

Table S1 Risk scores and risk groups in the TCGA database

Id	Risk score	Risk group
TCGA-AA-3524-01	0.11	Low
TCGA-A6-6780-01	0.12	Low
TCGA-CK-4950-01	0.19	Low
TCGA-AA-3681-01	0.22	Low
TCGA-CM-6674-01	0.23	Low
TCGA-A6-2678-01	0.25	Low
TCGA-AG-3578-01	0.25	Low
TCGA-AA-3555-01	0.26	Low
TCGA-AH-6903-01	0.26	Low
TCGA-AA-3494-01	0.27	Low
TCGA-AA-3947-01	0.27	Low
TCGA-CK-4952-01	0.27	Low
TCGA-D5-6920-01	0.27	Low
TCGA-A6-3809-01	0.28	Low
TCGA-CM-6675-01	0.28	Low
TCGA-AG-3731-01	0.29	Low
TCGA-DM-A28K-01	0.29	Low
TCGA-DM-A28M-01	0.31	Low
TCGA-AD-6890-01	0.32	Low
TCGA-AF-4110-01	0.32	Low
TCGA-CI-6622-01	0.32	Low
TCGA-A6-2674-01	0.33	Low
TCGA-A6-5656-01	0.33	Low
TCGA-AA-3986-01	0.33	Low
TCGA-AA-A01I-01	0.33	Low
TCGA-EI-6506-01	0.33	Low
TCGA-F5-6861-01	0.33	Low
TCGA-AA-3560-01	0.34	Low
TCGA-AA-3971-01	0.34	Low
TCGA-AG-3608-01	0.34	Low
TCGA-CI-6623-01	0.34	Low
TCGA-CM-6171-01	0.34	Low
TCGA-G5-6235-01	0.34	Low
TCGA-A6-5661-01	0.35	Low
TCGA-AZ-4682-01	0.35	Low
TCGA-CK-4947-01	0.36	Low
TCGA-AA-3495-01	0.37	Low
TCGA-AA-3519-01	0.37	Low
TCGA-AA-3968-01	0.37	Low
TCGA-AG-3902-01	0.37	Low
TCGA-AG-A00C-01	0.37	Low

TCGA-DC-6160-01	0.37	Low
TCGA-DM-A288-01	0.37	Low
TCGA-G4-6322-01	0.37	Low
TCGA-AA-3660-01	0.38	Low
TCGA-AD-6548-01	0.38	Low
TCGA-AH-6547-01	0.38	Low
TCGA-CM-6172-01	0.38	Low
TCGA-AG-A011-01	0.39	Low
TCGA-AG-A025-01	0.39	Low
TCGA-D5-6531-01	0.4	Low
TCGA-D5-6539-01	0.4	Low
TCGA-A6-4105-01	0.41	Low
TCGA-AA-3492-01	0.41	Low
TCGA-AA-3549-01	0.41	Low
TCGA-AA-3819-01	0.41	Low
TCGA-AG-3742-01	0.41	Low
TCGA-CI-6624-01	0.41	Low
TCGA-F4-6806-01	0.42	Low
TCGA-AA-3955-01	0.43	Low
TCGA-AF-2693-01	0.43	Low
TCGA-AG-4008-01	0.43	Low
TCGA-AY-A8YK-01	0.43	Low
TCGA-DC-4745-01	0.43	Low
TCGA-EI-6508-01	0.43	Low
TCGA-EI-6512-01	0.43	Low
TCGA-AA-3975-01	0.44	Low
TCGA-AA-A00J-01	0.44	Low
TCGA-AG-3592-01	0.44	Low
TCGA-AG-3612-01	0.44	Low
TCGA-CI-6620-01	0.44	Low
TCGA-AA-3688-01	0.45	Low
TCGA-AA-3875-01	0.46	Low
TCGA-AA-3980-01	0.46	Low
TCGA-G4-6293-01	0.46	Low
TCGA-G4-6309-01	0.46	Low
TCGA-AA-3509-01	0.47	Low
TCGA-AA-3950-01	0.47	Low
TCGA-AA-3982-01	0.47	Low
TCGA-AH-6897-01	0.47	Low
TCGA-AG-4022-01	0.48	Low
TCGA-AH-6644-01	0.48	Low
TCGA-CK-6747-01	0.48	Low
TCGA-D5-6931-01	0.48	Low
TCGA-F4-6854-01	0.48	Low

TCGA-G4-6307-01	0.48	Low
TCGA-A6-5657-01	0.49	Low
TCGA-A6-6650-01	0.49	Low
TCGA-AA-3520-01	0.49	Low
TCGA-AD-6889-01	0.49	Low
TCGA-AG-3586-01	0.49	Low
TCGA-NH-A8F7-01	0.49	Low
TCGA-AA-3526-01	0.5	Low
TCGA-AA-3854-01	0.5	Low
TCGA-AG-3896-01	0.5	Low
TCGA-CM-5864-01	0.5	Low
TCGA-DY-A1DG-01	0.5	Low
TCGA-A6-6648-01	0.51	Low
TCGA-AA-3511-01	0.51	Low
TCGA-AA-3713-01	0.51	Low
TCGA-AG-3725-01	0.51	Low
TCGA-G4-6315-01	0.51	Low
TCGA-AA-3685-01	0.52	Low
TCGA-AA-3846-01	0.52	Low
TCGA-AA-A01T-01	0.52	Low
TCGA-AG-A008-01	0.52	Low
TCGA-AZ-5403-01	0.52	Low
TCGA-DM-A1HB-01	0.52	Low
TCGA-EI-7002-01	0.52	Low
TCGA-A6-6138-01	0.53	Low
TCGA-AA-3856-01	0.53	Low
TCGA-AA-3979-01	0.53	Low
TCGA-AZ-4684-01	0.53	Low
TCGA-CM-4752-01	0.53	Low
TCGA-CM-6163-01	0.53	Low
TCGA-AG-3901-01	0.54	Low
TCGA-AG-A023-01	0.54	Low
TCGA-CM-6678-01	0.54	Low
TCGA-D5-6532-01	0.54	Low
TCGA-EI-6884-01	0.54	Low
TCGA-CK-4948-01	0.55	Low
TCGA-D5-5538-01	0.55	Low
TCGA-F5-6465-01	0.55	Low
TCGA-AA-3522-01	0.56	Low
TCGA-AA-3821-01	0.56	Low
TCGA-AA-3862-01	0.56	Low
TCGA-AA-A00K-01	0.56	Low
TCGA-AA-A01G-01	0.56	Low
TCGA-DM-A280-01	0.56	Low

TCGA-EI-6510-01	0.56	Low
TCGA-F4-6805-01	0.56	Low
TCGA-4T-AA8H-01	0.57	Low
TCGA-AG-3600-01	0.57	Low
TCGA-AG-3609-01	0.57	Low
TCGA-AG-3887-01	0.57	Low
TCGA-CK-5914-01	0.57	Low
TCGA-DC-6157-01	0.57	Low
TCGA-G4-6306-01	0.57	Low
TCGA-AA-3510-01	0.58	Low
TCGA-AA-3673-01	0.58	Low
TCGA-AA-A00D-01	0.58	Low
TCGA-AH-6544-01	0.58	Low
TCGA-AA-3844-01	0.59	Low
TCGA-AG-A015-01	0.59	Low
TCGA-CM-6677-01	0.59	Low
TCGA-A6-6781-01	0.6	Low
TCGA-AG-3598-01	0.6	Low
TCGA-AZ-5407-01	0.6	Low
TCGA-CA-5796-01	0.6	Low
TCGA-CM-6170-01	0.6	Low
TCGA-DC-5869-01	0.6	Low
TCGA-NH-A6GB-01	0.6	Low
TCGA-QG-A5Z2-01	0.6	Low
TCGA-AA-3815-01	0.61	Low
TCGA-CA-5254-01	0.61	Low
TCGA-CM-5862-01	0.61	Low
TCGA-D5-6932-01	0.61	Low
TCGA-EI-6513-01	0.61	Low
TCGA-AA-3831-01	0.62	Low
TCGA-AG-3594-01	0.62	Low
TCGA-D5-7000-01	0.62	Low
TCGA-A6-5665-01	0.63	Low
TCGA-AA-3678-01	0.63	Low
TCGA-AA-3861-01	0.63	Low
TCGA-AF-3911-01	0.63	Low
TCGA-AG-3581-01	0.63	Low
TCGA-AG-3893-01	0.63	Low
TCGA-CM-5860-01	0.63	Low
TCGA-DM-A28H-01	0.63	Low
TCGA-A6-A567-01	0.64	Low
TCGA-AA-A01V-01	0.64	Low
TCGA-AA-A02Y-01	0.64	Low
TCGA-AF-A56N-01	0.64	Low

TCGA-DC-6681-01	0.64	Low
TCGA-A6-6137-01	0.65	Low
TCGA-AA-3518-01	0.65	Low
TCGA-AA-3877-01	0.65	Low
TCGA-AA-A00L-01	0.65	Low
TCGA-AY-A69D-01	0.65	Low
TCGA-AA-3697-01	0.66	Low
TCGA-AA-3818-01	0.66	Low
TCGA-AA-3956-01	0.66	Low
TCGA-A6-A565-01	0.67	Low
TCGA-AG-3892-01	0.67	Low
TCGA-AG-3894-01	0.67	Low
TCGA-AG-4005-01	0.67	Low
TCGA-CA-5797-01	0.67	Low
TCGA-D5-6927-01	0.67	Low
TCGA-DM-A0X9-01	0.67	Low
TCGA-DM-A1D4-01	0.67	Low
TCGA-A6-6140-01	0.68	Low
TCGA-A6-6653-01	0.68	Low
TCGA-AA-3543-01	0.68	Low
TCGA-AA-3556-01	0.68	Low
TCGA-AA-3675-01	0.68	Low
TCGA-D5-5539-01	0.68	Low
TCGA-F5-6571-01	0.68	Low
TCGA-AA-A00R-01	0.69	Low
TCGA-AH-6643-01	0.69	Low
TCGA-DC-6154-01	0.69	Low
TCGA-DC-6158-01	0.69	Low
TCGA-EI-7004-01	0.69	Low
TCGA-CM-6680-01	0.7	Low
TCGA-DM-A0XD-01	0.7	Low
TCGA-AU-3779-01	0.71	Low
TCGA-AA-3939-01	0.72	Low
TCGA-AA-3534-01	0.73	Low
TCGA-CM-5863-01	0.73	Low
TCGA-DC-5337-01	0.73	Low
TCGA-G4-6321-01	0.73	Low
TCGA-A6-6142-01	0.74	Low
TCGA-AA-3662-01	0.74	Low
TCGA-D5-6530-01	0.74	Low
TCGA-AA-3532-01	0.75	Low
TCGA-AF-3913-01	0.75	Low
TCGA-AF-6136-01	0.75	Low
TCGA-CM-6165-01	0.75	Low

TCGA-D5-5540-01	0.75	Low
TCGA-EI-6881-01	0.75	Low
TCGA-EI-6885-01	0.75	Low
TCGA-A6-6141-01	0.76	Low
TCGA-AA-A00Z-01	0.76	Low
TCGA-AF-2692-01	0.76	Low
TCGA-AZ-6598-01	0.76	Low
TCGA-CI-6621-01	0.76	Low
TCGA-D5-6536-01	0.76	Low
TCGA-DC-6682-01	0.76	Low
TCGA-AG-A00H-01	0.77	Low
TCGA-CK-5913-01	0.77	Low
TCGA-G4-6317-02	0.77	Low
TCGA-AA-3860-01	0.78	Low
TCGA-AA-3977-01	0.78	Low
TCGA-AD-A5EK-01	0.78	Low
TCGA-AG-A036-01	0.78	Low
TCGA-F4-6856-01	0.78	Low
TCGA-NH-A50V-01	0.78	Low
TCGA-AG-A02X-01	0.79	Low
TCGA-DM-A1DA-01	0.79	Low
TCGA-RU-A8FL-01	0.79	Low
TCGA-A6-2682-01	0.8	Low
TCGA-A6-2684-01	0.8	Low
TCGA-AA-3970-01	0.8	Low
TCGA-AU-6004-01	0.8	Low
TCGA-AZ-6605-01	0.8	Low
TCGA-AA-3530-01	0.81	Low
TCGA-G4-6625-01	0.81	Low
TCGA-A6-A566-01	0.82	Low
TCGA-AA-3679-01	0.82	Low
TCGA-AA-3930-01	0.82	Low
TCGA-AA-A03J-01	0.82	Low
TCGA-AF-5654-01	0.82	Low
TCGA-CL-4957-01	0.82	Low
TCGA-CM-4746-01	0.82	Low
TCGA-G5-6641-01	0.82	Low
TCGA-AA-3664-01	0.83	Low
TCGA-AG-3580-01	0.83	Low
TCGA-AG-A02N-01	0.83	Low
TCGA-D5-6537-01	0.83	Low
TCGA-D5-6898-01	0.83	Low
TCGA-G4-6311-01	0.83	Low
TCGA-A6-5664-01	0.84	Low

TCGA-AG-3575-01	0.84	Low
TCGA-AG-3582-01	0.84	Low
TCGA-AF-6655-01	0.85	Low
TCGA-CM-6676-01	0.85	Low
TCGA-DC-6155-01	0.85	Low
TCGA-G4-6294-01	0.85	Low
TCGA-AA-3544-01	0.86	Low
TCGA-AA-A02W-01	0.86	Low
TCGA-CK-4951-01	0.86	Low
TCGA-D5-6924-01	0.86	Low
TCGA-G4-6317-01	0.86	Low
TCGA-A6-2677-01	0.87	Low
TCGA-D5-6922-01	0.88	Low
TCGA-G4-6323-01	0.88	Low
TCGA-QG-A5YV-01	0.88	Low
TCGA-AA-A00W-01	0.89	Low
TCGA-DM-A28E-01	0.89	Low
TCGA-AA-3663-01	0.9	Low
TCGA-AG-3611-01	0.9	Low
TCGA-AZ-4313-01	0.9	Low
TCGA-AA-A017-01	0.91	Low
TCGA-AG-A00Y-01	0.91	Low
TCGA-AY-A71X-01	0.91	Low
TCGA-CM-5341-01	0.91	Low
TCGA-F5-6811-01	0.91	Low
TCGA-G4-6586-01	0.91	Low
TCGA-AA-3561-01	0.92	Low
TCGA-AA-3672-01	0.92	Low
TCGA-QL-A97D-01	0.92	Low
TCGA-AA-3972-01	0.93	Low
TCGA-AD-6888-01	0.93	Low
TCGA-CK-6751-01	0.93	Low
TCGA-A6-2675-01	0.94	Low
TCGA-AA-3531-01	0.94	Low
TCGA-AA-3655-01	0.94	Low
TCGA-AA-3696-01	0.94	Low
TCGA-AG-3909-01	0.94	Low
TCGA-DC-6683-01	0.94	Low
TCGA-CM-5344-01	0.95	Low
TCGA-AA-3864-01	0.95	High
TCGA-AY-A54L-01	0.95	High
TCGA-AA-A029-01	0.96	High
TCGA-AG-3605-01	0.96	High
TCGA-AZ-4308-01	0.96	High

TCGA-CA-5256-01	0.96	High
TCGA-DY-A1H8-01	0.96	High
TCGA-DM-A28G-01	0.97	High
TCGA-G4-6627-01	0.97	High
TCGA-T9-A92H-01	0.97	High
TCGA-AG-3583-01	0.98	High
TCGA-AG-3599-01	0.98	High
TCGA-AZ-6606-01	0.98	High
TCGA-AA-3858-01	0.99	High
TCGA-AA-A03F-01	0.99	High
TCGA-AF-A56L-01	0.99	High
TCGA-CM-6679-01	0.99	High
TCGA-AA-3488-01	1	High
TCGA-AA-3949-01	1	High
TCGA-AF-6672-01	1	High
TCGA-AA-A01F-01	1.01	High
TCGA-CM-6162-01	1.01	High
TCGA-SS-A7HO-01	1.01	High
TCGA-AA-A02O-01	1.02	High
TCGA-A6-2680-01	1.03	High
TCGA-AA-3554-01	1.03	High
TCGA-AA-A024-01	1.04	High
TCGA-D5-6928-01	1.04	High
TCGA-3L-AA1B-01	1.05	High
TCGA-AA-3989-01	1.06	High
TCGA-AA-3710-01	1.07	High
TCGA-CM-6169-01	1.07	High
TCGA-AA-3994-01	1.08	High
TCGA-CI-6619-01	1.08	High
TCGA-NH-A50U-01	1.08	High
TCGA-AA-3502-01	1.09	High
TCGA-EI-6883-01	1.09	High
TCGA-G4-6297-01	1.09	High
TCGA-G4-6626-01	1.09	High
TCGA-AA-3870-01	1.1	High
TCGA-EI-6507-01	1.1	High
TCGA-EI-6882-01	1.11	High
TCGA-CA-6717-01	1.12	High
TCGA-CK-6748-01	1.12	High
TCGA-AA-3976-01	1.13	High
TCGA-AA-3542-01	1.14	High
TCGA-AA-3941-01	1.14	High
TCGA-D5-6540-01	1.14	High
TCGA-F5-6813-01	1.14	High

TCGA-A6-3807-01	1.15	High
TCGA-A6-5660-01	1.15	High
TCGA-A6-5667-01	1.15	High
TCGA-AA-3666-01	1.15	High
TCGA-AA-3833-01	1.15	High
TCGA-AA-3811-01	1.16	High
TCGA-AZ-4615-01	1.16	High
TCGA-AZ-6603-01	1.16	High
TCGA-DY-A0XA-01	1.16	High
TCGA-AA-3506-01	1.17	High
TCGA-AA-3842-01	1.17	High
TCGA-AA-3866-01	1.17	High
TCGA-A6-5662-01	1.18	High
TCGA-AD-6899-01	1.18	High
TCGA-QG-A5YX-01	1.19	High
TCGA-AA-3553-01	1.2	High
TCGA-AA-3966-01	1.2	High
TCGA-AG-3885-01	1.21	High
TCGA-CM-4748-01	1.21	High
TCGA-A6-4107-01	1.22	High
TCGA-CA-6715-01	1.22	High
TCGA-EI-6514-01	1.22	High
TCGA-AA-3489-01	1.23	High
TCGA-AA-3837-01	1.23	High
TCGA-AG-A016-01	1.23	High
TCGA-AY-6197-01	1.23	High
TCGA-AY-6386-01	1.24	High
TCGA-CM-4751-01	1.24	High
TCGA-A6-3808-01	1.25	High
TCGA-AG-3999-01	1.25	High
TCGA-D5-6923-01	1.26	High
TCGA-EI-6511-01	1.26	High
TCGA-G4-6314-01	1.26	High
TCGA-NH-A50T-01	1.26	High
TCGA-AA-3841-01	1.27	High
TCGA-AZ-4315-01	1.27	High
TCGA-AA-A02F-01	1.28	High
TCGA-DM-A282-01	1.28	High
TCGA-AA-A00A-01	1.29	High
TCGA-AA-A01D-01	1.29	High
TCGA-A6-5659-01	1.3	High
TCGA-DM-A1D7-01	1.3	High
TCGA-BM-6198-01	1.31	High
TCGA-DM-A1D6-01	1.31	High

TCGA-DM-A1DB-01	1.31	High
TCGA-AD-5900-01	1.32	High
TCGA-CL-5918-01	1.32	High
TCGA-A6-2672-01	1.33	High
TCGA-CM-5349-01	1.34	High
TCGA-EI-6509-01	1.34	High
TCGA-G5-6233-01	1.34	High
TCGA-AG-A01J-01	1.35	High
TCGA-F4-6704-01	1.36	High
TCGA-AF-2687-01	1.37	High
TCGA-CM-4744-01	1.37	High
TCGA-AG-A020-01	1.38	High
TCGA-CM-6161-01	1.38	High
TCGA-D5-5541-01	1.38	High
TCGA-D5-6930-01	1.38	High
TCGA-AA-A00N-01	1.39	High
TCGA-AA-A02H-01	1.39	High
TCGA-AA-A02J-01	1.4	High
TCGA-AG-3883-01	1.4	High
TCGA-CM-5868-01	1.4	High
TCGA-DC-4749-01	1.4	High
TCGA-CL-5917-01	1.42	High
TCGA-F4-6463-01	1.42	High
TCGA-F4-6569-01	1.42	High
TCGA-AA-A01Q-01	1.43	High
TCGA-AF-A56K-01	1.43	High
TCGA-AG-3591-01	1.43	High
TCGA-CM-6166-01	1.43	High
TCGA-NH-A8F7-06	1.43	High
TCGA-AA-3667-01	1.45	High
TCGA-F4-6703-01	1.45	High
TCGA-AA-A00U-01	1.46	High
TCGA-AY-5543-01	1.46	High
TCGA-CK-5916-01	1.47	High
TCGA-CM-6168-01	1.47	High
TCGA-EF-5830-01	1.47	High
TCGA-AG-A032-01	1.48	High
TCGA-AM-5820-01	1.48	High
TCGA-D5-6538-01	1.48	High
TCGA-NH-A6GC-01	1.48	High
TCGA-AA-3851-01	1.49	High
TCGA-AY-4071-01	1.49	High
TCGA-5M-AAT6-01	1.5	High
TCGA-AA-3538-01	1.5	High

TCGA-DM-A0XF-01	1.5	High
TCGA-DM-A28C-01	1.5	High
TCGA-G4-6628-01	1.5	High
TCGA-AA-A00O-01	1.51	High
TCGA-AA-3517-01	1.52	High
TCGA-AG-A002-01	1.52	High
TCGA-D5-6926-01	1.52	High
TCGA-DM-A1D0-01	1.52	High
TCGA-A6-6654-01	1.53	High
TCGA-CA-6719-01	1.55	High
TCGA-AG-3732-01	1.56	High
TCGA-AZ-6601-01	1.56	High
TCGA-G4-6303-01	1.56	High
TCGA-AZ-6608-01	1.57	High
TCGA-A6-6649-01	1.59	High
TCGA-AH-6549-01	1.59	High
TCGA-D5-6535-01	1.6	High
TCGA-DM-A28F-01	1.6	High
TCGA-DY-A1DC-01	1.61	High
TCGA-CM-5861-01	1.62	High
TCGA-F4-6807-01	1.62	High
TCGA-QG-A5YW-01	1.63	High
TCGA-AG-A01N-01	1.64	High
TCGA-G4-6588-01	1.65	High
TCGA-AA-3525-01	1.68	High
TCGA-AF-2691-01	1.69	High
TCGA-AA-3514-01	1.71	High
TCGA-AA-3562-01	1.71	High
TCGA-DY-A1DE-01	1.72	High
TCGA-AA-A00E-01	1.73	High
TCGA-AA-A01S-01	1.73	High
TCGA-AA-A01Z-01	1.73	High
TCGA-A6-2681-01	1.74	High
TCGA-AG-3878-01	1.75	High
TCGA-AA-3496-01	1.76	High
TCGA-AA-A01K-01	1.76	High
TCGA-AA-A02R-01	1.76	High
TCGA-AD-6965-01	1.78	High
TCGA-AA-A00F-01	1.81	High
TCGA-CA-6716-01	1.81	High
TCGA-AA-A02K-01	1.82	High
TCGA-F4-6459-01	1.82	High
TCGA-G4-6304-01	1.82	High
TCGA-A6-3810-01	1.84	High

TCGA-F5-6702-01	1.84	High
TCGA-A6-6652-01	1.85	High
TCGA-D5-6529-01	1.85	High
TCGA-WS-AB45-01	1.87	High
TCGA-A6-2679-01	1.9	High
TCGA-AA-3855-01	1.9	High
TCGA-AA-A010-01	1.9	High
TCGA-CM-4750-01	1.9	High
TCGA-A6-2676-01	1.92	High
TCGA-G5-6572-02	1.92	High
TCGA-AF-3400-01	1.94	High
TCGA-AG-3881-01	1.94	High
TCGA-AG-3727-01	1.95	High
TCGA-CA-5255-01	1.95	High
TCGA-AA-3812-01	1.96	High
TCGA-DM-A285-01	1.96	High
TCGA-AG-4001-01	1.99	High
TCGA-D5-6533-01	1.99	High
TCGA-AA-3680-01	2.01	High
TCGA-AA-A02E-01	2.01	High
TCGA-AD-6963-01	2.01	High
TCGA-AG-3728-01	2.01	High
TCGA-CM-4747-01	2.01	High
TCGA-AG-3890-01	2.02	High
TCGA-A6-A5ZU-01	2.03	High
TCGA-A6-5666-01	2.04	High
TCGA-AA-3516-01	2.05	High
TCGA-AA-A00Q-01	2.05	High
TCGA-AD-6895-01	2.06	High
TCGA-CK-5912-01	2.07	High
TCGA-AG-3726-01	2.08	High
TCGA-AA-A01C-01	2.1	High
TCGA-DM-A1D9-01	2.1	High
TCGA-AD-6964-01	2.11	High
TCGA-DM-A1HA-01	2.11	High
TCGA-AA-3848-01	2.18	High
TCGA-AG-3593-01	2.18	High
TCGA-AG-3898-01	2.18	High
TCGA-F5-6812-01	2.21	High
TCGA-G4-6320-01	2.21	High
TCGA-AZ-4323-01	2.22	High
TCGA-AZ-6599-01	2.23	High
TCGA-A6-2686-01	2.24	High
TCGA-F4-6461-01	2.24	High

TCGA-F5-6814-01	2.24	High
TCGA-F4-6808-01	2.25	High
TCGA-A6-A56B-01	2.26	High
TCGA-AM-5821-01	2.27	High
TCGA-DM-A1D8-01	2.27	High
TCGA-A6-6782-01	2.28	High
TCGA-A6-2685-01	2.29	High
TCGA-G4-6310-01	2.32	High
TCGA-G4-6302-01	2.34	High
TCGA-AA-3973-01	2.38	High
TCGA-AG-A014-01	2.39	High
TCGA-EF-5831-01	2.42	High
TCGA-EI-6917-01	2.42	High
TCGA-NH-A6GA-01	2.43	High
TCGA-CM-6164-01	2.44	High
TCGA-A6-2671-01	2.46	High
TCGA-AZ-6600-01	2.46	High
TCGA-F4-6855-01	2.46	High
TCGA-D5-5537-01	2.49	High
TCGA-AG-3882-01	2.52	High
TCGA-4N-A93T-01	2.55	High
TCGA-AA-3867-01	2.55	High
TCGA-AA-3869-01	2.55	High
TCGA-AG-3587-01	2.6	High
TCGA-F4-6570-01	2.6	High
TCGA-F4-6460-01	2.61	High
TCGA-NH-A8F8-01	2.61	High
TCGA-G4-6299-01	2.63	High
TCGA-AA-A01P-01	2.64	High
TCGA-AA-3548-01	2.68	High
TCGA-G4-6298-01	2.79	High
TCGA-CA-6718-01	2.81	High
TCGA-D5-6541-01	2.81	High
TCGA-AA-3692-01	2.86	High
TCGA-D5-6929-01	2.91	High
TCGA-DT-5265-01	2.93	High
TCGA-5M-AATE-01	3	High
TCGA-AG-4007-01	3.08	High
TCGA-G5-6572-01	3.1	High
TCGA-F5-6863-01	3.19	High
TCGA-5M-AAT4-01	3.28	High
TCGA-A6-6651-01	3.28	High
TCGA-AA-A01R-01	3.28	High
TCGA-AG-A02G-01	3.32	High

TCGA-AG-4021-01	3.45	High
TCGA-F5-6464-01	3.51	High
TCGA-NH-A5IV-01	3.57	High
TCGA-DM-A28A-01	3.69	High
TCGA-A6-2683-01	3.77	High
TCGA-QG-A5Z1-01	3.79	High
TCGA-F5-6864-01	3.89	High
TCGA-D5-6534-01	3.91	High
TCGA-CM-5348-01	4.07	High
TCGA-AG-3574-01	4.12	High
TCGA-CM-6167-01	4.14	High
TCGA-AA-A01X-01	4.16	High
TCGA-F4-6809-01	4.29	High
TCGA-AA-3552-01	4.32	High
TCGA-AF-2690-01	4.5	High
TCGA-DY-A1DD-01	4.52	High
TCGA-AZ-6607-01	4.55	High
TCGA-AD-6901-01	4.56	High
TCGA-CM-4743-01	4.63	High
TCGA-AG-A026-01	4.72	High
TCGA-AZ-4616-01	4.76	High
TCGA-AY-6196-01	4.78	High
TCGA-AA-3715-01	5.32	High
TCGA-DY-A1DF-01	5.62	High
TCGA-DC-6156-01	5.68	High
TCGA-AG-3584-01	5.98	High
TCGA-AA-3952-01	6.24	High
TCGA-AA-A004-01	6.67	High
TCGA-AZ-4614-01	7.81	High
TCGA-G4-6295-01	8.47	High
TCGA-AY-4070-01	9.09	High
