





Methylation profiling report

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Supplier information

Sample identifier: sampleName1540720713
Sentrix ID: 202273260119_R07C01

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	Warning, missmatch of predictio and supplier information	n

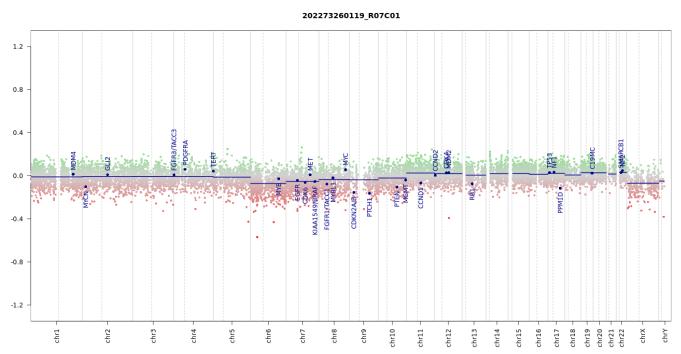
Brain tumor methylation classifier results (v11b4)

Methylation classes (MCs w	ith score >= 0.3)	Calibrated score	Interpretati	on
methylation class control tissu	e, cerebellar hemisphere	0.53	no match	X
Legend: ✓ Match (score >= 0.9)	X No match (score < 0.9): possibly still relevant for low tumor content and quality cases.	d low DNA • Match to (score >=	MC family mem	ber

Class descriptions

Methylation class control tissue, cerebellar hemisphere: The methylation class "control tissue, cerebellar hemisphere" is comprised of normal tissue samples from this specific anatomic structure. In case tumor samples display this molecular pattern, it is typically an indication of low tumor cell content in the analyzed material and indicates that the extracted DNA is likely not suitable for classification by methylation profiling.

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