

Study of origin



Profiled for copy number alterations



EZH2

8%



EZH1

2.2%



SUZ12

2.2%

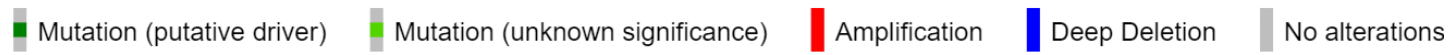


EED

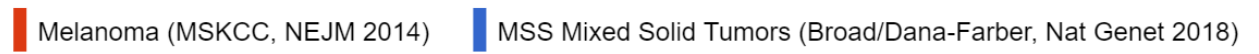
2.2%



Genetic Alteration



Study of origin



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