Table S1. Mutations identified in protein phosphatase 2A (PP2A) scaffolding subunits.

Subunit (Gene)	Tumor Type	Frequency	Mutation	Functional Consequence
Aα (PPP2R1A)	Melanoma [1]	1/14 (7%)	Arg-418-Trp	Defective binding to A and/or c subunits [2]. Decreased B regulatory subunit and cα stability [3]
	Lung Cancer [1]	1/16 (6%)	Glu-64-Asp	
	Breast Cancer [1]	2/43 (7%)	Glu-64-Gly Frameshift at AA170	
	MCF-7 breast cancer cell line [4]		Decreased expression	
	Gliomas [5]	25/58 (43%)	Decreased expression	
	Ovarian cancer [6]	3/42 (7%)	Arg-183-Gly Arg-183-Trp Arg-182-Trp	Mutations predicted to impair subunit binding
Aβ (PPP2R1B)	Lung Cancer [1]	2/16 (13%)	Exon 9 deletion	
	Breast Cancer [1]	4/43 (9%)	Exon deletion/abnormal splicing	
	Colorectal Cancers	4/30 (13%) [7]	Glyc-15-Ala & Leu-499-Iso Val-498-Glu Val-500-Gly Ser-365-Pro	Mutations predicted to impair subunit binding
		2/13 (15%) [8]		
		5/50 (10%) [9]	Missense: 4/50 (8%) Homozygous deletions: 1/50 (2%)	
	Lung Cancer [8]	5/33 (15%) – Primary 4/70 (6%) cell lines	Deletions, frameshifts, and point mutations	Decreased affinity for PP2Ac

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