

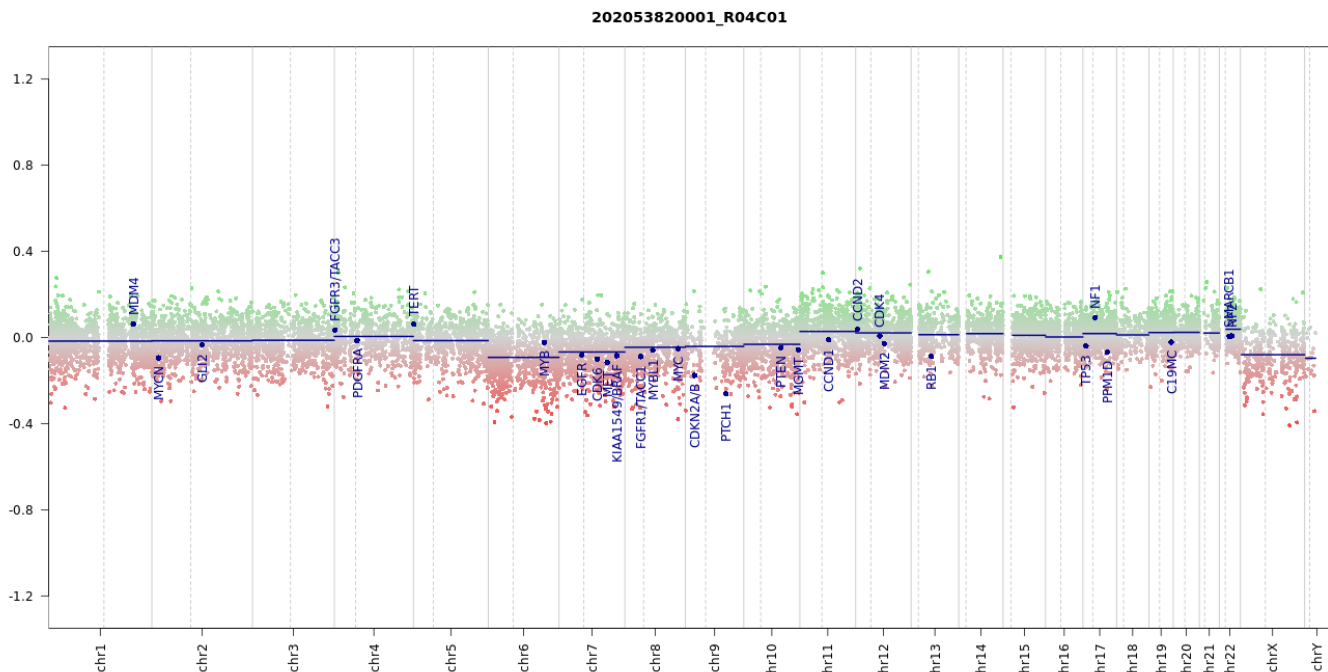
## Methylation profiling report

### General information

Sentrix ID: **202053820001\_R04C01**  
 Array type: **EPIC**  
 Material type: **KRYO DNA**  
 Gender: **male**

**Brain tumor methylation classifier results (v11b4) - No matching methylation classes with calibrated score  $\geq 0.3$**

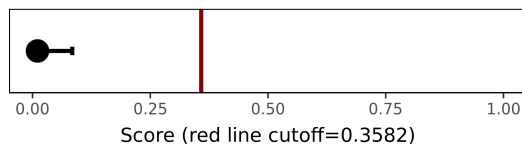
### Copy number variation profile



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



**Status**  
unmethylated

**Estimated**  
0.0105

**CI lower**  
0.00122

**CI upper**  
0.08431

(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 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