

# BV-BRC Test Report

## D4. Taxonomy Data Tab

<b>Item to test</b>	<b>Taxonomy Data Tab</b>
<b>URL</b>	<a href="https://www.bv-brc.org/view/Taxonomy/629#view_tab=taxontree">https://www.bv-brc.org/view/Taxonomy/629#view_tab=taxontree</a>
<b>Prerequisites</b>	None
<b>References</b>	<a href="https://www.bv-brc.org/docs/quick_references/organisms_taxon/taxonomy.html">https://www.bv-brc.org/docs/quick_references/organisms_taxon/taxonomy.html</a>
<b>Tester(s)</b>	Rebecca Wattam, Ron Kenyon
<b>Test date</b>	6-Jan-2022 and 4-Feb-22 (original-passed), 8-May-2022 (follow-up-passed)
<b>Test result</b>	<b>Passed</b>

### Overview

- Test the Taxonomy Tab with example bacterial and viral data.
- Verify the correct Taxonomy.
- Test branch expansion and collapse
- Test Taxon Overview action button.
- Test Genomes and Features action buttons.

### Test Results

- Test results were verified by examination of returned data via inspection.
- All operations performed as expected.

**Bacteria:**  
**Yersinia:**

**Taxonomy:**

 Taxon View

 Bacteria » Proteobacteria » Gammaproteobacteria » Enterobacterales » Yersiniaceae » **Yersinia** ( 1537 Genomes )

Overview	Phylogeny	<b>Taxonomy</b>	Genomes	AMR Phenotypes	Sequences	Proteins	Protein Structures	Specialty Genes	Domains and Motifs	Epitopes	Pathw
Name			Rank	Genomes							
<input type="checkbox"/>	▲	Yersinia	genus	1537							
<input type="checkbox"/>	▶	Yersinia aldovae	species	11							
<input type="checkbox"/>	▶	Yersinia aleksiciae	species	9							
<input type="checkbox"/>	▶	Yersinia bercovieri	species	19							
<input type="checkbox"/>	▶	Yersinia canariae	species	1							
<input type="checkbox"/>	▶	Yersinia enterocolitica	species	304							
<input type="checkbox"/>	▶	Yersinia entomophaga	species	2							
<input type="checkbox"/>	▶	Yersinia frederiksenii	species	54							
<input type="checkbox"/>	▶	Yersinia hibernica	species	2							
<input type="checkbox"/>	▶	Yersinia intermedia	species	47							
<input type="checkbox"/>	▶	Yersinia kristensenii	species	45							
<input type="checkbox"/>	▶	Yersinia massiliensis	species	20							
<input type="checkbox"/>	▶	Yersinia mollaretii	species	24							
<input type="checkbox"/>	▶	Yersinia nurmii	species	1							
<input type="checkbox"/>	▶	Yersinia pekkanenii	species	2							
<input type="checkbox"/>	▶	Yersinia pseudotuberculosis complex	species group	897							
<input type="checkbox"/>	▶	Yersinia rohdei	species	12							
<input type="checkbox"/>	▶	Yersinia ruckeri	species	80							
<input type="checkbox"/>	▶	unclassified Yersinia	no rank	7							

Branch expand/collapse:

- ▲ Yersinia
- ▶ Yersinia aldovae
- ▶ Yersinia aleksiciae
- ▶ Yersinia bercovieri
- ▶ Yersinia canariae
- ▲ Yersinia enterocolitica
- ▲ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica IP 10393
- ▲ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica (type O
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- ▶ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica W22703
- ▲ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica IP2222
- ▶ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica (type O
- ▶ Yersinia enterocolitica LC20
- ▶ Yersinia enterocolitica subsp. enterocolitica
- ▶ Yersinia enterocolitica subsp. palearctica
- ▶ Yersinia entomophaga

Taxon Overview Action Button:

- [Yersinia enterocolitica](#)
- [▲ Yersinia enterocolitica \(type O\)](#)
- [▶ Yersinia enterocolitica \(type O\)](#)
- [▶ Yersinia enterocolitica IP 10393](#)
- [▲ Yersinia enterocolitica \(type O\)](#)
- [▶ Yersinia enterocolitica \(type O\)](#)
- [▲ Yersinia enterocolitica \(type O\)](#)

Taxon View  
 Bacteria » Proteobacteria » Gammaproteobacteria » Enterobacterales » Yersiniaceae » Yersinia » Yersinia enterocolitica » **Yersinia enterocolitica IP 10393** ( 1 Genomes )

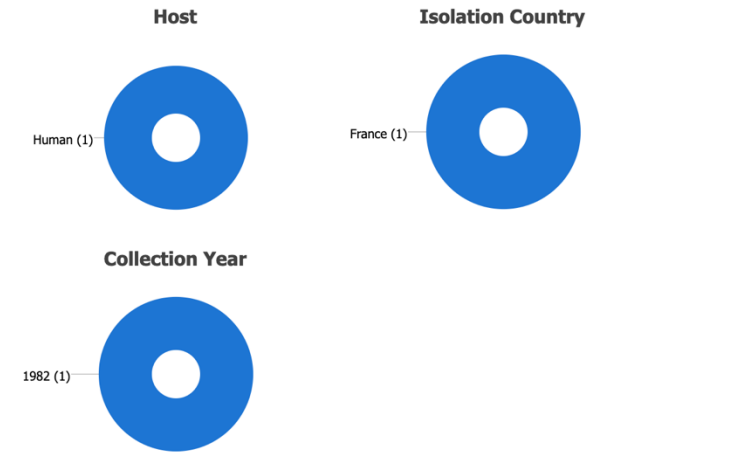
- Overview**
- Phylogeny
- Taxonomy
- Genomes
- AMR Phenotypes
- Sequences
- Proteins
- Protein Structures
- Specialty Genes
- Domains and Motifs
- Epitopes

Taxonomy Summary Genomes By Metadata

**Taxon ID** 1243181  
**Taxon Name** Yersinia enterocolitica IP 10393  
**Taxon Rank** strain  
**Genomes / Segments** 1  
**Protein Coding Genes (CDS)** 8452

Reference/Representative Genomes Genomes By Metadata

Type	Genome Name
Reference	Yersinia enterocolitica subsp. enterocolitica 8081
Representative	Yersinia enterocolitica LC20



## Genomes Action Button:

<input type="checkbox"/>	▶ <i>Yersinia bercovieri</i>	species	19
<input type="checkbox"/>	▶ <i>Yersinia canariae</i>	species	1
<input checked="" type="checkbox"/>	▶ <i>Yersinia enterocolitica</i>	species	304
<input type="checkbox"/>	▶ <i>Yersinia entomophaga</i>	species	2
<input type="checkbox"/>	▶ <i>Yersinia frederiksenii</i>	species	54
<input type="checkbox"/>	▶ <i>Yersinia hibernica</i>	species	2

### Taxon View

Bacteria » Proteobacteria » Gammaproteobacteria » Enterobacterales » Yersiniaceae » *Yersinia* » ***Yersinia intermedia*** ( 47 Genomes )

Overview | **Phylogeny** | Taxonomy | Genomes | AMR Phenotypes | Sequences | **Proteins** | Protein Structures | Specialty Genes | Domains and Motifs | Epitopes | Pathw

DOWNLOAD		KEYWORDS	ADV Search	CDS	PATRIC	FILTERS					
Genome Name	Accession	Feature Type	BRC ID	RefSeq Locus Tag	Start	End	Strand	Length (NA)	Gene Symbol	Product	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2892C	184	951	+	768		DedA family inner membrane protein Yabl	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2893C	973	1677	-	705		Thiamin ABC transporter, ATPase compon	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2894C	1664	3271	-	1608		Thiamin ABC transporter, transmembrane	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2895C	3247	4299	-	1053		Thiamin ABC transporter, substrate-binding	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2896C	4614	6275	-	1662		SgrR, sugar-phosphate stress, transcrip	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2897C	6344	7288	+	945		Pyruvate formate-lyase activating enzyme	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2898C	7293	9725	+	2433		Pyruvate formate-lyase (EC 2.3.1.54)	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2899C	10002	11183	+	1182		Sugar efflux transporter A	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2900C	11211	12473	-	1263		Tryptophan-specific transport protein	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2901C	12584	13984	-	1401		Tryptophanase (EC 4.1.99.1)	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2902C	14579	15181	-	603		3-isopropylmalate dehydratase small subu	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2903C	15196	16626	-	1431		3-isopropylmalate dehydratase large subu	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2904C	16628	17719	-	1092		2-isopropylmalate dehydrogenase (EC 1.1	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2905C	17722	19284	-	1563		2-isopropylmalate synthase (EC 2.3.3.13)	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2907C	20390	21346	+	957		LysR family transcriptional activator LeuO	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2908C	21640	23592	+	1953		Long-chain-fatty-acid-CoA ligase (EC 6.2	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2909C	23964	25739	+	1776		Acetolactate synthase large subunit (EC 2	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2910C	25742	26236	+	495		Acetolactate synthase small subunit (EC 2	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_29110	26531	27535	+	1005		Fructose repressor FruR, LacI family	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2912C	27683	28123	-	441		L-alanine exporter AlaE	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2913C	29046	29504	+	459		Transcriptional regulator MraZ	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2914C	29507	30469	+	963		16S rRNA (cytosine(1402)-N(4))-methyltra	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2915C	30466	30786	+	321		Cell division protein FtsL	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2916C	30847	32610	+	1764		Cell division protein FtsI [Peptidoglycan sy	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2917C	32597	34084	+	1488		UDP-N-acetylmuramoyl-dipeptide-2,6-dia	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2918C	34081	35445	+	1365		UDP-N-acetylmuramoyl-tripeptide-D-alan	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2919C	35439	36521	+	1083		Phospho-N-acetylmuramoyl-pentapeptide-	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2920C	36524	37840	+	1317		UDP-N-acetylmuramoyl-L-alanine-D-gluta	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2921C	37840	39042	+	1203		Peptidoglycan glycosyltransferase FtsW (E	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2922C	39039	40109	+	1071		UDP-N-acetylglucosamine-N-acetylmurar	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2923C	40192	41667	+	1476		UDP-N-acetylmuramate-L-alanine ligase (	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2924C	41660	42580	+	921		D-alanine-D-alanine ligase (EC 6.3.2.4)	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2925C	42582	43439	+	858		Cell division protein FtsQ	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2926C	43436	44692	+	1257		Cell division protein FtsA	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2927C	44765	45916	+	1152		Cell division protein FtsZ	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2928C	46001	46936	+	936		UDP-3-O-[3-hydroxymristoyl] N-acetylglu	
<input type="checkbox"/>	<i>Yersinia intermedia</i> ATCC 29909	NZ_AALF02000C	CDS	fig 349965.6.peg.yinte0001_2929C	47004	47528	-	525		Zn-ribbon-containing, possibly RNA-bindin	

1 - 200 of 227797 results

< 1 2 3 ... 1139 >

## Virus:

### Influenza

Taxon View

Viruses » Negarnaviricota » Insthoviricetes » Articulavirales » Orthomyxoviridae » Alphainfluzavirius » **Influenza A virus** ( 810384 Genomes )

Overview	Taxonomy	Strains	Genomes	Proteins	Protein Structures	Domains and Motifs	Epitopes	Experiments	Surveiller
Name		Rank	Genomes						
<input type="checkbox"/>	▲ Influenza A virus	species	854164						
<input type="checkbox"/>	▶ H10N1 subtype	serotype	416						
<input type="checkbox"/>	▶ H10N2 subtype	serotype	308						
<input type="checkbox"/>	▶ H10N3 subtype	serotype	941						
<input type="checkbox"/>	▶ H10N4 subtype	serotype	531						
<input type="checkbox"/>	▶ H10N5 subtype	serotype	1155						
<input type="checkbox"/>	▶ H10N6 subtype	serotype	507						
<input type="checkbox"/>	▶ H10N7 subtype	serotype	4710						
<input type="checkbox"/>	▶ H10N8 subtype	serotype	855						
<input type="checkbox"/>	▶ H10N9 subtype	serotype	344						
<input type="checkbox"/>	▶ H11N1 subtype	serotype	280						
<input type="checkbox"/>	▶ H11N2 subtype	serotype	962						
<input type="checkbox"/>	▶ H11N3 subtype	serotype	350						
<input type="checkbox"/>	▶ H11N4 subtype	serotype	21						
<input type="checkbox"/>	▶ H11N5 subtype	serotype	77						
<input type="checkbox"/>	▶ H11N6 subtype	serotype	171						
<input type="checkbox"/>	▶ H11N7 subtype	serotype	118						
<input type="checkbox"/>	▶ H11N8 subtype	serotype	134						
<input type="checkbox"/>	▶ H11N9 subtype	serotype	3388						
<input type="checkbox"/>	▶ H12N1 subtype	serotype	75						
<input type="checkbox"/>	▶ H12N2 subtype	serotype	33						
<input type="checkbox"/>	▶ H12N3 subtype	serotype	49						
<input type="checkbox"/>	▶ H12N4 subtype	serotype	84						
<input type="checkbox"/>	▶ H12N5 subtype	serotype	1422						
<input type="checkbox"/>	▶ H12N6 subtype	serotype	48						
<input type="checkbox"/>	▶ H12N7 subtype	serotype	40						
<input type="checkbox"/>	▶ H12N8 subtype	serotype	50						
<input type="checkbox"/>	▶ H12N9 subtype	serotype	42						
<input type="checkbox"/>	▶ H13N1 subtype	serotype	2						
<input type="checkbox"/>	▶ H13N2 subtype	serotype	642						
<input type="checkbox"/>	▶ H13N3 subtype	serotype	31						
<input type="checkbox"/>	▶ H13N4 subtype	serotype	1						
<input type="checkbox"/>	▶ H13N6 subtype	serotype	1896						
<input type="checkbox"/>	▶ H13N8 subtype	serotype	865						
<input type="checkbox"/>	▶ H13N9 subtype	serotype	135						
<input type="checkbox"/>	▶ H14N2 subtype	serotype	16						
<input type="checkbox"/>	▶ H14N3 subtype	serotype	90						
<input type="checkbox"/>	▶ H14N4 subtype	serotype	16						
<input type="checkbox"/>	▶ H14N5 subtype	serotype	38						
<input type="checkbox"/>	▶ H14N6 subtype	serotype	44						
<input type="checkbox"/>	▶ H14N7 subtype	serotype	16						



data

## Dengue

 Taxon View

 Viruses » [Kitrinoviricota](#) » [Flasuviricetes](#) » [Amarillovirales](#) » [Flaviviridae](#) » [Flavivirus](#) » **Dengue virus** ( 39517 Genomes )

Overview	Taxonomy	Genomes	Proteins	Protein Structures	Domains and Motifs	Epitopes	Experiments
Name		Rank	Genomes				
<input type="checkbox"/>	▾ Dengue virus	species	39517				
<input type="checkbox"/>	▸ Dengue virus 1	no rank	13701				
<input type="checkbox"/>	▸ Dengue virus 2	no rank	12087				
<input type="checkbox"/>	▸ Dengue virus 3	no rank	6473				
<input type="checkbox"/>	▸ Dengue virus 4	no rank	3342				

## Ebola

 Taxon View

 Viruses » [Negarnaviricota](#) » [Monjiviricetes](#) » [Mononegavirales](#) » [Filoviridae](#) » **Ebolavirus** ( 3969 Genomes )

Overview	Taxonomy	Genomes	Proteins	Protein Structures	Domains and Motifs	Epitopes	Experiments
Name		Rank	Genomes				
<input type="checkbox"/>	▾ Ebolavirus	genus	3969				
<input type="checkbox"/>	▸ Bombali ebolavirus	species	12				
<input type="checkbox"/>	▸ Bundibugyo ebolavirus	species	43				
<input type="checkbox"/>	▸ Reston ebolavirus	species	59				
<input type="checkbox"/>	▸ Sudan ebolavirus	species	44				
<input type="checkbox"/>	▸ Tai Forest ebolavirus	species	9				
<input type="checkbox"/>	▸ Zaire ebolavirus	species	3720				
<input type="checkbox"/>	▸ unclassified Ebolavirus	no rank	82				

## Enterovirus

 Taxon View

 Viruses »  Pisuviricota »  Pisoniviricetes »  Picornavirales »  Picornaviridae » **Enterovirus** ( 111358 Genomes )

Overview	<b>Taxonomy</b>	Genomes	Proteins	Protein Structures	Domains and Motifs	Epitopes	Experiments
Name		Rank	Genomes				
<input type="checkbox"/>	▲ Enterovirus	genus	111358				
<input type="checkbox"/>	▶ Enterovirus A	species	38978				
<input type="checkbox"/>	▶ Enterovirus B	species	28352				
<input type="checkbox"/>	▶ Enterovirus C	species	12109				
<input type="checkbox"/>	▶ Enterovirus D	species	5239				
<input type="checkbox"/>	▶ Enterovirus E	species	264				
<input type="checkbox"/>	▶ Enterovirus F	species	76				
<input type="checkbox"/>	▶ Enterovirus G	species	829				
<input type="checkbox"/>	▶ Enterovirus H	species	20				
<input type="checkbox"/>	▶ Enterovirus I	species	3				
<input type="checkbox"/>	▶ Enterovirus J	species	140				
<input type="checkbox"/>	▶ Enterovirus K	species	2				
<input type="checkbox"/>	▶ Enterovirus L	species	2				
<input type="checkbox"/>	▶ Rhinovirus A	species	8067				
<input type="checkbox"/>	▶ Rhinovirus B	species	1687				
<input type="checkbox"/>	▶ Rhinovirus C	species	5914				
<input type="checkbox"/>	▶ environmental samples	no rank	95				
<input type="checkbox"/>	▶ unclassified Enterovirus	no rank	2320				
<input type="checkbox"/>	▶ unclassified Rhinovirus	no rank	7210				

## Lassa


 Taxon View

 Viruses »  Negarnaviricota »  Ellioviricetes »  Bunyavirales »  Arenaviridae »  Mammarenavirus » **Lassa mammarenavirus** ( 2328 C

Overview	<b>Taxonomy</b>	Strains	Genomes	Proteins	Protein Structures	Domains and Motifs	Epitopes	Experiments
Name		Rank	Genomes					
<input type="checkbox"/>	▲ Lassa mammarenavirus	species	2328					
<input type="checkbox"/>	▶ Lassa virus Josiah	no rank	1					



## Rota

 Taxon View

 Viruses » [Duplornaviricota](#) » [Resentoviricetes](#) » [Reovirales](#) » **Reoviridae** ( 136480 Genomes )

	Overview	<b>Taxonomy</b>	Genomes	Proteins	Protein Structures	Domains and Motifs	Epitopes	Experiments
		Name	Rank	Genomes				
<input type="checkbox"/>	▾	Reoviridae	family	136480				
<input type="checkbox"/>	▾	Sedoreovirinae	subfamily	123822				
<input type="checkbox"/>	▶	Cardoreovirus	genus	27				
<input type="checkbox"/>	▶	Mimoreovirus	genus	22				
<input type="checkbox"/>	▶	Orbivirus	genus	18100				
<input type="checkbox"/>	▶	Phytoreovirus	genus	390				
<input checked="" type="checkbox"/>	▾	Rotavirus	genus	104625				
<input type="checkbox"/>	▶	Rotavirus A	species	93527				
<input type="checkbox"/>	▶	Rotavirus B	species	1736				
<input type="checkbox"/>	▶	Rotavirus C	species	4807				
<input type="checkbox"/>	▶	Rotavirus D	species	199				
<input type="checkbox"/>	▶	Rotavirus F	species	90				
<input type="checkbox"/>	▶	Rotavirus G	species	141				
<input type="checkbox"/>	▶	Rotavirus H	species	429				
<input type="checkbox"/>	▶	Rotavirus I	species	77				
<input type="checkbox"/>	▶	Rotavirus J	species	22				
<input type="checkbox"/>	▶	unclassified Rotavirus	no rank	3440				
<input type="checkbox"/>	▶	Seadornavirus	genus	658				
<input type="checkbox"/>	▶	Spinareovirinae	subfamily	11031				
<input type="checkbox"/>	▶	unclassified Reoviridae	no rank	1627				

## Herpes

 Taxon View

 Viruses » [Peploviricota](#) » [Herviviricetes](#) » [Herpesvirales](#) » **Herpesviridae** ( 63480 Genomes )

	Overview	<b>Taxonomy</b>	Genomes	Proteins	Protein Structures	Domains and Motifs	Epitopes	Experiments
		Name	Rank	Genomes				
<input type="checkbox"/>	▾	Herpesviridae	family	63480				
<input type="checkbox"/>	▶	Alphaherpesvirinae	subfamily	29091				
<input type="checkbox"/>	▶	Betaherpesvirinae	subfamily	14436				
<input type="checkbox"/>	▶	Gammaherpesvirinae	subfamily	19360				
<input type="checkbox"/>	▶	Iguanid herpesvirus 2	species	2				
<input type="checkbox"/>	▶	environmental samples	no rank	2				
<input type="checkbox"/>	▶	unclassified Herpesviridae	no rank	589				

