

AI reveals prostate cancer is not just one disease

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Artificial Intelligence has helped scientists reveal a new form of aggressive prostate cancer that could revolutionize how the disease is diagnosed and treated in the future.

The study [published](#) in *Cell Genomics* reveals that [prostate cancer](#), which affects one in eight men in their lifetime, includes two different subtypes

termed evotypes.

The discovery was made by an international team led by the University of Oxford, and the University of Manchester, who applied AI ([artificial intelligence](#)) on data from DNA to identify two different subtypes affecting the prostate.

The team hopes their findings could save thousands of lives in future and revolutionize how prostate cancer is diagnosed and treated. Ultimately, it could provide tailored treatments to each individual patient according to a genetic test which will also be delivered using AI.

According to Cancer Research UK, prostate cancer is the most common cancer affecting men in the UK with around 52,000 cases a year. Dr. Rupal Mistry, the charity's senior Science Engagement Manager, said,

"The work published ... by this global consortium of researchers has the potential to make a real difference to people affected by prostate cancer. The more we understand about cancer the better chance we have of developing treatments to beat it. We are proud to have helped fund this cutting-edge work, which has laid the foundations for personalized treatments for people with prostate cancer, allowing more people to beat their disease."

The research, which involved scientists from the University of Oxford, the University of Manchester, the University of East Anglia and the Institute of Cancer Research, London, highlights how a prostate cancer diagnosis can affect physical, emotional and mental well-being.

Lead researcher Dr. Dan Woodcock, of the Nuffield Department of Surgical Sciences at the University of Oxford, said, "Our research demonstrates that prostate tumors evolve along multiple pathways, leading to two distinct disease types.

"This understanding is pivotal as it allows us to classify tumors based on how the cancer evolves rather than solely on individual gene mutations or expression patterns."

The researchers worked together as part of international consortium, called The Pan Prostate Cancer Group, set up by scientists at The Institute of Cancer Research (ICR) and The University of East Anglia to analyze [genetic data](#) from thousands of prostate cancer samples across nine countries.

Crucially, the team's collaboration with Cancer Research UK (CRUK) aims to develop a genetic test that, when combined with conventional staging and grading, can provide a more precise prognosis for each patient, allowing tailored treatment decisions.

The researchers used AI to study changes in the DNA of prostate cancer samples (using whole genome sequencing) from 159 patients.

They identified two distinct cancer groups among these patients using an AI technique called neural networks. These two groups were confirmed by using two other mathematical approaches applied to different aspects of the data. This finding was validated in other independent datasets from Canada and Australia.

They went on to integrate all the information to generate an [evolutionary tree](#) showing how the two subtypes of prostate cancer develop, ultimately converging into two distinct disease types termed "evotypes."

Professor David Wedge of Manchester Cancer Research Center, who led the study, explained, "This realization is what enables us to distinguish the disease types. This hasn't been done before because it's more complicated than HER2+ in breast cancer, for instance.

"This understanding is pivotal as it allows us to classify tumors based on their evolutionary trajectory rather than solely on individual gene mutations or expression patterns."

Researcher Professor Colin Cooper, from UEA's Norwich Medical School, highlighted that while prostate cancer is responsible for a large proportion of all male cancer deaths, it is more commonly a disease men die with rather than from. This means that unnecessary treatment can often be avoided, sparing men from [side-effects](#) such as incontinence and impotence.

He added, "This study is really important because until now, we thought that prostate cancer was just one type of disease. But it is only now, with advancements in artificial intelligence, that we have been able to show that there are actually two different subtypes at play.

"We hope that the findings will not only save lives through better diagnosis and tailored treatments in the future, but they may help researchers working in other cancer fields better understand other types of cancer too."

Dr. Naomi Elster, Director of Research at Prostate Cancer Research, said, "We simply don't know enough about what a prostate cancer diagnosis means at present—there are many men who have disease which is or may become aggressive and being able to treat aggressive disease more effectively is critical. But on the other side of the coin are the too many men who live with side effects of cancer treatment they may never have needed.

"These results could be the beginning of us being able to take the same 'divide and conquer' approach to prostate cancer that has worked in other diseases, such as breast cancer."

Professor Ros Eeles, Professor of Oncogenetics at The Institute of Cancer Research, London, and Honorary Consultant in Clinical Oncology and Cancer Genetics at The Royal Marsden NHS Foundation Trust, said, "This study has utilized the enormous genomic dataset from The Pan Prostate Cancer Group—a powerhouse of information about prostate cancer from around the world. These results will hopefully lead to better treatments for patients, demonstrating the importance of data sharing and team science."

More information: Genomic evolution shapes prostate cancer disease type, *Cell Genomics* (2024). DOI: [10.1016/j.xgen.2024.100511](https://doi.org/10.1016/j.xgen.2024.100511). [www.cell.com/cell-genomics/ful ... 2666-979X\(24\)00038-7](https://www.cell.com/cell-genomics/fulltext/S2666-979X(24)00038-7)

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