
ReQON Documentation

Release 0.1.0

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August 11, 2016

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ReQON ([ikn], /rikn/, RE-kon) lets you build simple, read-only ReQL queries from a JSON data structure.

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Query descriptor

The ReQON query descriptor is an object with specific attributes:

```
{
  '$db': 'imdb',
  '$table': 'movies',
  '$query': [],
}
```

1.1 \$db

References a specific database.

ReQON	ReQL
{ '\$db': 'imdb' }	r.db('imdb')

This attribute is optional.

1.2 \$table

Return all of the documents in the specified table of the default database.

ReQON	ReQL
{ '\$table': 'movies' }	r.table('movies')
{ '\$db': 'imdb', '\$table': 'movies' }	r.db('imdb').table('movies')

This attribute is required.

1.3 \$query

The query attribute is a sequence of [terms](#) that filter, manipulate, or aggregate the document sequence in some way. Each term in the sequence is a list of 1 or 2 items, where the first item is the name of the term, followed (optionally, depending on the term) by a list of arguments.

This attribute is optional; omitting this attribute is the same as fetching every document in the specified \$table.

Terms

2.1 Selecting data

2.1.1 \$get

`get (key)`

See also: <http://rethinkdb.com/api/python/get/>

2.1.2 \$get_all

`get_all (keys[, index])`

See also: http://rethinkdb.com/api/python/get_all/

2.1.3 \$filter

`filter (predicate)`

Boolean operator	ReQL	See also:
<code>\$and</code>	<code>r.and_</code>	http://rethinkdb.com/api/python/and/
<code>\$or</code>	<code>r.or_</code>	http://rethinkdb.com/api/python/or/
<code>\$not</code>	<code>r.not_</code>	http://rethinkdb.com/api/python/not/

See also: <http://rethinkdb.com/api/python/filter/>

2.1.4 \$between

`between (lower_key, upper_key[, index, left_bound, right_bound])`

2.2 Transformations

2.2.1 \$has_fields

`has_fields (fields)`

See also: http://rethinkdb.com/api/python/has_fields/

2.2.2 \$with_fields

with_fields (*fields*)

See also: http://rethinkdb.com/api/python/with_fields/

2.2.3 \$order_by

order_by (*key* | *index*[, *ordering*])

See also: http://rethinkdb.com/api/python/order_by/

2.2.4 \$skip

skip (*n*)

See also: <http://rethinkdb.com/api/python/skip/>

2.2.5 \$limit

limit (*n*)

See also: <http://rethinkdb.com/api/python/limit/>

2.2.6 \$slice

slice (*start_offset*[, *end_offset*, *left_bound*, *right_bound*])

See also: <http://rethinkdb.com/api/python/slice/>

2.2.7 \$nth

nth (*n*)

See also: <http://rethinkdb.com/api/python/nth/>

2.2.8 \$sample

sample (*n*)

See also: <http://rethinkdb.com/api/python/sample/>

2.3 Manipulation

2.3.1 \$pluck

pluck (*fields*)

See also: <http://rethinkdb.com/api/python/pluck/>

2.3.2 \$without

without (*fields*)

See also: <http://rethinkdb.com/api/python/without/>

2.4 Aggregation

2.4.1 \$group

group (*field* | *index*[, *multi*])

See also: <http://rethinkdb.com/api/python/group/>

2.4.2 \$count

count ()

See also: <http://rethinkdb.com/api/python/count/>

2.4.3 \$sum

sum (*field*)

See also: <http://rethinkdb.com/api/python/sum/>

2.4.4 \$avg

avg (*field*)

See also: <http://rethinkdb.com/api/python/avg/>

2.4.5 \$min

min (*field*)

See also: <http://rethinkdb.com/api/python/min/>

2.4.6 \$max

max (*field*)

See also: <http://rethinkdb.com/api/python/max/>

2.5 Geospatial

2.5.1 \$get_intersecting

`get_intersecting (geometry, index)`

See also: http://rethinkdb.com/api/python/get_intersecting/

2.5.2 \$get_nearest

`get_nearest (geometry, index[, max_results, max_dist, unit, geo_system])`

See also: http://rethinkdb.com/api/python/get_nearest/

Selecting data

Term	Description
<code>\$get</code>	Get a single document by its primary key
<code>\$get_all</code>	Get all documents where the given value matches the requested index
<code>\$filter</code>	Get all the documents for which the specified sequence is true

Transformations

Term	Description
<code>\$with_fields</code>	Exclude documents that do not have the specified fields and return only those fields
<code>\$has_fields</code>	Test if a document has the specified fields, filtering out any that do not
<code>\$order_by</code>	Sort the documents by the specified field or index
<code>\$skip</code>	Skip a number of documents from the head of the sequence
<code>\$limit</code>	End the sequence after the given number of documents
<code>\$slice</code>	Return the documents within the specified range
<code>\$nth</code>	Get the <i>n</i> th document in the sequence
<code>\$sample</code>	Select a given number of elements from a sequence with uniform random distribution

Manipulation

Term	Description
<code>\$pluck</code>	Return only the specified fields
<code>\$without</code>	The opposite of <code>\$pluck</code> , return the documents without the specified fields

Aggregation

Term	Description
<code>\$group</code>	Partition the documents into multiple groups based on the specified field
<code>\$count</code>	Count the number of documents in the sequence
<code>\$sum</code>	Sum the specified field of the sequence
<code>\$avg</code>	Average the specified field of the sequence
<code>\$min</code>	Find the minimum value of the specified field in the sequence
<code>\$max</code>	Find the maximum value of the specified field of the sequence

Geospatial

Term	Description
<code>\$get_intersecting</code>	Get all documents where the given geometry object intersects with a geometry object of a geospatial index
<code>\$get_nearest</code>	Return the documents closest to the specified point based on a geospatial index

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