University of British Columbia, Vancouver

Department of Computer Science

CPSC 304 Project Cover Page

Milestone #: 4

Date: Nov. 23, 2022

Group Number: 20

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By typing our names and student numbers in the above table, we certify that the work in the attached assignment was performed solely by those whose names and student IDs are included above. (In the case of Project Milestone 0, the main purpose of this page is for you to let us know your e-mail address, and then let us assign you to a TA for your project supervisor.)

In addition, we indicate that we are fully aware of the rules and consequences of plagiarism, as set forth by the Department of Computer Science and the University of British Columbia

Project Description:

Repo link: https://github.students.cs.ubc.ca/CPSC304-2022W-T1/project b7e6r d3e2b n3q8v

The final project initializes a database according to the schema provided in earlier milestones and is populated with mostly real, sample data, which are epidemiological information of viruses. We've implemented a multi-page web application that uses our database. The app, through a GUI, is capable of running queries that modify the database (deleting, inserting, and updating entries). Also, it can show the result of queries that filter the data through projection and selection, joins data from two tables, and runs division and aggregation queries that reveal new information about the data. Most functionalities also allow for user input, apart from division and aggregation.

Most of the schema submitted earlier remains the same in the final project. Although, as Oracle does not support assertions, certain participation constraints are not enforced. These are indicated by a comment in the initialization script (<u>initialization.sql</u> in the root folder).

A copy of the schema can be found in schema.sql in the "Documents" folder, which contains only create table statements from the initialization script.

Screenshots of the data:

Virus1

FAMILY	STRANDEDNESS
Coronaviridae Poxviridae Retroviridae Orthomyxoviridae Picornaviridae Secoviridae Herpesviridae Microviridae Filoviridae Paramyxoviridae Hepadnaviridae Papillomaviridae	Single Double Single Single Single Single Single Single Single Double Single Single Single Single Single Single

Virus2

GENUS	FAMILY
Betacoronavirus Orthopoxvirus Lentivirus Alphainfluenzavirus Enterovirus Comovirus Simplexvirus Sinsheimervirus Ebolavirus Morbillivirus Orthohepadnavirus Chipapillomavirus	Coronaviridae Poxviridae Retroviridae Orthomyxoviridae Picornaviridae Secoviridae Herpesviridae Microviridae Filoviridae Paramyxoviridae Hepadnaviridae Papillomaviridae

Virus3

VIRUSCOMMONNAME	GENUS	STATUS	TRANSMISSIONTYPE
SARS-CoV	 Betacoronavirus	Active	Respiratory Droplets
Variola Virus	Orthopoxvirus	Eradicated	Airborne Particles
Human Immunodeficiency Virus	Lentivirus	Active	Bodily Fluids
Monkeypox Virus	Orthopoxvirus	Active	Direct Contact
Influenza A Virus	Alphainfluenzavirus	Active	Respiratory Droplets
Poliovirus	Enterovirus	Active	Fecal-oral
Vaccinia Virus	Orthopoxvirus	Active	Direct Contact
Cowpea Mosaic Virus	Comovirus	Active	Insects
Herpes Simplex Virus	Simplexvirus	Active	Bodily Fluids
Phi X 174	Sinsheimervirus	Active	Bacterial
Zaire Ebolavirus	Ebolavirus	Active	Bodily Fluids
Rinderpest morbillivirus	Morbillivirus	Eradicated	Direct Contact
Hepatitis B <u>Virus</u>	Orthohepadnavirus	Active	Bodily Fluids
Human Papillomavirus	Chipapillomavirus	Active	Direct Contact

RNAVirus

VIRUSCOMMONNAME	SENSE
SARS-CoV Human Immunodeficiency Virus Influenza A Virus Poliovirus Cowpea Mosaic Virus Zaire Ebolavirus Rinderpest morbillivirus	Positive Positive Negative Positive Positive Positive Negative Negative Negative

DNAVirus

VIRUSCOMMONNAME	GENOMESHAPE
Variola Virus Monkeypox Virus Vaccinia Virus Herpes Simplex Virus Phi X 174 Hepatitis B Virus Human Papillomavirus	Linear Linear Linear Linear Circular Linear

VaccineAgainst1

TYPE	IMMUNOCOMPROMISESAFETY
Attenuated Inactivated Subunit mRNA Viral Vector	Unsafe Safe Safe Safe Safe

VaccineAgainst2

VIRUSCOMMONNAME	TYPE	MANUFACTURE	VALENCE	DELIVERYMODE	YEAR
Zaire Ebolavirus SARS-CoV Hepatitis B Virus SARS-CoV Poliovirus Human Papillomavirus	Viral Vector mRNA Subunit Inactivated Attenuated Subunit	Merck & Co. Pfizer Merck & Co. Sinovac Biotech Cantacuzino Institute of Bucharest Merck & Co.	2 5 1 1	Injection Injection Injection Injection Injection Oral Injection	2014 2021 2019 2021 1961 2014

Host

HOSTCOMMONNAME	TYPE
Human E. coli Cowpea Deer Buffalo	Mammal Bacteria Plant Mammal Mammal

Infects

HOSTCOMMONNAME	VIRUSCOMMONNAME
HOSTCOMMONNAME	VIRUSCOMMONNAME
Human	Poliovirus
Human	Vaccinia Virus
Human Human	Variola Virus Zaire Ebolavirus

Receptor

RECEPTORNAME	CELLTYPE	TISSUETYPE
ACE2 MARC0 CCR5 GAG Sialic Acid CD155	Enterocytes Keratinocytes Immune Cells Keratinocytes All Immune Cells	Epithelia Epidermis All Epidermis All

Targets

RECEPTORNAME	VIRUSCOMMONNAME
ACE2	SARS-CoV
CCR5	Human Immunodeficiency Virus
CD155	Poliovirus
GAG	Monkeypox Virus
GAG	Vaccinia Virus
MARCO	Variola Virus
Sialic Acid	Influenza A Virus

ViralDisease

DISEASENAME	DISEASETYPE
Smallpox AIDS Hepatitis Herpes Labialis Influenza	Acute Chronic Acute Dormant Acute

Causes

DISEASENAME	VIRUSCOMMONNAME
AIDS Hepatitis Herpes Labialis Influenza Smallpox	Human Immunodeficiency Virus Hepatitis B Virus Herpes Simplex Virus Influenza A Virus Variola Virus

Symptom

SYMPTOMNAME	SPECIFICITY
Blisters Vomit Diarrhea Lymphadenopathy Macules	Non-specific Non-specific Non-specific Specific Specific

Has

DISEASENAME	SYMPTOMNAME	SEVERITY
Herpes Labialis Hepatitis Hepatitis AIDS Smallpox	Blisters Vomit Diarrhea Lymphadenopathy Macules	Mild Mild Mild Severe Severe

Country

COUNTRYNAME	POPULATIONDENSITY	CONTINENT	STATUS
United States China Mexico South Africa	153 66	North America Asia North America Africa	Developed Developing Developing Developed Developed Developed
Japan Congo Canada	16	Africa North America	Underdeveloped Developed

Outbreak

COUNTRYNAME	VIRUSCOMMONNAME	OUTBREAKSIZE	CASUALTY	YEAR	ORIGIN
United States	Influenza A Virus	Pandemic	100000000	 1346	Spain
China	SARS-CoV	Pandemic	25000000	2019	China
Mexico	Variola Virus	Epidemic	8000000	1519	Mexico
Japan	Variola Virus	Epidemic	2000000	735	Japan
Congo	Zaire Ebolavirus	Epidemic	55	2020	Congo
China	Influenza A Virus	Pandemic	4000000	1968	Hong Kong
Canada	Human Immunodeficiency Virus	Pandemic	40100000	1981	Central Africa

EndemicTo

COUNTRYNAME	VIRUSCOMMONNAME
Canada	Hepatitis B Virus
China	Influenza A Virus
Congo	Human Immunodeficiency Virus
South Africa	Human Immunodeficiency Virus
United States	Influenza A Virus

Application

APPLICATIONNAME	USAGE
Weapon Synthetic Virus Nanotechnology Virotherapy Vaccine	Biological Warfare Research Research Medicine Medicine

UsedIn

APPLICATIONNAME	VIRUSCOMMONNAME
Nanotechnology Synthetic Virus Vaccine Virotherapy Weapon	Cowpea Mosaic Virus Poliovirus Poliovirus Herpes Simplex Virus Vaccinia Virus

```
List of queries:
* "[]" indicates user input.
INSERT INTO Virus1 (Family, Strandedness) []
INSERT INTO Virus2 (Genus, Family) []
INSERT INTO Virus3 (virusCommonName, Genus, Status, TransmissionType) []
INSERT INTO RNAVirus (virusCommonName, Sense) []
INSERT INTO DNAVirus (virusCommonName, genomeShape) []
DELETE FROM Virus3
      WHERE virusCommonName = []
UPDATE RNAVirus SET Sense = []
      WHERE virusCommonName = []
UPDATE DNAVirus SET genomeShape = []
      WHERE virusCommonName = []
UPDATE RNAVirus SET Status = [], Genus = [], TransmissionType = []
      WHERE virusCommonName = []
* all insert, delete, and update queries can be found in insert, delete, update.php
SELECT *
      FROM []
      WHERE []
SELECT []
      FROM []
* selection and projection queries can be found in <u>select-project.php</u>
SELECT DNAVirus.virusCommonName, Genus, Status, TransmissionType, GenomeShape
       FROM DNAVirus, Virus3
      WHERE DNAVirus.virusCommonName = Virus3.virusCommonName AND []
SELECT RNAVirus.virusCommonName, Genus, Status, TransmissionType, Sense
       FROM RNAVirus, Virus3
      WHERE RNAVirus.virusCommonName = Virus3.virusCommonName AND []
SFLFCT DISTINCT D.virusCommonName
```

```
FROM DNAVirus D
```

WHERE NOT EXISTS ((SELECT ho.hostCommonName

FROM Host ho)

MINUS

(SELECT I.hostCommonName

FROM Infects I

WHERE D.virusCommonName = I.virusCommonName))

* all join and divide queries can be found in join-div.php

SELECT virusCommonName AS Virus, AVG(Casualty) AS Average_Deaths

FROM Outbreak

WHERE Casualty > 1000000

GROUP BY virusCommonName

HAVING COUNT(*) > 1

ORDER BY Average_Deaths DESC

SELECT TransmissionType AS Transmission, COUNT(*) AS Viruses Count

FROM Virus3

GROUP BY TransmissionType

ORDER BY Viruses_Count DESC

SELECT c.diseaseName AS Disease, SUM(o.Casualty) AS Death Count

FROM Causes c, Outbreak o

WHERE c.virusCommonName = o.virusCommonName AND c.diseaseName IN

(SELECT v.diseaseName

FROM ViralDisease v

WHERE v.diseaseType = 'Acute')

GROUP BY c.diseaseName

ORDER BY Death Count DESC

^{*} all aggregation queries can be found in <u>aggregations.php</u>

Graphical User Interface Screenshots:

1. Selection and Projection Queries:

a. Users' inputs and initial database:

Table used: Virus3

VIRUSCOMMONNAME	GENUS	STATUS	TRANSMISSIONTYPE
SARS-CoV	Betacoronavirus	 Active	Respiratory Droplets
Variola Virus	Orthopoxvirus	Eradicated	Airborne Particles
Human Immunodeficiency Virus	Lentivirus	Active	Bodily Fluids
Monkeypox Virus	Orthopoxvirus	Active	Direct Contact
Influenza A Virus	Alphainfluenzavirus	Active	Respiratory Droplets
Poliovirus	Enterovirus	Active	Fecal-oral
Vaccinia Virus	Orthopoxvirus	Active	Direct Contact
Cowpea Mosaic Virus	Comovirus	Active	Insects
Herpes Simplex Virus	Simplexvirus	Active	Bodily Fluids
Phi X 174	Sinsheimervirus	Active	Bacterial
Zaire Ebolavirus	Ebolavirus	Active	Bodily Fluids
Rinderpest morbillivirus	Morbillivirus	Eradicated	Direct Contact
Hepatitis B <u>Virus</u>	Orthohepadnavirus	Active	Bodily Fluids
Human Papillomavirus	Chipapillomavirus	Active	Direct Contact

Projection

Available Data:	Vaccine (3	
Choose			
	acture <mark>▼</mark> D	OeliveryM	ode V Year
Select Attributes			

Selection

Avaiable data: Vaccine	e 😊
Number of constraints	: 2
Create conditions	
DeliveryMode 😊 =	Injection
OR 😊 Year	② = ② 2021
Use The Condition	
Submit	

b. Results:

Selection:

VIRUSCOMMONNAME	ТҮРЕ	MANUFACTURE	VALENCE	DELIVERYMODE	YEAR
Zaire Ebolavirus	Viral Vector	Merck & Co.	1	Injection	2014
SARS-CoV	mRNA	Pfizer	2	Injection	2021
Hepatitis B Virus	Subunit	Merck & Co.	5	Injection	2019
SARS-CoV	Inactivated	Sinovac Biotech	1	Injection	2021
Human Papillomavirus	Subunit	Merck & Co.	9	Injection	2014

Projection:

TYPE	MANUFACTURE
Attenuated	Cantacuzino Institute of Bucharest
Inactivated	Sinovac Biotech
Subunit	Merck & Co.
Subunit	Merck & Co.
Viral Vector	Merck & Co.
mRNA	Pfizer

2. Join and Divide Queries:

a. Users Inputs:

Join:

Tabled used: RNAVirus, Virus3

VIRUSCOMMONNAME	SENSE
SARS-CoV Human Immunodeficiency Virus Influenza A Virus Poliovirus Cowpea Mosaic Virus Zaire Ebolavirus Rinderpest morbillivirus	Positive Positive Negative Positive Positive Negative Negative

VIRUSCOMMONNAME	GENUS	STATUS	TRANSMISSIONTYPE
SARS-CoV	Betacoronavirus	Active	Respiratory Droplets
Variola Virus	Orthopoxvirus	Eradicated	Airborne Particles
Human Immunodeficiency Virus	Lentivirus	Active	Bodily Fluids
Monkeypox Virus	Orthopoxvirus	Active	Direct Contact
Influenza A Virus	Alphainfluenzavirus	Active	Respiratory Droplets
Poliovirus	Enterovirus	Active	Fecal-oral
Vaccinia Virus	Orthopoxvirus	Active	Direct Contact
Cowpea Mosaic Virus	Comovirus	Active	Insects
Herpes Simplex Virus	Simplexvirus	Active	Bodily Fluids
Phi X 174	Sinsheimervirus	Active	Bacterial
Zaire Ebolavirus	Ebolavirus	Active	Bodily Fluids
Rinderpest morbillivirus	Morbillivirus	Eradicated	Direct Contact
Hepatitis B <u>Virus</u>	Orthohepadnavirus	Active	Bodily Fluids
Human Papillomavirus	Chipapillomavirus	Active	Direct Contact

RNA Virus And DNA Virus

Number of c	onstraints:	2				
Find The Inf	ormation:	RNA \	/irus	0		
Status	© =	©	Active	е		
AND 🔞 Se	nse	©	<>	•	Negative	
Use The Cond	dition					
Submit						

Divide:

```
SQL> select * from DNAvirus;
VIRUSCOMMONNAME
                                    GENOMESHAPE
Variola Virus
                                    Linear
Monkeypox Virus
                                   Linear
Vaccinia Virus
                                   Linear
Herpes Simplex Virus
Phi X 174
Hepatitis B Virus
Human Papillomavirus
                                    Linear
                                    Circular
                                    Linear
                                    Circular
Example Virus
                                    Updated Genome Shape
8 rows selected.
```

HOSTCOMMONNAME	ТҮРЕ
Human E. coli Cowpea Deer Buffalo	Mammal Bacteria Plant Mammal Mammal

HOSTCOMMONNAME	VIRUSCOMMONNAME
Buffalo Buffalo Buffalo Cowpea Cowpea Cowpea Deer Deer Deer E. coli	Rinderpest morbillivirus Vaccinia Virus Variola Virus Cowpea Mosaic Virus Vaccinia Virus Variola Virus Rinderpest morbillivirus Vaccinia Virus Vaccinia Virus Phi X 174
E. coli HOSTCOMMONNAME	Vaccinia Virus VIRUSCOMMONNAME
E. coli Human	Variola Virus Hepatitis B Virus Herpes Simplex Virus Human Immunodeficiency Virus Human Papillomavirus Influenza A Virus Monkeypox Virus Poliovirus SARS-CoV Vaccinia Virus Variola Virus
HOSTCOMMONNAME	VIRUSCOMMONNAME
Human	Zaire Ebolavirus

Find DNA Viruses infect all hosts Submit

b. Results:

Join:

VIRUSCOMMONNAME	GENUS	STATUS	TRANSMISSIONTYPE	SENSE
Cowpea Mosaic Virus	Comovirus	Active	Insects	Positive
Human Immunodeficiency Virus	Lentivirus	Active	Bodily Fluids	Positive
Poliovirus	Enterovirus	Active	Fecal-oral	Positive
SARS-CoV	Betacoronavirus	Active	Respiratory Droplets	Positive

Divide:

Find DNA Viruses infect all hosts

VIRUSCOMMONNAME		
Vaccinia Virus		
Variola Virus		

3. Insert, Delete, Update Queries:

initial database:

VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
51A1U5	TRANSMISSIONITYE
SARS-CoV	Betacoronavirus
Active	Respiratory Droplets
Variola Virus	Orthopoxvirus
Eradicated	Airborne Particles
Human Immunodeficiency Virus	Lentivirus
Active	Bodily Fluids
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Monkeypox Virus	Orthopoxvirus
Active	Direct Contact
Influenza A Virus	Alphainfluenzavirus
Active	Respiratory Droplets
Poliovirus	Enterovirus
Active	Fecal-oral
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Vaccinia Virus	Orthopoxvirus
Active	Direct Contact
Cowpea Mosaic Virus	Comovirus
Active	Insects
Herpes Simplex Virus	Simplexvirus
Active	Bodily Fluids
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Phi X 174	Sinsheimervirus
Active	Bacterial
Zaire Ebolavirus	Ebolavirus
Active	Bodily Fluids
Rinderpest morbillivirus	Morbillivirus
Eradicated	Direct Contact
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Hepatitis B Virus	Orthohepadnavirus

Insert input and results: Insertion, Deletion, Update

(if select RNA, leave Genome Shape empty. If select DNA, leave Sense empty.) $% \left(\frac{1}{2}\right) =\left(\frac{1}{2}\right) \left(\frac{1}{2}\right) \left$
Type: DNA ▼
Virus Common Name: Example Virus
Genome Shape: Example Genome Shape
Sense:
Strandedness: Example Strandedness
Genus: Example Genus
Family: Example Family
Status: Example Status
TransmissionType: Example TranmissionType
Insert
Delete To delete input the virus common name
Update

VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
 Monkeypox Virus Active	Orthopoxvirus Direct Contact
Influenza A Virus	Alphainfluenzavirus
Active	Respiratory Droplets
Poliovirus	Enterovirus
Active	Fecal-oral
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Vaccinia Virus	Orthopoxvirus
Active	Direct Contact
Cowpea Mosaic Virus	Comovirus
Active	Insects
Herpes Simplex Virus	Simplexvirus
Active	Bodily Fluids
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Phi X 174	Sinsheimervirus
Active	Bacterial
Zaire Ebolavirus	Ebolavirus
Active	Bodily Fluids
Rinderpest morbillivirus	Morbillivirus
Eradicated	Direct Contact
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Hepatitis B Virus	Orthohepadnavirus
Active	Bodily Fluids
Human Papillomavirus	Chipapillomavirus
Active	Direct Contact
Example Virus	Example Genus
Example Status	Example TranmissionType

SQL> select* from virus2;	
GENUS	FAMILY
Betacoronavirus Orthopoxvirus Lentivirus Alphainfluenzavirus Enterovirus Comovirus Simplexvirus Sinsheimervirus Ebolavirus Morbillivirus Orthohepadnavirus	Coronaviridae Poxviridae Retroviridae Orthomyxoviridae Picornaviridae Secoviridae Herpesviridae Microviridae Filoviridae Paramyxoviridae Hepadnaviridae
GENUS	FAMILY
Chipapillomavirus Example Genus	Papillomaviridae Example Family
13 rows selected.	

VIRUSCOMMONNAME GENOMESHAPE

Variola Virus Linear
Vaccinia Virus Linear
Herpes Simplex Virus Linear
Phi X 174 Circular
Hepatitis B Virus Linear
Human Papillomavirus Circular
Example Virus Example Genome Shape

FAMILY	STRANDEDNESS
Coronaviridae	Single
Poxviridae	Double
Retroviridae	Single
Orthomyxoviridae	Single
Picornaviridae	Single
Secoviridae	Single
Herpesviridae	Double
Microviridae	Single
Filoviridae	Single
Paramyxoviridae	Single
Hepadnaviridae	Double
FAMILY	STRANDEDNESS
Papillomaviridae	Double
Example Family	Example Strandedness

Update Input and Results:

Insertion, Deletion, Update

(if select RNA, leave Genome Shape empty. If select DNA, leave Sense empty.)
Type: DNA ✔
Virus Common Name: Example Virus
Genome Shape: Updated Genome Shape
Sense:
Strandedness:
Genus: Orthohepadnavirus
Family:
Status: Updated Status
TransmissionType: Updated TransmissionType
Insert
Delete To delete input the virus common name
Update

/IRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Monkeypox Virus Active	Orthopoxvirus Direct Contact
Influenza A Virus Active	Alphainfluenzavirus Respiratory Droplets
Poliovirus Active	Enterovirus Fecal-oral
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
	Orthopoxvirus Direct Contact
Cowpea Mosaic Virus Active	Comovirus Insects
Herpes Simplex Virus Active	Simplexvirus Bodily Fluids
VIRUSCOMMONNAME	GENUS
STATUS	TRANSMISSIONTYPE
Phi X 174 Active	Sinsheimervirus Bacterial
Zaire Ebolavirus Active	Ebolavirus Bodily Fluids
Rinderpest morbillivirus Eradicated	Morbillivirus Direct Contact
VIRUSCOMMONNAME	GENUS
VIROSCOMMONIVAME	
	TRANSMISSIONTYPE
STATUS 	TRANSMISSIONTYPE Orthohepadnavirus Bodily Fluids
STATUS 	Orthohepadnavirus

SQL> select * from DNAvirus; VIRUSCOMMONNAME GENOMESHAPE Variola Virus Monkeypox Virus Vaccinia Virus Herpes Simplex Virus Linear Linear Linear Linear Circular Phi X 174 Hepatitis B Virus Human Papillomavirus Linear Circular Example Virus Updated Genome Shape 8 rows selected.

Delete Input and results:

Insertion, Deletion, Update

(if select RNA, leave Genome Shape empty. If select DNA, leave Sense empty.)
Type: DNA ▼
Virus Common Name: Vaccinia Virus
Genome Shape:
Sense:
Strandedness:
Genus:
Family:
Status:
TransmissionType:
Insert
Delete To delete input the virus common name
Update

VIRUSCOMMONNAME TRANSMISSIONTYPE SARS-CoV Betacoronavirus Respiratory Droplets Variola Virus Eradicated Orthopoxvirus Airborne Particles Lentivirus Bodily Fluids Human Immunodeficiency Virus VIRUSCOMMONNAME GENUS STATUS TRANSMISSIONTYPE Orthopoxvirus Direct Contact Monkeypox Virus Influenza A Virus Active Alphainfluenzavirus Respiratory Droplets Enterovirus Fecal-oral Poliovirus Active VIRUSCOMMONNAME GENUS STATUS TRANSMISSIONTYPE owpea Mosaic Virus Comovirus Active Insects Simplexvirus Bodily Fluids Herpes Simplex Virus Active Sinsheimervirus Bacterial Phi X 174 Active VIRUSCOMMONNAME GENUS STATUS TRANSMISSIONTYPE Ebolavirus Bodily Fluids Zaire Ebolavirus Active Orthohepadnavirus Bodily Fluids Hepatitis B Virus Example Virus Updated Status Orthohepadnavirus Updated TransmissionType

ON DELETE CASCADE example:

SQL> select * from DNAvirus;	
VIRUSCOMMONNAME	GENOMESHAPE
Variola Virus Monkeypox Virus Herpes Simplex Virus Phi X 174 Hepatitis B Virus Example Virus	Linear Linear Linear Circular Linear Updated Genome Shape
6 rows selected	

4. Aggregation Queries:

Aggregation with GROUP BY:

Tables used: Virus3

VIRUSCOMMONNAME	GENUS	STATUS	TRANSMISSIONTYPE
SARS-CoV	Betacoronavirus	Active	Respiratory Droplets
Variola Virus	Orthopoxvirus	Eradicated	Airborne Particles
Human Immunodeficiency Virus	Lentivirus	Active	Bodily Fluids
Monkeypox Virus	Orthopoxvirus	Active	Direct Contact
Influenza A Virus	Alphainfluenzavirus	Active	Respiratory Droplets
Poliovirus	Enterovirus	Active	Fecal-oral
Vaccinia Virus	Orthopoxvirus	Active	Direct Contact
Cowpea Mosaic Virus	Comovirus	Active	Insects
Herpes Simplex Virus	Simplexvirus	Active	Bodily Fluids
Phi X 174	Sinsheimervirus	Active	Bacterial
Zaire Ebolavirus	Ebolavirus	Active	Bodily Fluids
Rinderpest morbillivirus	Morbillivirus	Eradicated	Direct Contact
Hepatitis B Virus	Orthohepadnavirus	Active	Bodily Fluids
Human Papillomavirus	Chipapillomavirus	Active	Direct Contact

Input and result:

Virus Data

Choose a query:	Count Viruses by Transmission Type	~
Submit Home		

Result:

TRANSMISSION	VIRUSES_COUNT
Bodily Fluids	4
Direct Contact	4
Respiratory Droplets	2
Insects	1
Bacterial	1
Fecal-oral	1
Airborne Particles	1

Aggregation with HAVING:

Tables used: Outbreak

COUNTRYNAME	VIRUSCOMMONNAME	OUTBREAKSIZE	CASUALTY	YEAR	ORIGIN
United States	Influenza A Virus	Pandemic	100000000	1346	Spain
China	SARS-CoV	Pandemic	25000000	2019	China
Mexico	Variola Virus	Epidemic	8000000	1519	Mexico
Japan	Variola Virus	Epidemic	2000000	735	Japan
Congo	Zaire Ebolavirus	Epidemic	55	2020	Congo
China	Influenza A Virus	Pandemic	4000000	1968	Hong Kong
Canada	Human Immunodeficiency Virus	Pandemic	40100000	1981	Central Africa

Input and result:

Virus Data

Choose a query: Average Casualty By Virus (Virus must have caused atleast two outbreaks) >

Submit

Home

Result:

VIRUS	AVERAGE_DEATHS
Influenza A Virus	52000000
Variola Virus	5000000

Nested aggregation with GROUP BY:

Tables used: Causes, Outbreak, ViralDisease

DISEASENAME	DISEASETYPE
Smallpox AIDS Hepatitis Herpes Labialis Influenza	Acute Chronic Acute Dormant Acute

COUNTRYNAME	VIRUSCOMMONNAME	OUTBREAKSIZE	CASUALTY	YEAR	ORIGIN
United States China Mexico Japan Congo China Canada	Influenza A Virus SARS-CoV Variola Virus Variola Virus Zaire Ebolavirus Influenza A Virus Human Immunodeficiency Virus	Pandemic Pandemic Epidemic Epidemic Epidemic Epidemic Pandemic Pandemic	100000000 25000000 8000000 2000000 55 4000000 40100000	2019 1519 735 2020 1968	Spain China Mexico Japan Congo Hong Kong Central Africa

DISEASENAME	VIRUSCOMMONNAME
AIDS Hepatitis	Human Immunodeficiency Virus Hepatitis B Virus
Herpes Labialis	Herpes Simplex Virus
Influenza	Influenza A Virus
Smallpox	Variola Virus

Input and result:

Virus Data

Choose a query: Total death count of each acute viral disease due to outbreaks

Submit Home

Result:

DISEASE	DEATH_COUNT		
Influenza	104000000		
Smallpox	10000000		