S4 Table. *PIK3CA* alteration analysis

	Chinese(n=62)	Caucasian(n=82)
***PIK3CA* alterations**		
All	23(37%)	30(37%)
Mutations	19(31%)	21(26%)
Amplification	6(10%)	9(11%)
***PIK3CA* mutant variants**		
E542K	5(8%)	5(6%)
E545K/Q	5(8%)	9(11%)
H1047R/L	4(6%)	2(2%)