

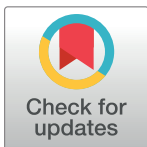
CORRECTION

Correction: Inherited variants in genes somatically mutated in thyroid cancer

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There is an error in the email address listed for corresponding author Chiara Campo. The email address should be chiaracampo28@gmail.com.

[Table 3](#), “Association of the 8 top SNPs in the two Italian sets with WDTC risk (Part II),” does not appear correctly. Please view [Table 3](#) here.



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Table 3. Association of the 8 top SNPs in the two Italian sets with WDTC risk (Part II).

Gene/SNP		Italian1*				Italian2*				Combined	
		Case	Control	OR [95% CI]	pValue	Case	Control	OR [95% CI]	pValue	OR [95% CI]	pValue
TSHR (rs2268477)	CC	1210	1236	1.0 [reference]		566	395	1.0 [reference]			
	AC	248	304	0.79[0.64–0.98]	0.03	115	76	1.19[0.86–1.65]	0.30	0.95[0.64–1.42]	0.81
	AA	10	27	0.42[0.18–0.95]	0.04	9	6	1.30[0.44–3.85]	0.63	0.64[0.33–1.23]	0.18
	AC + AA	258	331	0.76[0.62–0.94]	0.01	124	82	1.20[0.87–1.65]	0.27	0.94[0.60–1.47]	0.78
	Per allele			0.75[0.62–0.91]	3.96×10^{-3}			1.18[0.88–1.58]	0.26	0.93[0.59–1.44]	0.74
SMAD4 (rs7229678)	GG	566	542	1.0 [reference]		251	187	1.0 [reference]			
	CG	651	729	0.84[0.70–1.01]	0.07	337	229	1.05[0.81–1.37]	0.70	0.94[0.78–1.05]	0.19
	CC	225	258	0.75[0.58–0.96]	0.02	103	82	0.99[0.69–1.41]	0.93	0.82[0.67–1.01]	0.06
	CG + CC	876	987	0.82[0.69–0.97]	0.02	440	311	1.04[0.81–1.33]	0.77	0.88[0.77–1.02]	0.09
	Per allele			0.86[0.76–0.97]	0.01			1.01[0.85–1.20]	0.94	0.91[0.82–1.00]	0.06
GNAS (rs7121)	TT	571	678	1.0 [reference]		269	166	1.0 [reference]			
	CT	651	679	1.19[0.99–1.42]	0.06	322	237	0.81[0.62–1.06]	0.12	0.99[0.68–1.45]	0.98
	CC	240	202	1.51[1.18–1.95]	1.28×10^{-3}	100	95	0.63[0.44–0.89]	9.63×10^{-3}	0.98[0.42–2.32]	0.97
	CT + CC	891	881	1.26[1.06–1.49]	7.12×10^{-3}	422	332	0.76[0.59–0.97]	0.03	0.99[0.60–1.62]	0.96
	Per allele			1.23[1.09–1.39]	7.90×10^{-4}			0.79[0.66–0.94]	7.26×10^{-3}	0.99[0.64–1.53]	0.97
GNAS (rs1800900)	GG	447	491	1.0 [reference]		190	148	1.0 [reference]			
	AG	658	744	1.03[0.85–1.25]	0.76	354	260	1.04[0.79–1.37]	0.77	1.03[0.88–1.21]	0.69
	AA	360	309	1.36[1.08–1.70]	8.82×10^{-3}	144	90	1.27[0.90–1.81]	0.17	1.33[1.10–1.61]	3.00×10^{-3}
	AG + AA	1018	1053	1.13[0.94–1.34]	0.19	498	350	1.10[0.84–1.43]	0.49	1.12[0.97–1.30]	0.13
	Per allele			1.17[1.04–1.31]	0.01			1.11[0.94–1.32]	0.22	1.15[1.05–1.27]	4.00×10^{-3}

SNP: single nucleotide polymorphism, OR: odd ratio, CI: confidence interval

(*) The analysis was conducted adjusting for the covariates of sex and age

<https://doi.org/10.1371/journal.pone.0202208.t001>

Reference

1. Campo C, Köhler A, Figlioli G, Elisei R, Romei C, Cipollini M, et al. (2017) Inherited variants in genes somatically mutated in thyroid cancer. PLoS ONE 12(4): e0174995. <https://doi.org/10.1371/journal.pone.0174995>. PMID: 28410400