

Methylation profiling report

Supplier information

Sample identifier: GCGR-NS12ST_A-NF1-EGFRvIII-GFP-LUC
Sentrix ID: 205715840020_R08C01
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)

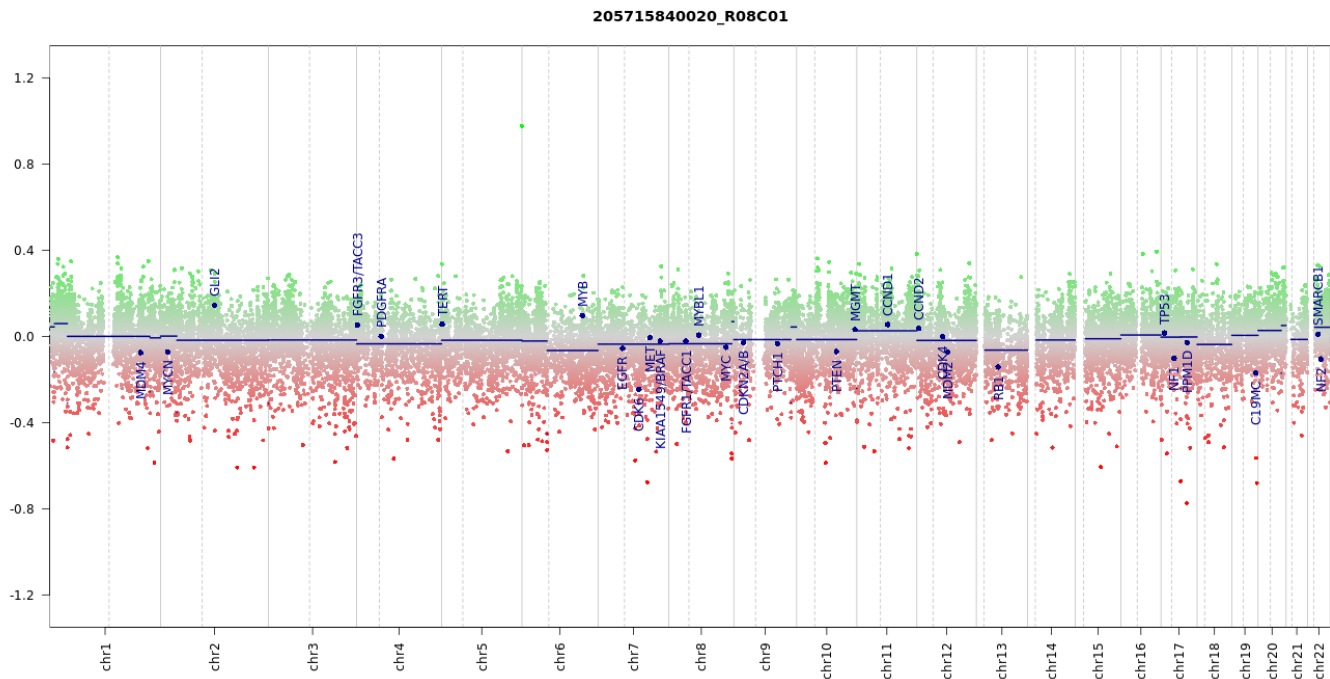
Methylation classes (MCs with score ≥ 0.3)

	Score	Interpretation
methylation class infantile hemispheric glioma	0.35	no match ✗
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

Methylation class infantile hemispheric glioma: The methylation class "infantile hemispheric glioma" comprises tumors with widely varying morphology, commonly including features more typical of higher grade lesions especially glioblastoma. These tumors are located supratentorially, all cases so far were observed in infants; median age is 0 years (age range 0 to 1). Copy number alterations are scarce, and typical molecular features of this class are not currently known. The name given here is provisional.

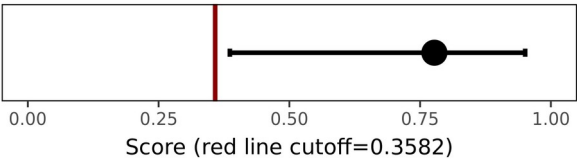
Copy number variation profile



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	0.77728	0.38624	0.95087

(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