ROSALIND

Research Reimagined

ROSALIND
nCounter Analysis

ROSALIND has collaborated with NanoString to combine the power of NanoString nCounter with the utility of ROSALIND'S leading cloud-based data visualization. Explore your nCounter data sets like never before with the most advanced discovery and collaboration tools right at your fingertips.

INNOVATIVE

COLLABORATIVE

SCALEABLE

ACTIONABLE

SECURE

ROSALIND NanoString Features

- NanoString Gene & miRNA expression
- Interactive Plots
- Interpretation from more than 200 leading knowledge bases
- Differential expressed genes & details for every gene
- Sharing of analysis across research teams*
- Muti-Omic Meta-Analysis*

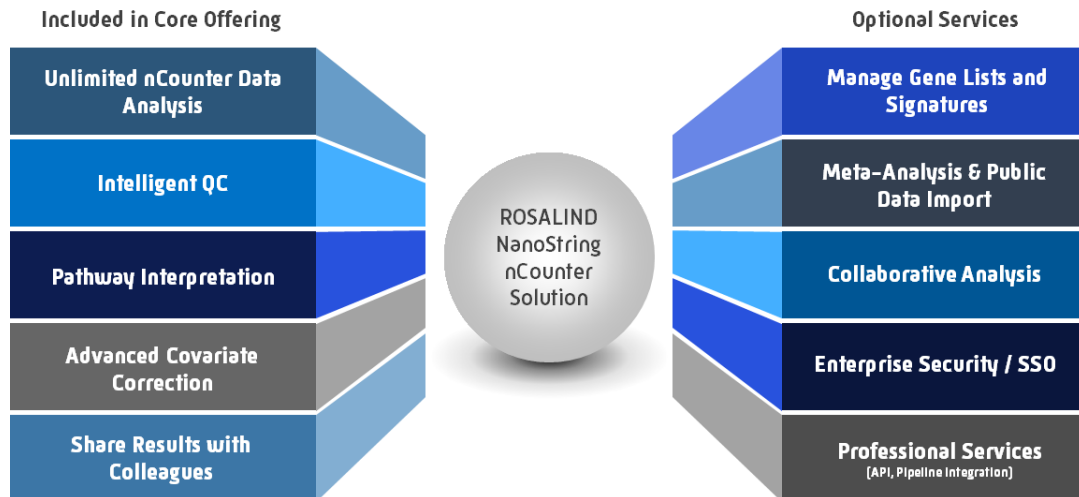** nCounter Plus+ Subscription Req'd (starting at $49/mo.)*

TO LEARN MORE, VISIT https://www.rosalind.bio/nanostring

1 (855) 766-7267   info@rosalind.bio   www.rosalind.bio

nCounter Starter Solution and Beyond

Included in Core Offering

- Unlimited nCounter Data Analysis
- Intelligent QC
- Pathway Interpretation
- Advanced Covariate Correction
- Share Results with Colleagues

ROSALIND NanoString nCounter Solution

Optional Services

- Manage Gene Lists and Signatures
- Meta-Analysis & Public Data Import
- Collaborative Analysis
- Enterprise Security / SSO
- Professional Services [API, Pipeline Integration]

Explore the Full ROSALIND Platform

ROSALIND enables scientists and researchers to analyze and interpret differential gene expression without the need for bioinformatics or programming skills.

- Record experiment design and custom attributes
- Capture metadata with NCBI BioSample attributes
- Perform covariate & batch corrections
- Setup comparisons using biological attributes
- Create gene filters to adjust cut-off parameters
- Download publication-ready figures and plots
- Discover gene signatures and create gene lists
- Explore pathway, disease & drug knowledge bases
- Real-time collaboration & results sharing*
- Re-use existing samples for new experiments*
- Import NCBI Short Read Archive Public Data*
- Find patterns across datasets with Meta-Analyses*
- Analyze multi-omic experiments & assays*

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