Methylation profiling report

Supplier information
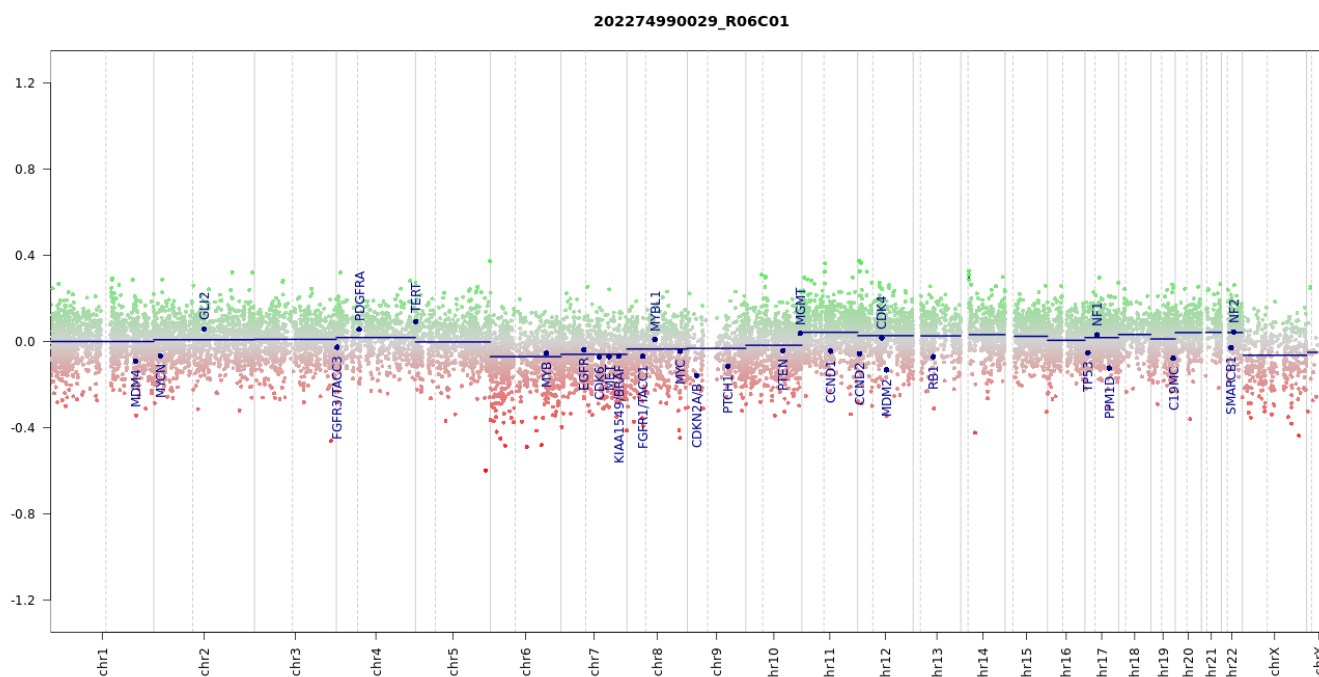
		Automatic prediction		
Sample identifier:	**sampleName1540748675**	Array type:	**EPIC**	
Sentrix ID:	**202274990029_R06C01**	Material type:	**KRYO DNA**	✔
Material type:	**KRYO DNA**	Gender:	**male**	❗
Gender:	**NA**			
Supplier diagnosis:	**-**	Legend: ✔ OK ❗ Supplier information or prediction not available	✖ Warning, missmatch of prediction and supplier information	

Brain tumor methylation classifier results (v11b4) - No matching methylation classes with calibrated score >= 0.3
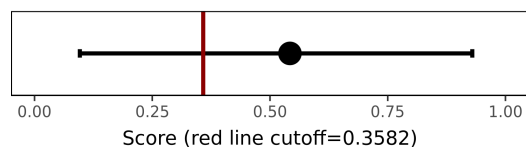
Copy number variation profile

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

Score (red line cutoff=0.3582)

Status	Estimated	CI lower	CI upper
methylated	0.54239	0.09619	0.92958

(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