

Erratum

Development and validation of a N6 methylation regulator-related gene signature for prognostic and immune response prediction in non-small cell lung cancer: Am J Cancer Res. 2023; 13(7): 2984-2997

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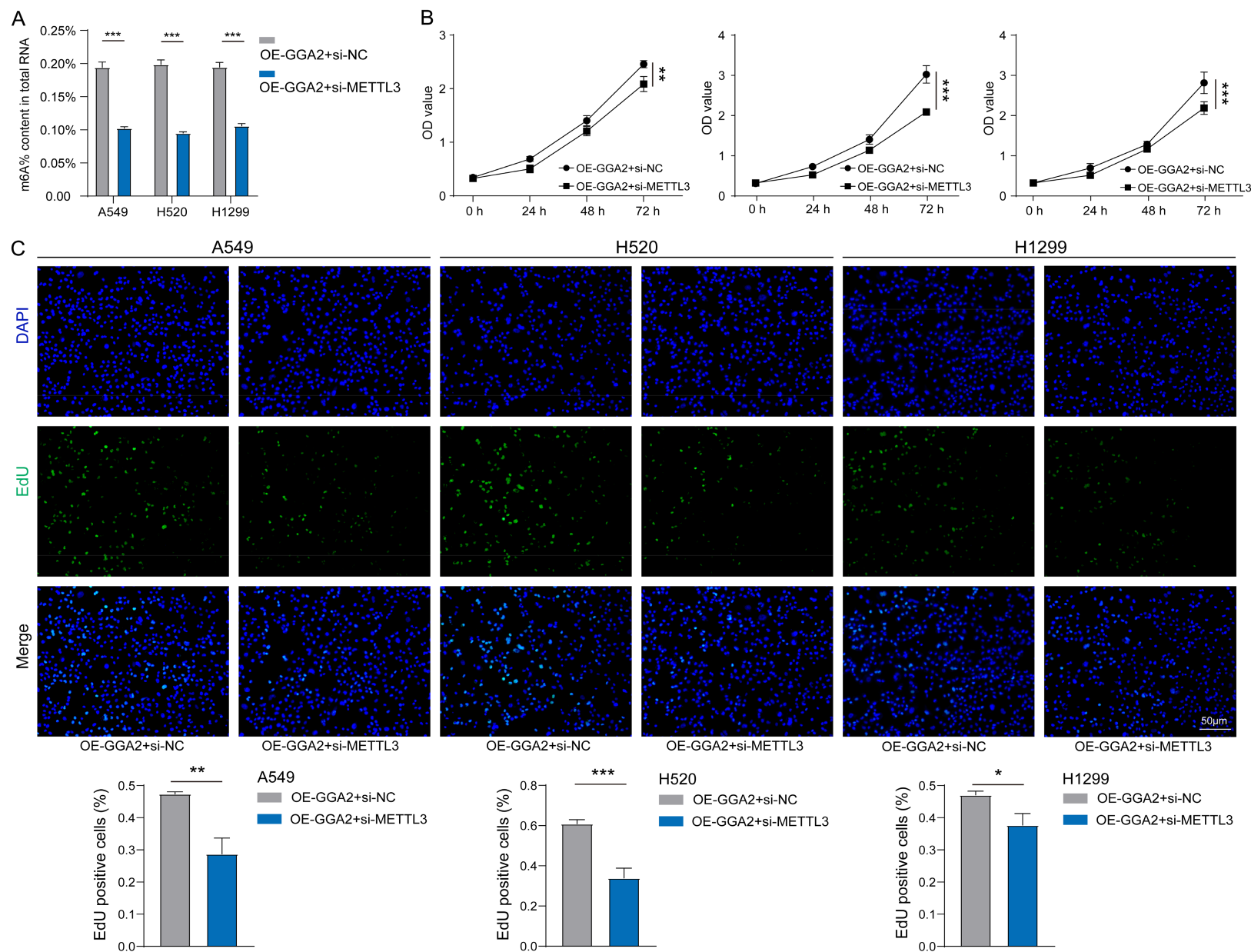
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Received March 26, 2026; Accepted May 8, 2026; Epub October 15, 2026; Published October 30, 2026

An unintentional duplication of images has been found in **Figure 7D**. Therefore, we provide the correct version to displace the wrong figure and reflect changes. We sincerely apologize for this oversight and any confusion it may have caused. The corrected **Figure 7** is shown below.

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A gene signature in non-small cell lung cancer



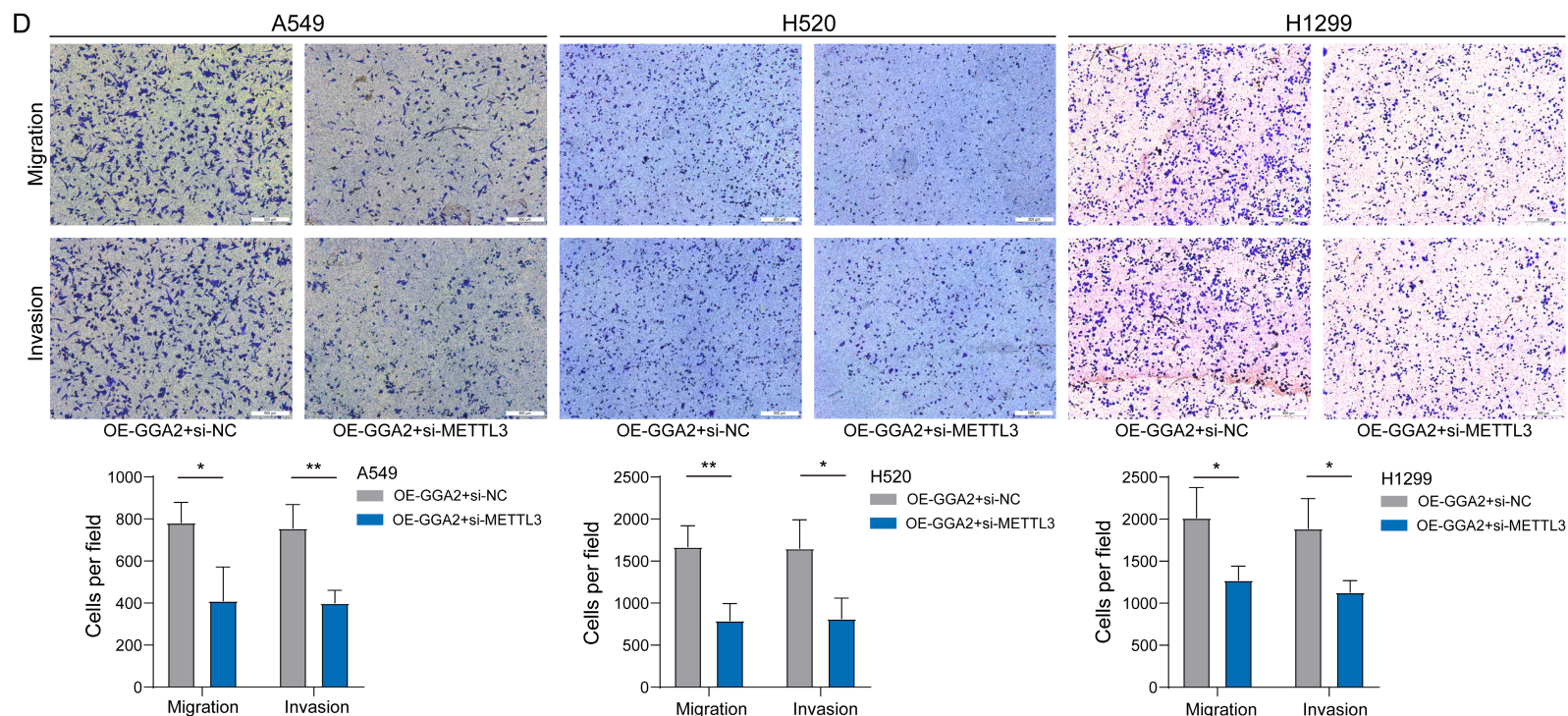


Figure 7. GGA2 inhibited NSCLC cells by downregulating METTL3. Detection of m6A content in total RNA, *P* value was calculated by 2-tailed Student *t* test (***, *P* < 0.001) (A). Cell proliferation was detected by measuring the optical density (OD) in 450 nm, *P* value was calculated by 2-tailed Student *t* test (**, *P* < 0.01; ***, *P* < 0.001) (B). EdU staining (green fluorescence) on cell proliferation 3 days after the construction of cell lines, nuclei were counterstained blue with DAPI (Scale bars, 50 μ m) (*, *P* < 0.05; **, *P* < 0.01; ***, *P* < 0.001) (C). Representative images of Transwell assays (migration and invasion) after 24 hours incubation (Scale bars, 500 μ m) (*, *P* < 0.05; **, *P* < 0.01) (D).