

# Unlock genomic insights

Increase efficiency and confidence with explainable-AI (XAI) and automation to support your tertiary analysis clinical research workflows

Emedgene streamlines and integrates variant interpretation through report generation workflows that you define.

## Trusted by leading institutions

**BAYLOR**  
GENETICS

**SickKids**



**fleury** medicina e saúde | Genômica



# Confidently scale tertiary analysis clinical research workflows

## Streamlined

Save valuable time through the application of XAI and customizable automation by as much as 75% per subject

## Integrated

Connect with LIMS and other health IT systems to simplify and secure the complete assay workflow

## Powered for growth

Confidently keep pace with evolving science, technology and demand supported by a team of experts

## Powerful core AI technology applied across diverse use cases

### Launch assays

Implement a high-throughput genome (WGS), exome (WES), virtual panel, or targeted panel workflow that is integrated into your lab's digital ecosystem.

### Expand menu

Broaden your analysis to WGS or WES or standardize panels on a backbone assay; analyze across a range of variant types—SNVs, indels, CNVs, mtDNA, SVs, and STR variants.

### Scale volume

Increase throughput without increasing headcount using XAI and automated workflows.

### Share curated knowledge

Leverage the power of collaboration to share knowledge across a private network of labs.

# The role of AI in genomics is to reduce noise and highlight data with the most evidence



## Efficiency

Support scale and efficiency gains



## Time per subject

Reduce turnaround time and costs



## Accuracy

Confidently prioritize variants across applications

## All the enterprise features needed for smooth adoption, integration, and operation

### Automated ACMG classification

Save as much as 90% of classification time with automated classification for SNVs, indels, and CNVs.

### Broad portfolio of tests and variant types

Support your interpretation of SNVs, indels, CNVs, SVs, STRs, and mtDNA from WGS, WES, targeted panels, or virtual panels.

### SOP and advanced filters

Implement your standard operating procedure (SOP) on the platform by creating sets of predefined custom filters or use our advanced filtering system.

### Knowledge management

Maintain a knowledge base of your organization's curated and tagged genomic data.

### Streamlined workflow and collaboration

Implement a customized workflow for your lab and facilitate team communications.

### Automated custom reporting

Customize, edit, and automatically populate reports; no HTML required and minimal manual work.

### Powerful API interoperability

Automate workflows with API integrations to leading LIMS, storage, pipelines, and more.

### Share curated knowledge

Securely share curated variants across a private network of labs and facilitate team communications.

# Integrate and streamline your research workflows from library prep, sequencing, and data analysis

Webinar



## Breaking the Interpretation Bottleneck

Dr. Linyan Meng, Baylor Genetics, presents a joint study demonstrating 97% accuracy in pinpointing causative variants with AI

Publication



## Nature Genomic Medicine

Boston Children's Hospital incorporates Emedgene in their clinical research genomics initiative

Publication



## American Journal of Medical Genetics

Brazilian healthcare giant Fleury increases yield with AI

**Emedgene serves your mission to unlock genomic insights by providing data-driven AI solutions for genomic discovery at scale.**

**illumina**<sup>®</sup>

1.800.809.4566 toll-free (US) | +1.858.202.4566 tel  
techsupport@illumina.com  
[www.illumina.com](http://www.illumina.com)

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## Abbreviation list

ACMG American College of Medical Geneticists  
AI Artificial Intelligence  
API Application Programming Interface  
CNV Copy Number Variant  
EHR Electronic Health Record  
GDPR General Data Protection Regulation  
HIPAA Health Insurance Portability and Accountability Act

LIMS Lab Information Management System  
mtDNA Mitochondrial DNA  
NLP Natural Language Processing  
SNV Single Nucleotide Variant  
SOC Service Organization Control  
SOP Standard Operating Procedure  
STR Short Tandem Repeat

SV Structural Variant  
TAT Turnaround Time  
WES Whole-Exome Sequencing  
WGS Whole-Genome Sequencing  
XAI Explainable Artificial Intelligence