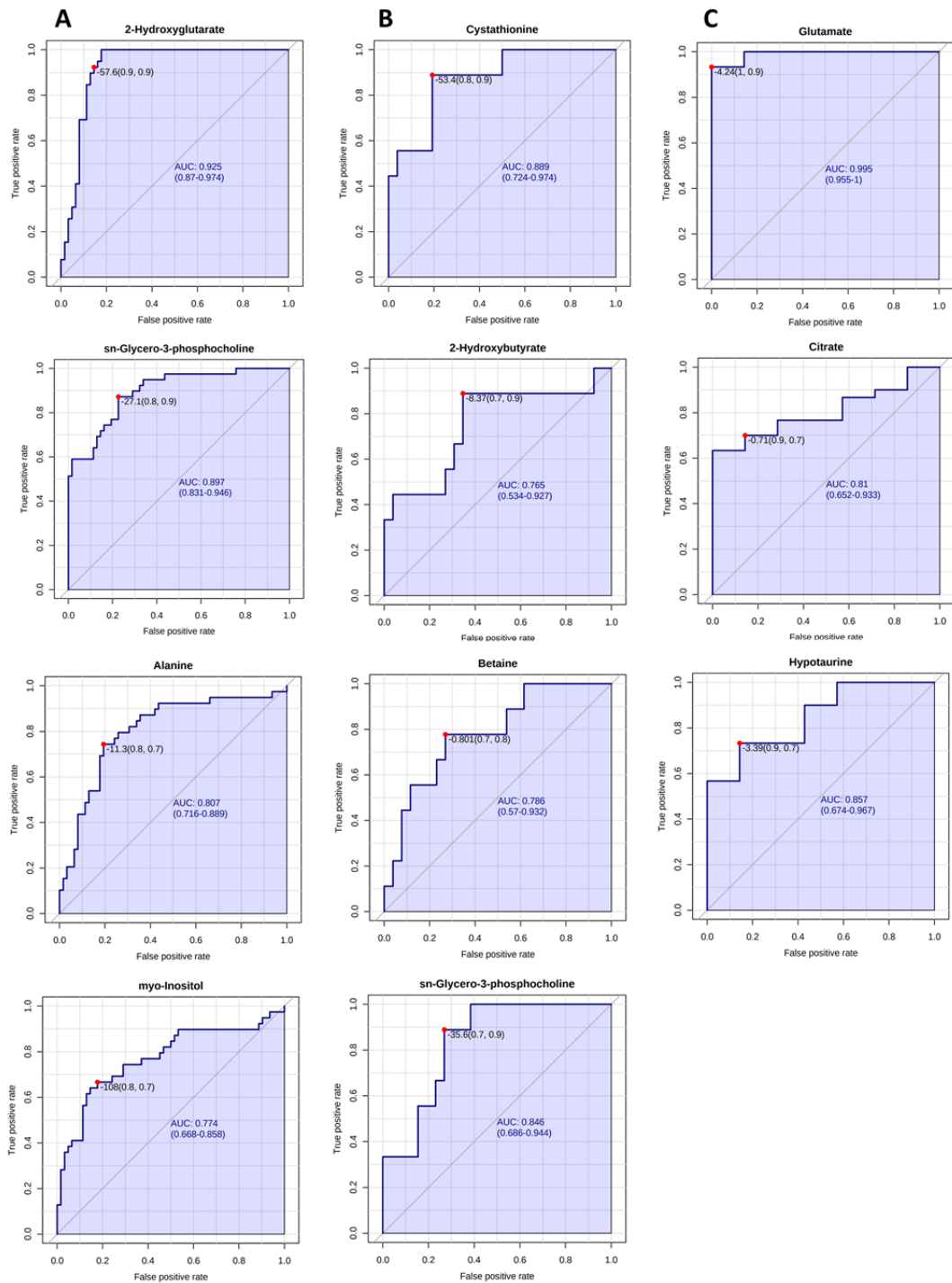


Supplementary information (Trautwein *et al.*)

Content

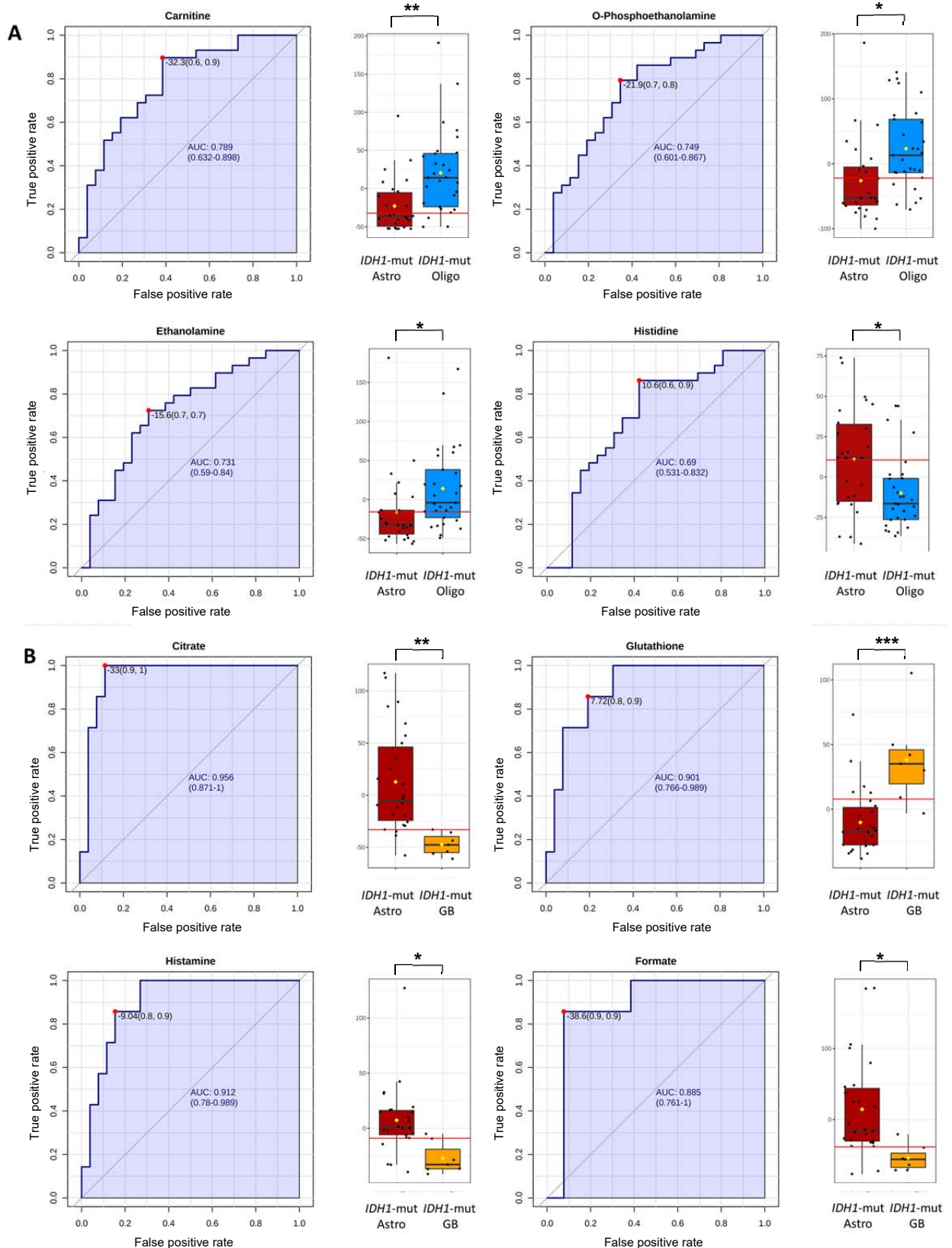
- 1 Supplementary figures S1-S5
- 2 Supplementary tables ST1-ST2

1 Supplementary Figures

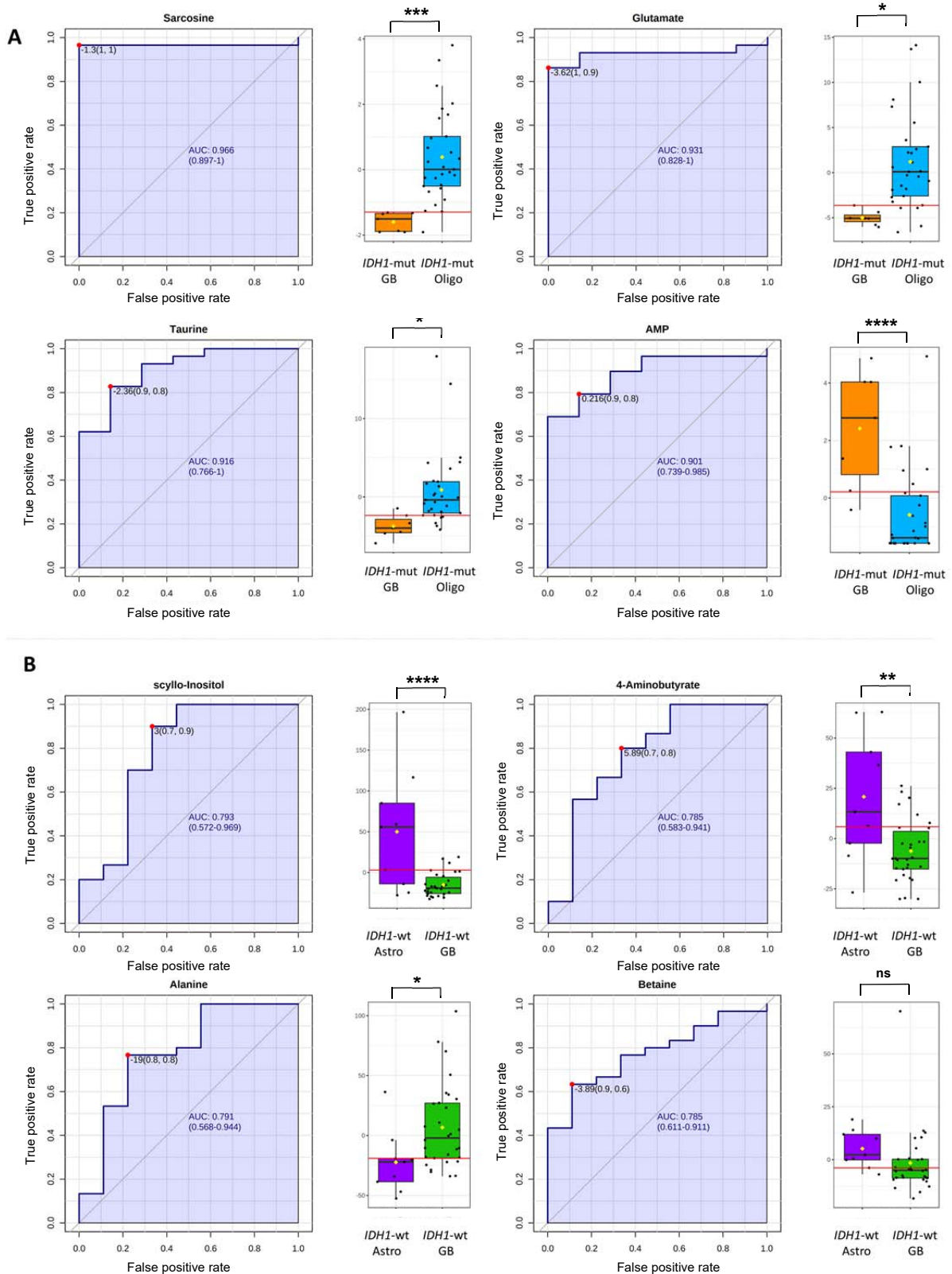


Supplementary Figure S1: Receiver operating characteristic (ROC) analysis.

Area under the curve (AUC) plots for *IDH1*-mutation (*IDH1*-mut, n = 62) vs. wildtype (*IDH1*-wt, n = 39) comparison I (A), astrocytoma (Astro) *IDH1*-mut (n = 26) vs. wt (n = 9) comparison II (B) and glioblastoma (GB) *IDH1*-mut (n = 7) vs. wt (n = 30) comparison III (C).



Supplementary Figure S2: Comparisons of tumor entities by receiver operating characteristic (ROC) analysis and significant metabolites. (A) Astro IDH1-mut (n = 26) vs. Oligo IDH1-mut (n = 29) comparison IV, and (B) Astro IDH1-mut (n = 26) vs. GB IDH1-mut (n = 7) comparison V. p-values determined by student's t-tests. Illustrated as box and whiskers plots showing all points with mean, min/max and outliers. Abbreviations: Astro, astrocytoma; GB, glioblastoma; Oligo, oligodendroglioma. p-values: * < 0.001, ** < 0.01, * < 0.05.**



Supplementary Figure S3: Comparisons of tumor entities by receiver operating characteristic ROC analysis and significant metabolites. (A) GB *IDH1*-mut (n = 7) vs. Oligo *IDH1*-mut (n = 29), comparison VI, and **(B)** Astro *IDH1*-wt (n = 9) vs. GB *IDH1*-wt (n = 30) comparison VII. *p*-values determined by student's t-tests. Illustrated as box and whiskers plots showing all points with mean, min/max and outliers. Abbreviations: Astro, astrocytoma; GB, glioblastoma; Oligo, oligodendroglioma. *p*-values: **** < 0.0001, *** < 0.001, ** < 0.01, * < 0.05.

A Progression-free survival

PFS <u>Univariate</u>	Thresholds (mM) [#]	Median (months)	
		<u>below</u>	<u>above</u>
2-Hydroxyglutarate	-87.44	09, CI 95% 06-12, n=23	54, CI 95% 31-60, n=43
Alanine	-43.06	56, CI 95% 31-68, n=22	14, CI 95% 10-30, n=44
Creatine phosphate	-20.52	12, CI 95% 06-23, n=24	54, CI 95% 30-66, n=42
Homocysteine	40.40	36, CI 95% 16-56, n=58	11, CI 95% 03-14, n=08
Isoleucine	-3.58	41, CI 95% 23-60, n=45	12, CI 95% 06-16, n=21
Leucine	56.06	36, CI 95% 16-56, n=56	08, CI 95% 04-12, n=10
Mannitol	-48.33	54, CI 95% 14-66, n=17	16, CI 95% 12-36, n=49
<i>myo</i> -Inositol	-23.51	14, CI 95% 09-23, n=34	60, CI 95% 32-76, n=32
<i>o</i> -Acetylcholine	18.75	36, CI 95% 16-56, n=53	11, CI 95% 04-16, n=13
<i>scyllo</i> -Inositol	-80.09	11, CI 95% 06-14, n=20	54, CI 95% 31-66, n=46
Serine	-61.86	56, CI 95% 14-76, n=11	16, CI 95% 12-36, n=55
Threonine	18.95	41, CI 95% 16-60, n=48	12, CI 95% 06-16, n=18
<i>sn</i> -Glycero-3-phosph.	-22.31	42, CI 95% 25-60, n=37	12, CI 95% 09-30, n=29

BBivariate pathology (*IDH1*-status, pathology)

Group 1 <i>IDH1</i> -wt	09, CI 95% 06-12,	n=27
Group 2 <i>IDH1</i> -mut, astrocytoma	54, CI 95% 50-76,	n=17
Group 3 <i>IDH1</i> -mut, oligodendroglioma	56, CI 95% 25-73,	n=17

IDH1-mut glioblastoma (n=5) were excluded, because 4 were censored.

Bivariate metabolites (2-Hydroxyglutarate -60,05 and *myo*-Inositol -23,51)

Group 1 (2-Hydroxyglutarate < -60,05)	10, CI 95% 06-12,	n=31
Group 2 (2-Hydroxyglutarate > -60,05 and < <i>myo</i> -Inositol 23,51)	31, CI 95% 16-60,	n=13
Group 3 (2-Hydroxyglutarate > -60,05 and > <i>myo</i> -Inositol 23,51)	68, CI 95% 42-76,	n=22

C Overall survival

OS <u>Univariate</u>	Thresholds (mM) [#]	Median (months)	
		<u>below</u>	<u>above</u>
2-Hydroxyglutarate	-42.99	30, CI 95% 18-48, n=36	103, CI 95% 73-152, n=30
Acetate	33.53	68, CI 95% 35-97, n=58	156, CI 95% 72-221, n=8
Alanine	-39.59	110, CI 95% 73-152, n=28	35, CI 95% 20-95, n=38
Formate	40.51	73, CI 95% 37-103, n=58	152, CI 95% 23-175, n=8
Glutamate	-32.71	97, CI 95% 68-152, n=33	37, CI 95% 19-103, n=33
Glycine	-44.75	110, CI 95% 68-156, n=29	45, CI 95% 20-83, n=37
Homocysteine	-15.84	133, CI 95% 35-133, n=28	73, CI 95% 30-103, n=38
Isoleucine	-11.02	95, CI 95% 52-110, n=42	35, CI 95% 18-open, n=24
Leucine	11.74	95, CI 95% 68-110, n=49	23, CI 95% 13-open, n=17
Mannitol	10.74	97, CI 95% 52-110, n=48	20, CI 95% 13-83, n=18
<i>myo</i> -Inositol	-163.2	24, CI 95% 13-97, n=11	95, CI 95% 52-110, n=55
Phenylalanine	-0.08	95, CI 95% 52-110, n=42	35, CI 95% 18-open, n=24
Threonine	18.95	95, CI 95% 68-110, n=48	39, CI 95% 18-open, n=18
Tyrosine	-5.68	97, CI 95% 68-116, n=39	31, CI 95% 18-open, n=27
<i>sn</i> -Glycero-3-phosph.	-11.20	37, CI 95% 18-83, n=39	110, CI 95% 68-152, n=27

DBivariate pathology

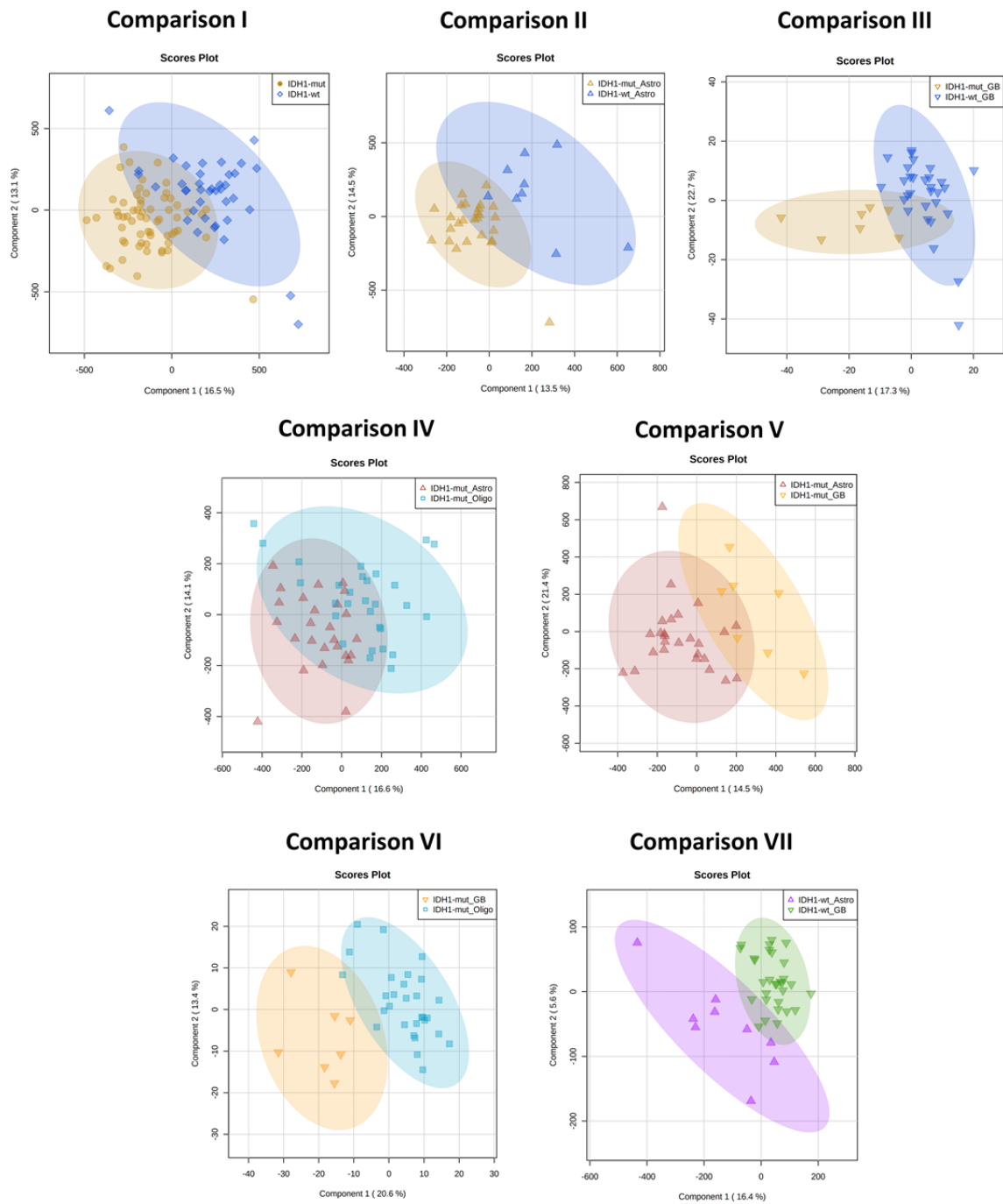
Group 1 <i>IDH1</i> -wt	19, CI 95% 15-30,	n=27
Group 2 <i>IDH1</i> -mut, astrocytoma	110, CI 95% 68-116,	n=1
Group 3 <i>IDH1</i> -mut, oligodendroglioma	110, CI 95% 52-156,	n=17

IDH1-mut glioblastoma (n=5) were excluded, because all were censored.

Bivariate metabolites (Acetate 33,53 and 2-Hydroxyglutarate -22,4)

Group 1 (Acetate < 33,53 and 2-Hydroxyglutarate < -22,4)	31, CI 95% 18-133,	n=37
Group 2 (Acetate < 33,53 and 2-Hydroxyglutarate > -22,4)	95, CI 95% 68-110,	n=21
Group 3 (Acetate > 33,53)	156, CI 95% 72-221,	n=8

Supplementary Figure S4: Tissue metabolite correlations with progression free and overall survival. Overview of metabolites and their univariate thresholds determined by one-step classification and regression tree analysis for optimal dichotomization of patients for progression-free survival (PFS, **A**) and overall survival (OS, **C**); Optimal trichotomization of patients by bivariate analyses of either pathological parameters or metabolites each by two-step classification and regression tree analysis for PFS (**B**) and OS (**D**). #Metabolite concentrations (mM) were normalized and Pareto-scaled for dilution effects. *sn*-Glycero-3-phosph. = *sn*-Glycero-3-phosphocholine.



Supplementary Figure S5: Partial Least Squares Discriminant Analysis (PLSDA) scores plots from the comparisons I – VII. PLSDA scores plots were integrated in main Figures 2 – 5.

2 Supplementary tables

Comparison I: <i>IDH1</i> -mut (n = 62) vs <i>IDH1</i> -wt (n = 39)				Comparison II: <i>IDH1</i> -mut Astro (n = 26) vs <i>IDH1</i> -wt Astro (n = 9)				Comparison III: <i>IDH1</i> -mut GB (n = 7) vs <i>IDH1</i> -wt GB (n = 30)			
Biomarker	AUC	t-test	log FC2	Biomarker	AUC	t-test	log FC2	Biomarker	AUC	t-test	log FC2
2-Hydroxyglutarate	0.923	8.95E-10	6.59	2-Hydroxyglutarate	0.880	0.003	3.47	Glutamate	0.990	7.30E-05	-3.55
sn-Glycero-3-phosphocholine	0.895	5.95E-11	2.96	Cystathionine	0.876	0.052	3.72	myo-Inositol	0.924	4.25E-06	-0.43
scyllo-Inositol	0.833	8.97E-04	2.96	sn-Glycero-3-phosphocholine	0.838	0.001	1.36	sn-Glycero-3-phosphocholine	0.905	1.32E-06	-0.62
Alanine	0.804	6.77E-07	0.37	Betaine	0.782	0.007	-0.88	scyllo-Inositol	0.876	5.57E-04	-1.03
myo-Inositol	0.770	0.020	2.57	Histamine	0.774	0.074	1.35	Hypotaurine	0.848	0.062	-4.80
Hypotaurine	0.758	7.30E-05	-0.34	2-Hydroxybutyrate	0.752	0.065	0.99	Citrate	0.805	0.061	-3.51
Sarcosine	0.729	0.002	3.47	Valine	0.735	0.438	0.75	AMP	0.800	0.012	2.02
Cystathionine	0.698	0.004	4.24	Hypoxanthine	0.701	0.021	-0.96	Formate	0.786	0.081	-3.34
Glutathione	0.693	2.72E-04	1.39	ADP	0.697	0.177	2.29	Creatine	0.779	0.008	-1.46
Mannitol	0.691	5.25E-04	0.78	O-Phosphocholine	0.692	0.082	-0.59	Alanine	0.776	0.038	-3.25
O-Acetylcholine	0.689	0.002	0.81	Leucine	0.688	0.900	0.42	Inosine	0.776	0.047	-3.49
Serine	0.686	7.96E-04	0.82	Hypotaurine	0.684	0.033	-0.68	Taurine	0.774	0.030	-3.46
O-Phosphocholine	0.683	0.011	0.87	O-Acetylcholine	0.679	0.048	-0.44	Acetate	0.767	0.195	-2.98
Creatine phosphate	0.678	0.333	2.30	Glutamate	0.671	0.174	-0.29	Guanosine	0.745	0.101	-3.10
Succinate	0.665	0.008	0.98	4-Aminobutyrate	0.662	0.082	-0.35	Carnitine	0.717	0.048	-1.43
Threonine	0.664	0.003	1.03	Homocysteine	0.658	0.093	-0.55	Glutamine	0.714	0.202	-1.89
Glycine	0.658	0.004	0.87	Serine	0.654	0.040	-0.52	Fumarate	0.712	0.056	-3.20
Creatine	0.658	0.036	1.88	Ascorbate	0.645	0.044	-0.32	Hypoxanthine	0.700	0.166	-3.04
Glutamate	0.653	0.040	1.01	Phenylalanine	0.645	0.853	0.49	Lactate	0.698	0.157	-1.88
Carnitine	0.649	0.002	1.83	Histidine	0.641	0.283	0.63	O-Phosphocholine	0.671	0.294	-2.75

Histamine	0.649	0.007	2.51	O-Phosphoethanolamine	0.637	0.430	-0.08	Histamine	0.657	0.190	-3.34
Aspartate	0.648	0.033	-0.95	Tyrosine	0.632	0.972	0.39	Phenylalanine	0.657	0.782	-1.98
Methionine	0.645	0.035	1.26	Glutathione	0.628	0.035	-0.35	Aspartate	0.652	0.151	-1.32
Histidine	0.643	0.143	2.43	Alanine	0.628	0.188	-0.13	Homocysteine	0.648	0.150	-2.97
Guanosine	0.635	0.021	0.66	Citrate	0.620	0.276	0.44	Cystathionine	0.645	0.316	-0.21

Comparison IV: <i>IDH1</i> -mut Astro (n = 26) vs. <i>IDH1</i> -mut Oligo (n = 29)				Comparison V: <i>IDH1</i> -mut GB (n = 7) vs. <i>IDH1</i> -mut Astro (n = 26)				Comparison VI: <i>IDH1</i> -mut GB (n = 7) vs. <i>IDH1</i> -mut Oligo (n = 29)				Comparison VII: <i>IDH1</i> -wt Astro (n = 9) vs. <i>IDH1</i> -wt GB (n = 30)			
Biomarker	AUC	t-test	log FC2	Biomarker	AUC	t-test	log FC2	Biomarker	AUC	t-test	log FC2	Biomarker	AUC	t-test	log FC2
Carnitine	0.786	0.001	-3.95	Citrate	0.951	0.004	5.90	Sarcosine	0.966	7.23E-04	-8.01	scyllo-Inositol	0.785	4.86E-05	2.89
O-Phospho-ethanolamine	0.744	0.005	-1.99	Histamine	0.901	0.008	5.74	Glutamate	0.926	0.004	-6.26	4-Amino-butyrate	0.778	0.001	2.88
Ethanolamine	0.728	0.032	-2.37	Glutathione	0.896	2.47E-04	2.72	Taurine	0.906	0.020	-6.94	Alanine	0.778	0.028	0.18
scyllo-Inositol	0.707	0.021	-2.12	Formate	0.879	0.015	5.06	AMP	0.892	9.62E-05	0.83	Betaine	0.774	0.227	2.37
Histidine	0.696	0.007	-0.77	Uridine	0.868	0.002	5.50	Uridine	0.892	4.16E-04	-7.83	Phenylalanine	0.759	0.113	0.36
Sarcosine	0.695	0.014	-3.00	AMP	0.863	3.46E-04	-1.31	Phenylalanine	0.887	1.57E-05	-3.84	Tyrosine	0.744	0.149	0.47
Guanosine	0.684	0.147	0.84	Acetate	0.857	0.062	4.80	Lactate	0.887	1.60E-04	-4.31	Valine	0.737	0.145	0.51
GSSG	0.682	0.010	-0.78	Histidine	0.846	0.006	5.51	Glutathione	0.887	8.05E-04	-3.77	myo-Inositol	0.722	7.54E-04	2.61
Nicotinurate	0.676	0.029	-0.23	Glutamate	0.830	0.023	4.66	Formate	0.882	0.133	-6.75	Creatine	0.711	0.006	2.03
Citrate	0.674	0.035	-0.77	Taurine	0.819	0.003	5.18	Citrate	0.872	0.019	-6.66	Hypotaurine	0.704	0.194	-0.28
Lactate	0.662	0.113	-1.08	Butyrate	0.813	0.009	2.60	Succinate	0.852	0.001	-4.33	Histidine	0.704	0.273	2.82
O-Acetylcholine	0.658	0.030	-1.93	Lactate	0.791	0.019	3.23	Butyrate	0.837	0.006	-4.11	Hypoxanthine	0.696	0.007	2.38
Mannitol	0.639	0.033	-0.91	Serine	0.780	0.046	3.00	Histidine	0.818	0.063	-7.10	Sarcosine	0.696	0.040	3.30
Acetate	0.634	0.954	-1.02	O-Acetylcholine	0.775	0.013	2.89	Tyrosine	0.808	3.91E-04	-4.22	Ethanolamine	0.685	0.797	2.76
Glutathione	0.633	0.727	-1.05	Phenylalanine	0.769	0.024	2.86	Mannitol	0.808	0.006	-4.10	Creatine	0.682	0.024	2.55

												phosphate			
Taurine	0.629	0.038	-1.76	Succinate	0.764	0.025	3.10	Methionine	0.803	0.120	-4.40	Acetate	0.682	0.102	2.06
O-Phosphocholine	0.627	0.272	-1.80	GTP	0.764	0.064	5.34	Histamine	0.793	0.020	-6.55	Cystathionine	0.678	0.060	-0.11
myo-Inositol	0.621	0.160	-0.93	4-Aminobutyrate	0.758	0.070	4.78	scyllo-Inositol	0.783	0.122	-6.25	Ascorbate	0.674	0.028	2.63
Phenylalanine	0.615	0.047	-0.98	N-Acetylaspartate	0.753	0.092	5.23	Isoleucine	0.778	7.36E-04	-3.88	Leucine	0.674	0.237	0.68
GTP	0.614	0.112	-0.25	Inosine	0.742	0.070	4.86	Glucose	0.768	0.033	-4.17	Isoleucine	0.667	0.317	0.45
2-Hydroxybutyrate	0.614	0.133	-1.01	Fumarate	0.736	0.028	4.68	Serine	0.768	0.039	-4.54	Glycine	0.663	0.171	0.81
Hypoxanthine	0.602	0.855	-0.45	Methionine	0.731	0.169	2.85	Fumarate	0.749	0.046	-5.88	O-Acetylcholine	0.656	0.268	0.87
3-Hydroxybutyrate	0.597	0.927	-1.60	Tyrosine	0.720	0.076	3.08	Carnitine	0.739	0.074	-5.93	Mannitol	0.652	0.116	0.97
Alanine	0.595	0.311	-1.57	Hypoxanthine	0.703	0.188	4.54	4-Aminobutyrate	0.739	0.078	-6.26	GTP	0.648	0.153	1.97
Glutamate	0.594	0.311	-1.60	Isoleucine	0.703	0.311	2.96	Valine	0.724	0.006	-4.55	Uridine	0.648	0.732	2.19

Supplementary table ST1: Top 25 AUC details for univariate ROC biomarker analysis of comparisons I-III in the “*IDH1* mutation status cohort”, and comparisons IV - VII for different histologies and identical *IDH1* status in the “*IDH1* mutation status cohort”. Abbreviations: AMP, adenosine monophosphate; Astro, astrocytoma; AUC, area under curve; Dex, dexamethasone; FC, fold change; GB, glioblastoma; GTP, guanosine triphosphate; GSSG, glutathione disulfide; Oligo, oligodendroglioma; ROC, receiver operator characteristics; Explanations: Bold are biomarkers with $p < 0.05$; Highlighted in grey are biomarkers shown and discussed in the manuscript and correlated with gene expression data.

Supplementary Table ST2: Basic clinical characteristics of study population

ID	IDH1 status	Treatment prior to tissue acquisition	Histo	WHO	Age (years)	Resection	PFS (months)	OS (months)
1201	IDH1-wt	Untreated	GB	IV	50	1	12	18
1202	IDH1-wt	Untreated	GB	IV	78	1	13	19
1203	IDH1-wt	Untreated	GB	IV	65	1	3	13
1204	IDH1-wt	Untreated	GB	IV	49	1	11	no information
1205	IDH1-wt	Untreated	GB	IV	45	1	6	10
1206	IDH1-wt	Untreated	GB	IV	70	1	7	15
1207	IDH1-wt	Untreated	GB	IV	45	1	4	28
1208	IDH1-wt	Untreated	GB	IV	55	1	11	35
1209	IDH1-wt	Untreated	GB	IV	70	1	4	12
1210	IDH1-wt	Untreated	Astro	III	49	1	6	18
1211	IDH1-wt	Untreated	GB	IV	47	1	4	37
1214	IDH1-wt	Untreated	GB	IV	48	1	> 53	n/a
1215	IDH1-wt	Untreated	GB	IV	58	1	14	20
1218	IDH1-wt	Untreated	GB	IV	71	1	4	12
1219	IDH1-wt	Untreated	GB	IV	72	1	12	no information
1220	IDH1-wt	Untreated	GB	IV	80	1	6	13
1221	IDH1-wt	Untreated	Astro	II	37	1	no information	no information
1222	IDH1-wt	Untreated	GB	IV	77	1	no information	no information
1224	IDH1-wt	Untreated	GB	IV	70	1	6	13
1225	IDH1-wt	Untreated	GB	IV	71	1	no information	no information
1226	IDH1-wt	Untreated	Astro	II	40	1	> 36	n/a
1227	IDH1-wt	Untreated	GB	IV	50	1	16	31
1228	IDH1-wt	Untreated	GB	IV	77	1	no information	no information
1401	IDH1-mut	Untreated	GB	IV	33	1	56	n/a
1402	IDH1-mut	Untreated	GB	IV	34	1	> 56	n/a
1403	IDH1-mut	Untreated	GB	IV	40	1	> 55	n/a
1404	IDH1-mut	Untreated	Astro	III	40	1	> 52	n/a
1405	IDH1-mut	Untreated	Astro	III	45	1	36	> 57
1406	IDH1-mut	Untreated	Astro	III	32	1	> 60	n/a
1407	IDH1-mut	Untreated	GB	IV	37	1	31	> 63
1408	IDH1-mut	Untreated	Oligo	II	56	1	11	> 27
1409	IDH1-mut	Untreated	GB	IV	48	1	no information	no information
1410	IDH1-mut	Untreated	GB	IV	34	1	> 48	no information
1411	IDH1-mut	Untreated	Astro	III	34	1	15	> 50
1412	IDH1-mut	Untreated	Oligo	II	71	1	> 52	n/a

1414	<i>IDH1</i> -mut	Untreated	Oligo	II	32	1	> 48	n/a
1416	<i>IDH1</i> -mut	Untreated	Oligo	II	57	1	> 39	n/a
1417	<i>IDH1</i> -mut	Untreated	Astro	II	63	1	> 57	no information
1418	<i>IDH1</i> -mut	Untreated	Astro	II	45	1	no information	no information
1419	<i>IDH1</i> -mut	Untreated	Astro	II	36	1	> 40	n/a
1420	<i>IDH1</i> -mut	Untreated	Astro	II	62	1	41	> 52
1424	<i>IDH1</i> -mut	Untreated	Astro	II	39	1	76	> 88
1427	<i>IDH1</i> -mut	Untreated	GB	IV	34	1	no information	no information
1429	<i>IDH1</i> -mut	Untreated	Oligo	II	47	1	> 36	n/a
1430	<i>IDH1</i> -mut	Untreated	Astro	III	28	1	15	no information
2726	<i>IDH1</i> -mut	Untreated	Oligo	III	43	1	25	156
3088	<i>IDH1</i> -mut	Untreated	Oligo	III	45	2	25	156
1506	<i>IDH1</i> -mut	Untreated	Oligo	II	66	1	73	103
2686	<i>IDH1</i> -mut	Untreated	Oligo	III	71	2	73	103
5096	<i>IDH1</i> -mut	Untreated	Oligo	II	45	1	32	48
5939	<i>IDH1</i> -mut	Untreated	Oligo	III	48	2	32	48
2287	<i>IDH1</i> -mut	Untreated	Oligo	II	29	1	23	175
2754	<i>IDH1</i> -mut	Untreated	Oligo	III	31	2	23	175
3509	<i>IDH1</i> -mut	Untreated	Astro	II	37	1	31	83
4133	<i>IDH1</i> -mut	Untreated	Astro	III	39	2	31	83
5120	<i>IDH1</i> -mut	Untreated	Astro	II	22	1	13	45
5475	<i>IDH1</i> -mut	Untreated	Astro	III	23	2	13	45
T01	<i>IDH1</i> -mut	Untreated	Astro	II	39	1	76	> 86
T02	<i>IDH1</i> -mut	Untreated	Astro	II	39	2	76	> 84
T05	<i>IDH1</i> -wt	Untreated	Astro	III	45	1	35	> 51
T06	<i>IDH1</i> -wt	Untreated	Astro	III	45	2	no information	no information
1534	<i>IDH1</i> -wt	RT	Astro	III	51	1	14	18
1774	<i>IDH1</i> -wt	RT	Astro	III	52	2	14	18
2252	<i>IDH1</i> -wt	RT -> PCV	Astro	III	47	1	16	18
2581	<i>IDH1</i> -wt	RT -> PCV	Astro	III	48	2	16	18
4589	<i>IDH1</i> -mut	RT	Astro	III	24	1	42	68
5647	<i>IDH1</i> -mut	RT	Astro	III	37	2	42	68
3071	<i>IDH1</i> -mut	RT	Astro	III	35	1	66	73
4218	<i>IDH1</i> -mut	RT	Astro	III	39	2	66	73
3186	<i>IDH1</i> -mut	RT	Astro	III	29	1	60	116
4306	<i>IDH1</i> -mut	RT	Astro	III	34	2	60	116
3648	<i>IDH1</i> -mut	RT -> TMZ	Astro	III	36	1	54	110

4942	<i>IDH1</i> -mut	RT ->TMZ	Astro	III	40	2	54	110
T03	<i>IDH1</i> -mut	RT -> TMZ	Astro	III	39	1	10	> 74
T04	<i>IDH1</i> -mut	RT -> TMZ	Astro	III	39	2	10	> 72
T07	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	31	1	9	30
T08	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	31	2	9	30
T09	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	39	1	5	39
T10	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	39	2	5	39
T11	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	54	1	6	12
T12	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	54	2	6	12
T13	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	53	1	9	18
T14	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	53	2	9	18
T15	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	50	1	9	20
T16	<i>IDH1</i> -wt	RT/TMZ->TMZ	GB	IV	50	2	9	20
1742	<i>IDH1</i> -mut	TMZ	Oligo	III	36	1	68	152
2875	<i>IDH1</i> -mut	TMZ	Oligo	III	41	2	68	152
3663	<i>IDH1</i> -mut	TMZ	Oligo	III	33	1	54	97
4943	<i>IDH1</i> -mut	TMZ	Oligo	III	38	2	54	97
3637	<i>IDH1</i> -mut	TMZ	Oligo	III	47	1	78	110
5446	<i>IDH1</i> -mut	TMZ	Oligo	III	53	2	78	110
3346	<i>IDH1</i> -mut	TMZ	Oligo	III	31	1	114	133
5837	<i>IDH1</i> -mut	TMZ	Oligo	III	40	2	114	133
2396	<i>IDH1</i> -mut	RT->TMZ	Oligo	III	33	1	30	52
2913	<i>IDH1</i> -mut	RT->TMZ	Oligo	III	35	2	30	52
1991	<i>IDH1</i> -mut	RT	Oligo	III	58	1	60	95
2930	<i>IDH1</i> -mut	RT	Oligo	III	63	2	60	95
5086	<i>IDH1</i> -mut	RT->PC	Oligo	III	57	1	16	23
5527	<i>IDH1</i> -mut	RT->PC	Oligo	III	58	2	16	23
2065	<i>IDH1</i> -mut	RT	Oligo	III	36	1	56	221
2996	<i>IDH1</i> -mut	RT	Oligo	III	40	2	56	221

Supplementary Table ST2: Basic clinical characteristics of study population. Patients have been characterized by their assigned identity number (ID), *IDH1* mutation status, treatment prior to tissue acquisition, histology and WHO grade (according to WHO classification 2016), age, resection status, PFS and OS. Abbreviations: Astro, Astrocytoma; GB, Glioblastoma; Histo, Histology; ID, Identifier; Oligo, oligodendroglioma; OS, overall survival; PC, procarbazine/lomustine; PCV, procarbazine/lomustine/vincristine; PFS, progression free survival; RT, radiotherapy; TMZ, temozolomide; WHO, World Health Organization; Further details on cohorts are outlined in Figure 1 and in the text.