

<b>Sample</b>	<b>Age</b>	<b>Sex</b>	<b>Primary/Recurrent (P/R)?</b>
BT241	66	F	R
BT972	53	M	R
BT594	50	M	P
BT618	67	F	R
BT566	55	F	R
BT155	66	F	P
BT162	57	F	P
BT163	56	M	P
BT168	46	F	P
BT189	67	F	P
MBT190	27	F	R
BT191	50	F	P
BT956	70	F	P
BT698	57	F	P
MBT225	27	M	R
MBT173	69	F	R
BT566	55	F	R
MBT103	52	F	P
MBT121	67	F	P
BT458	81	M	P
BT428	63	F	P
BT954	65	M	
BT799	77	F	P
MBT63	68	F	P
MBT27	56	M	P
BT935	53	F	P
BT459	60	M	P
BT778	54	F	P

**Sup Table 1**

Ensembl ID	Gene symbol	Log (FC)			Combined Differential Gene Expression	
		BT241	GBM8	GBM4	Log (FC)	P-value
ENSG00000175592	FOSL1	-3.0876903	-2.9685466	-1.4827605	-1.3302582	0.002764603
ENSG00000205927	OLIG2	0.99188628	1.51882472	2.02962725	0.95394172	0.027641632
ENSG00000112715	VEGFA	-1.079047	-2.993489	-0.2838893	-0.8847545	0.040829495
ENSG00000170558	CDH2	-1.8132171	-0.4377818	-1.1326002	-0.633147	0.148275349
ENSG00000011422	PLAUR	-2.0416545	-1.1341974	-2.2947466	-0.4965695	0.206493437
ENSG00000121966	CXCR4	-1.5615955	-2.8821719	-1.3521936	-0.361023	1
ENSG00000026508	CD44	-0.6256976	-0.0583328	-1.5159146	-0.3010297	0.482572636
ENSG00000181449	SOX2	-0.0175167	-0.8038392	-0.905513	-0.3279201	0.417025332

Sup Table 2