

# ER Model - An Example

## Chapter 2

in the Course Textbook

For material relating to ER Model

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# Problem

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## A Bioinformatics Application

- *Patient*: has a unique *MSP number*, a *Patient name*, a *Date of Birth*, a *Tissue Type* and an *indicator* denoting whether the tissue is cancerous or normal.
- A *patient library* associates a patient with multiple *tags*
- Each tag has a unique *tag number* and a unique *nucleotide sequence*.
- For each tag in the patient library, a *count* is given to record the number of times the tag occurs in the library. In general, the same tag can be associated with any number of patients.
- A tag may be mapped to a *gene*. Each gene has a unique *gene name* and a *type*.
- In general, multiple tags may be mapped to the same gene. However, two different genes cannot be mapped to the same tag.
- Finally, an *article* is identified by a unique *article number* and a *journal name*. An article may analyze multiple genes and a gene may be analyzed by multiple articles.

Let us construct an ER model for the above application.

# WARM UP

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## Entity Sets:

- Patients - Attributes are MSP Number, Name, DOB, Tissue Type and Indicator
- Tags - Attributes are Tag Number and Nucleotide Sequence
- Genes - Attributes are Gene Name and Type
- Articles - Attributes are Article Number and Journal Name

## Relationship Sets:

- Patient Library - Many to Many, Has an attribute Count
- Map - Many to 1 from Tags to Genes
- Analyzes - Many to Many

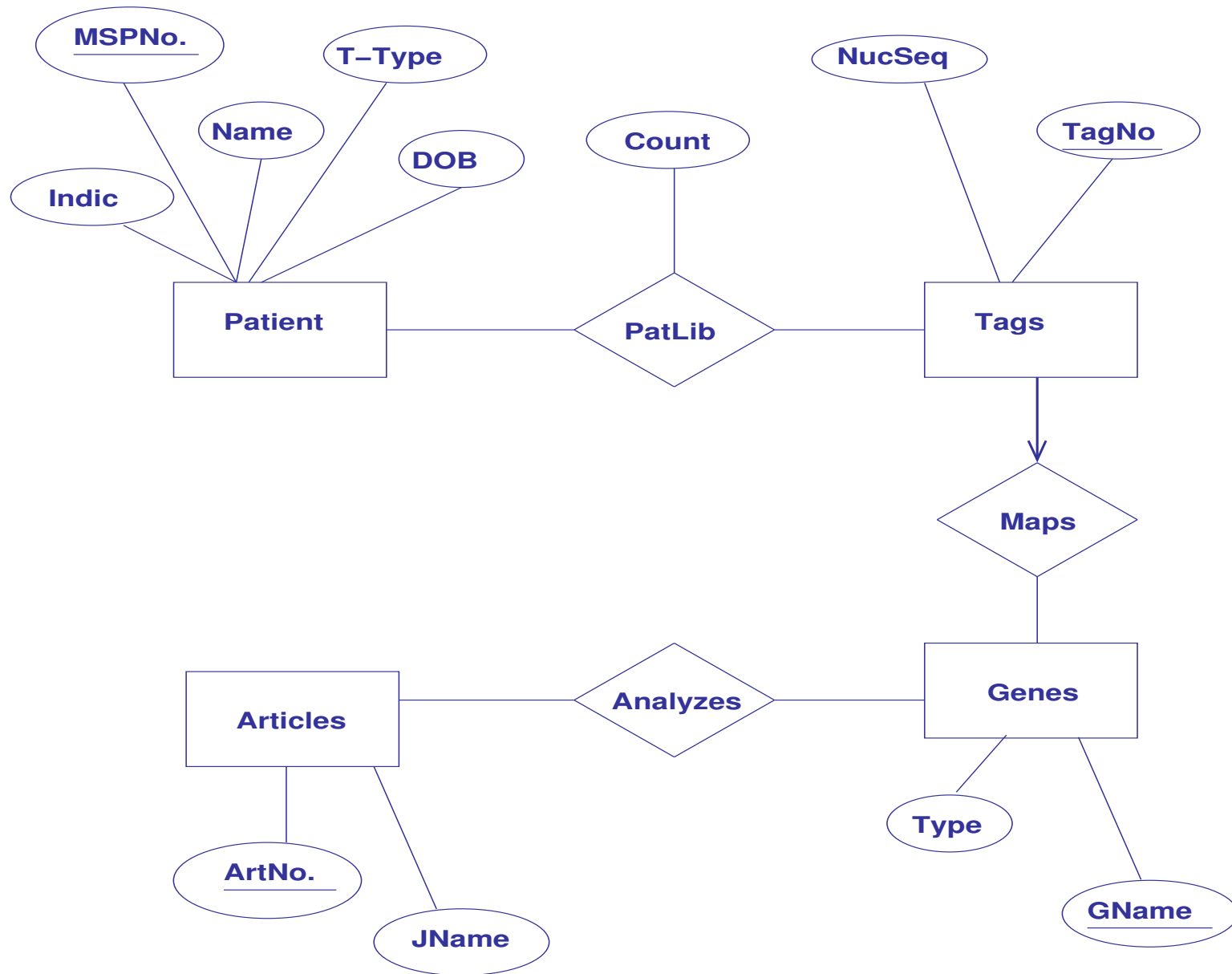


Figure 1: ER Diagram

## Relations for the Entities

**Patient Table:** Patients(MSPNo, Name, TType, DOB, Indic)

```
CREATE TABLE Patient(  
    MSPNo    CHAR(10),  
    Name     CHAR(30),  
    TType    CHAR(30),  
    DOB      DATE,  
    Indic    ENUM('Cancerous', 'Normal'),  
    PRIMARY KEY (MSPNo));
```

**Tags Table:** Tags(TagNo, NucSeq)

```
CREATE TABLE Tags(  
    TagNo    CHAR(10),  
    NucSeq    CHAR(30),  
    PRIMARY KEY (TagNo),  
    UNIQUE(NucSeq));
```

## Relations for the Entities

**Genes Table:** Genes(GName, Type)

```
CREATE TABLE Genes(  
    GName    CHAR(30),  
    Type     CHAR(30),  
    PRIMARY KEY (GName));
```

**Articles Table:** Articles(ArtNo, JName)

```
CREATE TABLE Articles(  
    ArtNo    CHAR(10),  
    JName    CHAR(30),  
    PRIMARY KEY (ArtNo));
```

## Relations for the Relationships

**Patients Library Table:** PatLib(MSPNo, TagNo, Count)

```
CREATE TABLE PatLib(  
    MSPNo    CHAR(10),  
    TagNo    CHAR(10),  
    Count    INTEGER,  
    PRIMARY KEY (MSPNo, TagNo)  
    FOREIGN KEY (MSPNo) REFERENCES Patient,  
    FOREIGN KEY (TagNo) REFERENCES Tags);
```

**Maps Table:** Maps(TagNo, GName)

```
CREATE TABLE Maps(  
    TagNo    CHAR(10),  
    GName    CHAR(30),  
    PRIMARY KEY (TagNo)  
    FOREIGN KEY (TagNo) REFERENCES Tags,  
    FOREIGN KEY (GName) REFERENCES Genes);
```

## Relations for the Relationships

### **Analyzes Table:** Analyzes(ArtNo, GName)

```
CREATE TABLE Analyzes(  
    ArtNo      CHAR(10),  
    GName      CHAR(30),  
    PRIMARY KEY (ArtNo, GName)  
    FOREIGN KEY (ArtNo) REFERENCES Articles,  
    FOREIGN KEY (GName) REFERENCES Genes);
```

Tags and Maps can also be combined into one table as they share the same primary key, as follows.

```
CREATE TABLE TagsMap(  
    TagNo      CHAR(10),  
    NucSeq     CHAR(30),  
    GName      CHAR(30),  
    PRIMARY KEY (TagNo)  
    FOREIGN KEY (GName) REFERENCES Genes,  
    UNIQUE (NucSeq));
```