

SUPPLEMENTARY TABLE S2. FREQUENCY DISTRIBUTIONS OF BRAF GENOTYPES BY EXTRATHYROIDAL EXTENSION IN PATIENTS WITH PAPILLARY THYROID CANCER

Variants	No, n (%)	Yes, n (%)	OR [CI] ^a
rs1042179: A>G			
AA	136 (70.5)	121 (70.3)	1.00
AG	51 (26.4)	47 (27.3)	0.95 [0.59–1.52]
GG	6 (3.1)	4 (2.3)	1.33 [0.36–4.93]
GG+AG	57 (29.5)	51 (29.7)	0.97 [0.62–1.54]
rs11762469: A>T			
AA	134 (69.1)	113 (65.7)	1.00
AT	55 (28.4)	55 (32.0)	0.85 [0.54–1.34]
TT	5 (2.6)	4 (2.3)	1.10 [0.28–4.23]
AT+TT	60 (30.9)	59 (34.3)	1.87 [0.56–1.35]
rs17161747: G>C			
GG	124 (63.9)	119 (69.2)	1.00
GC	59 (30.4)	51 (29.7)	1.14 [0.72–1.82]
CC	11 (5.7)	2 (1.2)	5.72 [1.22–26.77]^b
GC+CC	70 (36.1)	53 (30.8)	1.32 [0.84–2.06]
rs3748093: T>A			
TT	117 (60.6)	100 (58.1)	1.00
AT	68 (35.2)	60 (34.9)	0.98 [0.63–1.52]
AA	8 (4.1)	12 (7.0)	0.54 [0.21–1.37]
AT+AA	76 (42.7)	72 (36.4)	0.90 [0.59–1.37]

^aORs are adjusted for age, sex, cancer familial history, alcohol drinking, and smoking.

^bBoldface indicates statistical significance ($p < 0.05$).

CI, 95% confidence interval; OR, odds ratio.