



Stay alive in the digitized world

AI in Pediatric Infectious Diseases:

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บทบาทของ AI ในงานโรคติดเชื้อในเด็ก

Epidemic Forecasting

Predict and track the spread of infectious diseases → early intervention and prevention measures.

Dx & Tx Optimization

Analyze medical images, laboratory results, & patient data.

Drug & Vaccine Development

Analyze large amounts of data to identify potential targets & accelerate the research process.

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An AI Tool That Can Help Forecast Viral Variants

EVEscape predicts future viral mutations, new variants using evolutionary, biological information

By CATHERINE CARUSO | October 11, 2023 | Research
4 min read



At a glance:

- New AI tool called EVEscape uses evolutionary and biological information to predict how a virus could change to escape the immune system.
- The tool successfully predicted the most concerning new variants that occurred during the COVID-19 pandemic.
- Researchers say the tool can help inform the development of vaccines and therapies for SARS-CoV-2 and other rapidly mutating viruses.

NATURE INDEX | 13 March 2024

How AI is being used to accelerate clinical trials

From study design to patient recruitment, researchers are investigating ways that technology could speed up the process.

By Matthew Hutson



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Modern Clinical Trials



>\$1.2B
Average Cost

>5
Years

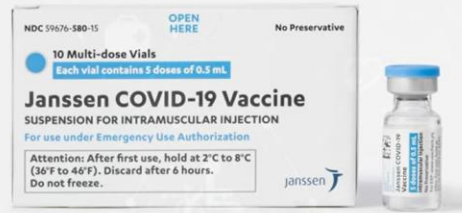
<20%
Success Rate

Improve the gold standard process with AI

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The First Successful AI-Driven Phase III Clinical Trial



<https://www.youtube.com/watch?v=ctRnh65cAel>

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The First Successful AI-Driven Phase III Clinical Trial



Time	4 - 6 months	1 - 2 years	1 - 3 years	6 months+
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>1M
Daily Cases

>10,000
Daily Deaths

4

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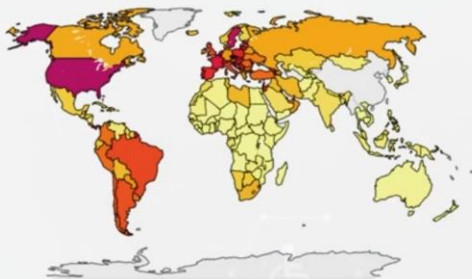
The First Successful AI-Driven Phase III Clinical Trial



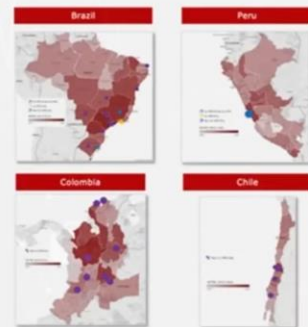
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Implementation for Janssen's Phase III Vaccine Trial



May 2020:
Select trial countries globally



June 2020:
Determining local site selections

Argentina, Brazil, Chile, Columbia, Mexico, Peru, South Africa, United States

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Can a fabled blue parrot flourish again in the wild? p. 1148 | Unlocking hidden physics with quantum computers pp. 1154 & 1182 | Inherited microbiomes are shaped by lifestyle p. 1220

Science

515
10 JUNE 2022
SPECIAL ISSUE
science.org

AAAS

NUCLEAR PORE

A gateway complex p. 1172

The cover features a large 3D model of a nuclear pore complex (NPC) in the center, showing its complex structure with yellow and blue components. To the right, a circular diagram compares protein structure prediction methods: Cryo-EM (Cryo-EM), AlphaFold (AlphaFold), and a 'Combined model'. The diagram shows the NPC structure being predicted by these methods, with labels for 'Cryo-EM', 'AlphaFold', 'Nup133', 'Nup159', 'Nup145-42', 'Nup157', 'Nup158', 'Nup159-42', 'Nup157', and 'Combined model'.

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LASKER FOUNDATION

AlphaFold—for predicting protein structures

2023 Albert Lasker Basic Medical Research Award

Demis Hassabis
Google DeepMind

John Jumper
Google DeepMind

15 September 2023

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