

SUPPLEMENTARY TABLES

Supplementary Table 1. Details of GPER1 ROC in pan-cancer.

Tumor type	Tumor(n)	Normal(n)	AUC(CI)	Cut-off	Sensitivity	Specificity	Positive predictive value	Negative predictive value	YI
ACC	128	77	0.687(0.598-0.776)	2.033	0.532	0.922	0.804	0.766	0.454
BLCA	407	28	0.896(0.835-0.958)	1.272	0.865	0.786	0.983	0.286	0.651
BRCA	1099	292	0.849(0.827-0.871)	1.763	0.768	0.832	0.945	0.488	0.6
CESC	306	13	0.843(0.770-0.917)	1.128	0.596	1.000	1.000	0.095	0.595
CHOL	36	9	0.951(0.886-1.000)	3.181	0.889	0.917	0.727	0.971	0.806
COAD	290	349	0.964(0.951-0.977)	1.707	0.900	0.928	0.913	0.918	0.828
DLBC	47	444	0.492(0.422-0.561)	1.373	0.596	0.523	0.117	0.924	0.118
ESCA	182	666	0.781(0.745-0.818)	1.578	0.742	0.686	0.392	0.907	0.428
GBM	689	1157	0.551(0.524-0.577)	2.410	0.787	0.310	0.404	0.709	0.097
HNSC	44	502	0.784(0.725-0.843)	1.037	0.745	0.791	0.228	0.968	0.495
KICH	66	53	0.902(0.835-0.970)	1.491	0.909	0.868	0.896	0.885	0.777
KIRC	531	100	0.690(0.635-0.745)	3.079	0.678	0.650	0.911	0.275	0.328
KIRP	289	60	0.568(0.494)0.641	3.299	0.478	0.750	0.902	0.230	0.228
LAML	173	70	0.959(0.936-0.983)	0.373	0.913	0.914	0.963	0.81	0.828
LGG	523	1152	0.548(0.519-0.577)	2.058	0.946	0.161	0.339	0.869	0.107
LIHC	371	160	0.548(0.497-0.598)	2.683	0.447	0.706	0.779	0.355	0.154
LUAD	535	59	0.925(0.897-0.952)	2.704	0.898	0.865	0.424	0.987	0.764
LUSC	498	338	0.919(0.899-0.939)	1.590	0.884	0.822	0.880	0.827	0.706
OSCC	329	32	0.753 (0.678-0.828)	1.627	0.906	0.520	0.155	0.983	0.426
OV	427	88	0.787(0.741-0.833)	1.046	0.621	0.830	0.946	0.311	0.450
PAAD	179	171	0.854(0.813-0.895)	1.014	0.777	0.836	0.832	0.781	0.613
PRAD	496	152	0.787(0.749-0.824)	2.272	0.659	0.822	0.924	0.425	0.482
READ	93	318	0.965(0.947-0.983)	1.676	0.882	0.947	0.828	0.965	0.828
SKCM	469	813	0.596(0.561-0.632)	0.884	0.337	0.899	0.658	0.702	0.236
STAD	414	210	0.940(0.919-0.960)	2.555	0.903	0.848	0.921	0.817	0.751
TGCT	154	165	0.831(0.779-0.883)	1.094	0.747	0.952	0.935	0.801	0.698
THCA	512	338	0.723(0.689-0.757)	2.880	0.611	0.766	0.798	0.566	0.378
THYM	119	446	0.469(0.420-0.517)	2.252	0.824	0.377	0.261	0.889	0.200
UCEC	181	101	0.911(0.877-0.945)	1.805	0.845	0.901	0.939	0.765	0.746
UCS	57	78	0.823 (0.742-0.905)	2.051	0.737	0.859	0.792	0.817	0.596

Abbreviations: ROC, Receiver Operator Characteristic curve; AUC, Area Under Curve; CI, Confidence Interval; YI, Youden's index.

Supplementary Table 2. Details of the prognostic K-M analysis of GPER1 in pan-cancer.

Cancer	N	HR (95% CI)	P value
ACC	79	0.64(0.30-1.36)	0.243
BLCA	433	1.19(0.89-1.59)	0.243
BRCA	1090	0.69(0.50-0.97)	0.03
CESC	309	1.04(0.66-1.65)	0.865
CHOL	45	0.70(0.27-1.85)	0.473
COAD	521	1.21(0.82-1.79)	0.33
DLBC	48	0.09(0.01-0.79)	0.029
ESCA	173	0.46(0.28-0.77)	0.003
GBM	174	0.90(0.64-1.27)	0.548
HNSC	500	0.75(0.57-0.99)	0.042
KICH	89	1.30(0.35-4.83)	0.698
KIRC	611	0.59(0.44-0.80)	0.001
KIRP	288	0.45(0.25-0.81)	0.008
LAML	151	1.18(0.78-1.80)	0.436
LGG	529	0.75(0.54-1.05)	0.093
LIHC	424	1.04(0.74-1.47)	0.825
LUAD	513	0.71(0.52-0.98)	0.036
LUSC	551	1.03(0.79-1.35)	0.826
MESO	86	0.69(0.43-1.11)	0.131
OS	101	1.26(0.68-2.35)	0.462
OSCC	361	0.93(0.67-1.29)	0.675
OV	379	1.16(0.89-1.50)	0.263
PAAD	177	0.59(0.38-0.93)	0.022
PCPG	186	0.63(0.15-2.65)	0.531
PRAD	551	0.78(0.22-2.80)	0.703
READ	177	2.15(0.93-4.95)	0.073
SKCM	472	1.04(0.80-1.36)	0.764
SARC	259	0.38(0.20-0.70)	0.002
STAD	375	1.50(1.06-2.12)	0.023
TGCT	156	1.24(0.17-9.35)	0.833
THCA	568	1.72(0.64-4.65)	0.284
THYM	121	0.52(0.13-2.09)	0.359
UCEC	587	0.53(0.34-0.80)	0.003
UCS	56	1.10(0.54-2.21)	0.798
UVM	80	1.52(0.67-3.47)	0.321