

Erratum to the association between the ERCC1/2 polymorphisms and radiotherapy efficacy in 87 patients with non-small cell lung cancer

Editorial Office

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Erratum to: J Thorac Dis 2021;13:3126-36

In the article (1) titled "the association between the ERCC1/2 polymorphisms and radiotherapy efficacy in 87 patients with nonsmall cell lung cancer" published in *Journal of Thoracic Disease* (J Thorac Dis 2021;13:3126-36. doi: 10.21037/jtd-21-755), there were some errors in the Results section.

(I) There is a typo in a confidence interval (CI) value.

Original sentence: The median PFS of patients with the G allele was 18.9 months (95% CI: 13.960–24.780), which was significantly higher than that of patients with the A allele (11.3 months, 95% CI: 8.655–14.005); the difference between the two groups was statistically significant (P=0.040).

Corrected sentence: The median PFS of patients with the G allele was 18.9 months (95% CI: 13.960–23.780), which was significantly higher than that of patients with the A allele (11.3 months, 95% CI: 8.655–14.005); the difference between the two groups was statistically significant (P=0.040).

(II) There were errors in Table 3. The corrected version of Table 3 is included below.

Gene	MST (months)	95% CI	Р	Median PFS (months)	95% CI	Р
rs11615						
AA	6.77	0-14.05	0.000	5.13	0–15.13	0.019
AG	19.77	11.14–28.40		11.33	5.76–16.91	
GG	30.90	12.83–48.97		18.87	13.96–23.78	
A allele	16.17	9.10-23.24	0.003	11.33	8.66–14.01	0.040
G allele	30.90	12.83–48.97		18.87	13.96–23.78	
rs3212961						
GG	24.77	4.75–44.79	0.735	11.53	6.41–16.66	0.910
GT	23.97	15.75–32.19		18.07	12.67–23.47	
TT	24.93	9.37–40.49		11.60	5.28–17.92	

Table 3 Association between the ERCC1/2 gene polymorphisms and prognosis in 87 NSCLC patients who received definitive radiotherapy

Table 3 (continued)

 Table 3 (continued)

Gene	MST (months)	95% CI	Р	Median PFS (months)	95% CI	Р
G allele	24.93	18.95–30.91	0.441	16.17	11.84–20.50	0.897
T allele	24.77	4.75-44.79		11.53	6.41–16.66	
rs3212986						
AA	71.13	0–160.416	0.092	15.53	0–38.01	0.718
AC	29.10	18.92–39.28		18.70	11.91–25.49	
CC	21.53	12.97–30.09		14.67	8.85–20.49	
A allele	30.90	15.68-46.12	0.086	18.07	8.85–20.49	0.423
C allele	21.53	12.97–30.09		14.67	11.89–24.25	
rs13181						
TT	24.93	16.19–33.67	0.692	14.67	10.17–19.17	0.921
TG	24.77	12.67–36.87		19.07	9.96–28.19	
rs238406						
TT	16.17	0–36.26	0.686	11.33	8.92-13.74	0.143
TG	28.67	22.72-34.62		21.50	16.86–26.14	
GG	19.80	13.76–25.84		11.30	6.67–15.93	
T allele	26.83	21.44–32.22	0.562	16.83	12.16–21.50	0.097
G allele	19.80	13.76–25.84		11.30	6.67–15.93	
rs1799793						
CC	25.40	18.74–32.05	0.222	15.53	11.79–19.27	0.411
СТ	19.07	10.48–27.66		16.13	2.89–29.37	

NSCLC, non-small cell lung cancer; CI, confidence interval; MST, median overall survival time; PFS, progression-free survival.

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References

1. Jiang C, Guo Y, Li Y, et al. The association between the ERCC1/2 polymorphisms and radiotherapy efficacy in 87 patients with non-small cell lung cancer. J Thorac Dis 2021;13:3126-36.

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