

COVID-19 is an emerging, rapidly evolving situation.
 Get the latest public health information from CDC: <https://www.coronavirus.gov>.
 Get the latest research from NIH: <https://www.nih.gov/coronavirus>.
 Find NCBI SARS-CoV-2 literature, sequence, and clinical content: <https://www.ncbi.nlm.nih.gov/sars-cov-2/>.

BLAST® » [blastn suite](#) » results for RID-T04UZR6B016

Job Title [Nucleotide Sequence ...](#)
 RID [T04UZR6B016](#) Search expires on 10-22 18:54 pm
 Program BLASTN
 Database nt
 Query ID lcl|Query_27527
 Description [None ...](#)
 Molecule type dna
 Query Length 10009

Descriptions

Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
Abalistes stellaris mitochondrial DNA, complete genome	18484	18484	100%	0.0	100.00%	AP009202.1
Abalistes stellaris mitochondrion, complete genome	12497	17328	100%	0.0	96.82%	MG515171.1
Balistapus undulatus mitochondrial DNA, complete genome	12338	12338	99%	0.0	88.96%	AP009203.1
Rhinecanthus aculeatus mitochondrial DNA, complete genome	12059	12059	100%	0.0	88.44%	AP009210.1
Pseudobalistes flavimarginatus mitochondrial DNA, complete genome	11897	11897	100%	0.0	88.17%	AP009209.1
Odonus niger mitochondrial DNA, complete genome	11865	11865	99%	0.0	88.11%	AP009208.1
Balistes vetula mitochondrial DNA, complete genome	11814	11814	100%	0.0	88.00%	AP009204.1
Pseudobalistes fuscus mitochondrion, complete genome	11786	11786	100%	0.0	87.95%	KU985150.1
Xanthichthys auromarginatus mitochondrial DNA, complete genome	11707	11707	100%	0.0	87.82%	AP009211.1
Xenobalistes tumidipectoris mitochondrial DNA, complete genome	11673	11673	100%	0.0	87.76%	AP009182.1
Balistoides conspicillum mitochondrial DNA, complete genome	11642	11642	99%	0.0	87.73%	AP009205.1
Sufflamen fraenatum mitochondrial DNA, complete genome	11638	11638	100%	0.0	87.69%	AP004456.1
Melichthys vidua mitochondrial DNA, complete genome	11585	11585	99%	0.0	87.61%	AP009207.1
Thamnaconus modestus mitochondrial DNA, complete genome	8835	8835	99%	0.0	82.72%	AP009185.1
Monacanthus chinensis mitochondrial DNA, complete genome	8820	8820	99%	0.0	82.71%	AP009219.1
Paramonacanthus choirocephalus mitochondrial DNA, complete genome	8770	8770	99%	0.0	82.62%	AP009223.1
Thamnaconus tessellatus mitochondrion, complete genome	8741	8741	99%	0.0	82.56%	KJ009561.1
Pterocaesio digramma mitochondrial DNA, complete genome	8696	8696	99%	0.0	82.49%	LC549803.1
Eubalichthys mosaicus mitochondrial DNA, complete genome	8696	8696	99%	0.0	82.47%	AP009217.1
Thamnaconus hypargyreus mitochondrion, complete genome	8674	8674	99%	0.0	82.45%	KJ743986.1