

The 25 patients (newly diagnosed + recurrent)

25 real, anonymized GBM cases pulled from public cancer-research databases — the people the tool is tested against. Values marked UNKNOWN are facts the databases don't track and are never guessed.

Columns 1 of 6 · anchored on patient_id

patient_id	sample_id	source_cohort	source_study	disease_stage	recurrence_number	age	sex	cancer_type_detailed	oncotree_code
TCGA-28-25-02	TCGA-28-2502-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	65	Male	Glioblastoma Multiforme	GBM
TCGA-41-25-75	TCGA-41-2575-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	75	Male	Glioblastoma Multiforme	GBM
TCGA-06-54-13	TCGA-06-5413-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	67	Male	Glioblastoma Multiforme	GBM
TCGA-02-00-24	TCGA-02-0024-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	35	Male	Glioblastoma Multiforme	GBM
TCGA-06-54-15	TCGA-06-5415-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	60	Male	Glioblastoma Multiforme	GBM
TCGA-32-19-78	TCGA-32-1978-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	57	Male	Glioblastoma Multiforme	GBM
TCGA-14-14-01	TCGA-14-1401-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	54	Male	Glioblastoma Multiforme	GBM
TCGA-32-19-82	TCGA-32-1982-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	76	Female	Glioblastoma Multiforme	GBM
TCGA-02-00-34	TCGA-02-0034-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	60	Male	Glioblastoma Multiforme	GBM
TCGA-14-18-25	TCGA-14-1825-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	70	Male	Glioblastoma Multiforme	GBM
TCGA-02-00-54	TCGA-02-0054-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	44	Female	Glioblastoma Multiforme	GBM
TCGA-06-10-86	TCGA-06-1086-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	42	Male	Glioblastoma Multiforme	GBM
TCGA-06-25-70	TCGA-06-2570-01	TCGA-newlyDx	gbm_tcga (Firehose) + gbm_tcga_pan_can_atlas_2018	newly_diagnosed	0	21	Female	Glioblastoma Multiforme	GBM
GLSS-HF-30-81	GLSS-HF-3081-R2	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	2	62	Female	Glioblastoma, IDH-mutant	GB
GLSS-HF-4B-46	GLSS-HF-4B46-R1	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	1	32	Male	Oligodendroglioma, IDH-mutant and 1p/19q-codeleted	ODG
GLSS-HF-1-CCD	GLSS-HF-1CC-D-R1	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	1	50	Male	Glioblastoma, IDH-wildtype	GB
GLSS-HF-1-CCD	GLSS-HF-1CC-D-R2	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	2	50	Male	Glioblastoma, IDH-wildtype	GB

patient_id	sample_id	source_cohort	source_study	disease_stage	recurrence_number	age	sex	cancer_type_detailed	oncotree_code
GLSS-HF-2829	GLSS-HF-2829-R1	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	1	44	Female	Glioblastoma, IDH-wildtype	GB
GLSS-HF-4FBF	GLSS-HF-4FBF-R2	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	2	52	Male	Glioblastoma, IDH-wildtype	GB
GLSS-HF-50F3	GLSS-HF-50F3-R2	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	2	46	Male	Glioblastoma, IDH-wildtype	GB
GLSS-HF-4F0A	GLSS-HF-4F0A-R1	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	1	35	Male	Oligoastrocytoma, IDH-mutant	AOAST
GLSS-HF-4F0A	GLSS-HF-4F0A-R2	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	2	35	Male	Glioblastoma, IDH-mutant	GB
GLSS-HF-57AE	GLSS-HF-57AE-R2	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	2	59	Female	Glioblastoma, IDH-wildtype	GB
GLSS-HF-3016	GLSS-HF-3016-R1	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	1	45	Male	Glioblastoma, IDH-wildtype	GB
GLSS-HF-4D9A	GLSS-HF-4D9A-R1	GLASS-recurrent	difg_glass (GLASS Consortium)	recurrent	1	43	Male	Glioblastoma, IDH-wildtype	GB

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patient_id	subtype	grade	tumor_tissue_site	IDH_status	MGMT_methylation	1p19q_codeletion	EGFR_amp	EGFR_mut	CDKN2A_B_del	PTEN_alt
TCGA-28-2502	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	no	none	yes	PTEN C71Y
TCGA-41-2575	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	AMP	none	yes	none
TCGA-06-5413	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	AMP	none	yes	none
TCGA-02-0024	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	no	none	yes	none
TCGA-06-5415	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	AMP	none	yes	none
TCGA-32-1978	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	AMP	none	yes	none
TCGA-14-1401	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	AMP	none	yes	none
TCGA-32-1982	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	AMP	EGFR G598V	yes	PTEN C105R
TCGA-02-0034	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	no	none	yes	HOMDEL
TCGA-14-1825	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	no	none	yes	HOMDEL
TCGA-02-0054	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	no	none	yes	none

patient_id	subtype	grade	tumor_tissue_site	IDH_status	MGMT_methylation	1p19q_codeletion	EGFR_amp	EGFR_mut	CDKN2A_B_del	PTEN_alt
TCGA-06-1086	IDH-wildtype	IV	Brain	IDH-wildtype	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	no	none	yes	none
TCGA-06-2570	IDH-mutant	IV	Brain	IDH-mutant	UNKNOWN (not in TCGA cBioPortal; external assay needed)	No (GBM; non-codeleted by definition)	no	none	no	none
GLSS-HF-3081	IDH-mutant	IV	Brain	IDH-mutant	Methylated	No	no	yes	HOMDEL	none
GLSS-HF-4B46	IDH-mutant	III	Brain	IDH-mutant	Methylated	Yes (1p/19q co-deleted)	no	no	none	none
GLSS-HF-1CCD	IDH-wildtype	IV	Brain	IDH-wildtype	Unmethylated	No	no	yes	none	none
GLSS-HF-1CCD	IDH-wildtype	IV	Brain	IDH-wildtype	Unmethylated	No	no	no	HOMDEL	none
GLSS-HF-2829	IDH-wildtype	IV	Brain	IDH-wildtype	Methylated	No	no	yes	HOMDEL	mut
GLSS-HF-4FBF	IDH-wildtype	IV	Brain	IDH-wildtype	Methylated	No	no	no	none	none
GLSS-HF-50F3	IDH-wildtype	IV	Brain	IDH-wildtype	Unmethylated	No	yes	yes	none	none
GLSS-HF-4F0A	IDH-mutant	III	Brain	IDH-mutant	Unmethylated	No	no	no	none	none
GLSS-HF-4F0A	IDH-mutant	IV	Brain	IDH-mutant	Methylated	No	no	no	none	mut
GLSS-HF-57AE	IDH-wildtype	IV	Brain	IDH-wildtype	Unmethylated	No	no	no	none	none
GLSS-HF-3016	IDH-wildtype	IV	Brain	IDH-wildtype	Unmethylated	No	yes	yes	none	mut
GLSS-HF-4D9A	IDH-wildtype	IV	Brain	IDH-wildtype	Unmethylated	No	no	no	none	none

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patient_id	TP53_mut	ATRX_mut	NF1_mut	BRAF_mut	actionable_other	all_marker_mutations
TCGA-28-2502	none	none	NF1 C167Qfs*10	none	none	NF1 C167Qfs*10; PTEN C71Y
TCGA-41-2575	TP53 X225_splice	none	none	none	none	TP53 X225_splice
TCGA-06-5413	none	none	none	none	PIK3CA E39K	PIK3CA E39K
TCGA-02-0024	none	none	none	none	none	none
TCGA-06-5415	none	none	none	none	none	none

patient_id	TP53_mut	ATRX_mut	NF1_mut	BRAF_mut	actionable_other	all_marker_mutations
TCGA-32-1978	none	none	none	none	none	none
TCGA-14-1401	none	none	none	none	none	none
TCGA-32-1982	none	none	none	none	none	EGFR G598V; PTEN C105R
TCGA-02-0034	none	none	none	none	none	none
TCGA-14-1825	TP53 R196*; TP53 R282W	none	none	none	none	TP53 R196*; TP53 R282W
TCGA-02-0054	none	none	none	none	none	none
TCGA-06-1086	none	none	none	none	none	none
TCGA-06-2570	TP53 Q136E	ATRX K1424Gfs*65	none	none	PIK3CA E542V	ATRX K1424Gfs*65; IDH1 R132H; PIK3CA E542V; TP53 Q136E
GLSS-HF-3081	yes	no	yes	yes	BRAF P141L; EGFR A92V,V876M; IDH1 R132H; NF1 G1842D; TP53 L111P	BRAF P141L; EGFR A92V,V876M; IDH1 R132H; NF1 G1842D; TP53 L111P
GLSS-HF-4B46	no	no	no	no	none	IDH1 R132H
GLSS-HF-1CCD	no	no	no	no	EGFR V774M	EGFR V774M
GLSS-HF-1CCD	no	no	no	no	none	none
GLSS-HF-2829	no	yes	yes	no	ATRX A1064T; EGFR M766I; NF1 M1035*,P678Rfs*10; PTEN R335*	ATRX A1064T; EGFR M766I; NF1 M1035*,P678Rfs*10; PTEN R335*
GLSS-HF-4FBF	no	no	no	no	none	none
GLSS-HF-50F3	no	no	no	no	EGFR D46H	EGFR D46H
GLSS-HF-4F0A	yes	yes	no	no	none	ATRX S458*; IDH1 R132H; TP53 K120E
GLSS-HF-4F0A	yes	yes	yes	no	none	ATRX S458*; CDKN2A V82Cfs*64; IDH1 R132H; NF1 G1869S; PTEN *404Wext*8; TP53 K120E
GLSS-HF-57AE	no	no	no	no	none	none
GLSS-HF-3016	yes	no	no	no	EGFR I744T; PTEN I122S; TP53 M133T	EGFR I744T; PTEN I122S; TP53 M133T
GLSS-HF-4D9A	no	no	no	no	none	none

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patient_id	all_marker_cnas	mutation_count	fraction_genome_altered	KPS_performance_status	ECOG_derived	extent_of_resection	prior_lines_of_therapy
TCGA-28-2502	CDKN2A:HOMDEL; CDKN2B:HOMDEL	43	0.1588	20	4	UNKNOWN (not in TCGA)	0
TCGA-41-2575	PDGFRA:AMP; TERT:AMP; CDKN2A:HOMDEL; CDKN2B:HOMDEL; EGFR:AMP	58	0.2403	60	2	UNKNOWN (not in TCGA)	0
TCGA-06-5413	CDKN2A:HOMDEL; CDKN2B:HOMDEL; EGFR:AMP	41	0.1798	60	2	UNKNOWN (not in TCGA)	0
TCGA-02-0024	MET:AMP; PDGFRA:AMP; CDKN2A:HOMDEL; CDKN2B:HOMDEL		0.3196	100	0-1	UNKNOWN (not in TCGA)	0
TCGA-06-5415	CDKN2A:HOMDEL; CDKN2B:HOMDEL; EGFR:AMP	43	0.2468	100	0-1	UNKNOWN (not in TCGA)	0
TCGA-32-1978	MDM2:AMP; CDKN2A:HOMDEL; CDKN2B:HOMDEL; EGFR:AMP		0.3329	UNKNOWN	UNKNOWN	UNKNOWN (not in TCGA)	0
TCGA-14-1401	MDM4:AMP; CDKN2A:HOMDEL; CDKN2B:HOMDEL; EGFR:AMP		0.109	80	0-1	UNKNOWN (not in TCGA)	0
TCGA-32-1982	TERT:AMP; CDKN2A:HOMDEL; CDKN2B:HOMDEL; EGFR:AMP	66	0.1315	80	0-1	UNKNOWN (not in TCGA)	0
TCGA-02-0034	PTEN:HOMDEL; CDKN2A:HOMDEL; CDKN2B:HOMDEL		0.2146	80	0-1	UNKNOWN (not in TCGA)	0
TCGA-14-1825	PTEN:HOMDEL; CDKN2A:HOMDEL; CDKN2B:HOMDEL	46	0.2814	80	0-1	UNKNOWN (not in TCGA)	0
TCGA-02-0054	CDKN2A:HOMDEL; CDKN2B:HOMDEL		0.1997	80	0-1	UNKNOWN (not in TCGA)	0
TCGA-06-1086	CDKN2A:HOMDEL; CDKN2B:HOMDEL		0.374	80	0-1	UNKNOWN (not in TCGA)	0
TCGA-06-2570	IDH2:AMP; CDK4:AMP	35	0.0786	100	0-1	UNKNOWN (not in TCGA)	0
GLSS-HF-3081	CDKN2A:HOMDEL; CDKN2B:HOMDEL	5083	0.6405	UNKNOWN (not collected in GLASS)	UNKNOWN	Total	2
GLSS-HF-4B46	none	36	0.095	UNKNOWN (not collected in GLASS)	UNKNOWN	Subtotal	1
GLSS-HF-1CCD	none	13	0.0001	UNKNOWN (not collected in GLASS)	UNKNOWN	Total	1
GLSS-HF-1CCD	CDKN2A:HOMDEL; CDKN2B:HOMDEL	75	0.1743	UNKNOWN (not collected in GLASS)	UNKNOWN	Total	2
GLSS-HF-2829	CDKN2A:HOMDEL; CDKN2B:HOMDEL	1327	0.3219	UNKNOWN (not collected in GLASS)	UNKNOWN	Subtotal	1
GLSS-HF-4FBF	none	UNKNOWN	UNKNOWN	UNKNOWN (not collected in GLASS)	UNKNOWN	Total	2
GLSS-HF-50F3	EGFR:AMP	70	0.0854	UNKNOWN (not collected in GLASS)	UNKNOWN	Biopsy	2
GLSS-HF-4FOA	none	50	0.0131	UNKNOWN (not collected in GLASS)	UNKNOWN	Total	1

patient_id	all_marker_cnas	mutation_count	fraction_genome_altered	KPS_performance_status	ECOG_derived	extent_of_resection	prior_lines_of_therapy
GLSS-HF-4F0A	none	1741	0.2262	UNKNOWN (not collected in GLASS)	UNKNOWN	Subtotal	2
GLSS-HF-57AE	none	UNKNOWN	UNKNOWN	UNKNOWN (not collected in GLASS)	UNKNOWN	Total	2
GLSS-HF-3016	EGFR:AMP	91	0.5197	UNKNOWN (not collected in GLASS)	UNKNOWN	Subtotal	1
GLSS-HF-4D9A	none	UNKNOWN	UNKNOWN	UNKNOWN (not collected in GLASS)	UNKNOWN	Subtotal	1

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patient_id	prior_radiation_tmz_completed	radiation_therapy	measurable_disease	ihc_biomarkers_CD70_GPC3_B7H3	os_status	os_months	pfs_status	pfs_months
TCGA-28-2502	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	0:LIVING	0.6575	0:CENSORED	0.6575
TCGA-41-2575	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	9.534	1:PROGRESSION	4.307
TCGA-06-5413	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	0:LIVING	8.811	1:PROGRESSION	6.411
TCGA-02-0024	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	53.1	1:PROGRESSION	46.06
TCGA-06-5415	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	0:LIVING	8.548	0:CENSORED	8.548
TCGA-32-1978	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	15.85	1:PROGRESSION	15.85
TCGA-14-1401	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	3.748	1:PROGRESSION	3.748
TCGA-32-1982	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	4.668	1:PROGRESSION	4.668
TCGA-02-0034	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	14.14	1:PROGRESSION	12.69
TCGA-14-1825	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	7.627	1:PROGRESSION	2.696
TCGA-02-0054	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	6.542	1:PROGRESSION	2.367
TCGA-06-1086	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	1:DECEASED	6.838	1:PROGRESSION	6.838
TCGA-06-2570	No (treatment-naive primary resection sample)	UNKNOWN	UNKNOWN (RANO not assessed in TCGA)	not assayed (IHC required)	0:LIVING	31.5	0:CENSORED	31.5
GLSS-HF-3081	Partial/UNKNOWN	No	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	39	UNKNOWN	UNKNOWN
GLSS-HF-4B46	Yes	Yes	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	0:LIVING	50	UNKNOWN	UNKNOWN

patient_id	prior_radiation_tmz_completed	radiation_therapy	measurable_disease	ihc_biomarkers_CD70_GPC3_B7H3	os_status	os_months	pfs_status	pfs_months
GLSS-HF-1 CCD	UNKNOWN	No	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	21	UNKNOWN	UNKNOWN
GLSS-HF-1 CCD	UNKNOWN	No	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	21	UNKNOWN	UNKNOWN
GLSS-HF-28 29	UNKNOWN	No	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	13	UNKNOWN	UNKNOWN
GLSS-HF-4F BF	Partial/UNKNOWN	No	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	56	UNKNOWN	UNKNOWN
GLSS-HF-50 F3	Partial/UNKNOWN	Yes	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	36	UNKNOWN	UNKNOWN
GLSS-HF-4F 0A	Partial/UNKNOWN	No	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	139	UNKNOWN	UNKNOWN
GLSS-HF-4F 0A	Yes	Yes	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	139	UNKNOWN	UNKNOWN
GLSS-HF-57 AE	UNKNOWN	No	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	11	UNKNOWN	UNKNOWN
GLSS-HF-30 16	Partial/UNKNOWN	Yes	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	21	UNKNOWN	UNKNOWN
GLSS-HF-4 D9A	UNKNOWN	No	UNKNOWN (RANO not assessed in GLASS)	not assayed (IHC required)	1:DECEASED	17	UNKNOWN	UNKNOWN

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patient_id	provenance_notes
TCGA-28-25 02	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-41-25 75	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-06-54 13	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-02-00 24	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-06-54 15	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-32-19 78	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-14-14 01	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA

patient_id	provenance_notes
TCGA-32-19 82	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-02-00 34	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-14-18 25	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-02-00 54	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-06-10 86	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
TCGA-06-25 70	molecular+outcomes real (TCGA); KPS/age/sex real (Firehose); MGMT/resection/measurable not available in TCGA
GLSS-HF-30 81	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-4B 46	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-1 CCD	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-1 CCD	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-28 29	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-4F BF	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-50 F3	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-4F 0A	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-4F 0A	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-57 AE	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed

patient_id	provenance_notes
GLSS-HF-3016	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed
GLSS-HF-4D9A	MGMT/resection/prior-therapy/recurrence/IDH/co del real (GLASS); KPS not collected; measurable not assessed

Fields the source cohorts don't carry (MGMT on newly-diagnosed, RANO measurable disease, IHC markers) are shown as UNKNOWN / not assayed and are treated as "not enough information," never as a measurement.