Calibrated scores Heidelberg vs. New York

- Identical classification: 96% (51/53)
- Discrepant classification: 0% (0/53)
- No match in one center: 4% (2/53)

b

- Analyzed in Heidelberg
- Analyzed in New York

- Amplification EGFR, deletion CDKN2A/B, gain Chr 7, loss Chr 10 in glioblastoma
- Loss of Chr 22 with homozygous deletion of SMARCB1 in ATRT
- Isochromosome 17q in medulloblastoma

c

- Illumina 450k vs Illumina EPIC
- Raw score EPIC vs Raw score 450k

d

- Illumina 450k vs High-coverage WGBS
- Raw score WGBS vs Raw score 450k