		
	Stage III	12 3	0.33±0.92	0.67	2.38
	Stage IV	82	0.36±1.01		
Survival status	Unknown	3	0.22±0.20		
	Alive	36 6	0.77±1.95	0.02 *	2.39
	Dead	17 2	0.45±1.19		