## Thioredoxin system protein expression is associated with poor clinical outcome in adult and paediatric gliomas and medulloblastomas.

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Variable	Number of Patients	Percentage (%)
Age (years)		
≤59	10	55.6
>59	8	44.4
Gender		
Male	8	44.4
Female	10	55.6
Tumour Site		
Parietal	8	44.4
Temporal	5	27.8
Occipital	1	5.6
Frontal	4	22.2
Resection status		
Complete	14	77.8
Partial	4	22.2
Radiotherapy		
60 Gy	17	94.4
< 60 Gy	1	5.6
Temozolomide		
Yes	16	88.9
No	2	11.1
Gliadel wafer		
Yes	9	50.0
No	9	50.0
IDH-1		
Wild-type	17	94.4
Mutant	1	5.6
MGMT status		
Hypomethylated	9	90.0
Hypermethylated	1	10.0
N/A	8	
Survival status		
Living	4	28.6
Deceased	14	77.8

Supplementary Table 1: Clinicopathological variables of the adult GBM cohort.	
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Abbreviations: GBM, glioblastoma; IDH1, isocitrate dehydrogenase 1; MGMT, O-6-methylguanine DNA methyltransferase; N/A, not available.

	pLGG cohort	pHGG cohort
Variable	N (%)	N (%)
Age (years)		
≤3	25 (23.1)	23 (20.2)
>3	83 (76.9)	91 (79.8)
ND	18	23
Gender		
Male	57 (52.3)	80 (63.0)
Female	52 (47.7)	47 (37.0)
ND	17	10
Tumour site		
Supratentorial	38 (35.2)	84 (71.8)
Infratentorial	70 (64.8)	33 (28.2)
ND	18	20
Tumour grade		
Ι	92 (80.7)	0 (0.0)
II	22 (19.3)	0 (0.0)
III	0 (0.0)	35 (26.7)
IV	0 (0.0)	96 (73.3)
ND	12	6
Resection status		
Complete	NA	16 (15.8)
Partial	NA	44 (43.6)
Biopsy	NA	41 (40.6)
ND	126	36
Recurrent status		
Yes	43 (40.6)	96 (88.9)
No	63 (59.4)	12 (11.1)
ND	20	29
Survival status		
Living	77 (73.3)	19 (17.3)
Deceased	28 (26.7)	91 (82.7)
ND	21	27

Supplementary Table 2: Clinicopathological variables of the paediatric LGG and HGG cohorts.

Abbreviations: pLGG, paediatric low-grade glioma; pHGG, paediatric high-grade glioma; ND, not determined.

Variable	Number of Patients	Percentage (%)
Age (years)		
≤3	20	18.7
>3	87	81.3
ND	7	
Gender		
Male	78	72.2
Female	30	27.8
ND	6	
Histological Classification		
Classical	57	61.3
Desmoplastic	15	16.1
Large cell/anaplastic variant	16	17.2
Extensive nodularity	1	1.1
myogenic	4	4.3
ND	21	
Resection status		
Complete	52	61.2
Partial	33	38.8
ND	29	
Recurrent status		
Yes	42	44.7
No	52	55.3
ND	20	
Metastatic stage		
M0	56	60.9
M1	6	6.5
M2	10	10.9
M3	17	18.5
M4	3	3.3
ND	22	
Survival status		
Living	44	45.4
Deceased	53	54.6
ND	17	

Supplementary Table 3: Clinicopathological variables of the MB cohort.

Abbreviations: MB, medulloblastoma; ND, not determine



**Supplementary Figure 1: The specificity of anti-TrxR, anti-Trx and anti-TxNIP antibodies.** (a) TrxR, (b) Trx and (c) TxNIP expression across the panel of 6 brain (representing different tumour types) and 2 breast cancer cell lines were detected by Western blotting analysis. A single band corresponding to the molecular mass of the intended target was observed in the Western blotting studies which indicated the specificity of all antibodies. Figures present the representative blots of two independent experiments with lysates from cells with different passage numbers.

Moultong		Tumour	types	
Markers	Adult GBM	Paediatric LGG	Paediatric HGG	MB
Cytoplasmic TrxR	0.701	0.724	0.702	0.712
Nuclear TrxR	0.796	0.741	0.836	0.900
Cytoplasmic Trx	0.835	0.734	0.874	0.897
Nuclear Trx	0.790	0.700	0.925	0.892
Cytoplasmic TxNIP	0.808	0.740	0.873	0.956

Supplementary Table 4: Intra-class correlation coefficient analysis of Trx, TrxR and TxNIP scoring across different brain tumour cohorts.

Abbreviations: GBM, glioblastoma; LGG, low-grade glioma; HGG, high-grade glioma; MB, medulloblastoma.

Markers		Adult GBN	1	Paediatric	Paediatric	MB
Warkers	Core	Core Rim Invasive		LGG	HGG	
Cutonlasmia TauD	78	100	86	120	67	130
Cytopiasinic Trxk	(30-187)	(77-190)	(50-105)	(5-235)	(0-217)	(0-270)
Nuclear $\mathbf{T}_{\mathbf{m}}\mathbf{D}(0)$	47	72	79	55	36	64
Nuclear Trxk (%)	(7-95)	(15-100)	(45-98)	(0-95)	(0-100)	(0-100)
Cutonlasmia Tau	122	105	105	97	110	137
Cytopiasinic 11x	(5-185)	(0-198)	(90-140)	(0-273)	(0-283)	(0-280)
Note that $T_{max}(0/1)$	35	34	52	12	30	38
Nuclear Trx (%)	(0-90)	(0-90)	(10-85)	(0-88)	(0-98)	(0-99)
	175	140	170	173	188	150
Cytopiasinic TXNIP	(45-228)	(30-222)	(115-195)	(0-280)	(10-280)	(12-270)

Supplementary Table 5: The median scores and ranges of TrxR, Trx and TxNIP expression in all four brain tumour cohorts.

Abbreviations: GBM, glioblastoma; LGG, low-grade glioma; HGG, high-grade glioma; MB, medulloblastoma.

Supplementary Table 6: X-tile cut points for stratification of TrxR, Trx and TxNIP expression in all four brain tumour cohorts.

Markers		Adult GB	Μ	Paediatric	Paediatric	MB	
	Core	Rim	Invasive	LGG	HGG		
Cytoplasmic TrxR	78	100	86	167	160	180	
Nuclear TrxR (%)	63	80	85	77	58	58	
Cytoplasmic Trx	143	150	105	27	160	172	
Nuclear Trx (%)	38	23	60	14	57	60	
Cytoplasmic TxNIP	175	173	175	140	195	125	

Abbreviations: GBM, glioblastoma; LGG, low-grade glioma; HGG, high-grade glioma; MB, medulloblastoma.

Variable	Nucle	ear TrxR (Co	re)	Cytoplas	Cytoplasmic TrxR (Invasive)				
Variable	Low	High	<i>P</i> -value	Low	High	<i>P</i> -value			
Age (years)									
$\leq$ 59	7 (50.0%)	2 (14.3%)	0.580	5 (41.7%)	3 (25.0%)	0.545			
> 59	3 (21.4%)	2 (14.3%)		1 (8.3%)	3 (25.0%)				
Gender									
Male	6 (42.9%)	1 (7.1%)	0.559	0 (0.0%)	5 (41.7%)	0.015			
Female	4 (28.6%)	3 (21.4%)		6 (50.0%)	1 (8.3%)				
Tumour site									
Parietal	7 (50.0%)	0 (0.0%)	0.008	1 (8.3%)	3 (25.0%)	0.096			
Temporal	0 (0.0%)	3 (21.4%)		4 (33.3%)	0 (0.0%)				
Occipital	1 (7.1%)	0 (0.0%)		0 (0.0%)	1 (8.3%)				
Frontal	2 (14.3%)	1 (7.1%)		1 (8.3%)	2 (16.7%)				
Resection status									
Complete	6 (42.9%)	7 (50.0%)	1.000	2 (16.7%)	7 (58.3%)	0.236			
Partial	1 (7.1%)	0 (0.0%)		2 (16.7%)	1 (8.3%)				
IDH-1 status									
Wild-type	10 (71.4%)	3 (21.4%)	0.286	5 (41.7%)	6 (50.0%)	1.000			
Mutant	0 (0.0%)	1 (7.1%)		1 (8.3%)	0 (0.0%)				

Supplementary Table 7: Associations between TrxR expression and clinicopathological variables in adult glioblastoma (Core + Invasive margin).

The *P* values are resultant from the Pearson  $\chi^2$  test of association or Fisher's Exact test if a cell count was less than five. Significant *P* values are indicated in bold.

	Cyto	plasmic T	rxR	Nu	Nuclear TrxR			Cytoplasmic TxNIP		
variable	Low	High	Р	Low	High	Р	Low	High	Р	
Age										
≤3	14 (17.9%)	3 (3.8%)	0.682	9 (11.5%)	8 (10.3%)	0.072	5 (6.1%)	12 (14.6%)	0.644	
>3	54 (69.2%)	7 (9.0%)		46 (59%)	15 (19.2%)		23 (28.0%)	42 (51.2%)		
Gender										
Male	31 (46.3%)	5 (7.5%)	0.437	26 (38.8%)	10 (14.9%)	0.498	15 (21.1%)	22 (31.0%)	0.474	
Female	29 (43.3%)	2 (3.0%)		20 (29.9%)	11 (16.4%)		11 (15.5%)	23 (32.4%)		
Tumour site										
Supratentorial	14 (21.2%)	6 (9.1%)	0.002	17 (25.8%)	3 (4.5%)	0.083	5 (7.1%)	17 (24.3%)	0.125	
Infratentorial	45 (68.2%)	1 (1.5%)		28 (42.4%)	18 (27.3%)		20 (28.6%)	28 (40.0%)		
Recurrence										
No	43 (66.2%)	1 (1.5%)	0.004	25 (38.5%)	19 (29.2%)	0.001	12 (17.4%)	34 (49.3%)	0.013	
Yes	15 (23.1%)	6 (9.2%)		20 (30.8%)	1 (1.5%)		13 (18.8%)	10 (14.5%)		

Supplementary	Table	8:	Associations	between	TrxR	and	TxNIP	expression	and	clinicopathological
variables in pae	diatric l	low	-grade glioma	ı.						

The *P* values are resultant from the Pearson  $\chi^2$  test of association or Fisher's Exact test if a cell count was less than five. Significant *P* values are indicated in bold. The table does not include the number of observations where clinicopathological data or scores were not available.

Variable	Cyte	oplasmic T	'nx	N	Nuclear Trx			
variable	Low	High	Р	Low	High	Р		
Age								
≤3	2 (2.6%)	15 (19.2%)	0.643	8 (10.3%)	9 (11.5%)	0.526		
>3	5 (6.4)	56 (71.8%)		34 (43.6%)	27 (34.6%)			
Gender								
Male	4 (5.9%)	32 (47.1%)	1.000	17 (25.0%)	19 (27.9%)	0.316		
Female	3 (4.4%)	29 (42.6%)		19 (27.9%)	13 (19.1%)			
Tumour site								
Supratentorial	2 (30.%)	19 (28.4%)	1.000	12 (17.9%)	9 (13.4%)	0.587		
Infratentorial	5 (7.5%)	41 (61.2%)		23 (34.3%)	23 (34.3%)			
Recurrence								
No	2 (3.0%)	43 (65.2%)	0.029	22 (33.3%)	23 (34.8%)	0.324		
Yes	5 (7.6%)	16 (24.2%)		13 (19.7%)	8 (12.1%)			

Supplementary Table 9: Associations between Trx expression and clinicopathological variables in paediatric low-grade glioma.

The *P* values are resultant from the Pearson  $\chi^2$  test of association or Fisher's Exact test if a cell count was less than five. Significant *P* values are indicated in bold. The table does not include the number of observations where clinicopathological data or scores were not available.

X7	Cyte	plasmic T	rx	N	uclear Trx		Cytoplasmic TxNIP		
Variable	Low	High	Р	Low	High	Р	Low	High	Р
Age									
≤3	16 (15.0%)	2 (1.9%)	0.145	17 (15.7%)	1 (0.9%)	0.188	6 (6.3%)	11 (11.6%)	0.114
>3	62 (57.9%)	27 (25.2%)		72 (66.7%)	18 (16.7%)		44 (46.3%)	34 (35.8%)	
Gender									
Male	41 (51.3%)	9 (11.3%)	0.564	41 (51.3%)	9 (11.3%)	0.520	21 (30.0%)	25 (35.7%)	0.314
Female	23 (28.8%)	7 (8.8%)		27 (33.8%)	3 (3.8%)		14 (20.0%)	10 (14.3%)	
Histological									
subtypes	16	2		16	2		12	3	
AA	(20.8%)	(2.6%)	0.008	(20.8%)	(2.6%)	0.189	(17.9%)	(4.5%)	0.009
GBM	42	9		44	7		17	28	
	(54.5%)	(11.7%)		(57.1%)	(9.1%)		(25.4%)	(41.8%)	
AO	(3.9%)	(6.5%)		(6.5%)	(3.9%)		(7.5%)	(3.0%)	
Tumour site									
Supratentorial	52 (67.5%)	14 (18.2%)	1.000	55 (71.4%)	11 (14.3%)	1.000	25 (36.8%)	32 (47.1%)	0.021
Infratentorial	9 (11.7%)	2 (2.6%)		10 (13.0%)	1 (1.3%)		9 (13.2%)	2 (2.9%)	
WHO grade	. ,	. ,			. ,		. ,		
III	19 (24.7%)	6 (7.8%)	0.545	21 (27.3%)	4 (5.2%)	0.741	16 (23.9%)	5 (7.5%)	0.005
IV	43	9		45	7		18	28	
Extent of resection	(55.8%)	(11.7%)		(58.4%)	(9.1%)		(26.9%)	(41.8%)	
Gross total	12 (18.5%)	1 (1.5%)	0.446	13 (20.0%)	0 (0.0%)	0.141	8 (13.3%)	4 (6.7%)	0.333
Subtotal	25 (38.5%)	8 (12.3%)		25 (38.5%)	8 (12.3%)		13 (21.7%)	18 (30.0%)	
Biopsy	15 (23.1%)	4 (6.2%)		16 (24.6%)	3 (4.6%)		9 (15.0%)	8 (13.3%)	
Recurrence									
No	10 (14.5%)	0 (0.0%)	0.189	9 (13.0%)	1 (1.4%)	1.000	7 (11.3%)	3 (4.8%)	0.167
Yes	46 (66.7%)	13 (18.8%)		50 (72.5%)	9 (13.0%)		24 (38.7%)	28 (45.2%)	

Supplementary Table 10: Associations between Trx and TxNIP expression and clinicopathological variables in paediatric high-grade glioma.

The *P* values are resultant from the Pearson  $\chi^2$  test of association or Fisher's Exact test if a cell count was less than five. Significant *P* values are indicated in bold. Abbreviations: AA, anaplastic astrocytoma; GBM, glioblastoma multiforme; AO, anaplastic oligodendroglioma.

Variable	Cytoplasmic Trx			Nuclear Trx		
	Low	High	Р	Low	High	Р
Age						
≤3	7 (8.8%)	5 (6.3%)	0.495	9 (11.3%)	3 (3.8%)	0.711
>3	49 (61.3%)	19 (23.8%)		54 (67.5%)	14 (17.5%)	
Gender						
Male	35 (46.7%)	15 (20.0%)	0.859	40 (53.3%)	10 (13.3%)	0.690
Female	17 (22.7%)	8 (10.7%)		19 (25.3%)	6 (8.0%)	
Histological subtypes						
Classic	28 (45.2%)	15 (24.2%)	0.149	35 (56.5%)	8 (12.9%)	0.041
Desmoplastic/nodular	7 (11.3%)	1 (1.6%)		7 (11.3%)	1 (1.6%)	
MB with extensive nodularity	0 (0.0%)	2 (3.2%)		0 (0.0%)	2 (3.2%)	
Large cell/ anaplastic	5 (8.1%)	2 (3.2%)		4 (6.5%)	3 (4.8%)	
Medullomyoblastoma	2 (3.2%)	0 (0.0%)		2 (3.2%)	0 (0.0%)	
Extent of resection						
Complete	26 (44.1%)	11 (18.6%)	0.559	20 (33.9%)	2 (3.4%)	0.179
Partial	17 (28.8%)	5 (8.5%)		27 (45.8%)	10 (16.9%)	
Recurrence						
Yes	18 (28.1%)	14 (21.9%)	0.031	21 (32.8%)	11 (17.2%)	0.016
No	26 (40.6%)	6 (9.4%)		29 (45.3%)	3 (4.7%)	
Metastasis						
Absence	28 (45.9%)	10 (16.4%)	0.482	31 (50.8%)	7 (11.5%)	1.000
Presence	15 (24.6%)	8 (13.1%)		19 (31.1%)	4 (6.6%)	

Supplementary Table 11: Associations between Trx expression and clinicopathological variables in paediatric medulloblastoma.

The *P* values are resultant from the Pearson  $\chi^2$  test of association or Fisher's Exact test if a cell count was less than five. Significant *P* values are indicated in bold.



Supplementary Figure 2: Box plots of protein expression in different WHO grades of paediatric gliomas. The expression of cytoplasmic TrxR (a) is significantly higher in grade I gliomas than both grade III (*P*=0.010) and grade IV gliomas (*P*=0.001). The expression of nuclear TrxR (b) is also significantly greater in grade I gliomas than grade IV gliomas (*P*=0.001), but not grade III gliomas (*P*=0.085). No significant difference of both cytoplasmic Trx (c) (*P*=0.876) and nuclear Trx (d) (*P*=0.448) expression is detected between tumour grades. A significantly higher expression of TxNIP (e) is noted in grade IV gliomas compared to grade I gliomas (*P*=0.001). \*\* represents  $P \leq 0.01$ .