

id	futime	fustat	AL136084.3	AL731567.1	AC006160.1	AC012568.1	AC034236.2
riskScore	risk						
TCGA-ZF-AA51	4.6958904109589	0	0.429602124	2.517325863	0.083820148		
	0.066594397	0.507606484	0.718707588	702786	high		
TCGA-BT-A42C	2.39178082191781	0	0.1415353	2.2526940	0.04142267		
	0.03290996	0.7814992	0.661283220	147806	high		
TCGA-DK-A3IT	1.77534246575342	0	0.7856667	0.9539293	0.1532923		
	0.04059648	0.5950788	0.734896798	418092	high		
TCGA-DK-A3IV	0.805479452054795	1	0.1753007	5.0971520	2.052187		
	0.0326089	1.3383830	3.978213002	30122	low		
TCGA-2F-A9KW	0.695890410958904	1	5.6066411	11.354620	0.09944709	0	
	0.9574099	0.68047816154325	high				
TCGA-GU-A762	0.635616438356164	1	0.374333255	0.400159819	0.082166149	0	
	0.625177224	0.757345493557312	high				
TCGA-XF-AAME	7.74794520547945	1	2.0436241	1.1327660	0.1246043	0	
	0.7545911	0.813978747341553	high				
TCGA-XF-AAN0	4.70684931506849	1	0.177699203	3.478365734	0.166421227	0	
	1.143499887	0.479678355006248	low				
TCGA-DK-A1A7	1.53424657534247	0	0.293680625	0.383707722	0.021487632		
	0.034143484	1.811767764	0.529259198	278955	low		
TCGA-FD-A3SR	1.64931506849315	1	0.2754449	8.2445680	0.1074847	0	
	0.7510579	0.455183942161924	low				
TCGA-GU-AATP	2.74794520547945	0	0.2027134	9.439460	0.1186547	0.06284677	
	1.2897250	4.491652565566	low				
TCGA-DK-AA6X	1.27945205479452	0	0.5220974	7.1935310	0.6621355	0	
	0.9253447	0.270861304324913	low				
TCGA-XF-A9T3	0.186301369863014	0	1.7451440	0.2211019	0.06809939	0	
	1.7130590	0.622688176831978	high				
TCGA-DK-A3WW	1.73424657534247	0	0.0885820	0.6312905	0.3370250		
	0.7970729	0.529717422723838	low				
TCGA-UY-A9PA	2.93698630136986	0	0.06564991	3.8052790	0.2305625		
	0.06105998	0.5907269	0.538597731569975	low			
TCGA-G2-AA3F	2.44657534246575	0	0.3181678	7.9901050	0.05320978	0	
	1.5615910	0.367817406075917	low				
TCGA-FJ-A871	0.745205479452055	1	0.1270220	0.08046562	0.1239171	0	
	1.1660520	0.593657697545079	high				
TCGA-GC-A3RB	1.59452054794521	0	0.0690156	2.4920310	0.08079426	0	
	0.6304211	0.651119068726288	high				
TCGA-YC-A89H	1.56986301369863	0	0.3775560	0.2242249	0.1381226	0	
	1.0294850	0.627982342415597	high				
TCGA-E7-A3X6	2.47671232876712	1	0.313302421	1.275879063	0.02619805	0	
	0.61020303	0.761246947620839	high				
TCGA-DK-A6AW	4.44109589041096	0	0.2128533	5.3598020	0.09344258	0	
	1.9152830	0.35585788626034	low				

TCGA-GC-A3BM	1.78356164383562	1	0.04469052	2.4630090.2092708				
	0.04156597	1.2795090.460901716042963	low					
TCGA-GV-A6ZA	1.89315068493151	0	0.25201	2.7777830.2065138	0.1875125			
	1.5942060.432467850572748	low						
TCGA-UY-A8OD	9.4027397260274	0	0.8751733	5.8212290.2241173	0.4069927			
	1.0887690.48975645391547	low						
TCGA-2F-A9KT	6.44383561643836	0	0.6506223	2.2256350.03808309				
	0.1210268	1.3128020.598784018681502	high					
TCGA-BT-A20Q	1.62465753424658	1	0.2955651	0.5148933	0.3171742			
	0.09163354	0.5104159	0.630786586647795	high				
TCGA-4Z-AA7Q	1.3972602739726	1	0.246401069	0.643869459	0.144226759			
	0.057293469	0.772643351	0.667042243911873	high				
TCGA-CF-A27C	1.16438356164384	0	0.03812694	8.5500080.2231697				
	0.2482291.7361490.292343967910717	low						
TCGA-DK-AA77	1.69315068493151	0	0.2074157	4.1881580.2428146	0			
	0.5514230.529215453671437	low						
TCGA-ZF-AA5N	0.46027397260274	1	0.3575789	0.3397773	0	0		
	0.6500088	0.814879385810466	high					
TCGA-XF-A9SZ	2.35342465753425	1	0.4645447	1.60014	0.1699459	0		
	0.6966727	0.641459240376475	high					
TCGA-4Z-AA82	4.26301369863014	1	0.3082251	0.6345746	0.09020722			
	0.04777925	0.5602933	0.762151418884154	high				
TCGA-DK-AA76	1.0027397260274	0	0.2885105	4.7153350.2701997	0			
	0.6922818	0.480979336941529	low					
TCGA-DK-A3IL	1.13150684931507	1	0.1423685	6.0425460.1666660.04413825				
	1.7210080.342581932723559	low						
TCGA-ZF-A9R1	2.11780821917808	0	0.05150933	3.3282610.1206005				
	0.04790804	0.9410211	0.541646506154689	low				
TCGA-C4-A0F0	0.161643835616438	0	0.32752934	0	0	0	0.107169227	
	1.00137678785437	high						
TCGA-FD-A3N5	1.87671232876712	1	0.3840633	0.03041194	0.05620127	0		
	0.2967148	0.890965287293286	high					
TCGA-BT-A2LA	1.43013698630137	0	0.2893862	0.3971927	0.2634915	0		
	2.6477740.307863355938074	low						
TCGA-BT-A2LB	1.34794520547945	1	0.440135003	6.133944956	0.021468807			
	0.102340713	1.120111688	0.510760543660896	low				
TCGA-XF-A8HB	3.75342465753425	0	0.07658528	0.0363863	0.2241396	0		
	0.3445624	0.715898200852212	high					
TCGA-FD-A5BX	0.473972602739726	1	0.6784212	0.1074412	0	0	0.81702	
	0.808537716432847	high						
TCGA-PQ-A6FI	1.01917808219178	0	0.03467674	0	0.0811898	0.0322523		
	0.6713287	0.738005866484698	high					
TCGA-4Z-AA7S	2.91506849315068	1	0.3436348	4.4314440.05746883	0			
	1.1779330.510686799243884	low						

TCGA-4Z-AA86	0.852054794520548	1	1.225670227	0.705849704	0.130441025	0
	0.891211974	0.716033844163389	high			
TCGA-5N-A9KI	0.208219178082192	1	0.588706345	1.333950157	0	0
	0.703831766	0.778546143000379	high			
TCGA-FD-A6TG	0.254794520547945	1	0.7193308	1.7467730.04678314	0	
	0.5666281	0.775008729211867	high			
TCGA-GV-A3QI	3.04109589041096	0	0.2363602	1.7325770.7708044	0	
	4.0878780.104095595245908	low				
TCGA-DK-A6AV	5.34794520547945	0	0.2614097	8.4454560.0765059	0	
	0.2851152	0.545954362147736	low			
TCGA-DK-A1A3	1.82191780821918	1	0.7957413	0.4536759	0.03105159	0
	0.9836219	0.735972833854692	high			
TCGA-GC-A3RD	1.17260273972603	0	0.2809227	1.4236660.1370279		
	0.04354699	1.1489910.56425178355351	low			
TCGA-BT-A20R	0.421917808219178	1	1.1479240.3231931	0.0746576		
	0.03954322	0.6607893	0.838161273851625	high		
TCGA-XF-A9T0	2.18904109589041	0	0.3458225	1.5334960.3036321	0	
	2.1845170.329666960393473	low				
TCGA-4Z-AA7Y	4.16986301369863	0	0.05125684	4.4321640.09000702		
	0.04767321	0.4192874	0.631447779722039	high		
TCGA-BL-A3JM	0.561643835616438	1	1.0906060.06095949	0.1126531	0	
	1.0495630.702400709663092	high				
TCGA-FJ-A3ZF	1.43561643835616	0	0.1833060.5515712	0.1788252		
	0.02841502	1.6160910.474431989784243	low			
TCGA-CF-A5UA	1	0	0	1.0348090.08314465	0	3.2147540.28082352150536
	low					
TCGA-FD-A5C0	1.50684931506849	1	0.2789463	3.5783010.05442552	0	
	0.3929793	0.702023106927056	high			
TCGA-FD-A5BS	4.49041095890411	0	0.60815076	1.73362456	0.15255896	0
	0.66329983	0.666759341805199	high			
TCGA-FD-A5BZ	2.28767123287671	1	0.4114713	1.6756580.2408479		
	0.05467188	0.7693454	0.585163146191091	high		
TCGA-UY-A9PE	0.517808219178082	0	0.4306632	3.2737860.1512489	0	
	1.8084110.401523872801239	low				
TCGA-GV-A40E	0.715068493150685	1	0.1701137	1.5760380.02489331	0	
	0.8117384	0.686631364201497	high			
TCGA-BT-A2LD	1.70684931506849	1	0.1522144	0	0	0.4565485
	0.867121868119438	high				
TCGA-G2-AA3B	5.5013698630137	0	0	3.4562950.1860359	0	1.973983
	0.344270990725829	low				
TCGA-CU-A5W6	0.153424657534247	1	0.1463046	1.6219120.08563695		
	0.3628684	1.7419940.508201646369691	low			
TCGA-DK-AA6R	13.8109589041096	0	0.2037009	0.5322899	0.2086577	0
	0.2221732	0.749903780398246	high			

TCGA-BL-A13I	0.610958904109589	1	2.09453333333333	1.05985753333333	
1.4731587	9.8443550.5542824	2.8181417152013	high		
TCGA-CF-A47T	1.05479452054795	1	0.1004256	4.0556080.1175649	0
2.1495760.339036743840254	low				
TCGA-DK-A3IS	4.18904109589041	0	0.1038549	2.0723780.1013163	0
1.1704860.54209989148118	low				
TCGA-XF-A9SJ	0.268493150684932	1	0.6621481.8875520	0	0.6388637
0.779660865372619	high				
TCGA-DK-AA6T	1.56712328767123	0	0.1644315	2.1353560.09624725	0
1.3600150.511572683033715	low				
TCGA-BT-A20N	2.17808219178082	1	0.06021935	2.803849412	0.035248393 0
0.426921525	0.718375494624064	high			
TCGA-CF-A7I0	1.00821917808219	0	0.5840422	13.369650.03107812	0
1.6649	0.278964553521669	low			
TCGA-GC-A3YS	2.07671232876712	0	0.1923035	0.5938723	0.3376849 0
1.1798	0.469357340613281	low			
TCGA-XF-A9T2	1.57534246575342	1	0.8752303	0.1848129	0.2846119 0
1.1932490.545994467990134	low				
TCGA-UY-A78O	6.33424657534247	0	0.1541814	3.1742910.2105776	
0.0956012	1.63959	0.40061962555441	low		
TCGA-GC-A3OO	1.31780821917808	0	0.9694677	1.7963480.02837309	0
0.5419084	0.817962058074444	high			
TCGA-XF-AAN5	6.28219178082192	0	0.2057931	0.5866441	0 0.09570253
1.2064460.664928895646788	high				
TCGA-FD-A3SN	2.43013698630137	0	0.276871134	0.789262519	0.054020635 0
0.440385608	0.804035498371061	high			
TCGA-E7-A4IJ	1.84657534246575	1	0.1772954	9.1534850.06918462	0
1.2085820.378373546854732	low				
TCGA-XF-AAN3	7.19178082191781	0	0.1500121.1759870.1536623	0	
0.7413833	0.632437049511428	high			
TCGA-FD-A3B8	1.05205479452055	0	4.0601420.7643235	0.2914620	
0.5326528	0.969871529024179	high			
TCGA-XF-A9ST	0.350684931506849	1	0.2327826	0.2764923	0.06812771 0
0.5712572	0.774667901680331	high			
TCGA-DK-A6B5	4.22465753424658	0	0.1123323	2.5261790.1972555	
0.03482618	1.09246	0.499580859040383	low		
TCGA-GV-A3QK	2.27945205479452	0	0.4026494	13.391140.2946051	
0.03120817	0.7593859	0.291849697424195	low		
TCGA-XF-AAMR	7.64383561643836	0	2.2394062.0039640.1908930	1.114531	
0.655860600561125	high				
TCGA-DK-A3IQ	1.47671232876712	1	1.4153332.0365220.04733953	0.03761085	
0.6395246	0.81521523401679	high			
TCGA-E7-A4XJ	0.186301369863014	1	0.2602329	4.8837310	0 1.898124
0.405026908116262	low				

TCGA-4Z-AA7M	1.35616438356164	0	0.05858858	7.29301	0.3772323		
	0.05449235	1.59754	0.269782806342414	low			
TCGA-C4-A0EZ	0.747945205479452	1	0.3445695	1.0231740	0.2016880		
	3.0535070	0.275861446478909	low				
TCGA-FD-A62P	0.523287671232877	1	0.6332648	1.3128840	0	0.6906913	
	0.787233991517779	high					
TCGA-CF-A47V	1.03835616438356	0	0.2137792	11.578780	0.2252378	0	
	1.5388910	0.254260518317085	low				
TCGA-XF-A9SI	6.63835616438356	0	0.9388348	1.8833140	0.1221181	0	
	0.9528623	0.640133744254448	high				
TCGA-GU-A42Q	0.942465753424658	1	0.1180760	0.0747985	0.1382276		
	0.03660689	0.7941649	0.672954808454003	high			
TCGA-KQ-A41N	4.39452054794521	0	0.1577661	0	0.1231277	0.04891195	
	1.6490320	0.511542483512615	low				
TCGA-GD-A3OQ	0.26027397260274	0	0.2024366	0.6411952	0.1184929	0	
	0.7175811	0.683986217690773	high				
TCGA-BT-A20U	1.24657534246575	1	0	0.2747121	0.08461133	0	
	0.2956141	0.81697314923374	high				
TCGA-XF-A9SW	0.991780821917808	1	4.12878601	0.84069437	0.34524515	0	
	0.83094719	0.83160624303636	high				
TCGA-FD-A3B3	2.66849315068493	1	0.9339805	0.3882741	0.0854203	0	
	0.6764651	0.793173849033195	high				
TCGA-FT-A3EE	0.271232876712329	1	0.3094538	1.0291680	0.04528341		
	0.02398486	0.5554953	0.776303436291715	high			
TCGA-ZF-A9RE	0.29041095890411	1	0.1696018	1.1281090	0.1323648		
	0.05258135	0.5241152	0.709966352775871	high			
TCGA-DK-A6B1	5.61369863013699	0	0.3637027	2.3500550	0.2341761		
	0.03382744	0.9421256	0.528388175812686	low			
TCGA-GU-A767	0.394520547945205	1	0.6712025	4.6011820	0.08418797	0	
	1.0981040	0.527148985484497	low				
TCGA-4Z-AA81	3.47945205479452	1	0	3.79853776	0.033911583	0.053884931	
	0.458122323	0.6782012776746	high				
TCGA-SY-A9G0	2.76164383561644	1	2.3956994	0.9955050	0.1168569	0	
	1.2157470	0.588977818409157	high				
TCGA-GV-A40G	1.58904109589041	0	0.2539045	6.4796680	0.1061563		
	0.0337361	1.17695	0.434552711202644	low			
TCGA-FD-A43Y	1.2986301369863	1	0	1.178669411	0.276594422	0.054937981	
	1.578392081	0.415709312957121	low				
TCGA-BT-A42E	3.03561643835616	0	0.03815633	0.6526219	0.06700251		
	0.07097724	0.6970758	0.722984864467625	high			
TCGA-FD-A43X	0.301369863013699	0	0.1168263	4.0333730	0.2279411		
	0.1086584	3.1748940	0.218619023376753	low			
TCGA-ZF-AA4R	2.83835616438356	1	0	1.395724531	0.074762288	0	
	0.835851666	0.641542053436944	high				

TCGA-BT-A20T	1.24109589041096	1	0.4821607	2.4435040.09407491		
	0.1494834	0.5916201	0.712682779086665	high		
TCGA-XF-AAN8	0.323287671232877	1	1.8288680.7965009	0	0.07087508	
	0.4155657	1.04758324298903	high			
TCGA-KQ-A41P	2.9972602739726	0	0.295293874	3.507413712	0.086422674	0
	0.993055569	0.553955859378021	low			
TCGA-UY-A9PH	4.27671232876712	0	1.3979744.9228140.3850734	0		
	2.8925350.223517015478403	low				
TCGA-FD-A5BY	0.687671232876712	0	0.42431828	0.3135955	0.0275964	0
	0.38566403	0.878680402205266	high			
TCGA-K4-A3WU	0.287671232876712	0	1.7522483.8103240.2958604	0.06268229		
	2.1500420.356863741534973	low				
TCGA-FD-A3SJ	2.02465753424658	1	0.8939428	0.9101134	0.3924409	0
	0.6529074	0.572292020504036	low			
TCGA-FD-A3B5	0.745205479452055	1	0.2402263	0.03260956	0.08034996	0
	0.87025	0.698795605400625	high			
TCGA-GC-A6I3	0.003	0	0	1.1695860	0	2.6849620.364505305119887
	low					
TCGA-ZF-A9R0	1.86301369863014	1	0.1846451	7.4859870.07205262		
	0.1144904	1.4265080.394304663505043	low			
TCGA-K4-A83P	1.35616438356164	0	1.43042001	1.6475256	0.15223137	0
	1.26465498	0.598184241441121	high			
TCGA-FJ-A3ZE	0.887671232876712	1	0.04036162	11.428980.378	0	
	1.4747280.221061439789339	low				
TCGA-DK-A2I1	1.4958904109589	0	0.9244188	1.26031	0.1176289	0.1121462
	1.4356580.576332123139901	low				
TCGA-DK-A6B0	6.38356164383562	0	0	3.4727310.22387	0.03952507	
	1.9466960.337975579135924	low				
TCGA-FD-A5BR	2.22739726027397	0	0.5475135	5.4626920.3204779		
	0.0509234	0.6270220.469860125537579	low			
TCGA-LT-A5Z6	1.2986301369863	0	0.1709399	8.1756390.3335228	0.1059924	
	1.3361630.302114875473059	low				
TCGA-GC-A4ZW	0.0410958904109589	0	0	0.7740573	0.03178795	0
	0.4738577	0.786390905429094	high			
TCGA-DK-AA74	4.67945205479452	0	0.6718609	0.5586116	0	0
	1.2594780.674539148978428	high				
TCGA-G2-A2EO	4.94246575342466	1	0.3389732	0.9202797	0.02834461	0
	1.3336050.602213552328668	high				
TCGA-UY-A78K	1.46849315068493	1	0.2599608	2.1408320.1521637	0	
	1.0396280.548128304748415	low				
TCGA-CF-A3MG	1.01095890410959	0	0.1074166	4.7972460	0	0.380762
	0.681769920586151	high				
TCGA-DK-A1A6	5.53424657534247	0	0.1837554.7143890.2366274	0.1025446		
	1.2926990.407812574211814	low				
TCGA-XF-AAMG	9.21643835616438	0	0.7591987	0.9275183	0.1904504	0

2.5137090.356283667864571 low

TCGA-XF-A8HE 10.4575342465753 0 0.286128536 1.042222161 0.16748057
0.044353964 0.455110244 0.714485995259846 high

TCGA-K4-A4AC 0.761643835616438 1 0.1621284 0.3081142 0.06326612
0.0502644 0.5894359 0.775364827025879 high

TCGA-FD-A3SL 1.95068493150685 1 1.3925640.2205395 0.0815114
0.04317341 1.3163330.68498358043671 high

TCGA-2F-A9KO 2.01095890410959 1 0.2393212.1035120.7354334 0.1112944
0.8972679 0.334174330001603 low

TCGA-DK-A3IM 0.679452054794521 1 0.03403159 0.09701212 0.09959911 0
1.0485740.626115546606368 high

TCGA-4Z-AA7N 3.74520547945205 1 1.2210046.8720620.3848355 0.1747135
0.6402543 0.454566601441347 low

TCGA-UY-A8OC 0.003 1 0.7375504 0.5005947 0 0 0.4740414
0.899279650220717 high

TCGA-XF-A9T8 1.14520547945205 1 1.0353560.5988422 0.1317453 0
0.7610131 0.736128711610721 high

TCGA-KQ-A41S 0.0958904109589041 0 1.452892997 2.384608517 0.103082024 0
0.576234919 0.770470954885053 high

TCGA-E7-A7XN 1.17260273972603 0 0.2977497 0.2829267 0.6099899 0
0.8389391 0.417214819130064 low

TCGA-DK-A2HX 3.89041095890411 1 0.9889884 0.7518024 0.09648131 0
0.5483255 0.809995890185174 high

TCGA-CF-A8HY 0.945205479452055 0 0.05512337 0.05237913 0.06453109 0
0.7966183 0.711683027073551 high

TCGA-4Z-AA7W 2.3013698630137 0 0 0.2824805 0 0 0.6160531
0.792683900506312 high

TCGA-BT-A0S7 0.547945205479452 1 1.0268690.1393926 0 0 0.4799954
0.947435455033982 high

TCGA-ZF-A9R3 2.6 1 0.3767636 14.439610.3920574 0.1946789 0.9245919
0.245512663927465 low

TCGA-4Z-AA89 2.81917808219178 0 0.06009191 6.6236370.0351738 0
1.2780540.432200543666718 low

TCGA-FD-A3NA 5.05479452054795 0 0.802372751 0.831739404 0.170783824
0.033921575 0.477346093 0.758232299067108 high

TCGA-ZF-AA54 1.61643835616438 1 6.69562 0.8938753 0.2267288 0
0.7845984 1.28644653929049 high

TCGA-DK-A1AB 1.39178082191781 1 0.2126145 0.8369807 0.03555725
0.1412497 0.4058164 0.850075114610904 high

TCGA-UY-A9PB 2.46301369863014 0 0.4857915 0.5193080.1421750
0.5795176 0.730542533065174 high

TCGA-XF-AAML 0.635616438356164 1 0.07075487 3.4960870.1242456 0
0.5980764 0.597545382145561 high

TCGA-ZF-AA58 4.51780821917808 0 0.623073864 0.54976539 0.208403372 0

	0.691711504	0.66780272469167	high					
TCGA-K4-A6FZ	0.150684931506849	0	0.768963024	1.913689158	0.214333186			
	0.10217154	0.339471816	0.728487411962227	high				
TCGA-KQ-A41R	3.6986301369863	0	0.2214161.8935390.08640146	0.06864523				
	2.2137020.396986967546678	low						
TCGA-DK-AA6M	4.33424657534247	0	0.1419003	0.7415978	0.1245884	0		
	0.8125332	0.649246271026389	high					
TCGA-HQ-A5ND	0.750684931506849	1	0.049125709	0.186720225	0.143774574			
	0.091382144	0.388459252	0.771679105347788	high				
TCGA-UY-A78P	6.52054794520548	0	0.5922352	0.8441275	0.1155517			
	0.1377072	0.2825992	0.857669796464118	high				
TCGA-ZF-A9RD	1.11780821917808	1	0.3620451	0.2580159	0.05297926	0		
	0.7897530.740123943210056	high						
TCGA-FD-A62S	1.11232876712329	1	0.7028408	0.7464216	0.07259932			
	0.07690606	0.6425716	0.790631544470491	high				
TCGA-E5-A2PC	3.63287671232877	0	0.1559728	3.3494990.01825922				
	0.1160543	1.0121960.600971892277555	high					
TCGA-GD-A76B	0.613698630136986	0	0.36465604	0.6545041	0.11858074	0		
	0.90592742	0.652087449375005	high					
TCGA-XF-A9SK	1.33150684931507	1	0.2954053	4.7718820.1383284				
	0.1099008	1.6270780.40408748974807	low					
TCGA-DK-A1AD	9.36986301369863	0	0.3593105	2.7313820.1051582	0			
	0.4212860.699251462858791	high						
TCGA-FD-A3SQ	3.8986301369863	1	1.53445	1.8440160.2641661	0.08395109			
	1.1198340.577026364866409	low						
TCGA-XF-AAN1	2.57808219178082	1	0.1469384	5.6547420.1720159	0			
	1.9231590.320745728963001	low						
TCGA-G2-A2EL	2.24383561643836	1	0.2704639	0.8352476	0.1978894	0		
	0.6729469	0.6411324432114	high					
TCGA-XF-AAMZ	3.69315068493151	1	0.09048109	3.5250420.0529616	0			
	1.6036510.450125097220473	low						
TCGA-UY-A8OB	5.77808219178082	0	0	0.3618930.03715435	0	0.7615488		
	0.722981553964644	high						
TCGA-CU-A3KJ	1.53972602739726	0	0.08238408	0.4436020.2250367	0			
	1.14565	0.527980901063932	low					
TCGA-DK-AA6S	13.8356164383562	0	0.4740423	1.0698020.1387364	0			
	1.43799	0.525024937296474	low					
TCGA-E7-A8O8	0.0356164383561644	0	0.2482925	6.1814090.02906678	0			
	1.2998810.452057309101351	low						
TCGA-LT-A8JT	1.75616438356164	0	0.08420597	8.6014940.1971542				
	0.1174781.3087490.338202249199795	low						
TCGA-DK-A1AC	10.9068493150685	0	0.1091241	2.5922870.2129132	0			
	0.2578762	0.651627150714644	high					
TCGA-C4-A0F1	0.243835616438356	0	0.7586546	0.08009846	0.02467032	0		

	0.4826803	0.897668879277434	high					
TCGA-4Z-AA83	5.54520547945206	0	0.1324319	6.8582240.2713088	0			
	1.6069090.304022565158099	low						
TCGA-ZF-AA4V	4.94794520547945	0	0.1248848	1.5822340.1949312	0			
	0.9194156	0.556243291176431	low					
TCGA-ZF-A9RM	3.98630136986301	0	0.09374483	11.491050.08230796				
	0.04359532	0.7540636	0.386250054283651	low				
TCGA-E7-A3Y1	0.446575342465753	0	0.2457575	12.189890.1150801	0			
	2.4660010.198450154663189	low						
TCGA-FD-A6TC	0.512328767123288	0	0.3525501	0.5862480.1031797	0			
	0.6128295	0.735668178673947	high					
TCGA-DK-A2I4	10.5068493150685	0	1.0259791.4151810.07748899	0				
	0.7309715	0.749627755744176	high					
TCGA-UY-A78M	1.89041095890411	1	0.1402134.8407890.05471424	0				
	1.4910480.43772054054378	low						
TCGA-4Z-AA7R	1.43013698630137	1	0.6877326	3.8276130.2012764	0			
	1.9288250.366582349887397	low						
TCGA-E7-A519	1.39178082191781	0	0.1445825	1.3738470	0.134474			
	2.6413690.385274456736236	low						
TCGA-BL-A5ZZ	1.03287671232877	0	2.398470651	0.46425424	0.207985899	0		
	0.423884306	0.911756313109545	high					
TCGA-FD-A62N	0.224657534246575	0	0.4623191	0.4881147	0	0		
	0.3921895	0.896259488580041	high					
TCGA-DK-AA71	1.13698630136986	0	0.1859	0.9421079	0.07254231	0		
	1.0813760.618044415194885	high						
TCGA-KQ-A41O	4.21369863013699	0	0.05421813	15.816320.1586784	0			
	0.6504829	0.288931034472181	low					
TCGA-DK-A1A5	0.178082191780822	1	0.6301206	1.6386870.09706068				
	0.06169111	0.4792748	0.76900011591428	high				
TCGA-XF-AAMT	0.246575342465753	1	0.7066927	0.4740078	0.1703268	0		
	1.0314820.623819613539527	high						
TCGA-XF-A9SU	0.498630136986301	1	0.9191511	3.4311850.03842927	0			
	1.7185760.486580278003087	low						
TCGA-FD-A6TK	0.904109589041096	0	0.9093543	0.3142122	0.04838867	0		
	0.7438631	0.804137112349486	high					
TCGA-UY-A78L	3.08767123287671	0	0.280937855	0.953399183	0.023491756	0		
	0.251697384	0.877942358005164	high					
TCGA-S5-A6DX	0.153424657534247	1	0.3742135	0.5333757	0	0		
	2.7720080.38271853659473	low						
TCGA-XF-A8HI	1.49041095890411	1	0.4159055	5.6316030.1825825	0			
	1.5309710.376811539210453	low						
TCGA-XF-A9T5	5.55342465753425	0	0.5497581	0.6268670.1608959	0			
	1.9037680.450746106935621	low						
TCGA-H4-A2HQ	1.61643835616438	0	0	5.4783180.1163670	0.9621964			

0.471407740221611 low
TCGA-XF-A9SP 1.24383561643836 1 1.5310071.6106580.03200533 0
1.1927450.700137682460827 high
TCGA-DK-A3IK 0.4 1 0.2618614 1.8839610.0437932 0.06958665 1.621844
0.512714820835897 low