

# Biomarker Ontology

Alteration Type	Description
Mutation	A genomic alteration resulting in the substitution, insertion, or deletion of nucleotides
Amplification	Copy Number Amplification
Deletion	Copy Number Deletion
Fusion	Rearrangement of chromosomal material that results in complete or partial sequences of two or more separate genes fused to create a novel transcript
Overexpression	Protein Overexpression
Loss	Protein Loss
Any	Any Type of Altered Status
Wildtype	Wildtype
Positive	Positive for expression of the protein. Dependent on the biomarker, a certain threshold of detection may be required for classification of the sample as "positive"
Negative	Lack of protein expression
Non-Amplified	Lack of copy number amplification
Instability Intermediate	Instability of microsatellite loci was detected at an intermediate level
High	Level of the biomarker is considered "high" by the testing platform. Currently used to describe the level of tumor mutation burden
Intermediate	Level of the biomarker is considered "intermediate" by the testing platform. Currently used to describe the level of tumor mutation burden
Low	Level of the biomarker is considered "low" by the testing platform. Currently used to describe the level of tumor mutation burden
Instability High	Instability of microsatellite loci was detected at a high level
Instability Low	Instability of microsatellite loci was detected at a low level
Stable	Instability of microsatellite loci was not detected
Tumor Positive	Expression of the biomarker was detected specifically within tumor cells
Immunocyte Positive	Expression of the biomarker was detected specifically within immunocytes
Tumor Negative	Expression of the biomarker was not detected specifically within tumor cells
Immunocyte Negative	Expression of the biomarker was not detected specifically within immunocytes
Unknown	Biomarker Status is Unknown
Uninterpretable	Biomarker Status was tested but results are uninterpretable
Clonal	Expression of the biomarker was clonal
Equivocal	Biomarker status could not be unequivocally determined
Promoter Methylation Positive	DNA methylation was detected at the promoter region of the gene
Promoter Methylation Negative	DNA methylation was not detected at the promoter region of the gene
Alternative Transcript	Expression of an alternative transcript lacking official nomenclature was detected
Phospho Overexpression	Overexpression of a phosphorylated form of the protein was detected