

Table S6 | Pan-cancer analysis of LETM2 expression in GEPIA dataset.

Cancer Type	Tumor	Normal	P value
ACC	1.88	0.6	
BLCA	1.95	1.48	
BRCA	2.51	1.52	
CESC	3.13	2.48	
CHOL	5.42	0.21	<0.05
COAD	1.3	1.41	
DLBC	1.64	1.34	
ESCA	3.47	1.32	
GBM	3.43	3.2	
HNSC	4.44	1.4	<0.05
KICH	3.24	3.55	
KIRC	2.68	3.6	
KIRP	6.37	3.1	
LAML	3.52	2.27	
LGG	3.27	3.2	
LIHC	0.35	0.24	
LUAD	3.16	2.11	
LUSC	2.25	2.13	
OV	2.23	2.36	
PDAC	8.11	1.18	<0.05
PCPG	6.66	0.48	<0.05
PRAD	0.7	1.45	
READ	1.18	1.61	
SARC	2.93	1.52	
SKCM	1.63	0.93	
STAD	2.31	1.2	
TGCT	1.39	38.71	<0.05
THCA	1.04	1.32	
THYM	1.95	1.35	
UCEC	3.65	2.15	
UCS	2.61	2.2	

The value represents the median expression of certain tumor type or normal tissue.