

# Etapa 4





# Banco de dados

## 1) Disease-Symptom Knowledge Database

### Disease-Symptom Knowledge Database

This table below is a knowledge database of disease-symptom associations generated by an automated method based on information in textual discharge summaries of patient Presbyterian Hospital admitted during 2004. The first column shows the disease, the second the number of discharge summaries containing a positive and current mention of and the associated symptom. Associations for the 150 most frequent diseases based on these notes were computed and the symptoms are shown ranked based on the strength. The method used the MedLEE natural language processing system to obtain UMLS codes for diseases and symptoms from the notes; then statistical methods based on frequency occurrences were used to obtain the associations. A more detailed description of the automated method can be found in Wang X, Chused A, Elhadad N, Friedman C, Markate. Automated knowledge acquisition from clinical reports. AMIA Annu Symp Proc. 2008; p. 783-7. PMID: PMC2656103.

Please contact friedman@dbmi.columbia.edu for any questions regarding the knowledge database.

Disease	Count of Disease Occurrence	Symptom
UMLS:C0020538_hypertensive disease	3363	UMLS:C0008031_pain chest
		UMLS:C0392660_shortness of breath
		UMLS:C0012833_dizziness
		UMLS:C0004093_asthenia
		UMLS:C0065636_fat
		UMLS:C0039070_syncope
		UMLS:C0042571_vertigo
		UMLS:C0036992_sweatUMLS:C0700590_sweating increased
		UMLS:C0030252_palpitation
		UMLS:C0027497_nausea
		UMLS:C0002962_angina pectoris
		UMLS:C0439716_pressure chest
UMLS:C0011847_diabetes	1421	UMLS:C0032617_polyuria
		UMLS:C0065602_polydipsia
		UMLS:C0392660_shortness of breath
		UMLS:C0009031_rain chest

## 2) Diseases Database

[Home](#) | [Index](#) | [Disclaimer](#) | [Contact](#) | [Previous page](#) | [Diseases Database Ver 2.0 : Medical lists and links](#) | [Search](#)

**Subject Index:**

ABCDEFGHIJKLM  
NOPQRSTUVWXYZ

- [Search](#) for information on a disease, symptom, physical sign, non-brand drug or common laboratory abnormality.
- Please understand our [disclaimer](#) and [what is in the Diseases Database](#)
- [Reviews](#)
- [Authorship and funding](#)
- [SNOMED, CTR, JAK, UMLS extensions](#) | [browser](#)

Start

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The Diseases Database is not a diagnostic or clinical decision-making tool.  
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Diseases updated: 2016-10-12  
This [YAML](#) [HTML](#) [1.0](#) | [revised 2016-10-10 06:04:21](#) | [EMOSQL Technology](#)

## 3) National Health Interview Survey

Summary Health Statistics: National Health Interview Survey, 2016

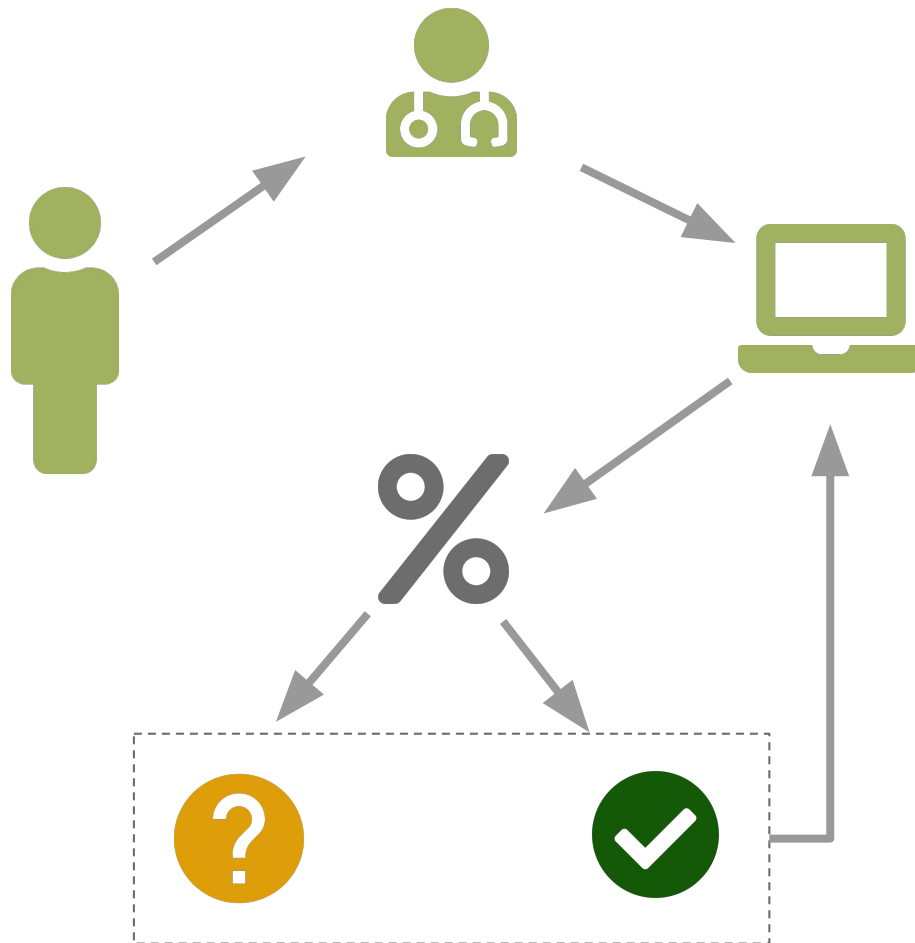
Table A-4a, page 1 of 9

Table A-4a. Age-adjusted percentages (with standard errors) of selected diseases and conditions among adults aged 18 and over, by selected characteristics: United States, 2016

Selected characteristic	Diabetes <sup>1</sup>	Ulcers <sup>1</sup>	Kidney disease <sup>1</sup>	Liver disease <sup>1</sup>	Arthritis diagnosis <sup>1</sup>	Chronic joint symptoms <sup>2</sup>
Total	8.8 (0.26)	5.7 (0.17)	1.9 (0.09)	1.9 (0.10)	21.6 (0.27)	28.3 (0.36)
Sex						
Male	9.4 (0.28)	5.7 (0.24)	1.8 (0.13)	1.9 (0.15)	18.5 (0.38)	26.7 (0.52)
Female	8.4 (0.26)	5.8 (0.22)	2.0 (0.12)	1.9 (0.14)	24.4 (0.38)	29.6 (0.45)
Age (years)						
18-44	2.8 (0.20)	3.4 (0.20)	0.7 (0.09)	1.2 (0.14)	7.3 (0.32)	16.4 (0.48)
45-64	12.5 (0.45)	7.5 (0.34)	2.1 (0.17)	2.8 (0.22)	31.1 (0.58)	38.8 (0.61)
65-74	23.0 (0.81)	8.8 (0.55)	3.9 (0.36)	2.6 (0.29)	47.6 (0.92)	46.1 (0.94)
75 and over	19.4 (0.85)	10.7 (0.68)	6.5 (0.50)	2.0 (0.33)	51.9 (1.04)	47.5 (1.07)
Race						
One race <sup>3</sup>	8.8 (0.21)	5.7 (0.17)	1.8 (0.09)	1.9 (0.10)	21.5 (0.27)	28.1 (0.36)
White	8.2 (0.22)	6.0 (0.19)	1.8 (0.10)	2.0 (0.12)	22.0 (0.30)	29.0 (0.39)
Black or African American	13.1 (0.67)	4.4 (0.38)	2.5 (0.31)	1.6 (0.26)	22.7 (0.80)	27.4 (0.95)
American Indian or Alaska Native	15.7 (2.53)	8.7 (1.84)	2.4 (0.99)	2.5 (1.12)	25.7 (2.91)	35.1 (3.98)
Asian	8.0 (0.36)	3.6 (0.67)	0.7 (0.22)	2.0 (0.42)	12.5 (1.10)	16.9 (1.28)
Native Hawaiian or Other Pacific Islander	14.4 (3.89)	-	0.7 (0.47)	-	18.3 (3.80)	25.9 (4.97)
Two or more races <sup>3</sup>	8.5 (1.38)	8.3 (1.32)	3.2 (0.78)	2.1 (0.61)	26.8 (2.09)	36.4 (2.50)
Black or African American, white	-	-	-	-	18.6 (5.62)	33.7 (5.81)
American Indian or Alaska Native, white	13.8 (3.38)	11.9 (2.32)	-	2.7 (0.88)	31.9 (3.81)	38.6 (3.96)

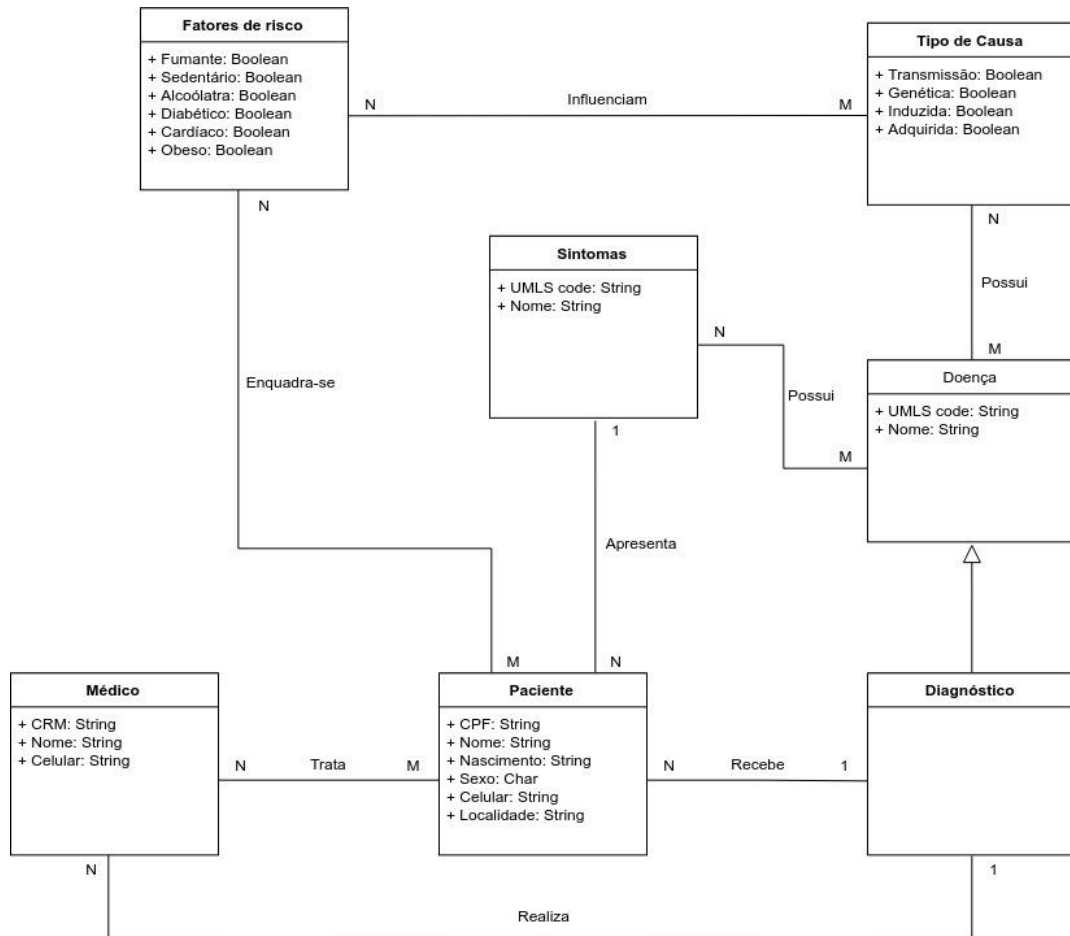


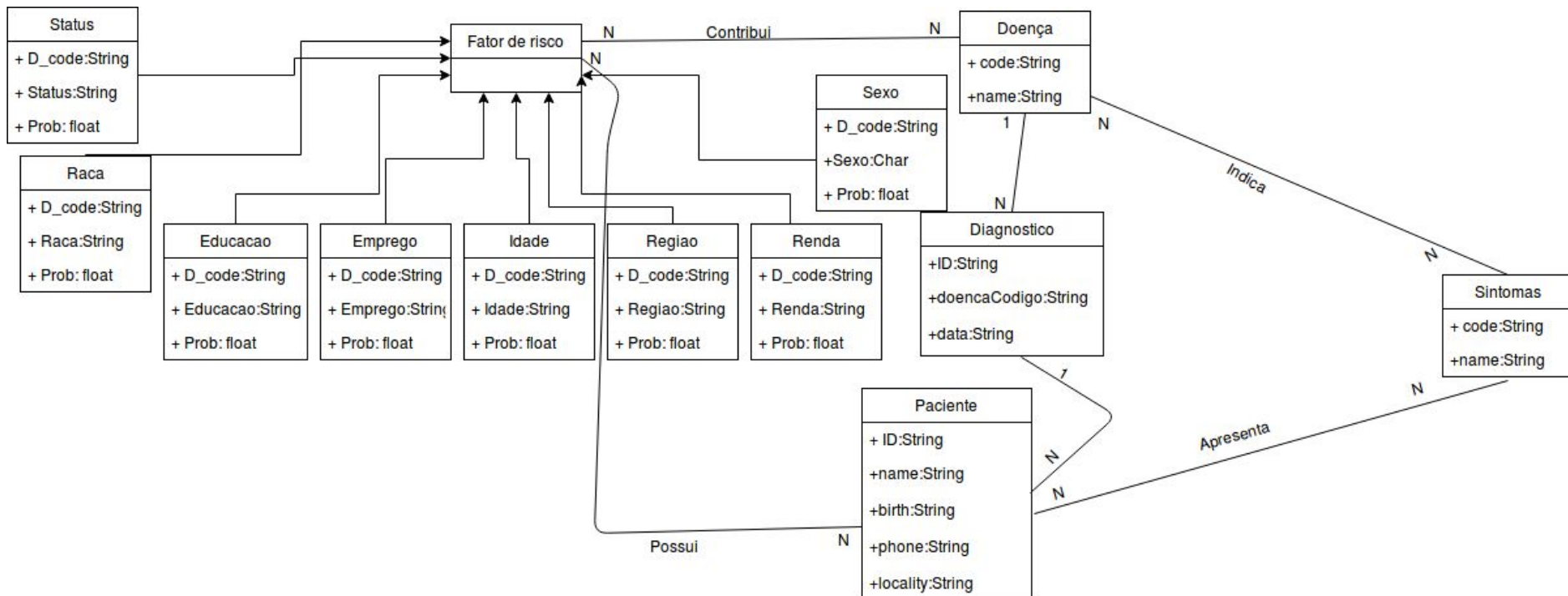
# Requisitos





# Modelo inicial







# Queries

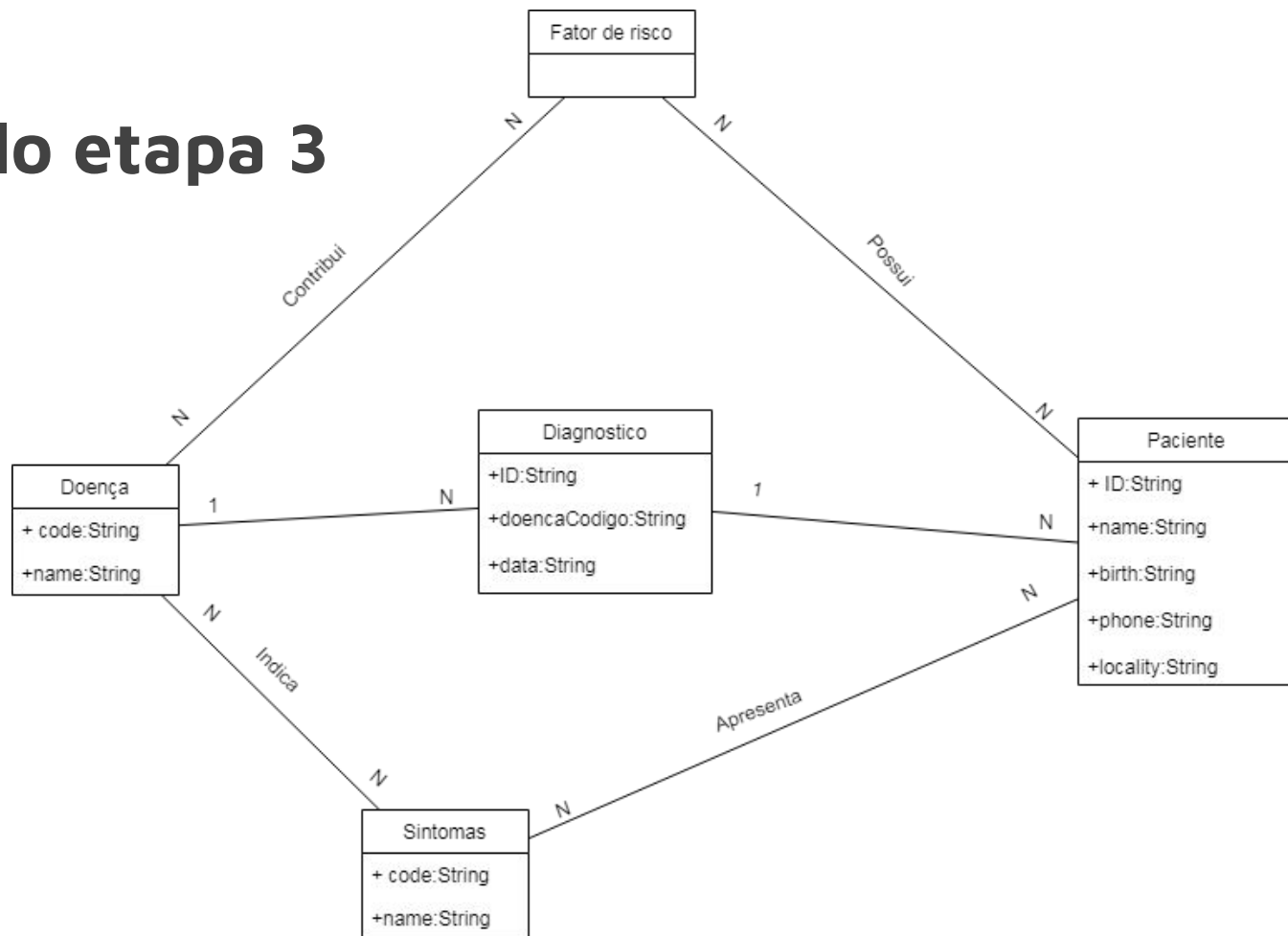
## Sintoma mais relatado por pacientes

```
In [ ]: SELECT nome
FROM (SELECT S.nome, COUNT(*) contagem
      FROM Sintomas S, DoencaSintomas DS, Doenca D
      WHERE DS.sintomaCodigo = S.codigo AND DS.doencaCodigo = D.codigo GROUP BY S.nome)
WHERE contagem = (SELECT MAX(contagem) FROM (SELECT S.nome, COUNT(*) contagem
      FROM Sintomas S, DoencaSintomas DS, Doenca D
      WHERE DS.sintomaCodigo = S.codigo AND DS.doencaCodigo = D.codigo GROUP BY S.nome))
```

## Nome dos constatados com a doença X (no caso abaixo: diabetes)

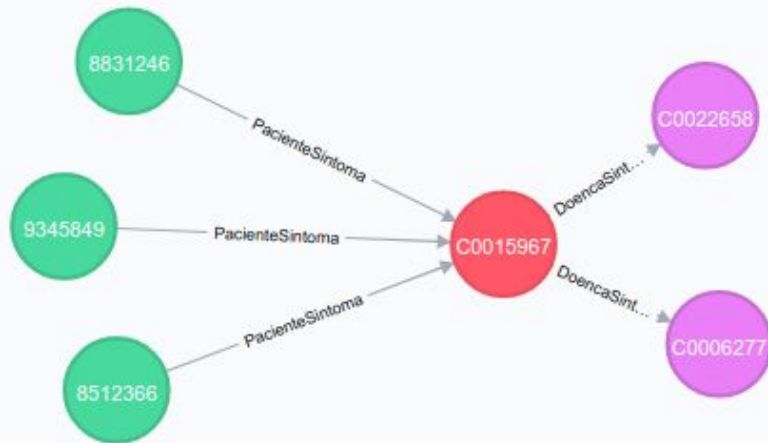
```
In [ ]: SELECT P.nome
FROM PacienteDiagnostico PD, Diagnostico D, Doenca Do, Paciente P
WHERE PD.pacienteID = P.ID AND PD.diagnosticoID = D.ID AND D.doencaCodigo = Do.codigo AND Do.nome = 'diabetes'
```

## Modelo etapa 3



# Queries

```
// retorna o caminho para um paciente que apresenta como sintoma 'febre'  
MATCH p = (l:Paciente)-[:PacienteSintoma]->(m:Sintomas {nome: 'fever'})-[t:DoencaSintoma]->(d:Doenca)  
RETURN p
```







# Queries

```
// Dado um sintoma X, retorna a probabilidade de um paciente ter a doença Y
MATCH p = (l:Paciente)-[:PacienteSintoma]->(m:Sintomas {nome: 'pain chest'})-[t:DoencaSintoma]->
(d:Doenca{nome: 'hypertensive disease'})
WITH COUNT(l) AS prob
MATCH q = (l:Paciente)-[:PacienteSintoma]->(m:Sintomas {nome: 'pain chest'})-[t:DoencaSintoma]->(d:Doenca)
RETURN toFloat(prob)/COUNT(l)
```

---

**toFloat(prob)/COUNT(l)**

---

0.2

---



# Queries

```
// Dado um Paciente X, retorna a probabilidade do paciente ter a doença Y levando em conta os
// seus fatores de risco e os seus sintomas
MATCH p = (l:Paciente {ID: 'patKMPV11'})-[:PacienteSintoma]->(m:Sintomas)-[t:DoencaSintoma]->
(d:Doenca{nome: 'hypertensive disease'})
WITH COUNT(l) AS prob
MATCH q = (l:Paciente {ID: 'patKMPV11'})-[:PacienteSintoma]->(m:Sintomas)-[t:DoencaSintoma]->
(d:Doenca) WITH COUNT(l)/prob AS tot
MATCH o = (n:Paciente{ID: 'patKMPV11'})-[FatorPaciente]->(Fator)-[t:FatorDoenca]->
(d:Doenca{nome: 'hypertensive disease'})
RETURN AVG(toFloat(replace(t.Probabilidade, ",", "."))) * toFloat(1) / tot
```

**AVG(toFloat(replace(t.Probabilidade, ",", "."))) \* toFloat(1) / tot**

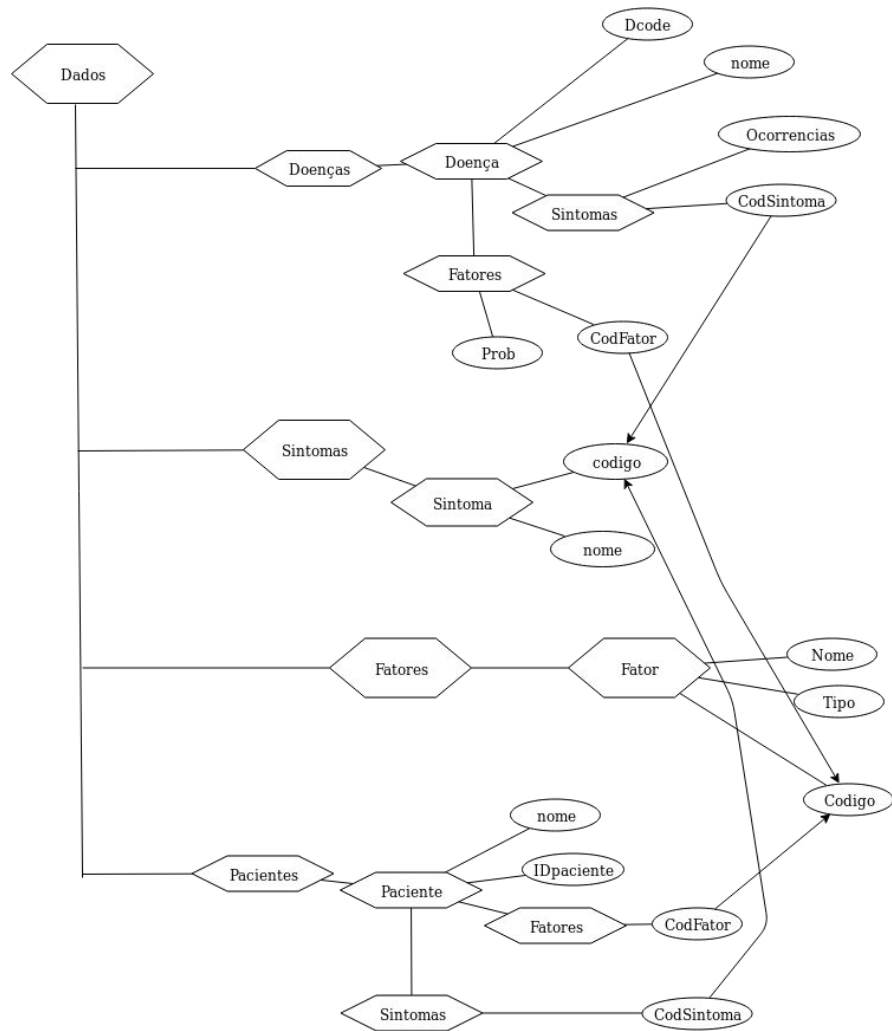
---

5.4175

---



# Modelo etapa 4





# Queries

```
# Get definitions for Neoplasms
```

```
PREFIX rdf: <http://www.w3.org/1999/02/22-rdf-syntax-ns#>
```

```
PREFIX rdfs: <http://www.w3.org/2000/01/rdf-schema#>
```

```
PREFIX xsd: <http://www.w3.org/2001/XMLSchema#>
```

```
PREFIX owl: <http://www.w3.org/2002/07/owl#>
```

```
PREFIX meshv: <http://id.nlm.nih.gov/mesh/vocab#>
```

```
PREFIX mesh: <http://id.nlm.nih.gov/mesh/>
```

```
PREFIX mesh2015: <http://id.nlm.nih.gov/mesh/2015/>
```

```
PREFIX mesh2016: <http://id.nlm.nih.gov/mesh/2016/>
```

```
PREFIX mesh2017: <http://id.nlm.nih.gov/mesh/2017/>
```

```
SELECT *
```

```
FROM <http://id.nlm.nih.gov/mesh>
```

```
WHERE {
```

```
    mesh:D009369 rdfs:label ?DescriptorLabel .
```

```
    mesh:D009369 meshv:preferredConcept ?concept .
```

```
    ?concept meshv:scopeNote ?ScopeNote
```

```
}
```



# Queries

*# Descendants of Heart Diseases (D006331)*

```
PREFIX mesh: <http://id.nlm.nih.gov/mesh/>
PREFIX mesh2015: <http://id.nlm.nih.gov/mesh/2015/>
PREFIX mesh2016: <http://id.nlm.nih.gov/mesh/2016/>
PREFIX mesh2017: <http://id.nlm.nih.gov/mesh/2017/>
```

```
SELECT DISTINCT ?descriptor ?label
FROM <http://id.nlm.nih.gov/mesh>
```

```
WHERE {
  mesh:D006331 meshv:treeNumber ?treeNum .
  ?childTreeNum meshv:parentTreeNumber+ ?treeNum .
  ?descriptor meshv:treeNumber ?childTreeNum .
  ?descriptor rdfs:label ?label .
}
```

```
ORDER BY ?label
```



# Queries

-----

Sintomas em comum entre duas doenças específicas

```
let $documento := doc('mydoc.xml')

for $i in ($documento//Doenca[nome = "diabetes"]/Sintomas)
for $j in ($documento//Doenca[nome = "hypertensive disease"]/Sintomas)
where $i/CodSintoma = $j/CodSintoma
for $k in ($documento//Sintoma)
where $k/codigo = $j/CodSintoma
return data($k/nome)
```

-----

Classificacao dos sintomas mais frequentes (com maior numero de Ocorrencias)  
segundo nosso banco de dados

```
let $documento := doc('mydoc.xml')

return
<classificacao>
{
  for $i in ($documento//Doenca/Sintomas)
  where $i[Ocorrencias > 2000] order by $i/@Ocorrencias
  return <frequente>{data($i/CodSintoma)}</frequente>
}
</classificacao>
```



# Queries

-----

Avalia qual doenca tem mais propensao a ocorrer em mulheres ou homens

```
let $documento := doc('mydoc.xml')
```

```
for $i in ($documento//Doenca)
```

```
return if ($i/Fatores[CodFator="24"]/Prob < $i/Fatores[CodFator="25"]/Prob)
```

```
then <Masculino>{distinct-values($i/nome)}</Masculino>
```

```
else <Feminino>{distinct-values($i/nome)}</Feminino>
```