





Methylation profiling report

Supplier information

Sample identifier: sampleName1540553397
Sentrix ID: 202274990030_R03C01

Material type: KRYO DNA

Gender: NA

Supplier diagnosis: -

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

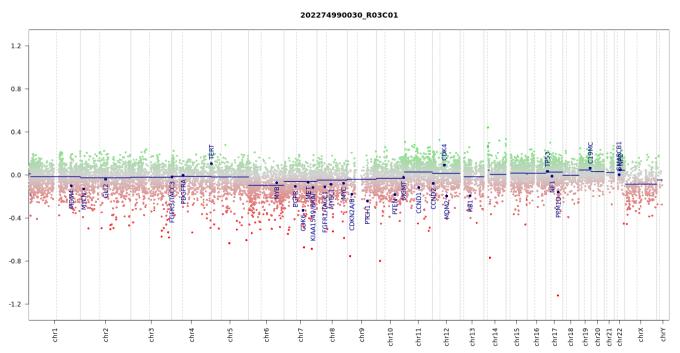
Brain tumor methylation classifier results (v11b4)

Methylation classes (MCs v	vith score >= 0.3)	Calibrated score	Interpretati	on
methylation class infantile he	mispheric glioma	0.63	no match	X
Legend: ✓ Match (score >= 0.9)	X No match (score < 0.9): possibly still relevant for low tumor content an quality cases.	d low DNA • Match to (score >=	MC family mem	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.

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