





## **Methylation profiling report**

**Supplier information** 

Sample identifier: sampleName1540574349 **Automatic prediction** Sentrix ID: 202273260142\_R05C01 Array type: **EPIC** Material type: **KRYO DNA** Material type: **KRYO DNA** Gender: Gender: male NA Supplier diagnosis: Warning, missmatch of prediction prediction not available and supplier information

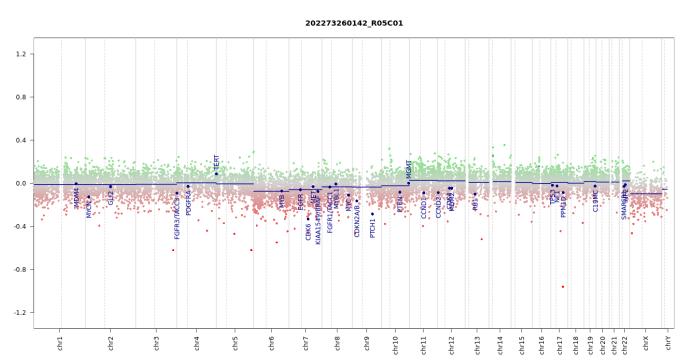
#### Brain tumor methylation classifier results (v11b4)

Methylation classes (MCs with score >= 0.3)	Calibrated score	Interpretati	on
methylation class diffuse midline glioma H3 K27M mutant	0.32	no match	X
Legend: ✓ Match (score >= 0.9) × No match (score < 0.9): possibly still relevant for low tumor content and	d low DNA • Match to (score >=	MC family mem : 0.5)	ber

#### Class descriptions

Methylation class diffuse midline glioma H3 K27M mutant: The methylation class "diffuse midline glioma H3 K27M mutant" is comprised of tumors with a histological diagnosis of diffuse midline glioma, H3 K27M mutant. The tumors are located in midline structures (thalamus, cerebellum, brainstem, spine). Median age is 12 years (range 1 to 54). Tumors of this class harbor mutations of codon 27 in one of the genes encoding histone 3 (most commonly H3F3A, but also the H3.1 genes HIST1H3A/B/C and very rarely H3.2). Additional mutations in ATRX and TP53/PPM1D are common. In brainstem tumors, ACVR1 mutations are also recurrently observed. Prognosis of this class is generally very poor, although rare lower grade lesions with K27M mutation have also been reported.

#### 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