

NYU cohort
22 samples

Sanger Sequencing
MLPA
DNA copy number
RNA sequencing
Pyrosequencing
IHC

63.6% of BAP1 alterations detected by
at least one of the molecular
techniques utilized
63.6% nuclear BAP1 loss by IHC

NMVB cohort
70 samples

IHC

67.1% nuclear BAP1 loss