

Proteome Comparison tool

Bidirectional BLAST analysis of up to 9 genomes

First description of tool

Nucleic Acids Research

[Nucleic Acids Res.](#) 2014 Jan 1; 42(Database issue): D206–D214.

PMCID: PMC3965101

Published online 2013 Nov 29. doi: [10.1093/nar/gkt1226](https://doi.org/10.1093/nar/gkt1226)

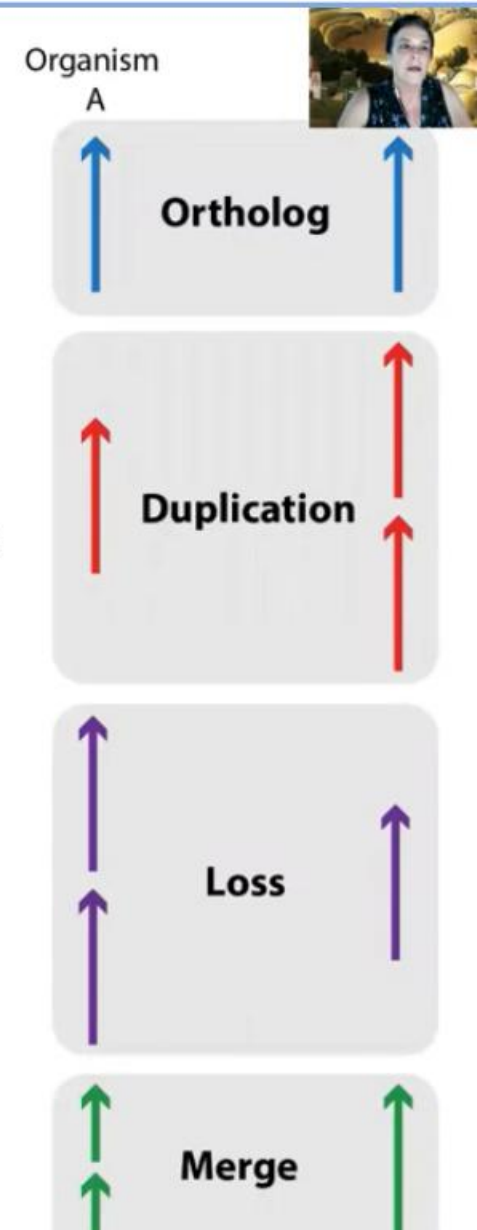
PMID: [24293654](https://pubmed.ncbi.nlm.nih.gov/24293654/)

The SEED and the Rapid Annotation of microbial genomes using Subsystems Technology (RAST)

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BLAST hits


- Best Bidirectional BLAST hits
 - Also known as reciprocal best hit
 - Can be called as orthologs between two species that are more similar
- Unidirectional BLAST hits
 - Paralogs
 - Indicative of duplication event, a loss or even merged genes














Input data

One reference and 9 comparison genomes

- Different types of reference data
- Different types up comparison data

Parameters ⓘ
ADVANCED PARAMETERS (OPTIONAL) ▼
OUTPUT FOLDER
1 2 
OUTPUT NAME

Reference Genome ⓘ
Select 1 reference genome from the following options.
SELECT A GENOME
1 2 
OR A FASTA FILE
1 2 
OR A FEATURE GROUP
1 2 

Comparison Genomes ⓘ
Add up to 9 genomes to compare (use plus buttons to add).
SELECT GENOME
1 2  
AND/OR SELECT PROTEIN FASTA FILE
1 2  
AND/OR SELECT FEATURE GROUP
1 2  
AND/OR SELECT GENOME GROUP
1 2  
SELECTED GENOME TABLE

Reset

Submit



Downloadable files

Bacteria SRR1955831-2-3_Unicycler_0_0												
	A	B	C	D	E	F	G	H	I	J	K	L
1	Bacteria SRR1955831-2-3_Unicycler_0_0											
2	ref_genom	ref_genom	ref_genom	ref_genom	ref_genom	ref_genom	ref_genom	ref_genom	ref_genom	ref_genom	ref_genom	ref_genom
3	2.8282.cor	1	584	fqi2.8282.peg.1				PGF_0705 Phage tem		1	1695	+
4	2.8282.cor	2	67	fqi2.8282.peg.2				PGF_1285 Phage pro		1709	1912	+
5	2.8282.cor	3	418	fqi2.8282.peg.3				PGF_0012 Phage por		1915	3171	+
6	2.8282.cor	4	338	fqi2.8282.peg.4				PGF_0656 Phage hes		3158	4174	+
7	2.8282.cor	5	411	fqi2.8282.peg.5				PGF_0289 Phage maj		4261	5496	+
8	2.8282.cor	6	52	fqi2.8282.peg.6				PGF_1292 Phage pro		5533	5691	+
9	2.8282.cor	7	110	fqi2.8282.peg.7				PGF_0347 Phage DN		5700	6032	+
10	2.8282.cor	8	110	fqi2.8282.peg.8				PGF_0683 Phage pro		6022	6354	+
11	2.8282.cor	9	125	fqi2.8282.peg.9				PGF_0347 Phage pro		6354	6731	+
12	2.8282.cor	10	126	fqi2.8282.peg.10				PGF_0291 Phage pro		6728	7108	+
13	2.8282.cor	11	317	fqi2.8282.peg.11				PGF_0562 Phage maj		7109	8062	+
14	2.8282.cor	12	148	fqi2.8282.peg.12				PGF_0347 Phage pro		8127	8573	+

