





Methylation profiling report

Sample identifier: sampleName1573854502

Sentrix ID: 203057570095_R08C01

Material type: KRYO DNA

Gender: NA

Supplier diagnosis: -

Supplier information

Automatic prediction		
Array type:	EPIC	
Material type:	KRYO DNA	~
Gender:	male	
Legend: ✔ OK Supplier information or prediction not available		

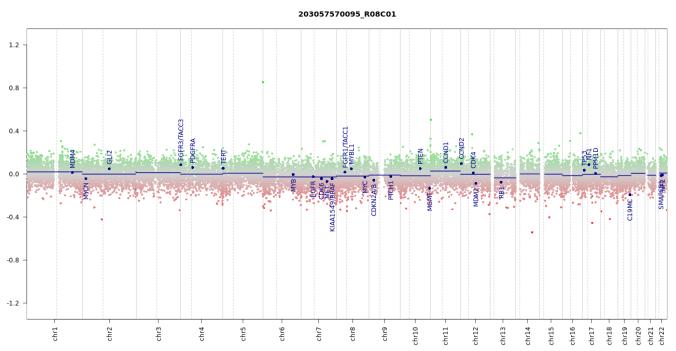
Brain tumor methylation classifier results (v11b4)

Methylation classes (MCs with score >= 0.3)	Calibrated score	Interpretation	on
methylation class infantile hemispheric glioma	0.8	no match	X
Legend: ✓ Match (score >= 0.9) × No match (score < 0.9): possibly still relevant for low tumor content and quality cases.	d low DNA • Match to (score >=	MC family meml : 0.5)	ber

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



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

Disclaimer

Classification using methylation profiling is a research tool under development, it is not verified and has not been clinically validated. Implementation of the results in a clinical setting is in the sole responsibility of the treating physician. Intended for non-commercial use only.

CI lower

0.06796

CI upper

0.39894

Run information

Report: idat_reportBrain_v11b4_sample Version 2.0 Task version:

Task	Version
idat_qc	2.0
idat_predictBrain	2.1
idat_rs_gender	2.0
idat_predictMGMT	2.0
idat_cnvp	3.0
idat_reportBrain_v11b4	2.0