

# Team 0 – D&A

## Project – Phase III

Aditya Nair (2023111029)

Shreyas Deb (2023113024)

Vishak Kashyap K (2023113012)

(total 8 pages)

Here is the draw.io link to our model diagrams - [team 0 phase 3](#)

### 1. ERD to Relational Model

The changes made during this transition are as follows:

- All the entity types are converted to relations. The arrow marks between the relations indicate the foreign key referencing.
- We have used the option 8A (given in the textbook, pg no. 299) to represent the subclass-superclass relations.
- All the composite and multivalued attributes have been kept as it is (We have assumed the traditional relational model, as the modern relational model is equivalent to 1NF).
- All the 1:N relationship types have been represented using the foreign key approach (textbook, pg 294, step 4).
- For the M:N binary relationships, the ternary and quaternary relationships, we have used the approach given in the textbook (step 5, pg 294 & pg 296).
- New attributes have been created or have been made a foreign key to represent the two recursive relationships in our database design. One is in the pathogen relation, where the **predecessor** key is made a foreign key to the Pathogen relation itself. Another instance is in the Global Vaccine Distribution where **importer\_id** and **exporter\_id** attributes have been added which reference the country relation.

Pathogen

id	scientific_name	type	lethality_rate	transmission_method	predecessor	incubation_period	mutation_probability	resistance
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Research Focus

lab_id	project_id	pathogen_id
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Mutation (Subclass of Pathogen)

id	parent_pathogen_id	date_discovered
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Infection

pathogen_id	country_id	first_case_date	total_infected	total_death
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Country

country_id	name	population	health_quality_index	climate	borders_open
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Country Response to Pathogen

country_id	project_id	pathogen_id
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Research Lab

lab_id	name	location	total_funding
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Participates In

lab_id	project_id	fund_allocated
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Research Project

project_id	labs	pathogen_id	duration	research_aim	milestones	current_status
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Vaccine Development

vaccine_id	project_id	pathogen_id
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Government Response

response_id	pathogen_id	country_id	date_implemented	response_type	response_severity	effectiveness_rate
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Vaccine

vaccine_id	pathogen_id	project_id	name	date_discovered	effectiveness	number_of_administrations	number_of_doses	side_effect
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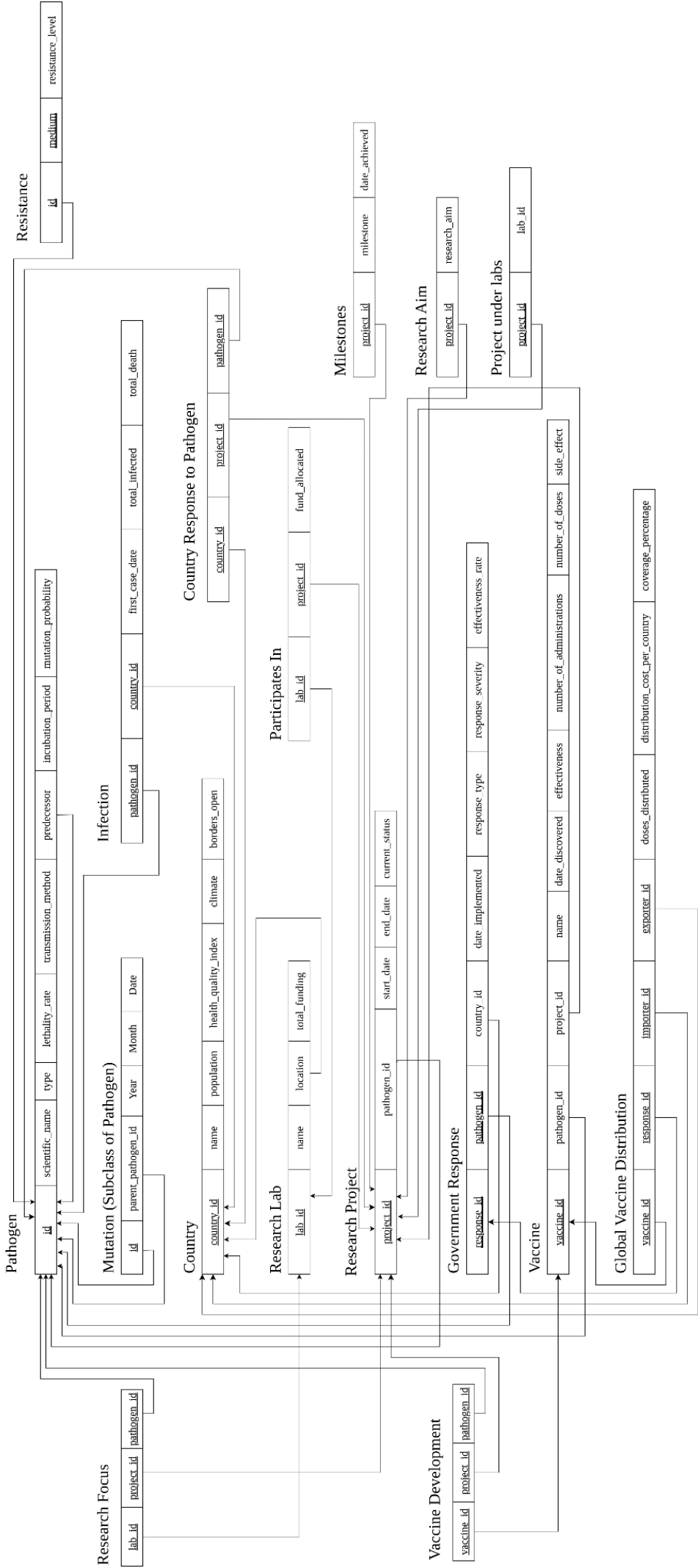
Global Vaccine Distribution

vaccine_id	response_id	importer_id	exporter_id	doses_distributed	distribution_cost_per_country	coverage_percentage
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## 2. Relational Model to 1NF

The following changes have been made during this transition:

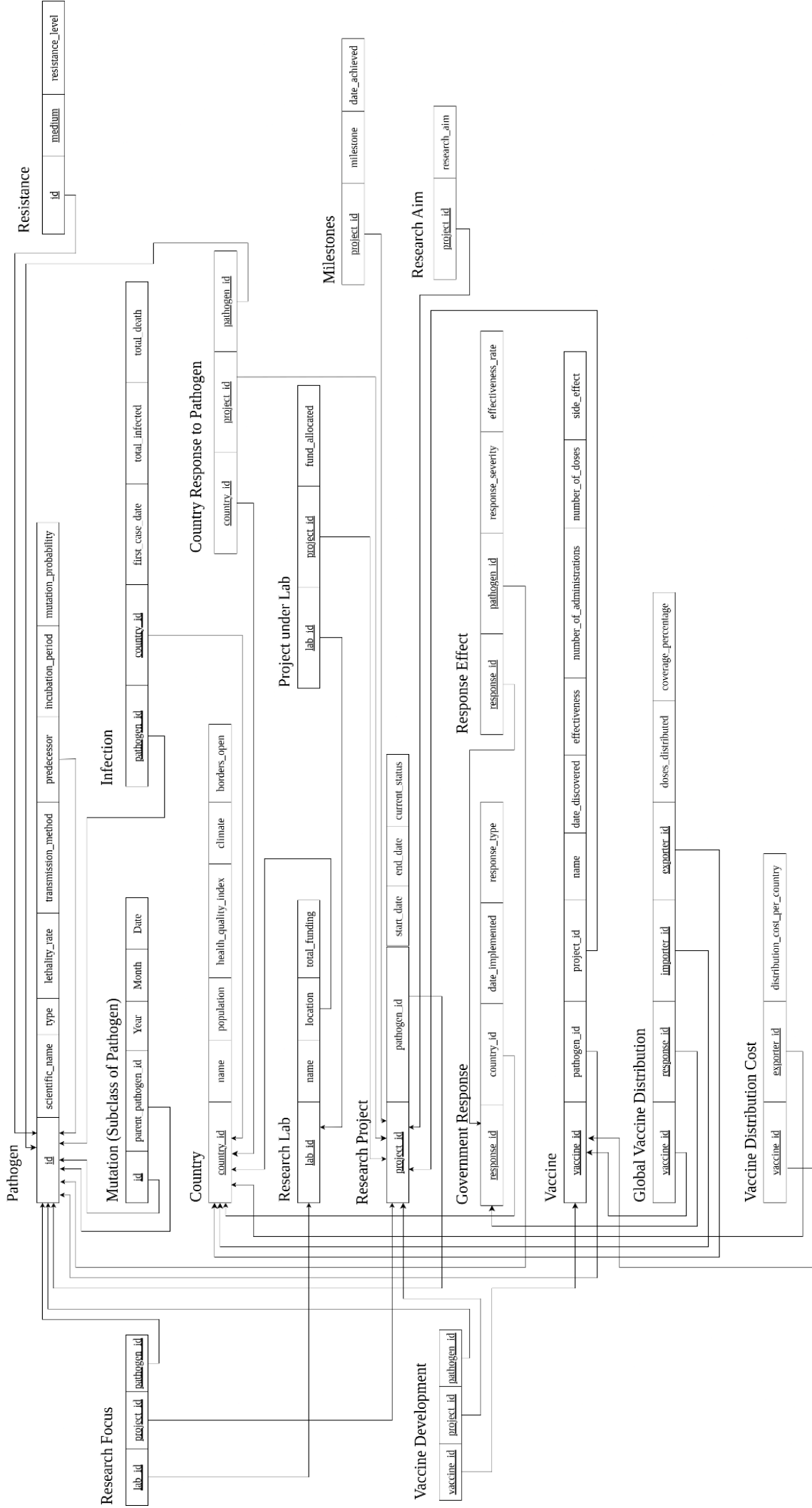
- All the composite attributes have been represented in terms of their simple components.
- All the multivalued attributes have been represented in a new relation.
- There were two composite multivalued attributes, one is **resistance** in the Pathogen relation and the other is **milestones** in the Research Project relation.



### 3. 1NF to 2NF

The following changes have been made during this transition:

- Government Response relation has `date_implemented` and `response_type` dependent only on the `response_id` and the others dependent on both `response_id` and `pathogen_id` both, and are broken off into a new relation Response Effect.
- Global Vaccine Distribution has `distribution_cost_per_country` depending only on the `vaccine_id` and `exporter_id` as it depends only on the exporting country for a vaccine, and refers to the customs and such costs excluding cost of actual vaccine units. Hence, we broke it off into a new relation Vaccine Distribution Cost.
- While creating a new relation for the multivalued attribute `labs` of Research Project, it turns out equivalent to an existing relationship type, based on which the relation Participates In was created. Hence we kept the relations Participates In and renamed it to `Project Under Lab` as well, to better reflect its purpose, and further remove redundancy.
- Also, to be noted that, the relations `Research Focus`, `Vaccine Development`, and `Country Response to Pathogen` can, and should, be implemented as Views in RDBMS implementation. However, we are retaining them in the relational model as they represent important relationship types.



#### 4. 2NF to 3NF

The following (and only) change has been made during this transition:

- It is observed that `pathogen_id` in Vaccine relation can be inferred from the non-prime attribute `project_id` as each Project, as well as Vaccine, focuses on only one Pathogen in our universe/mini-world. Since `pathogen_id` is already present in the Project that a given Vaccine emerges out of, it was simply removed from Vaccine.

