Methylation profiling report

Supplier information

Sample identifier:	GCGR-NS9FB_B-NF1-GFP-LUC
Sentrix ID:	205715840095_R07C01
Material type:	DNA-KRYO
Gender:	NA
Supplier diagnosis:	GBM

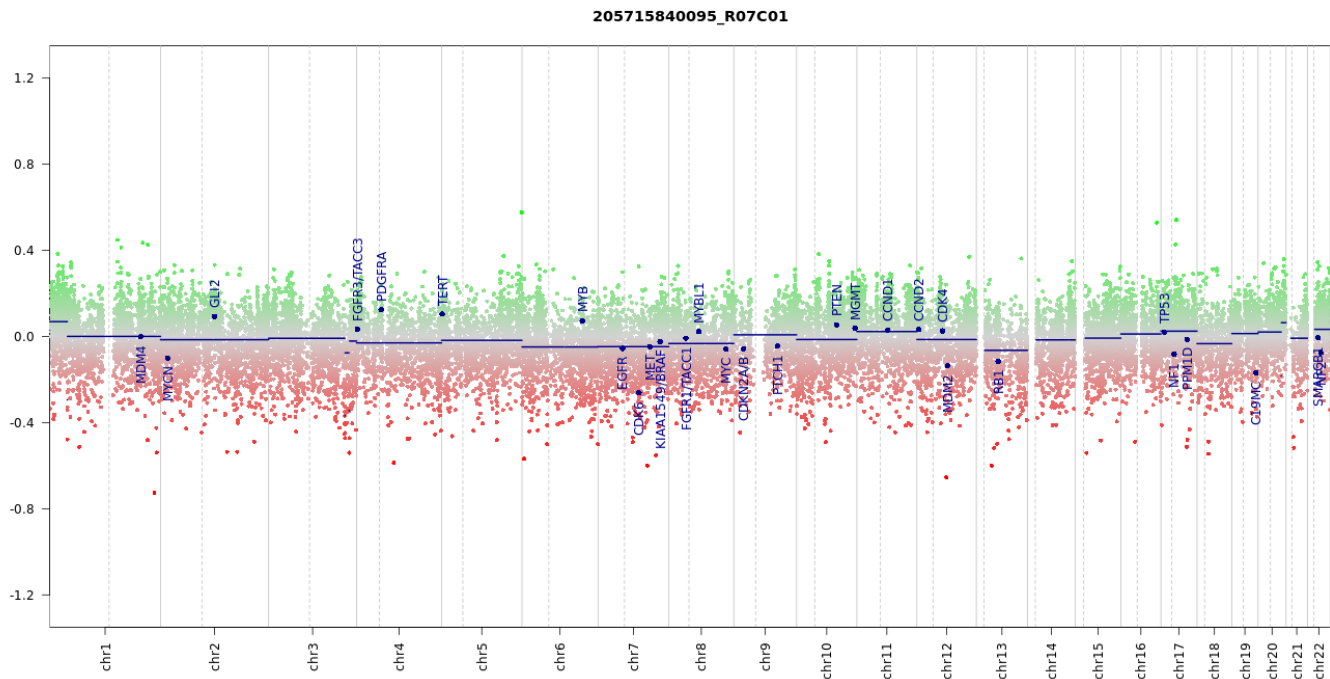| Automatic prediction		
Array type:	EPIC	
Material type:	DNA-KRYO	✔
Gender:	female	▯
Legend: ✔ Ok	▯ Supplier information or prediction not available	✗ Warning, missmatch of prediction and supplier information

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)
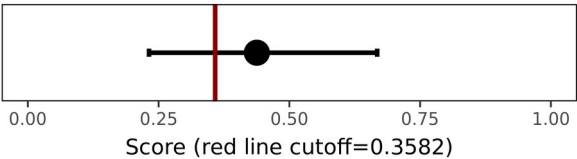
Class descriptions

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
methylated	0.43787	0.23175	0.66793

(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