

Soha Khalid

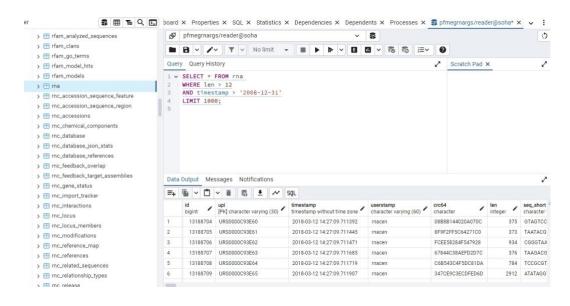
BWT - Data Engineering

Task 02 - Exercise

You will be using Rnacen (RNACentral) schema to find and explore the data available in all tables that will help you to recognise the potential tables you can query to answer the following questions.

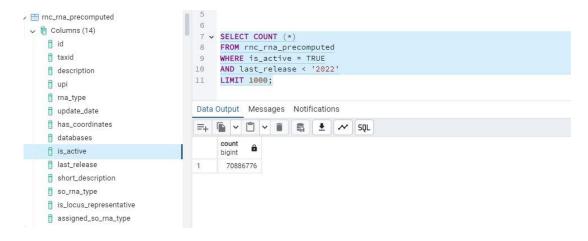
1. Write a query to get data having length of Rna structures more than 12 with them being added after 2008.

SELECT * FROM rna
WHERE len > 12
AND timestamp > '2008-12-31'
LIMIT 1000;



2. How many pre computed RNA are present that are still active and got their last release update before 2022.

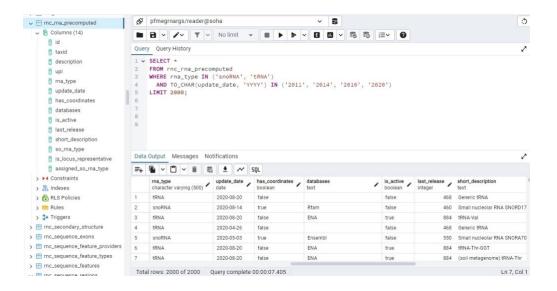
```
SELECT COUNT (*)
FROM rnc_rna_precomputed
WHERE is_active = TRUE
AND last_release < '2022'
LIMIT 1000;
```

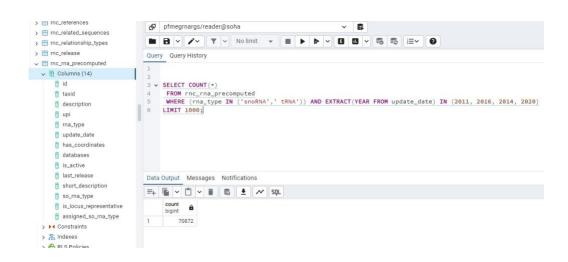


3. How many total pre computed RNA records for snoRNA and tRNA were recorded in 2011, 2016, 2014, and 2020.

```
SELECT *
FROM rnc_rna_precomputed
WHERE rna_type IN ('snoRNA', 'tRNA')
AND TO_CHAR(update_date, 'YYYY') IN ('2011', '2014', '2016', '2020')
LIMIT 1000;

SELECT COUNT(*)
FROM rnc_rna_precomputed
WHERE (rna_type IN ('snoRNA',' tRNA')) AND EXTRACT(YEAR FROM update_date) IN (2011, 2016, 2014, 2020)
LIMIT 1000;
```



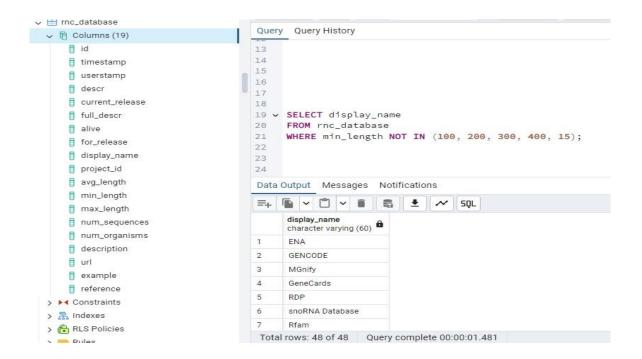


4. Can you give me the names of all databases built for RNA with minimum length other than 100, 200, 300, 400, and 15.

SELECT display_name

FROM rnc_database

WHERE min length NOT IN (100, 200, 300, 400, 15);



5. Can you get complete 500 records of sequences for active regions and name your column as myregions in which you are getting the region name column value. Then tell me what different chromosomes with exon_count we have for regions including center, east and north using the name you set for your column.

SELECT region name AS myregions

FROM rnc_sequence_regions

LIMIT 500;

For last part no result can be retrieved from data because no such value named "center, north and east" existed in the given database.

