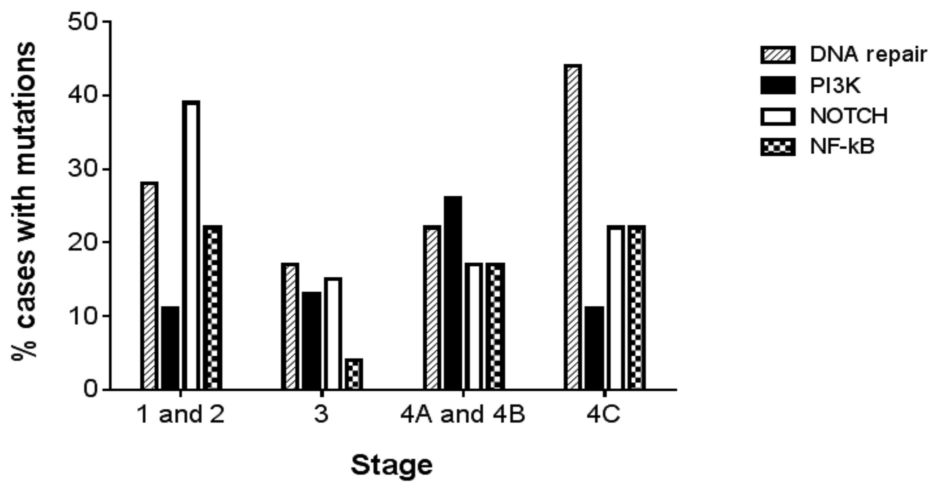


Title: Exome Sequencing Identifies Potentially Druggable Mutations in Nasopharyngeal Carcinoma

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Supplemental Figure legends:

Supplemental Figure 1. Correlation between NPC tumor stages and the frequencies of mutations detected in the 4 enriched druggable pathways.



Stage	Number of Cases (Percentage)				
	Total (n=98)	DNA repair	PI3K	NOTCH	NF-κB
1 and 2	18	5/18 (28%)	2/18 (11%)	7/18 (39%)	4/18 (22%)
3	48	8/48 (17%)	6/48 (13%)	7/48 (15%)	2/48 (4%)
4A and 4B	23	5/23 (22%)	6/23 (26%)	4/23 (17%)	4/23 (17%)
4C	9	4/9 (44%)	1/9 (11%)	2/9 (22%)	2/9 (22%)