

Methylation profiling report

Supplier information

Sample identifier: GCGR-NS12ST_A-CDKN2AB-EGFRVIII-GFP-LUC
Sentrix ID: 205715840125_R02C01
Material type: DNA-KRYO
Gender: NA
Supplier diagnosis: GBM

Automatic prediction

Array type:	EPIC	
Material type:	DNA-KRYO	✓
Gender:	male	!
Legend:	✓ Ok	! Supplier information or prediction not available
		✗ Warning, mismatch of prediction and supplier information

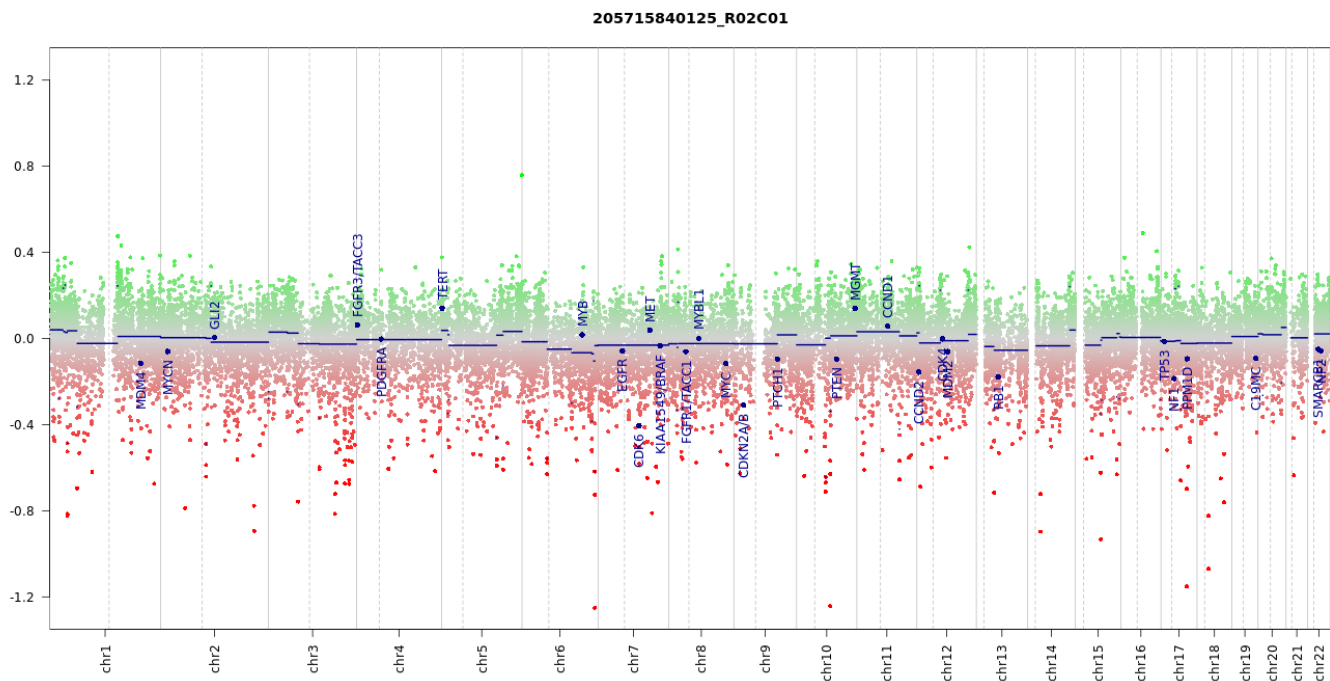
Brain tumor classifier results (11b4)

Not classifiable. No score is greater 0.3

Legend: ✓ Match (score ≥ 0.9) ✗ No match (score < 0.9): possibly still relevant for low tumor content and low DNA quality cases. ● Match to MC family member (score ≥ 0.5)

Class descriptions

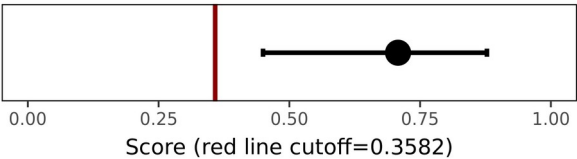
Copy number variation profile



Depiction of chromosome 1 to 22 (and X/Y if automatic prediction was successful). Gains/amplifications represent positive, losses negative deviations from the baseline. 29 brain tumor relevant gene regions are highlighted for easier assessment. (see Hovestadt & Zapatka, <http://www.bioconductor.org/packages/devel/bioc/html/conumee.html>)

MGMT promotor methylation (MGMT-STP27)

MGMT promotor status prediction



Status	Estimated	CI lower	CI upper
methyated	0.70782	0.44969	0.87777

(see Bady et al, J Mol Diagn 2016; 18(3):350-61)

Disclaimer

Classification using methylation profiling is a tool/website for research use only, it is not verified and has not been clinically validated and, therefore, must not be used for diagnostic purposes. This tool/website is not HIPAA compliant.

Run information

Report: report_website_mnp_brain_v11b4_sample (Version 2.1)

Task version:

Task	Version
idat_preprocess	2.0.1
idat_qc	2.0.1
idat_predictBrain	2.0.1
idat_rs_gender	2.0.1
idat_cnvp	3.0.1
idat_mgmt	2.0.1
report_website_mnp_brain_v11b4_research	2.1
report_website_mnp_brain_v11b4_sample	2.1
idat_predictBrain	12.5