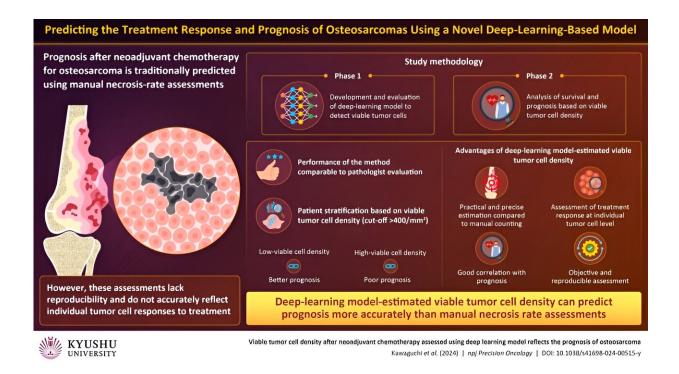


AI's ability to detect tumor cells could be key to more accurate bone cancer prognoses

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Researchers from Kyushu University, Japan have developed a novel deeplearning model which can accurately estimate viable tumor cell density from pathological images of osteosarcoma patients, and predict their prognosis following chemotherapy. Credit: Dr. Makoto Endo/Kyushu University

Researchers at Kyushu University have developed and validated a machine-learning model that can accurately evaluate the density of surviving tumor cells after treatment in pathological images of



osteosarcoma—the most prevalent malignant bone tumor. The model can assess how individual tumor cells respond to treatment and can predict overall patient prognosis more reliably than conventional methods.

Surgery and chemotherapy have significantly improved the outcomes of patients with localized osteosarcoma. However, patients with advanced metastatic disease (the stage where <u>cancerous cells</u> have spread to distant tissues) have a low survival rate. After a standard treatment of surgery and chemotherapy, assessing the prognosis of patients is essential for determining their subsequent individual treatment plans. However, predicting patient outcomes has many challenges.

Currently, prognosis relies on necrosis rate assessment, which involves pathologists evaluating the proportion of dead tissue within a tumor. Unfortunately, these methods are limited by variability between pathologists' assessments and may not accurately predict treatment response.

Recognizing the need for faster and more accurate prognoses, co-first authors Dr. Kengo Kawaguchi and Dr. Kazuki Miyama, from the Department of Orthopedic Surgery, Graduate School of Medical Sciences, Kyushu University, Japan, and Dr. Makoto Endo, a lecturer of Orthopedic Surgery at Kyushu University Hospital, along with collaborators, turned to artificial intelligence (AI) for a more nuanced evaluation. The multidisciplinary was team headed by Dr. Endo included Kyushu University's Professor Ryoma Bise, Professor Yoshinao Oda, and Professor Yasuharu Nakashima.

Explaining the rationale behind their research, which was <u>published</u> in *npj Precision Oncology*, Dr. Endo says, "In the traditional method, the necrosis rate is calculated as a necrotic area rather than individual cell counts, which is not sufficiently reproducible between assessors and does



not adequately reflect the effects of anticancer drugs. We therefore considered using AI to improve the estimation."

In phase 1 of the study, the team trained a type of AI, called a deeplearning model, to detect surviving <u>tumor cells</u> and validated its detection performance using patient data. The AI model showed proficiency in detecting viable tumor cells in pathological images, aligning with expert pathologists' capabilities.

In phase 2, the researchers analyzed two key measures: disease-specific survival, which tracks the duration after diagnosis or treatment without death directly caused by the disease, and metastasis-free survival, which monitors the time post-treatment without cancer cells spreading to distant body parts. They also explored the correlation between AI-estimated viable tumor cell density and prognosis. Notably, the AI model demonstrated comparable detection performance and precision to that of the pathologist, with good reproducibility.

Next, the researchers sorted the patients into groups based on whether the viable tumor cell density was above or below 400/mm². The survival analysis revealed that the high-density group showed a worse prognosis, while the low-density group showed a better prognosis for diseasespecific survival and metastasis-free survival.

Necrosis rate, on the other hand, was not associated with disease-specific survival or metastasis-free survival. Furthermore, analysis of individual cases revealed that AI-estimated viable tumor cell density was a more reliable predictor of prognosis than necrosis rate.

Overall, these findings suggest that the AI-based measurement of viable tumor cells reflects the inherent malignancy (ability of the cancer to spread) and individual tumor cell response of osteosarcomas. Incorporating AI in the analysis of pathological images improves



detection accuracy, reduces inter-assessor variability, and enables timely assessment.

Moreover, the estimation of viable tumor cells, which reflects their ability to keep multiplying following chemotherapy, is a more reliable predictor of treatment response than cell death. Large-scale validation of the AI model developed in this study can aid its wider application in reallife clinical settings.

"This new approach has the potential to enhance the accuracy of prognoses for osteosarcoma patients treated with chemotherapy. In the future, we intend to actively apply AI to rare diseases such as osteosarcoma, which have seen limited advancements in epidemiology, pathogenesis, and etiology.

"Despite the passage of decades, particularly in <u>treatment</u> strategies, substantial progress remains elusive. By putting AI to the problem, this might finally change," concludes Dr. Endo.

More information: Kengo Kawaguchi et al, Viable tumor cell density after neoadjuvant chemotherapy assessed using deep learning model reflects the prognosis of osteosarcoma, *npj Precision Oncology* (2024). DOI: 10.1038/s41698-024-00515-y

Provided by Kyushu University

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