

Table S5. Comparative analysis of VAF(%) observed in our analysis with that reported in previous published data of the indicated genes.

COMPARATIVE ANALYSIS OF VAF% ACCORDING TO THE PUBLISHED DATA			
MUTATED GENES	VAF% OBSERVED	VAF% EXPECTED (RANGE%)	REFERENCES (doi)
GNA11	59.1%	24.2%-60%	10.3390/cancers14133066, 10.1038/bjc.2013.299, 10.1007/s12253-017-0371-7, 10.1136/bjo.2009.174417, 10.1111/aos.14760, 10.21873/anticancerres.12010
GNAQ	29.8%	24.2%-53.3%	
BAP1	41%	40%-47%	10.1038/s41598-021-88390-6, 10.1126/science.1194472, 10.1038/s41467-020-15606-0, 10.3390/cancers14174105