

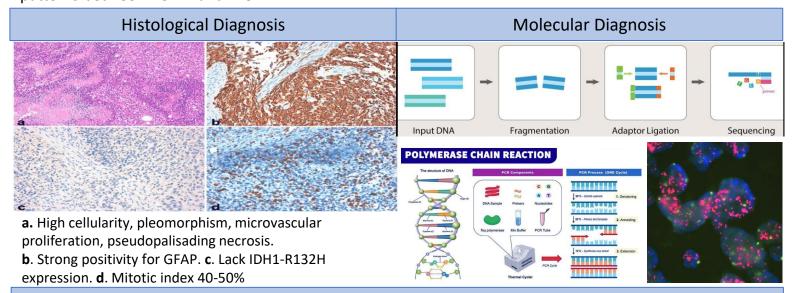




Understanding the differences between glioblastoma patients based on histological or molecular criteria

Background & Methods

WHO brain tumour diagnostic criteria changed in 2021 to incorporate advances in understanding gliomagenesis. With the incorporation of molecularly diagnosed GBM (mGBM) to histologically diagnosed GBM (hGBM) we sought to investigate whether treatment and prognosis had changed as a result. mGBM criteria includes EGFR amplification, TERT promoter mutation or combined gain of chromosome 7 and loss of chromosome 10. Using the Histo-Mol GBM Collaborative database, a large individual patient database of patients diagnosed with pathologically confirmed glioblastoma in 2021 we investigated differences and patterns between hGBM and mGBM.



Results

1828 patients across 51 participating centres diagnosed between 1 January 2021 - 31 December 2021. 3 centres have local approval with data collection due to start. Molecular testing performed according to

local practice				hGBM	mGBM
	•	Data collection complete	Total nations	1752	75
		Local approvals gained	Total patients	1753	75
		Not yet involved	Time from MRI → surgery	14 days	23 days
			Time from surgery → radiotherapy	40 days	46 days
		A ser of the	% alive at time of data analysis	13%	15%
			Chemotherapy	79%	80%
			Biopsy	30.3%	69.3%
	Sang.	who for the state of the state	Resection	69.3%	30.7%
	En /		EGFR mutation	37.3%	42.4%
	The state of the s		Chromosome +7/-10	36.9%	52.8%
			pTERT mutation	78.4%	89.4%

Conclusion

The data for this study was collected from 2021, as WHO recommendations being implemented, there were delays to treatment due to processing and additional tests, it would be interesting to see if these have improved. Ultimately the overall treatment regime remains largely unchanged, but this opens the possibility for further biomarkers to be identified and the development of targeted / adjuvant immunotherapies.