EVAXION

Evaxion Develops Method to Enhance AI Drug Development with Deep Probabilistic Programming

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COPENHAGEN, Denmark, June 25, 2021 (GLOBE NEWSWIRE) -- Evaxion Biotech A/S (Nasdaq: EVAX), a clinical-stage biotechnology company specializing in the development of Al-driven immunotherapies to improve the lives of patients with cancer and infectious diseases, announced today the acceptance of a new scientific paper by the International Conference on Machine Learning (ICML 2021). A draft of the article is available on the open-access scientific server bioRxiv.org.

The paper is entitled "Efficient Generative Modelling of Protein Structure Fragments using a Deep Markov Model", and was written and developed by Evaxion personnel in collaboration with Assoc. Prof. Thomas Hamelryck's probabilistic programming group at the University of Copenhagen. The paper describes BIFROST, a novel predictive system based on deep probabilistic programming that enables the rapid conversion of sequence data into structural information on protein fragments, which we believe may be useful for drug or vaccine design. Deep probabilistic programming is a new development in machine learning that combines the principled treatment of uncertainty provided by Bayesian statistics with the capabilities of deep learning. Compared to existing protein structure prediction approaches, BIFROST appears to be computationally more efficient, only requires sequence information and, importantly, incorporates an assessment of the reliability of its own predictions.

Lars Wegner, CEO of Evaxion, said: "This work is an exciting development by the collaborative team that we believe has the potential to make vaccine development more efficient. We intend to apply our expertise to further the development of Bayesian machine learning and to integrate these methods fully into Evaxion's AI platforms, including both our EDEN and RAVEN platforms for vaccine development."

Protein structure prediction methods such as BIFROST have the potential to facilitate Al-driven pharmaceutical design by indicating the likely conformation that components of immunotherapies or vaccines and their target might adopt. Existing methods for predicting the conformation of protein fragments do not explicitly evaluate the probability of conformations given the sequence which can make it difficult to dissect the reliability of subsequent calculations. By including estimates of uncertainty in predictions, BIFROST's Bayesian approach may be particularly useful in drug development datasets that, typically, are incomplete and relatively small.

Anders B. Sørensen, Evaxion Director, Research and Discovery, said: "We are excited to share this first-time application of Deep Markov Models within the field of protein structure prediction. This has significant potential to improve how we develop medicines and showcases the power harnessed when we combine academic research with industrial application."

About Evaxion

Evaxion Biotech A/S is a clinical-stage Al-immunology™ platform company decoding the human immune system to discover and develop novel immunotherapies to treat cancer, and vaccines against bacterial diseases and viral infections. Based on its proprietary and scalable Al-immunology core technology, Evaxion is developing a broad pipeline of novel product candidates which currently includes three patient-specific cancer immunotherapies, two of which are in Phase 1/2a clinical development. In addition, Evaxion is advancing a portfolio of vaccines to prevent bacterial and viral infections currently in preclinical development.

For more information

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Source: Evaxion Biotech

Forward-looking statement

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Forward-looking statements are subject to inherent risks and uncertainties beyond the Company's control that could cause the Company's actual results, performance, or achievements to be materially different from the expected results, performance, or achievements expressed or implied by such forward-looking statements. For a further description of the risks and uncertainties that could cause actual results to differ from those expressed in these forward-looking statements, as well as risks relating to the Company's business in general, see the risks described in the "Risk Factors" section included in the Company's Form 20-F for the year end December 31, 2020 and the Company's current and future reports filed with, or submitted to, the U.S. Securities and Exchange Commission (SEC). Any forward-looking statements contained in this announcement speak only as of the date hereof, and except as required by law, the Company assumes no obligation to update these forward-looking statements publicly, or to update the reasons actual results could differ materially from those anticipated in the forward-looking statements, even if new information becomes available in the future.



Source: Evaxion Biotech