

CORRECTION

# Correction: Genetic Variants Associated with Increased Risk of Malignant Pleural Mesothelioma: A Genome-Wide Association Study

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There are errors in [Table 1](#). The values in the row labeled Age (mean  $\pm$  s.e) are incorrect. Please see the corrected [Table 1](#) here.



## OPEN ACCESS

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**Table 1. Summary statistics of all the subjects included in the Italian GWAS.**

	CASALE M.		TURIN		GENOA		ALL SAMPLE	
	Cases N (%)	Controls N (%)	Cases N (%)	Controls N (%)	Cases N (%)	Controls N (%)	Cases N (%)	Controls N (%)
Eligible	241 (48.88)	252 (51.12)	91 (61.9)	56 (38.1)	75 (48.08)	81 (51.92)	407 (51.13)	389 (48.87)
After QC filtering	230 (49.25)	237 (50.75)	89 (61.81)	55 (38.19)	73 (49.32)	75 (50.68)	392 (51.65)	367 (48.35)
<b>GENDER</b>								
<b>Males</b>	155 (67.39)	162 (68.35)	62 (69.66)	38 (69.09)	67 (91.78)	56 (74.67)	284 (72.45)	256 (69.75)
<b>Females</b>	75 (32.61)	75 (31.65)	27 (30.34)	17 (30.91)	6 (8.22)	19 (25.33)	108 (27.55)	111 (30.25)
<b>BIRTH PLACE</b>								
<b>North Italy</b>	204 (90.27)	186 (78.81)	62 (69.66)	35 (63.64)	54 (78.26)	53 (76.81)	320 (83.33)	274 (76.11)
<b>Center Italy</b>	6 (2.65)	12 (5.08)	5 (5.62)	1 (1.82)	8 (11.59)	5 (7.25)	19 (4.95)	18 (5)
<b>South Italy</b>	14 (6.19)	33 (13.98)	16 (17.98)	17 (30.91)	4 (5.8)	4 (5.8)	34 (8.85)	54 (15)
<b>Sardinia</b>	0 (0)	2 (0.85)	3 (3.37)	2 (3.64)	1 (1.45)	3 (4.35)	4 (1.04)	7 (1.94)
<b>Other Caucasians</b>	2 (0.88)	3 (1.27)	3 (3.37)	0 (0)	2 (2.9)	4 (5.8)	7 (1.82)	7 (1.94)
<b>ASBESTOS EXPOSURE</b>								
<b>Non exposed</b>	4 (2.06)	54 (22.78)	3 (3.37)	18 (32.73)	10 (13.7)	41 (54.67)	17 (4.87)	113 (30.79)
<b>Medium exposed</b>	106 (54.64)	103 (43.46)	33 (37.08)	25 (45.45)	7 (9.59)	22 (29.33)	146 (41.01)	150 (40.87)
<b>High exposed</b>	84 (43.3)	80 (33.76)	53 (59.55)	12 (21.82)	56 (76.71)	12 (16)	193 (54.21)	104 (28.34)
<b>Age (mean±s.e.)</b>	67.61±11.14	63.36±11.06	68.74±8.84	68.31±8.80	69.64±9.64	58.59±15.03	68.25±10.39	63.11±12.01

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## Reference

1. Matullo G, Guarrera S, Betti M, Fiorito G, Ferrante D, Voglino F, et al. (2013) Genetic Variants Associated with Increased Risk of Malignant Pleural Mesothelioma: A Genome-Wide Association Study. PLoS ONE 8(4): e61253. doi: [10.1371/journal.pone.0061253](https://doi.org/10.1371/journal.pone.0061253) PMID: [23626673](https://pubmed.ncbi.nlm.nih.gov/23626673/)