Bethesda classification

No mutation	■ None
NBNR mutation	■ EZH1
	■ IDICER1
	■ EIF1AX
	■ PTEN
RAS mutation	■ NRAS
	■ KRAS
	■ HRAS
BRAF mutation	■ BRAF^{V600E}
Double mutation	■ Double mutation

Supplemental Fig. S3. Percentage distribution of detected mutations of nodules by Bethesda classification groups. NBNR, non-*BRAF*-non-*RAS*; EZH1, enhancer of zeste 1 polycomb repressive complex 2 subunit; DICER1, dicer 1, ribonuclease III; EIF1AX, eukaryotic translation initiation factor 1A X-linked; PTEN, phosphatase and tensin homolog.