Calculation of TMBPI for 530 ccRCC patients

ID	futime	fustat	TMBPI	Risk level
TCGA-3Z-A93Z	1.054794521	0	0.676488073	Low
TCGA-6D-AA2E	0.991780822	0	1.013515811	High
TCGA-A3-3306	3.068493151	0	0.967935076	Low
TCGA-A3-3307	3.934246575	0	0.731806162	Low
TCGA-A3-3308	0.043835616	0	1.252756664	High
TCGA-A3-3311	3.263013699	1	1.346109068	High
TCGA-A3-3313	2.01369863	1	1.443831067	High
TCGA-A3-3316	4.090410959	0	1.533201356	High
TCGA-A3-3317	4.084931507	0	1.44386027	High
TCGA-A3-3319	3.095890411	0	2.853270264	High
TCGA-A3-3320	4.131506849	0	1.009853379	High
TCGA-A3-3322	4.049315068	0	1.103001704	High
TCGA-A3-3323	3.030136986	0	1.280760914	High
TCGA-A3-3324	3.249315068	0	0.952110968	Low
TCGA-A3-3325	3.205479452	1	0.940472613	Low
TCGA-A3-3326	3.115068493	0	1.040472312	High
TCGA-A3-3328	3.794520548	0	0.214373555	Low
TCGA-A3-3329	4.449315068	0	0.698376577	Low
TCGA-A3-3331	3.443835616	0	0.692838926	Low
TCGA-A3-3335	5.167123288	0	0.94108278	Low
TCGA-A3-3343	2.589041096	0	0.753007265	Low
TCGA-A3-3346	0.375342466	1	1.014404733	High
TCGA-A3-3347	4.410958904	1	2.04941545	High
TCGA-A3-3349	3.794520548	0	0.706837458	Low
TCGA-A3-3351	2.493150685	0	1.001816569	High
TCGA-A3-3352	1.536986301	1	0.735529978	Low
TCGA-A3-3357	7.364383562	0	1.288093013	High
TCGA-A3-3358	3.580821918	0	1.011723565	High
TCGA-A3-3359	6.860273973	0	0.750794105	Low
TCGA-A3-3362	4.271232877	0	1.013207209	High
TCGA-A3-3363	0.873972603	0	0.942068254	Low
TCGA-A3-3365	2.391780822	0	0.785492083	Low
TCGA-A3-3367	6.219178082	0	0.828985026	Low
TCGA-A3-3370	6.230136986	0	0.74307254	Low
TCGA-A3-3372	2.01369863	0	0.980552964	High
TCGA-A3-3373	4.44109589	0	0.497292822	Low
TCGA-A3-3374	3.6	0	0.511135798	Low
TCGA-A3-3376	4.646575342	1	0.730372922	Low

TCGA-A3-3378	1.726027397	0	1.211607372	High
TCGA-A3-3380	1.553424658	0	0.791792107	Low
TCGA-A3-3382	1.57260274	0	2.006103881	High
TCGA-A3-3383	2.35890411	0	0.853121574	Low
TCGA-A3-3385	5.460273973	0	0.792379997	Low
TCGA-A3-3387	1.690410959	0	0.584400338	Low
TCGA-A3-A6NI	2.789041096	0	1.649277632	High
TCGA-A3-A6NJ	1.282191781	0	0.713819631	Low
TCGA-A3-A6NL	1.887671233	0	0.821151721	Low
TCGA-A3-A6NN	0.008219178	0	0.695705669	Low
TCGA-A3-A8CQ	0.008219178	0	0.752173753	Low
TCGA-A3-A8OU	0	0	1.298619496	High
TCGA-A3-A8OV	0.931506849	0	0.972979172	Low
TCGA-A3-A8OW	0.884931507	0	0.678180728	Low
TCGA-A3-A8OX	0	0	0.711893567	Low
TCGA-AK-3425	9.15890411	0	0.981705721	High
TCGA-AK-3426	2.424657534	1	1.568715892	High
TCGA-AK-3427	7.030136986	0	0.38631503	Low
TCGA-AK-3428	6.090410959	0	0.30538419	Low
TCGA-AK-3429	5.526027397	0	0.422261958	Low
TCGA-AK-3431	5.076712329	0	0.50363119	Low
TCGA-AK-3433	6.005479452	0	0.523026697	Low
TCGA-AK-3434	5.717808219	0	0.741194123	Low
TCGA-AK-3436	5.6	0	1.252362668	High
TCGA-AK-3440	4.780821918	0	1.049527001	High
TCGA-AK-3443	3.898630137	0	0.521963762	Low
TCGA-AK-3445	3.506849315	0	2.152738662	High
TCGA-AK-3447	3.334246575	0	2.259728286	High
TCGA-AK-3450	4.131506849	0	0.546772673	Low
TCGA-AK-3451	4.057534247	0	0.810519612	Low
TCGA-AK-3453	3.82739726	0	1.910526369	High
TCGA-AK-3454	2.394520548	0	2.344947714	High
TCGA-AK-3455	1.871232877	1	1.027165462	High
TCGA-AK-3456	3.131506849	0	0.858958508	Low
TCGA-AK-3458	3.2	0	1.020267353	High
TCGA-AK-3460	2.605479452	0	0.874782182	Low
TCGA-AK-3461	2.336986301	0	0.932776319	Low
TCGA-AK-3465	1.010958904	0	0.483965927	Low
TCGA-AS-3777	3.391780822	0	2.518490337	High
TCGA-AS-3778	0.117808219	0	1.045315276	High

TCGA-B0-4688	0.276712329	1	1.950716792	High
TCGA-B0-4690	0.117808219	1	1.774276033	High
TCGA-B0-4691	0.380821918	1	2.733786299	High
TCGA-B0-4693	0.210958904	1	0.909280868	Low
TCGA-B0-4694	0.290410959	1	0.758593519	Low
TCGA-B0-4696	2.37260274	1	2.971074624	High
TCGA-B0-4697	1.583561644	1	1.622853186	High
TCGA-B0-4698	0.115068493	1	2.627258013	High
TCGA-B0-4699	0.301369863	1	0.882765155	Low
TCGA-B0-4700	5.424657534	1	0.723335699	Low
TCGA-B0-4701	0.652054795	1	1.320705437	High
TCGA-B0-4703	0.498630137	1	2.397685098	High
TCGA-B0-4706	0.178082192	1	1.723072305	High
TCGA-B0-4707	1.643835616	1	3.934344805	High
TCGA-B0-4710	0.263013699	0	0.992723786	High
TCGA-B0-4712	3.663013699	1	1.377807672	High
TCGA-B0-4713	0.553424658	1	4.003203093	High
TCGA-B0-4714	0.271232877	1	1.265590193	High
TCGA-B0-4718	1.687671233	0	1.669344011	High
TCGA-B0-4810	1.309589041	1	1.529393679	High
TCGA-B0-4811	3.882191781	1	1.86761589	High
TCGA-B0-4813	0.049315068	1	6.121024641	High
TCGA-B0-4814	0.460273973	1	1.121453704	High
TCGA-B0-4815	4.350684932	1	2.147786365	High
TCGA-B0-4816	3.756164384	1	0.879074285	Low
TCGA-B0-4817	2.791780822	1	1.893114274	High
TCGA-B0-4818	1.397260274	1	0.844131133	Low
TCGA-B0-4819	0.501369863	1	2.190825568	High
TCGA-B0-4821	3.369863014	1	3.110178471	High
TCGA-B0-4822	3.043835616	1	2.335817767	High
TCGA-B0-4823	1.243835616	1	0.439244591	Low
TCGA-B0-4824	4.539726027	1	1.037412518	High
TCGA-B0-4827	2.424657534	1	1.067042155	High
TCGA-B0-4828	0.84109589	1	1.694983701	High
TCGA-B0-4833	6.536986301	1	1.298035285	High
TCGA-B0-4834	5.726027397	1	0.210822763	Low
TCGA-B0-4836	3.391780822	1	1.469444758	High
TCGA-B0-4837	3.775342466	1	1.376566569	High
TCGA-B0-4838	2.284931507	1	1.231795351	High
TCGA-B0-4839	4.490410959	1	0.624786601	Low

TCGA-B0-4841	0.55890411	1	0.96949104	Low
TCGA-B0-4842	4.723287671	1	2.903916955	High
TCGA-B0-4843	0.876712329	1	1.456818978	High
TCGA-B0-4844	0.857534247	1	2.047480324	High
TCGA-B0-4845	5.44109589	1	1.153198602	High
TCGA-B0-4846	3.287671233	1	0.7622647	Low
TCGA-B0-4847	2.17260274	1	1.935876603	High
TCGA-B0-4848	2.419178082	1	1.091248002	High
TCGA-B0-4849	0.189041096	1	1.209510094	High
TCGA-B0-4852	3.071232877	1	1.007401732	High
TCGA-B0-4945	5.876712329	1	0.96897609	Low
TCGA-B0-5075	1.745205479	1	1.989565486	High
TCGA-B0-5077	3.608219178	1	1.840728283	High
TCGA-B0-5080	0.936986301	1	1.084536911	High
TCGA-B0-5081	0.991780822	1	1.879777947	High
TCGA-B0-5083	2.863013699	1	1.325625112	High
TCGA-B0-5084	0.608219178	1	1.363348825	High
TCGA-B0-5085	2.109589041	1	1.538612131	High
TCGA-B0-5088	1.542465753	1	1.360169208	High
TCGA-B0-5092	1.257534247	1	2.925078822	High
TCGA-B0-5094	0.912328767	1	4.495789851	High
TCGA-B0-5095	0.671232877	1	1.983999035	High
TCGA-B0-5096	0.18630137	1	0.957524791	Low
TCGA-B0-5097	1.821917808	0	1.156956509	High
TCGA-B0-5098	4.339726027	1	2.196060944	High
TCGA-B0-5099	1.328767123	1	0.792716468	Low
TCGA-B0-5100	5.24109589	1	1.33275063	High
TCGA-B0-5102	7.57260274	1	1.032371185	High
TCGA-B0-5104	0	1	0.857553045	Low
TCGA-B0-5106	4.378082192	1	1.057629412	High
TCGA-B0-5107	2.539726027	1	4.444674872	High
TCGA-B0-5108	2.495890411	0	0.838720947	Low
TCGA-B0-5109	1.608219178	1	0.94939733	Low
TCGA-B0-5110	2.991780822	0	0.683504926	Low
TCGA-B0-5113	0.983561644	0	1.014573532	High
TCGA-B0-5115	2.183561644	0	1.07931099	High
TCGA-B0-5116	1.8	0	1.084546099	High
TCGA-B0-5117	1.465753425	0	0.794631789	Low
TCGA-B0-5119	0.161643836	0	0.491989055	Low
TCGA-B0-5120	1.350684932	0	0.691203686	Low

TCGA-B0-5121	1.517808219	0	0.818848851	Low
TCGA-B0-5399	1.78630137	0	0.890908025	Low
TCGA-B0-5400	3.101369863	0	2.056708449	High
TCGA-B0-5402	1.230136986	0	0.427835442	Low
TCGA-B0-5690	6.597260274	0	1.129545108	High
TCGA-B0-5691	9.4	0	0.681947377	Low
TCGA-B0-5692	4.073972603	0	0.692501702	Low
TCGA-B0-5693	8.42739726	0	0.526696972	Low
TCGA-B0-5694	1.315068493	1	0.475895522	Low
TCGA-B0-5695	3.890410959	0	0.745976791	Low
TCGA-B0-5696	4.731506849	0	1.034336457	High
TCGA-B0-5697	5.02739726	0	0.894477343	Low
TCGA-B0-5698	7.076712329	0	0.516847931	Low
TCGA-B0-5699	7.509589041	0	0.850891095	Low
TCGA-B0-5700	2.964383562	0	0.552040844	Low
TCGA-B0-5701	4.745205479	0	0.917672647	Low
TCGA-B0-5702	4.397260274	0	1.858030741	High
TCGA-B0-5703	3.295890411	0	1.103902241	High
TCGA-B0-5705	10.04931507	0	1.225877225	High
TCGA-B0-5706	6.61369863	0	1.414744305	High
TCGA-B0-5707	7.747945205	0	1.680718426	High
TCGA-B0-5709	8.539726027	0	0.718880805	Low
TCGA-B0-5710	3.997260274	0	0.615644326	Low
TCGA-B0-5711	8.030136986	0	1.460657023	High
TCGA-B0-5712	7.457534247	0	0.709310411	Low
TCGA-B0-5713	5.109589041	0	0.548202086	Low
TCGA-B0-5812	8.117808219	0	0.426216118	Low
TCGA-B2-3923	0.991780822	0	0.243019396	Low
TCGA-B2-3924	1.016438356	0	0.676832162	Low
TCGA-B2-4098	0.139726027	1	1.589095231	High
TCGA-B2-4099	1.024657534	0	0.692766331	Low
TCGA-B2-4101	0.515068493	0	1.353670572	High
TCGA-B2-4102	0.553424658	0	0.958956778	Low
TCGA-B2-5633	0.980821918	0	1.018227852	High
TCGA-B2-5635	0.863013699	0	0.963813083	Low
TCGA-B2-5636	0.726027397	0	0.750701447	Low
TCGA-B2-5639	1.142465753	0	1.306326689	High
TCGA-B2-5641	0.887671233	0	1.109136516	High
TCGA-B2-A4SR	1.389041096	0	0.931497568	Low
TCGA-B4-5377	1	0	0.589005701	Low

TCGA-B4-5378	0.479452055	0	0.592618708	Low
TCGA-B4-5832	0.424657534	0	0.980765568	High
TCGA-B4-5834	0.104109589	0	0.543238972	Low
TCGA-B4-5835	0.043835616	0	2.126455679	High
TCGA-B4-5836	0.38630137	0	0.666369236	Low
TCGA-B4-5838	0.454794521	0	0.462869124	Low
TCGA-B4-5843	0.030136986	0	0.510024118	Low
TCGA-B4-5844	0.019178082	0	0.378678921	Low
TCGA-B8-4143	1.942465753	1	4.073333876	High
TCGA-B8-4146	1.4	0	0.410591469	Low
TCGA-B8-4148	1.038356164	0	0.874312218	Low
TCGA-B8-4151	0.767123288	0	0.889697803	Low
TCGA-B8-4153	1.109589041	0	0.71249668	Low
TCGA-B8-4154	0.698630137	0	0.501937913	Low
TCGA-B8-4619	0.052054795	0	0.83737528	Low
TCGA-B8-4620	0.619178082	0	0.635778308	Low
TCGA-B8-4621	1.180821918	0	1.053162954	High
TCGA-B8-4622	0.495890411	0	0.434001216	Low
TCGA-B8-5158	0.802739726	0	1.778606976	High
TCGA-B8-5159	0.657534247	0	0.345754253	Low
TCGA-B8-5162	0.060273973	0	0.316767525	Low
TCGA-B8-5163	0.010958904	0	1.153702246	High
TCGA-B8-5164	0.063013699	0	0.521112348	Low
TCGA-B8-5165	0.021917808	0	0.530280043	Low
TCGA-B8-5545	1.430136986	0	0.434395864	Low
TCGA-B8-5546	1.383561644	0	0.618689779	Low
TCGA-B8-5549	0.531506849	0	1.419672689	High
TCGA-B8-5550	1.189041096	0	1.004152384	High
TCGA-B8-5551	0.04109589	0	1.104156916	High
TCGA-B8-5552	1.073972603	0	0.358848488	Low
TCGA-B8-5553	1.191780822	0	0.590967886	Low
TCGA-B8-A54D	2.273972603	0	1.847536662	High
TCGA-B8-A54E	2.490410959	0	0.977955855	Low
TCGA-B8-A54F	1.421917808	0	1.352204966	High
TCGA-B8-A54G	0.145205479	0	1.269337124	High
TCGA-B8-A54H	0.701369863	0	0.879183719	Low
TCGA-B8-A54I	0.410958904	0	1.262830587	High
TCGA-B8-A54J	1.446575342	0	0.645220427	Low
TCGA-B8-A54K	1.284931507	0	0.748631232	Low
TCGA-B8-A7U6	1.356164384	0	0.953696318	Low

TCGA-B8-A8YJ	1.180821918	0	0.837312471	Low
TCGA-BP-4158	9.252054795	0	0.836488325	Low
TCGA-BP-4159	7.126027397	1	0.484614908	Low
TCGA-BP-4160	7.893150685	0	0.565610304	Low
TCGA-BP-4161	7.523287671	0	1.189303941	High
TCGA-BP-4162	8.421917808	0	1.609962413	High
TCGA-BP-4163	7.778082192	0	1.651202486	High
TCGA-BP-4164	2.717808219	1	0.94852864	Low
TCGA-BP-4165	8.320547945	0	0.929360363	Low
TCGA-BP-4166	0.035616438	0	0.357492314	Low
TCGA-BP-4167	7.446575342	0	2.316709967	High
TCGA-BP-4169	1.920547945	1	1.01797489	High
TCGA-BP-4170	6.419178082	1	0.494215503	Low
TCGA-BP-4173	5.18630137	0	0.980353006	Low
TCGA-BP-4174	5.147945205	0	0.750260768	Low
TCGA-BP-4176	5.356164384	0	1.535762673	High
TCGA-BP-4177	4.575342466	0	0.42702663	Low
TCGA-BP-4325	8.120547945	0	1.093191895	High
TCGA-BP-4326	4.452054795	1	2.32979845	High
TCGA-BP-4327	0.298630137	1	1.531308703	High
TCGA-BP-4329	2.315068493	1	0.744240617	Low
TCGA-BP-4330	5.17260274	0	0.725985882	Low
TCGA-BP-4331	6.723287671	1	1.061776802	High
TCGA-BP-4332	3.104109589	0	1.014065338	High
TCGA-BP-4334	1.767123288	1	0.937229814	Low
TCGA-BP-4335	1.301369863	1	1.263505843	High
TCGA-BP-4337	0.005479452	1	2.590328143	High
TCGA-BP-4338	7.832876712	0	2.259172908	High
TCGA-BP-4340	1.539726027	1	0.995143841	High
TCGA-BP-4341	4.353424658	1	1.367033136	High
TCGA-BP-4342	6.180821918	1	1.135090459	High
TCGA-BP-4343	5.238356164	1	0.67797181	Low
TCGA-BP-4344	4.564383562	0	0.636743243	Low
TCGA-BP-4345	4.153424658	0	0.726965621	Low
TCGA-BP-4346	4.090410959	1	1.222832735	High
TCGA-BP-4347	3.745205479	0	0.703391389	Low
TCGA-BP-4349	1.019178082	0	1.091401379	High
TCGA-BP-4351	2.657534247	0	1.672220153	High
TCGA-BP-4352	0.942465753	1	8.451159071	High
TCGA-BP-4353	1.02739726	1	0.918873176	Low

TCGA-BP-4354	2.832876712	1	1.846046261	High
TCGA-BP-4355	2.610958904	1	1.08505563	High
TCGA-BP-4756	1.024657534	0	0.926205864	Low
TCGA-BP-4758	6.049315068	0	0.783939121	Low
TCGA-BP-4759	6.498630137	0	0.52715476	Low
TCGA-BP-4760	6.468493151	0	0.477219205	Low
TCGA-BP-4761	0.498630137	0	3.036724199	High
TCGA-BP-4762	3.679452055	1	0.378475325	Low
TCGA-BP-4763	3.479452055	1	1.039573706	High
TCGA-BP-4765	5.983561644	0	0.880595478	Low
TCGA-BP-4766	4.005479452	0	0.800167821	Low
TCGA-BP-4768	1.095890411	0	3.037293059	High
TCGA-BP-4769	5.139726027	0	0.526111503	Low
TCGA-BP-4770	0.901369863	1	0.526183222	Low
TCGA-BP-4771	0.443835616	1	1.986288988	High
TCGA-BP-4774	5.164383562	0	1.069409902	High
TCGA-BP-4775	5.049315068	0	0.836086769	Low
TCGA-BP-4776	1.126027397	0	0.895873751	Low
TCGA-BP-4777	4.742465753	0	1.01259562	High
TCGA-BP-4781	5.698630137	0	0.907560776	Low
TCGA-BP-4782	0.969863014	0	1.636695252	High
TCGA-BP-4784	5.079452055	0	0.348477651	Low
TCGA-BP-4787	1.315068493	1	3.965550418	High
TCGA-BP-4789	4.079452055	0	0.522667225	Low
TCGA-BP-4790	3.043835616	1	0.699148282	Low
TCGA-BP-4795	1.698630137	0	0.471766728	Low
TCGA-BP-4797	3.032876712	0	0.639572799	Low
TCGA-BP-4798	0.915068493	1	2.052960068	High
TCGA-BP-4799	3.104109589	1	1.063728471	High
TCGA-BP-4801	3.079452055	0	0.508401479	Low
TCGA-BP-4803	0.55890411	0	0.642158095	Low
TCGA-BP-4804	3.997260274	0	1.67089813	High
TCGA-BP-4807	0.578082192	0	0.531952624	Low
TCGA-BP-4959	7.287671233	0	0.645325388	Low
TCGA-BP-4960	5.950684932	0	1.444945929	High
TCGA-BP-4961	5.301369863	0	0.618968512	Low
TCGA-BP-4962	4.890410959	0	0.837081135	Low
TCGA-BP-4963	5.024657534	0	1.826927589	High
TCGA-BP-4964	5.101369863	0	1.275368163	High
TCGA-BP-4965	5.126027397	0	0.874718202	Low

TCGA-BP-4967	0.561643836	0	1.760027178	High
TCGA-BP-4968	4.783561644	0	1.115512999	High
TCGA-BP-4969	4.915068493	0	1.251531816	High
TCGA-BP-4970	1.18630137	0	0.907985571	Low
TCGA-BP-4971	4.073972603	0	1.706185108	High
TCGA-BP-4972	4.115068493	0	0.569109284	Low
TCGA-BP-4973	3.791780822	0	1.020488384	High
TCGA-BP-4974	0.578082192	1	0.949490532	Low
TCGA-BP-4975	3.926027397	0	0.667375149	Low
TCGA-BP-4976	4.471232877	0	0.382815204	Low
TCGA-BP-4977	1.243835616	0	1.639362256	High
TCGA-BP-4981	3.005479452	1	1.333439966	High
TCGA-BP-4982	2.778082192	0	0.673190774	Low
TCGA-BP-4983	3.871232877	0	2.27843151	High
TCGA-BP-4985	2.608219178	1	0.637139388	Low
TCGA-BP-4986	2.150684932	0	2.654042159	High
TCGA-BP-4987	3.079452055	0	0.833252155	Low
TCGA-BP-4989	0.323287671	0	2.44629168	High
TCGA-BP-4991	3.871232877	0	0.809220395	Low
TCGA-BP-4992	1.37260274	0	2.861854989	High
TCGA-BP-4993	0.484931507	0	0.620696274	Low
TCGA-BP-4994	3.583561644	0	0.948280989	Low
TCGA-BP-4995	3.756164384	0	0.990012081	High
TCGA-BP-4998	2.553424658	0	1.006396332	High
TCGA-BP-4999	3.468493151	0	0.663251116	Low
TCGA-BP-5000	1.542465753	0	1.874897392	High
TCGA-BP-5001	3.224657534	0	1.503349178	High
TCGA-BP-5004	3.084931507	0	1.185697665	High
TCGA-BP-5006	2.301369863	0	0.626973105	Low
TCGA-BP-5007	3.123287671	0	1.252614556	High
TCGA-BP-5008	2.934246575	0	0.634644927	Low
TCGA-BP-5009	2.991780822	1	1.235117371	High
TCGA-BP-5010	2.405479452	1	1.571985201	High
TCGA-BP-5168	4.008219178	1	0.404354693	Low
TCGA-BP-5169	0.528767123	0	2.516974064	High
TCGA-BP-5170	6.608219178	0	0.684634849	Low
TCGA-BP-5173	0.169863014	1	0.691297958	Low
TCGA-BP-5174	6.183561644	0	2.069789714	High
TCGA-BP-5175	2.553424658	0	3.76700542	High
TCGA-BP-5176	4.356164384	1	0.612330196	Low

TCGA-BP-5177	0.802739726	0	0.898072296	Low
TCGA-BP-5178	5.238356164	1	1.472851344	High
TCGA-BP-5180	6.2	0	1.244165783	High
TCGA-BP-5181	4.095890411	0	0.41562776	Low
TCGA-BP-5182	3.191780822	0	0.792593909	Low
TCGA-BP-5183	3.536986301	0	1.010332256	High
TCGA-BP-5184	3.104109589	0	0.475270268	Low
TCGA-BP-5185	3.101369863	0	0.984839945	High
TCGA-BP-5186	1.898630137	0	0.701749887	Low
TCGA-BP-5187	1.112328767	0	0.631629914	Low
TCGA-BP-5189	2.252054795	1	0.902530774	Low
TCGA-BP-5190	2.769863014	0	1.063440487	High
TCGA-BP-5191	2.649315068	0	2.405258363	High
TCGA-BP-5192	1.956164384	0	0.503973783	Low
TCGA-BP-5194	1.117808219	0	0.503913161	Low
TCGA-BP-5195	2.052054795	0	0.575683495	Low
TCGA-BP-5196	2.789041096	0	2.249440789	High
TCGA-BP-5198	1.652054795	0	0.839571	Low
TCGA-BP-5199	3.712328767	0	1.179921514	High
TCGA-BP-5200	2.912328767	0	0.545403767	Low
TCGA-BP-5201	2.605479452	0	1.531276428	High
TCGA-BP-5202	0.079452055	0	0.671934284	Low
TCGA-CJ-4634	4.98630137	0	0.83374555	Low
TCGA-CJ-4635	3.879452055	0	1.350416673	High
TCGA-CJ-4636	5.271232877	0	2.066083012	High
TCGA-CJ-4637	6.101369863	1	2.118656148	High
TCGA-CJ-4638	1.180821918	1	3.027615487	High
TCGA-CJ-4639	6.323287671	0	0.563011808	Low
TCGA-CJ-4640	5.473972603	0	0.66299333	Low
TCGA-CJ-4641	4.550684932	1	1.183254151	High
TCGA-CJ-4642	4.460273973	0	1.392126825	High
TCGA-CJ-4643	4.912328767	0	0.766790955	Low
TCGA-CJ-4644	0.920547945	1	0.385608618	Low
TCGA-CJ-4868	1.769863014	1	1.138207768	High
TCGA-CJ-4869	6.997260274	0	1.197708836	High
TCGA-CJ-4870	4.104109589	0	0.550924394	Low
TCGA-CJ-4871	6.638356164	0	0.549574596	Low
TCGA-CJ-4872	3.931506849	0	0.962715284	Low
TCGA-CJ-4873	4.865753425	0	3.126508947	High
TCGA-CJ-4874	6.254794521	0	0.33885542	Low

TCGA-CJ-4875	6.446575342	0	1.660754362	High
TCGA-CJ-4876	5.356164384	0	1.029197603	High
TCGA-CJ-4878	5.989041096	0	1.26157536	High
TCGA-CJ-4881	5.517808219	0	2.933915811	High
TCGA-CJ-4882	5.15890411	0	3.042139148	High
TCGA-CJ-4884	4.819178082	0	2.079602162	High
TCGA-CJ-4885	5.821917808	0	0.775644692	Low
TCGA-CJ-4886	5.347945205	0	0.569996458	Low
TCGA-CJ-4887	2.553424658	1	1.752419325	High
TCGA-CJ-4888	4.293150685	1	1.177758736	High
TCGA-CJ-4889	5.331506849	0	0.654249857	Low
TCGA-CJ-4890	5.712328767	0	1.199799309	High
TCGA-CJ-4891	2.243835616	1	3.23425769	High
TCGA-CJ-4892	4.167123288	0	1.056558555	High
TCGA-CJ-4893	2.054794521	0	0.490879357	Low
TCGA-CJ-4894	2.304109589	1	1.023738149	High
TCGA-CJ-4895	3.287671233	1	1.63068241	High
TCGA-CJ-4897	4.953424658	0	0.568220471	Low
TCGA-CJ-4899	4.18630137	0	0.670805734	Low
TCGA-CJ-4900	4.695890411	1	1.909387582	High
TCGA-CJ-4901	3.97260274	0	0.86771988	Low
TCGA-CJ-4902	4.164383562	0	1.043849819	High
TCGA-CJ-4903	4.273972603	0	1.2114934	High
TCGA-CJ-4904	4.909589041	0	0.31362923	Low
TCGA-CJ-4905	4.098630137	0	0.454345128	Low
TCGA-CJ-4907	4.106849315	0	0.895688107	Low
TCGA-CJ-4908	4.194520548	0	0.805367865	Low
TCGA-CJ-4912	4.539726027	0	3.642309098	High
TCGA-CJ-4916	3.761643836	0	1.206686472	High
TCGA-CJ-4918	0.254794521	1	1.030511684	High
TCGA-CJ-4920	0.380821918	1	1.622635578	High
TCGA-CJ-5671	5.323287671	0	1.088036393	High
TCGA-CJ-5672	5.402739726	1	1.32985047	High
TCGA-CJ-5675	6.657534247	0	0.463468405	Low
TCGA-CJ-5676	7.054794521	0	0.597150586	Low
TCGA-CJ-5677	2.142465753	1	0.629713288	Low
TCGA-CJ-5678	1.57260274	1	0.738923862	Low
TCGA-CJ-5679	1.860273973	1	0.5947735	Low
TCGA-CJ-5680	2.104109589	1	0.538654143	Low
TCGA-CJ-5681	1.512328767	1	0.859286134	Low

TCGA-CJ-5682	5.15890411	0	1.396047024	High
TCGA-CJ-5683	5.175342466	0	0.9425863	Low
TCGA-CJ-5684	6.112328767	0	0.665436854	Low
TCGA-CJ-5686	5.583561644	0	1.040979731	High
TCGA-CJ-5689	0	1	0.846516484	Low
TCGA-CJ-6027	5.082191781	0	1.31178235	High
TCGA-CJ-6028	4.452054795	1	1.214927384	High
TCGA-CJ-6030	6.298630137	1	0.786206036	Low
TCGA-CJ-6031	5.221917808	0	0.946211446	Low
TCGA-CJ-6032	6.980821918	0	0.524659998	Low
TCGA-CJ-6033	0.61369863	1	2.172397533	High
TCGA-CW-5580	5.380821918	1	1.505187607	High
TCGA-CW-5581	7.668493151	0	0.774762879	Low
TCGA-CW-5583	6.819178082	0	0.596565805	Low
TCGA-CW-5584	0.449315068	1	0.347135509	Low
TCGA-CW-5585	7.147945205	0	0.724404309	Low
TCGA-CW-5587	6.098630137	0	0.82943955	Low
TCGA-CW-5588	5.526027397	0	0.988965253	High
TCGA-CW-5589	6.515068493	0	0.960373823	Low
TCGA-CW-5590	2.945205479	1	1.095304221	High
TCGA-CW-5591	6.221917808	0	0.914716836	Low
TCGA-CW-6087	0.112328767	1	1.082851812	High
TCGA-CW-6088	8.82739726	0	1.496133124	High
TCGA-CW-6090	6.991780822	0	0.778823542	Low
TCGA-CW-6093	8.619178082	0	0.554649838	Low
TCGA-CW-6097	1.564383562	1	1.315140306	High
TCGA-CZ-4853	2.120547945	0	0.771643193	Low
TCGA-CZ-4854	3.846575342	1	2.328010703	High
TCGA-CZ-4856	0.049315068	0	0.527714299	Low
TCGA-CZ-4857	3.923287671	1	2.21698822	High
TCGA-CZ-4858	5.323287671	0	1.568088507	High
TCGA-CZ-4859	4.895890411	0	0.349142806	Low
TCGA-CZ-4860	0.564383562	1	1.370428472	High
TCGA-CZ-4861	1.221917808	1	0.983364848	High
TCGA-CZ-4862	5.049315068	0	1.145056402	High
TCGA-CZ-4863	4.632876712	0	0.692377192	Low
TCGA-CZ-4864	3.602739726	1	0.728942964	Low
TCGA-CZ-4865	0.454794521	1	1.498967112	High
TCGA-CZ-4866	4.843835616	0	1.205385627	High
TCGA-CZ-5451	4.569863014	0	0.764047339	Low

TCGA-CZ-5452	4.263013699	0	1.167535231	High
TCGA-CZ-5453	0.068493151	0	1.4727069	High
TCGA-CZ-5454	1.978082192	1	1.400538622	High
TCGA-CZ-5455	1.536986301	1	1.564995596	High
TCGA-CZ-5456	4.268493151	0	1.477386793	High
TCGA-CZ-5457	4.238356164	0	1.447436582	High
TCGA-CZ-5458	4.268493151	0	1.047386252	High
TCGA-CZ-5459	4.150684932	0	0.798512894	Low
TCGA-CZ-5460	3.917808219	0	0.874458203	Low
TCGA-CZ-5461	0.904109589	1	1.306246527	High
TCGA-CZ-5462	0.852054795	1	1.602029961	High
TCGA-CZ-5463	1.81369863	0	0.828537521	Low
TCGA-CZ-5464	4.087671233	0	0.845654808	Low
TCGA-CZ-5465	3.77260274	0	1.113494057	High
TCGA-CZ-5466	1.876712329	0	0.487105775	Low
TCGA-CZ-5467	0.2	1	1.357331505	High
TCGA-CZ-5468	0.161643836	1	0.82247415	Low
TCGA-CZ-5469	2.591780822	1	1.290849528	High
TCGA-CZ-5470	0	0	1.782546744	High
TCGA-CZ-5982	5.594520548	0	0.514155748	Low
TCGA-CZ-5984	4.084931507	0	0.557871088	Low
TCGA-CZ-5985	4.463013699	0	1.007373809	High
TCGA-CZ-5986	1.021917808	0	0.967794784	Low
TCGA-CZ-5987	1.219178082	1	0.61896647	Low
TCGA-CZ-5988	1.898630137	0	0.859561988	Low
TCGA-CZ-5989	4.380821918	0	1.34512295	High
TCGA-DV-5565	3.64109589	0	0.551504711	Low
TCGA-DV-5566	3.830136986	0	0.618524267	Low
TCGA-DV-5567	2.493150685	0	0.410877948	Low
TCGA-DV-5568	1.01369863	0	1.24612501	High
TCGA-DV-5569	0.97260274	0	0.694469108	Low
TCGA-DV-5573	3.095890411	0	0.81128023	Low
TCGA-DV-5574	1.980821918	0	0.500749652	Low
TCGA-DV-5575	2.756164384	0	0.621442976	Low
TCGA-DV-5576	1.991780822	1	0.582043038	Low
TCGA-DV-A4VX	4.454794521	1	2.636558185	High
TCGA-DV-A4VZ	1	0	1.27777777	High
TCGA-DV-A4W0	5.501369863	0	0.755144229	Low
TCGA-EU-5904	1.509589041	0	0.457187704	Low
TCGA-EU-5905	0.326027397	0	1.542657675	High

TCGA-EU-5906	0.564383562	0	0.434516804	Low
TCGA-EU-5907	0.347945205	0	0.658634005	Low
TCGA-G6-A5PC	0.663013699	1	2.310970348	High
TCGA-G6-A8L6	0.857534247	1	1.044233987	High
TCGA-G6-A8L7	5.843835616	0	1.995293949	High
TCGA-G6-A8L8	2.989041096	1	1.273315239	High
TCGA-GK-A6C7	0.167123288	0	0.682813148	Low
TCGA-MM-A563	1.619178082	0	1.140053384	High
TCGA-MM-A564	1.663013699	0	1.088704785	High
TCGA-MM-A84U	1.917808219	0	2.27993673	High
TCGA-MW-A4EC	1.364383562	0	0.792086937	Low
TCGA-T7-A92I	0.975342466	0	1.223088296	High