

**Supplementary Table 2. The prevalence of each genotype subgroup when combining *BRAF* and *TERT* promoter mutations**

Author	Total patients	Mean age	No. of patients			
			<i>BRAF</i> -/ <i>TERT</i> -	<i>BRAF</i> + only	<i>TERT</i> + only	<i>BRAF</i> +/ <i>TERT</i> +
de Biase 2015 (16)	374	47.4	131	224	6	13
Gandolfi 2015 (19)	121	48.1	58	42	9	12
Lee 2016 (59)	207	46.0	29	148	3	27
Liu 2016 (31)	1051	46.0	629	292	64	66
Song 2016 (50)	418	45.3	162	239	5	12
Xing 2014 (8)	507	45.9	287	159	26	35
Overall (%)	2678		1296 (48.4%)	1104 (41.2%)	113 (4.2%)	165 (6.2%)