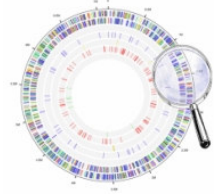


Genome Analysis by Ribocon - the Science Service Company

BIOINFORMATIC ANALYSIS OF MICROBIAL (META)GENOMIC SEQUENCE DATA - NOW AVAILABLE AS A CONVENIENT SERVICE!

- for scientific genome projects – done by scientists for scientists
- highest standards, fully transparent and supported
- based on a well-established bioinformatics pipeline



HOW DOES IT WORK?

Sending us your genomic sequence data is as easy as sending out DNA for sequencing!

1. Send us your sequence data in a standard format (up to 1,000 contigs and 10 Mb)
2. We do all the necessary analyses for you (gene prediction, annotation)
3. Get back your processed data in a common database format (a file) plus an intuitive software tool for subsequent local access, data exploration, and mining + support!

You define the level of support - from data processing to final submission – it's your project!

WHAT ARE YOUR BENEFITS?

- No need to take care about technological aspects of bioinformatics
- Keep full control compared to web-based solutions
- Individual contact person at Ribocon - continuous feedback
- Support for further analyses (comparative genomics, genome linguistics, etc.)



FACTS ...

- Sequencing of (meta)genomes is routine, the bioinformatic processing of (meta)genomic data is not - Ribocon has now closed this gap ...
- An **order of magnitude more cost-efficient** compared to setting up your own pipeline
- Getting your results done is a matter of days and not months!
- Individual Support - Maximum Experience - Maximum Outcome

Visit www.ribocon.com/genomics for more information
or contact us directly at genomics@ribocon.com.



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