

CSCI 3901 Algorithm, Database ERD, Code Design, Data Structure

Objective : The course project is the opportunity to demonstrate all of the concepts from the course in one body of work.

Report by:

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Algorithm:

PersonIdentity addPerson(String name)

1. Insert new entry in person with the new given name
2. Return true

Boolean recordAttributes(PersonIdentity person, Map<String, String> attributes)

1. Fetch PersonIdentity object
2. Select * from person where PersonIdentityperson.id = p_id
3. If null, return false
4. If the person exists, fetch the p_id.
5. for (Map.Entry<String,String> entry : Map.entrySet())
 - a. Validation the key value with the columns in person table
 - b. If key==column in person table, store the value in DB

Boolean recordReference(PersonIdentity person, String reference)

1. Select * from person where PersonIdentityperson.id = p_id
2. If null, return false
3. Insert into person_reference values (p_id, reference);
4. Return True

Boolean recordNote(PersonIdentity person, String note)

1. Select * from person where PersonIdentityperson.id = p_id
2. If null, return false
3. Insert into person_notes values (p_id, notes)
4. Return True;

Boolean recordChild(PersonIdentity parent, PersonIdentity child)

1. Check whether parent and child exist in person
2. If null, return false

3. Insert into parentchild_relation (parent_id, child_id)
4. Return false

Boolean recordPartnering(PersonIdentity partner1, PersonIdentity partner2)

1. Check whether partner1 and partner2 exist in person
2. If null, return false
3. Insert into partner_relation (partner 1, partner 2)
4. Return true

Boolean recordDissolution(PersonIdentity partner1, PersonIdentity partner2)

1. Check whether relation exist in partner_relation
2. If null, return false
3. Delete relation from partner_relation

FileIdentifier addMediaFile(String fileLocation)

1. Insert new entry in media archive with the new given name
2. Return true

Boolean recordMediaAttributes(FileIdentifier fileIdentifier, Map<String, String> attributes)

1. Fetch FileIdentifier object
2. Select * from media_archieve where FileIdentifier.id = p_id
3. If null, return false
4. If the file exists, fetch the mediaId.
5. for (Map.Entry<String,String> entry: Map.entrySet())
 - a. Validation the key value with the columns in person table
 - b. If key==column in file table, store the value in DB

Boolean peopleInMedia(FileIdentifier fileIdentifier, List<PersonIdentity> people)

1. Fetch the media_id from FileIdentifier object.
2. Find p_id from a person.
3. If media_id or person not found, then return false

4. Insert into people_in_media (mediaId, person_Id)
5. Return True

Boolean tagMedia(FileIdentifier, fileIdentifier, String tag)

1. Fetch the media_id from FileIdentifier object.
2. If null, return false
3. Insert into media_tags values (mediaId, tag);
4. Return True

PersonIdentity findPerson(String name)

1. Select query from person table with name in the argument
2. If null, return null;
3. Else, create PersonIdentity object
4. Assign column values from person table to each attributes of PersonIdentity object and return it.

FileIdentifier findMediaFile(String name)

1. Select query from media_archive table with name in the argument
2. If null, return null;
3. Else, create fileIdentifier object
4. Assign column values from the media_archive table to each attributes of file identifier object and return it.

BiologicalRelation findRelation(PersonIdentity person1, PersonIdentity person2)

1. Check if any of the person 1 or person2 has no ancestors
 - a. If yes, then check whether person2's ancestor is person 1 or person1's ancestor is person2;
 - b. If yes, find the generation gap in the same family line
2. Find common ancestors for person1 and person2

Set<PersonIdentity> descendants(PersonIdentity person, Integer generations)

1. Validate person in person table
2. If null, return false or null

3. If person found, create view of parentchild_relation where parentId=person_id
 - a. Select from parentchild_relation view where parentId = personId
 - b. Delete particular entry
4. Run step in for loop, for number of interactions mentioned in 'generation'
5. OR until select query data becomes null;

Set<PersonIdentity> ancestor(PersonIdentity person, Integer generations)

1. Validate person in person table
2. If null, return false or null
3. If person found, create view of parentchild_relation where childId=person_id
 - a. Select from parentchild_relation view where childId = personId
 - b. Delete particular entry
4. Run step in for loop, for number of interactions mentioned in 'generation'
5. OR until select query data becomes null;

List<String> notesAndReferences(PersonIdentity person)

1. Select p_id from person where name = person.name
2. Store it in a variable ID.
3. Select notes from person_notes where p_id = ID
4. Select references from person_reference where p_id = ID

Data Structure

1. Map

- a. Record attributes person
- b. Record attribute media file

2. List

- a. List will be used in findrelation
- b. It will compare the ancestors of partner1 and partner2 find relations
- c. List will also be used fetch multi value attributes
 - i. Example: Tags, notes, references and Individuals seen in picture.

Code Design

CRC method:

PersonIdentity Class	
Responsibilities: PersonIdentity addPerson(String name) Boolean recordAttributes(PersonIdentity person, Map<String, String> attributes) Boolean recordReference(PersonIdentity person, String reference) Boolean recordNote(PersonIdentity person, String note)	Collaboration: Genealogy BiologicalRelation

FileIdentifier	
Responsibilities: FileIdentifier addMediaFile(String fileLocation) Boolean recordMediaAttributes(FileIdentifier fileIdentifier, Map<String, String> attributes) Boolean peopleInMedia(FileIdentifier fileIdentifier, List<PersonIdentity> people) Boolean tagMedia(FileIdentifier, fileIdentifier, String tag)	Collaboration: Genealogy

Genealogy

Responsibilities:

PersonIdentity findPerson(String name)

FileIdentifier findMediaFile(String name)

String findName(PersonIdentity id)

String findMediaFile(FileIdentifier fileId)

Collaboration:

PersonIdentity

FileIdentifier

BiologicalRelation

Responsibilities:

Boolean recordChild

Boolean recordPartnering(PersonIdentity partner1, PersonIdentity partner2)

Boolean recordDissolution(PersonIdentity partner1, PersonIdentity partner2)

Collaboration:

PersonIdentity

DataBase Schema

ERD:



