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Getting started with next-generation sequencing

Next-generation sequencing (NGS) is a powerful tool for answering your research questions, and it's easier than you think.

Expand the scope of your research with a broader view

Compared to targeted approaches, NGS identifies variants across thousands of genes, enhancing discovery power when studying signaling pathways, disease markers, or novel drug targets.[1-3]

Compared to Sanger sequencing

Up to 1,000x more genes

sequenced in a single assay[4]

Compared to qPCR

>1,000 target regions profiled

in a single assay[5]

Compared to flow cytometry

Up to 530x more proteins

analyzed in a single experiment[6,7]

For us, sequencing changed everything. If we wouldn't have been able to use bulk RNA sequencing, we wouldn't have been able to see what's going on at the chromosomal level in our mutants.

Mercedes Carro, PhD, Postdoctoral Associate
Cohen Lab, Cornell University

RNA-Seq: A good place to start

Examine expression of thousands of genes in a single assay to characterize complex pathways.[8,9]

Get help from NGS professionals

You don't need to be an expert, get started by collaborating with core or service labs during experimental design phase.

No bioinformatics expertise required

Don't be intimidated by large NGS datasets, analysis tools and applications help you understand and interpret your data.

87%

increase in publications using NGS since 2013.[10]

The time for NGS is now

NGS is more accessible than ever, with costs decreasing and publications on the rise.[10] There has never been a better time to start using NGS.

We're ready when you are. Let's get you started!

Learn more about getting started with NGS by downloading our new eBook.

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