

# Temporal Query Language

version 2.2

TI = time interval (for example [0 – 2, 1 – 3, 1 – 5, 5 – 10])  
CTI = computed time interval (for example [0 – 10])  
TP = time point (internally a time interval with same time of start and end)  
CPT = examples: 12345 or 1234F  
ICD9 = examples: 022, 022.2, 022.22, E123, E123.1, E123.12, V01, V01.1, V01.01  
NOTE\_TP = TP with payload information [TP, NOTE\_ID]  
ICD9\_TI = TI with payload information [TI, ICD9\_ID]  
MAX = Integer.MAXIMUM (maximum positive number)  
MIN = Integer.MINIMUM (minimum negative number)

## Automatic Result type conversions

TP => TI[TP, TP]  
TP => BOOLEAN[is TP empty]  
TI => BOOLEAN[is TI empty]  
TI => TP (undefined)  
TI => CTI  
BOOLEAN => TIMELINE  
BOOLEAN => TP (undefined)  
NOTE\_TP => TP  
ICD9\_TI => TP

## Skipped patients

A query can skip patients. Reasons for skipping a patient are:

- FAIL PATIENT command
- querying for a feature that is flagged as incomplete (if at least one event for a feature does not have a valid time (dates before birth, dates after death, etc.), whole feature will be flagged as incomplete and querying for it will skip the patient
- querying negation commands (NOT, INVERT, NOTE(NOT(, BEFORE(X, Y)-()) if the whole feature set is missing in patient's records. For example querying for RX=1192 will skip patient if patient has no RX codes.

## Comments

Comments are portions of the query that are not evaluated.

1. Line comment is a comment starting with // and ending with end of line
2. Multi line comment is a comment starting with /\* and ending with \*/
3. Line starting with a # character denotes a comment

Examples:

```
var a = ICD9=250.00 // this is a comment
// also a comment
/* multi line comment
another multi line comment */
#comment
$a
```

## Variables

Variable definition:

```
var variable=UNION(ICD9=200.2, ICD9=100.1)
```

Variable usage:

```
INTERSECT($variable1, $variable2)
```

## Saving variables to server

```
var domain_name.variable_name=UNION(ICD9=200.2, ICD9=100.1);
```

Domain name is an identifier string that allows grouping variables into cohesive units.

Variables that contain other variables will be stored after evaluating the nested variables, so that there are no dependencies between stored variables.

Example:

```
var variable1=UNION(ICD9=100.1, ICD9=200.2)
```

```
var variable2=INTERSECT(ICD9=200.3, ICD9=300.3)
```

```
var john.variable3=UNION($variable1, $variable2)
```

john.variable3 will be stored as UNION(UNION(ICD9=100.1, ICD9=200.2), INTERSECT(ICD9=200.3, ICD9=300.3))

## Accessing saved variables

```
$my_workspace.variable_name
```

When declaring a variable and not using it in the same query, an error is thrown (no query), but the variable will be persisted.

Example:

```
var john.variable = ICD9=200.5
```

This will define a global variable, but will not store it. Variable will be stored only after you call the variable with .finalize, .store or .save suffix. Already defined variables cannot be overwritten. Once the finalize was called, the variable is read only. This also means that when you attempt to redefine a variable that was already saved, this will throw an error.

Example:

```
var john.variable = ICD9=200.5
```

```
$john.variable.finalize
```

Defining variable without storing it:

```
var john.variable = ICD9=200.5
```

```
$john.variable
```

Stored variables cannot be redefined.

## Basic temporal operations

### TIMELINE

Returns a single TI containing patient's whole timeline.

Example: TIMELINE

Returns TI

### ATC

Returns TI during which the patient had at least one RXNorm code belonging to the specified ATC

Example: ATC="XXX"

Returns TI

## ENCOUNTERS

A day during the patient had at least one encounter (only a visit counts as an encounter).

Returns a list of TI that are computed in such a way that each TI is exactly 1 day long and there is no overlap between TI.

## NOTES

Returns all time instances when the patient had text notes.

## PATIENTS(X, Y, Z)

Returns patients with the specified patient IDs

Example: AND(PATIENTS(1, 3, 5, 7, 12), ICD9=250.00)

Returns BOOLEAN

## SNOMED=

Returns a list of TI during which the patient had the specified SNOMED code

Example: SNOMED=12345

Returns TI

## SNOMED

Returns a list of all SNOMED TI for a patient

Example: SNOMED

Returns TI

## DEPARTMENT=

Returns a list of all TI for a patient during which he was in the specified department

Example: DEPARTMENT="xxxxx"

Returns TI

## CPT=

Returns a list of TI during which the patient had the specified CPT code

Example: CPT=12345

Returns TI

## CPT

Returns a list of all TI for all CPT codes of a patient. Is equivalent to UNION(CPT=.....)

Example: CPT

Returns TI

## VISIT TYPE="xxxx"

Returns time interval during which the patient had the specified visit type.

Returns TI

## ICD9=

Returns a list of ICD9\_TI during which the patient had the specified ICD9 code

Example: ICD9=E123.22

Returns TI

## ICD9

Returns a list of all ICD9\_TI for a patient. Equivalent to UNION(ICD9=...)

Example: ICD9

Returns TI

## ICD10=

Returns a list of ICD10\_TI during which the patient had the specified ICD10 code

Example: ICD10=E123.22

Returns TI

## ICD10

Returns a list of all ICD10\_TI for a patient. Is equivalent to UNION(ICD10=...)

Example: ICD10

Returns TI

## NOTE TYPE=

Returns a time interval during which the patient had note with the specified note type.

Example: NOTE TYPE="Progress Note, Outpatient"

NOTE command allows querying text commands within a note with the specified note type.

Example: NOTE(NOTE TYPE="Progress Note, Outpatient", TEXT="diabetes")

Returns NOTE\_TP

## PRIMARY

Only usable on a single ICD9 or ICD10 code. Returns TI for which the patient had the ICD9 or ICD10 code mentioned as primary diagnosis

Example: PRIMARY(ICD9=E123.22)

To return times when the code was mentioned but was not a primary code:

INTERSECT(ICD9=E123.22, INVERT(PRIMARY(ICD9=123.22)))

Returns TI

## RX=

Returns a list of RXNORM\_TI during which the patient had the specified RXNORM code

Example: RX=122

Returns RXNORM\_TI

## RX

Returns a list of all RXNORM\_TI for a patient. Equivalent to UNION(RX=...)

Example: RX

Returns RXNORM\_TI

## DRUG

Allows querying additional properties of a drug (drug status and drug route)

Example: DRUG(RX=122, STATUS="discontinued", ROUTE="intravenous")  
DRUG(RX=122, STATUS="discontinued")

DRUG(RX=122, ROUTE="intravenous")

Returns TI

## **TEXT=" "**

Returns a list of NOTE\_TP where the patient had specified text mentioned without negation or family history  
Returns NOTE\_TP

## **!TEXT=" "**

Negated text was present in notes.  
Returns NOTE\_TP

## **~TEXT=" "**

Family history text was present in notes  
Returns NOTE\_TP

## **Text field limitations**

Whenever using the text fields (for example: TEXT="something", ATC="something"), the text field between the quotation marks can contain any character, except for quotation mark character and a dollar sign. These are reserved for the internal use in the language.

## **NOTE(TEXT=" ", TEXT=" ")**

All the text specified has to occur in one note.

NOTE also accepts boolean operators (AND, NOT, OR) as their parameters.

NOTE itself behaves in a same way as the command NOTE(AND(TEXT=" ", TEXT=" "))

NOTE(OR(TEXT=" ", TEXT=" ")) returns all patients that have any of the specified texts. This is identical to OR(TEXT=" ", TEXT=" ")

NOTE(NOT(AND(TEXT=" ", TEXT=" ))) is identical to NOT(NOTE(TEXT=" ", TEXT=" "))

Boolean notes allow more complex queries such as:

NOTE(AND(TEXT="diabetes", OR(TEXT="treated", "treatment", "controlled"), NOT(TEXT="uncontrolled")))

Returns NOTE\_TP

## **YEAR**

Returns a TI during which the patient had any code during the specified year

Example: YEAR=2008

Caution: Year only contains the intervals explicitly defined in visits and notes. There is no computation of missing intervals, so using EXTEND BY could lead to an interval that has an undefined YEAR.

Returns TI

## **GENDER**

Example: GENDER="MALE"

Return BOOLEAN

## **RACE**

Example: RACE="WHITE"

Return BOOLEAN

## **ETHNICITY**

Example: ETHNICITY="LATINO"  
Return BOOLEAN

## **DEATH / DEAD**

Returns the TP of patient's death  
Example: DEATH  
Returns TP

## **NULL**

Returns an empty TI

## **VITALS**

Returns the time points at which the vitals had specified values  
Example: VITALS("Pulse", 60, 100)  
VITALS("Pulse")

If no parameters are used, all times when the patient had vitals taken are returned  
Example: VITALS  
Returns TP

## **LABS**

Returns the time points at which the labs had specified values.  
Example: LABS("WBC", "NORMAL")  
LABS("WBC")  
LABS("WBC", 2.5, MAX)

If no parameters are used, all times when the patient had labwork done are returned  
Example: LABS  
Returns TP

## **AGE**

Returns the time interval during which the patient was of a specified age range.  
Example: AGE(30 years, 35 years)

The difference between INTERVAL(30 years, 35 years) and AGE(30 years, 35 years) is, that AGE will return only the intervals during which the patient had some event recorded, whereas INTERVAL will construct the intervals regardless of whether there were some events or not.

Returns TI

## **Boolean operations**

Boolean operations operate on other boolean operations or basic temporal operations. If boolean operations operate on a mixture of basic temporal operations and boolean operations, appropriate temporal operation is performed first, then evaluation of the boolean operations is performed.

## AND

Performs a boolean AND operation on other boolean operations or returns TRUE if the input is at least one TI

Example: AND(CPT=1234,CPT=222,OR(CPT=2345,CPT=435))

Results:

CPT=1234 = 1-5

CPT=222 = 3-7

CPT=2345 = 2-3

CPT=435 =5-7

result = AND(1-5,3-7,OR(2-3,5-7))

result = AND(true,true,OR(true,true))

result = AND(true,true)

result = true

Returns BOOLEAN

## OR

Performs a boolean OR operation on other boolean operations or true if the input is at least one interval

Example:OR(CPT=222,CPT=333,AND(CPT=444,CPT=555))

Results:

CPT=222 = 1-5

CPT=333 = 2-8

CPT=444 = 2-4

CPT=555 = 6-9

result = OR(1-5,2-8,AND(2-4,6-9))

result = OR(1-8,AND(true,true))

result = OR(1-8,false)

result = OR(true,false)

result = true

Returns BOOLEAN

## NOT

Performs a boolean NOT operation on another single boolean operation or returns true if the input is an empty TI

Patient's record: CPT 3 = 1-5

CPT 4 = 10-12

CPT 1 = 7-11

Example:NOT(CPT=1)

result = false

Example: NOT(AND(CPT=4,CPT=1))

result = NOT(AND(true,true))

result = NOT(true)

result = false

Returns BOOLEAN

## IDENTICAL

Tests two commands whether they are completely identical (same number of intervals having the same starts and ends)

The difference between EQUAL and IDENTICAL is that EQUAL returns any intervals of same starts and ends,

but IDENTICAL returns all the intervals provided there are the same numbers of intervals with the same starts and ends for both arguments.

Example: IDENTICAL(INTERVAL(100, 200), UNION(INTERVAL(100, 200), INTERVAL(250, 300)))  
result = false

EQUAL(INTERVAL(100, 200), UNION(INTERVAL(100, 200), INTERVAL(250, 300)))  
result = [100, 200]

## TI operations

TI operations operate on TI, if the input is a boolean operation, true is converted to the TI of the whole patient's timeline and false is converted to an empty interval.

### START

Prerequisites: Performs TI => CTI

Returns a set of TI with the time coordinates [START, START]

Patient's record: CPT 3 = 1-5, 2-7, 9-10

Example: START(CPT=3)

result = [1-1, 2-2, 9-9]

Returns TP

### END

Prerequisites: Performs TI => CTI

Returns a set of TI with the time coordinates [END, END]

Patient's record: CPT 3 = 1-5, 2-7, 9-10

Example: END(CPT=3)

result = [5-5, 7-7, 10-10]

Returns TP

### COUNT

Does not perform computation of intervals before evaluation. Computation is performed on the result if both COMMAND\_EVALUATE and COMMAND\_INTERVAL are specified as intervals to return.

Returns a TI for which the number of specified commands was in the specified range.

Command accepts following configurations of parameters:

COUNT(COMMAND\_EVALUATE, MIN, MAX)

COUNT(COMMAND\_EVALUATE\*, COMMAND\_INTERVAL\*, TYPE, MIN, MAX)

Asterisk denotes which interval to return in the second case (command\_interval or command\_evaluate). Default is to return command\_evaluate.

COMMAND\_EVALUATE = Condition in which count we are interested in

MIN / MAX = interval of min / max counts. MIN and MAX values are accepted

COMMAND\_INTERVAL = section of patient's timeline for which we want to evaluate the COUNT command

TYPE = SINGLE (a single interval in patient's timeline must have the corresponding counts with double counting) / ALL (cumulative counts for all intervals with no double counting)

Example: COUNT(OR(~TEXT="diabetes", ~TEXT="cancer"))\*, INTERSECT(ICD9=250), ALL, 5, MAX)

Takes TI for which the patient had ICD9=250 and counts the occurrences of "diabetes" or "cancer" in family history and returns all the ICD9=250 intervals if cumulatively there were at least 5 counts.

## INTERVAL(X, Y)

Given two time points, constructs a TI. If the parameters are TI, all combinations of START of TI1, END of TI2 will be performed for which it is true, that START of TI1 <= END of TI2.

Interval parameters can be numeric, in which case it constructs an interval from the time points specified.

Interval command is also compatible with time commands.

Example: INTERVAL(3 years, 4 years)  
result = [(3\*365\*24\*60), (4\*365\*24\*60)]

Patient's record: CPT 3 = 1-5  
CPT 4 = 10-12

Example: INTERVAL(START\_OF(CPT=4), END\_OF\_RECORD)

result = INTERVAL(4-4, 12-12)

result = 4 – 12

Returns TI

## INTERVAL(X, Y, PAIRS)

Evaluates X and Y as series of pairs.

Result = UNION(INTERVAL(START(X1), START(Y1)), INTERVAL(START(X2), START(Y2)), INTERVAL(START(X3), START(Y3))) for all X1 <= Y1 and X2 <= Y2 and X3 <= Y3 etc.

Without the PAIRS parameter, the following evaluation is made:

Result = UNION(INTERVAL(START(X1), START(Y1)), INTERVAL(START(X1), START(Y2)), INTERVAL(START(X1), START(Y3)), INTERVAL(START(X2), START(Y1)), INTERVAL(START(X2), START(Y2)), INTERVAL(START(X2), START(Y3)), etc.)

Returns TI

## INTERSECT

Prerequisites: Performs TI => CTI

Returns an intersection of multiple TI. If there is no intersection, returns an empty TI.

Patient's record: CPT 3 = 1-5  
CPT 4 = 10-12  
CPT 1 = 7-11

Example: INTERSECT(CPT=1, CPT=4)

result = INTERSECT(7-11, 10-12)

result = 10-11

Returns CTI

## UNION

Prerequisites: Performs TI => CTI

Performs an union of multiple TI

Patient's record: CPT 3 = 1-5  
CPT 4 = 10-12  
CPT 1 = 7-11

Example: UNION(CPT=3, CPT=4)

result = UNION(1-5, 10-12)

result = 1-5, 10-12

Returns CTI

## FIRST MENTION

Returns a first time interval for a given expression

Patient's record: CPT 3 = 1-5, 10-12  
CPT 4 = 10-12  
CPT 1 = 7-11

Example: FIRST\_MENTION(CPT=3)

result = 1-5

Returns Single TI with min start and min end

First mention can work also in a particular context returning first mention of a term in a particular interval.

Patient's record: CPT 2 = 5-10, 20-30  
CPT 3 = 1-3, 15-22

Example: FIRST\_MENTION(CPT=3, CPT=2) (find first mention of CPT3 in CPT 2)

result = 20-22

Returns first mention (if any found of CPT=3 within CPT=2 time interval)

## LAST MENTION

Returns the last time interval for a given expression

Patient's record: CPT 3 = 1-5, 10-12  
CPT 4 = 10-12  
CPT 1 = 7-11

Example: LAST\_MENTION(CPT=3)

result = 10-12

Returns Single TI with max start and max end

Last mention can work also in a particular context returning last mention of a term in a particular interval.

Patient's record: CPT 2 = 5-10, 20-30  
CPT 3 = 2-8, 31-35

Example: LAST\_MENTION(CPT=3, CPT=2) (find first mention of CPT3 in CPT 2)

result = 5-8

Returns first mention (if any found of CPT=3 within CPT=2 time interval)

## EXTEND BY (RESIZE)

Prerequisites: Performs TI => CTI

Extend the start and end of each TI by a specified amount of time units.

EXTEND BY(TI, time\_start, time\_end)

EXTEND BY(TI, 0, START + 10)

EXTEND BY (TI, END – 10, 0)

If the intervals overlap after the extension, they will be merged. Intervals that would extend patient's time-line will be truncated so that they are contained in the time line.

Extends start of the interval by time\_start (negative towards past, positive towards future) and end of the interval by time\_end.

Patient's record: CPT 3 = 1-5, 10-12

Example: EXTEND\_BY(CPT=3,-2, 2)

result = (1-2)-(5+2),(10-2)-(12+2)

result = (-1)-(7),(8)-(14)  
result = 0-7, 8-12  
Returns TI

## EVALUATE / EVAL

Parameters: EVALUATE(TYPE, COMMAND)

TYPE:

INTEGER evaluates the first N patients and returns how many were true  
TIME returns as many patients as possible in the specified time  
CACHED evaluates cached patients only

## LIMIT

Parameters: LIMIT(TYPE, COMMAND)

TYPE:

INTEGER returns the specified number of patients LIMIT(1000, ICD9=200.00) => returns first 1000 pids  
TIME returns as many patients as possible in the specified time  
CACHED evaluates cached patients only

## ESTIMATE

Parameters: ESTIMATE(TYPE, COMMAND)

Only returns the positively evaluated PIDs. Estimates the total cohort statistics based on evaluated pids.

TYPE:

INTEGER returns the specified number of patients ESTIMATE(1000, ICD9=200.00) => returns first 1000 pids  
TIME returns as many patients as possible in the specified time  
CACHED evaluates cached patients only

## EQUAL

Parameters: EQUAL(TI1, TI2)

Returns only the intervals that have the same starts and ends.

## BEFORE STRUCTURE

Before command can be invoked in the following structure:

(X) BEFORE Y\*

returns every Y which exists at least one minute after the start of X

(X\*) BEFORE Y

returns every X which exists at least 1 minute before the start of Y

(X AND Y AND Z) BEFORE A\*

returns every A that has at least 1 minute before its start at least 1 minute of X, Y and Z

(X\* AND Y AND Z\*) BEFORE A

returns every X, Z for which it's true that it exists before a start of A

(X) AND NO (Y) BEFORE A\*

returns A that has X and no Y before its start

(X\*) AND NO (Y) BEFORE A

returns X that exists before A and where there is no Y before A

(X AND Y) AND NO (Z AND B) 3 MONTHS\* BEFORE A

returns 3 months before A for which it is true that there was a X and Y but no Z and B

Note that it is not possible to return any data points from the negative mentions NO (...). Otherwise it is possible to tag any elements of the expression including the time element.

## AFTER STRUCTURE

After command can be invoked in the same structure as before with the same syntax, except that the keyword AFTER is used.

## BEFORE (SEQUENCE)

Performs  $TI \Rightarrow CTI$  on  $TI1$  and  $TI2$

Returns  $TI$  (does not perform  $CTI$  on results, so if results are  $[2, 5]$   $[4, 8]$  it will NOT compute it into  $[2, 8]$ )

Parameters: BEFORE( $TI1$ ,  $TI2$ )

Asterisk denotes which parameter to return is evaluation is successful.

Returns  $TI1$  before  $TI2$ . Can be followed by multiple parameters.

$TI1$  and  $TI2$  can be followed by asterisk which designates whether to return  $TI1$  and/or  $TI2$  as a result of a successful query.

When no asterisk is specified and no return type is specified in other parameters, command cannot be executed.

Parameters: BEFORE( $TI1$ ,  $TI2$ )>>(-100, -1)\*<>(END+1, END+100)

+ sign before a parameter means that the  $TI1$  has to occur in that range. If the  $TI1$  does not happen, then the  $TI2$  for which we evaluate will fail

- sign before a parameter means that the  $TI1$  cannot occur in that range. If  $TI1$  occurs, the  $TI2$  fails

To return the range of the condition (-100, -1), asterisk needs to be added before the range:

BEFORE( $TI1$ ,  $TI2$ )>>\*(-100, -1)

This command returns the range of (-100, -1) before the  $TI2$ , if  $TI1$  was located in that range

If parameters start with <  $TI1$  had to begin in the specified range, if >  $TI1$  had to end in that range, if <> same  $TI1$  had to both start and end in the interval (contained by the interval).

The time range parameter >>(100, 200) is always evaluated based on the  $TI2$ , so BEFORE( $TI1$ ,  $TI2$ )>>(100, 200) will take the  $TI2$ , extend the START of  $TI2$  by 100 towards the future, extend the END by 200 into the future and evaluates whether there is  $TI1$  in that range.

BEFORE( $X$ ,  $X$ ) is not a valid command since  $TI$  compression takes place in BEFORE command

### Example 1

$X1 = [5, 5]$

$Y1 = [10, 20]$

$Y2 = [35, 40]$

BEFORE( $X$ ,  $Y$ )>>\*(-6, -1) = [29, 39]

As we can see parameter >>\*(-6, -1) is false for  $Y1$  ( $X1$  is within the specified range even though it should not be), but is true for  $Y2$ .

BEFORE( $X$ ,  $Y$ )>>\*(-6, -1) = [35, 40]

$Y2$  is returned since it is evaluated to true even though  $Y1$  was evaluated to false.

### Example 2

If we want to return only the intervals for which it is true that they were ALWAYS in a certain range before  $Y$

(there is no instance of X that was not in the range), we should use:

A = BEFORE(X, Y)+\*(R1, R2)

B = BEFORE(X, Y)-\*(R1, R2)

INTERSECT(NOT(\$B), \$A)

If there is at least one interval X that is not in range R1, R2, nothing will be returned.

Y1 = [10, 20]

Y2 = [30, 40]

X1 = [25, 27]

A = BEFORE(X, Y)+\*(MIN, -1) = [25, 27]

B = BEFORE(X, Y)-\*(MIN, -1) = [0, 9]

INTERSECT(NOT(\$B), \$A) = []

## DURATION

Performs computation of intervals first.

If COMMAND\_EVALUATE and COMMAND\_INTERVAL are followed by an asterisk, computation of intervals is performed on the result.

Returns a TI for which the duration of specified commands was in the specified range.

Command accepts following configurations of parameters:

DURATION(COMMAND\_EVALUATE, TYPE, MIN, MAX)

DURATION(COMMAND\_EVALUATE\*, COMMAND\_INTERVAL\*, TYPE, MIN, MAX)

Asterisk denotes which interval to return in the second case (command\_interval or command\_evaluate). Default is to return command\_evaluate.

COMMAND\_EVALUATE = Condition in which duration we are interested in

MIN / MAX = interval of min / max counts. MIN and MAX values are accepted

COMMAND\_INTERVAL = section of patient's timeline for which we want to evaluate the DURATION command.

Each time interval in the section is evaluated separately and then next time interval is iterated and evaluated.

ALL operator in this case will return if the patient had at desired durations cumulatively WITHIN the single time interval specified by the command.

TYPE = SINGLE (a single interval in patient's timeline must have the corresponding durations – double counting is permissible) / ALL (cumulative durations for all intervals – double counting is not permissible)

## INVERT

Prerequisites: Performs TI => CTI

Inverts a TI or TP by using patient's time line as a reference. Inversion of an empty interval is patient's complete interval.

Patient's record: CPT 3 = [1-3], [7-10]

CPT 2 = []

CPT 4 = [5 – 8]

Example: INVERT(CPT=2)

result = 1-10

Example: INVERT(INTERVAL(START\_OF\_RECORD, FIRST\_MENTION(CPT=4)))

This is equivalent to no history of

result = INVERT(1-5)

result = 6-10

Returns CTI

# Macro expressions

Macro expressions are sets of expression encapsulating common expression for convenience sake

## RECORD START

START(TIMELINE)

## RECORD END

END(TIMELINE)

## HISTORY OF(X)

Patient had a history of X. Returns TI from the first mention of X until the end of record. If patient never had X, empty interval will be returned.

INTERVAL(START(FIRST MENTION(X)), RECORD END)

## NO HISTORY OF(X)

Patient did not have a history of X. This will return a TI of the whole patient's time line if the patient never had X, or an interval from the beginning of the patient's time line to the development of X.

UNION(NOT(X), INVERT(HISTORY\_OF(X)))

## CONTAINS(X, Y)

Returns Y only if it's fully contained in X.

BEFORE(X, Y)+\*<>(START, END)

CONTAINS(X, X) is invalid as there is compression in BEFORE command

Returns TI of X that which are fully contained in Y

## RETURN X INTERSECTING Y

Returns full interval X if it intersects Y

BEFORE(Y, X\*)+(START, END)

## RETURN X INTERSECTING ANY (A, B, C)

Returns full interval X if it intersects at any point either A, B, or C.

BEFORE(UNION(A, B, C), X\*)+(START, END)

## RETURN X INTERSECTING ALL (A, B, C)

Returns full interval X if it intersects at any point A, B, and C. The intervals A, B, C do not have to intersect each other, but they have to intersect X

1 = BEFORE(A, X\*)+(START, END)

2 = BEFORE(B, X\*)+(START, END)

3 = BEFORE(C, X\*)+(START, END)

4 = INTERSECT(X, 1, 2, 3)

## RETURN X ALWAYS INTERSECTING Y

Returns full interval X if in patient's timeline all X intervals intersect Y

1 = BEFORE(Y, X\*)-(START, END)

2 = NOT(1)

3 = INTERSECT(X, 2)

## **RETURN X ALWAYS INTERSECTING ANY (A, B, C)**

Returns full interval X only if every X interval in patient's timeline intersects at some point A, B or C

1 = BEFORE(UNION(A, B, C), X\*)-(START, END)

2 = NOT (1)

3 = INTERSECT(2, X)

## **RETURN X ALWAYS INTERSECTING ALL (A, B, C)**

Returns full interval X if every X in patient's timeline always intersects A, B and C

1 = BEFORE(A, X\*)-(START, END)

2 = BEFORE(B, X\*)-(START, END)

3 = BEFORE(C, X\*)-(START, END)

4 = INTERSECT(1, 2, 3)

5 = NOT (4)

6 = INTERSECT(5, X)

## **RETURN X NOT INTERSECTING Y**

Returns full interval X if it does not intersect Y

BEFORE(Y, X\*)-(START, END)

## **RETURN X NOT INTERSECTING ANY (A, B, C)**

Returns full interval X if it does not intersect A, B or C. Intervals intersecting either A, B, or C will be removed

BEFORE(UNION(A, B, C), X\*)-(START, END)

## **RETURN X NOT INTERSECTING ALL (A, B, C)**

Returns full interval X if it does not intersect Y

1 = BEFORE(A, X\*)+(START, END)

2 = BEFORE(B, X\*)+(START, END)

3 = BEFORE(C, X\*)+(START, END)

4 = INTERSECT(1, 2, 3)

5 = INVERT(4)

6 = INTERSECT(X, 5)

## **RETURN X NEVER INTERSECTING Y**

Returns full interval X if it does not intersect Y

1 = BEFORE(Y, X\*)+(START, END)

2 = NOT (1)

3 = INTERSECT(X, 2)

## **RETURN X NEVER INTERSECTING ANY (A, B, C)**

Returns full interval X if it never intersected in patient's timeline A, B or C

1 = BEFORE(A, X\*)+(START, END)

2 = BEFORE(B, X\*)+(START, END)

3 = BEFORE(C, X\*)+(START, END)

4 = UNION(1, 2, 3)

5 = NOT(4)

6 = INTERSECT(5, X)

## **RETURN X NEVER INTERSECTING ALL (A, B, C)**

Returns full interval X if it is true that in patient's timeline it never intersected all of A, B, and C

1 = BEFORE(A, X\*)+(START, END)

2 = BEFORE(B, X\*)+(START, END)  
3 = BEFORE(C, X\*)+(START, END)  
4 = INTERSECT(1, 2, 3)  
5 = NOT(4)  
6 = INTERSECT(X, 5)

## NEVER HAD(X)

Patient never had a condition X. Returns patient's whole time line (true) if he never had X or an empty interval / false if he had.

NOT(HAS(X))

## EVENT FLOW (X)

Exports the current cohort into an EventFlow / CoCo file formats.

## OUTPUT (X)

When generating the list of PIDs in the export, the PID information will also contain time intervals for when the evaluated query was true in format PID [TAB] START\_TIME [TAB] END\_TIME

## CSV

Generates a csv file with specified list of columns.

CSV(INTERSECT(ICD9=250.00, CPT=24560), "METFORMIN"=AND(RX=1235), "DEATH TIME"=DEATH)

Apart from labeled columns, following commands will output all codes of the specified type:

ICD9, ICD10, LABS, VITALS, CPT, RX

If the column result would specify multiple time points / time intervals, they will be delimited with a semicolon

Example:

CSV(ICD9=250.00, "SURGERY"=CPT=22456)

CSV command returns all of patient's data for the entire timeline. It is possible to specify a subset of timeline using the TIME or START / END modifiers.

Example:

CSV(ICD9=250.50, TIME=VISIT TYPE="INPATIENT", CPT)

For all patients that had ICD9 code 250.50 output all CPT codes they had while being inpatient.

CSV(ICD9=250.50, START=EXTEND BY(DEATH, START-30 DAYS, START), END=DEATH, ICD9, LABS)

For all patients that had ICD9 code 250.50 output all ICD9 and LABS they had 30 days before death

## Cohort operations

### SAME

Returns list of patients that are in both compared cohorts.

VAR cohort1= INTERSECT(ICD9=250.0, NOTES)

VAR cohort2= AND(GENDER="MALE", RACE="WHITE")

SAME(\$cohort1, \$cohort2)

### DIFF

Returns list of patients that are represented in first parameter but not in second

DIFF(\$cohort1, \$cohort2)

# MERGE

Returns list of patients added from both cohorts

MERGE(\$cohort1, \$cohort2)

Examples:

History of disease A

HISTORY OF(A)

No history of disease A

NO HISTORY OF(A)

Patient with disease A taking drug B after procedure C

BEFORE(START(INTERSECT(A, B)), START(C))

Patient with disease A never had disease B

INTERSECT(NEVER HAD(B), HAS(A))

Patient with a history of disease B taking medication C

INTERSECT(HISTORY OF(B), C)

Patient with no history of disease A having disease B

INTERSECT(NO HISTORY OF(A), B)

Patient who had A maximum 6 months before developing B

BEFORE(A, B)+\*(-6\*30\*24\*60, -1)

Patient who had A at least 6 months before developing B

BEFORE(A, B)-(6\*30\*24\*60, -1)+\*(MIN, -6\*30\*24\*60-1)

Patient who had A exactly 6 months before developing B

BEFORE(A, B)+\*(-6\*30\*24\*60, -6\*30\*24\*60)

Patient hospitalized (H) for more than 6 months following procedure A (hospitalization had to occur 24 hours after procedure A)

DURATION(BEFORE(H, BEFORE(A, H)+\*(-24\*60, -1))+\*(START, END), 6\*30\*24\*60, MAX)

Define diabetes as at least 2 mentions of ICD9=250.\*, then extend it from the first mention to the end of life

INTERVAL(FIRST MENTION(COUNT(ICD9=250.\*, 2, MAX, ALL)), RECORD END)

Patient that had at least 2 mentions of family history of diabetes, take first mention of cancer and extend until end of recorded

INTERVAL(FIRST\_MENTION(INTERSECT(TEXT="cancer", AND(COUNT(~TEXT="diabetes", 2, MAX, ALL))))), RECORD\_END)

Patients who took medication X within 3 months before they died

INTERSECT(X, EXTEND BY(DEATH, -3\*30\*24\*60, 0))

Patients who underwent surgery A and they did not take medication X for at least 1 month prior to surgery

BEFORE(X, A\*)-(-30\*24\*60, -1)

Patients who started taking medication X 3 months before procedure A and stopped 1 month before the procedure and they never took the medication since. We need at least 6 months of continuous follow-up data afterwards

COUNT(ENCOUNTERS, EXTEND BY(BEFORE(X, BEFORE(X, A\*)+\*(-3\*30\*24\*60, -3\*30\*24\*60)\*->(-30\*24\*60, -1), 6\*30, MAX, ALL)

Patients that had diabetes(D) and during diabetes were not taking metformin(M).

INTERSECT(D, INVERT(M))

Patients who did not take medication X prior to event Y

BEFORE(X, Y\*)-(MIN, 0)

Patients who had surgery X while not being on a medication Y (X not intersecting Y)

INTERSECT(Y, INVERT(INTERSECT(X, Y)))

Patients that had lab A, B and C within 24 hours. Let's assume that labs are a single time points without duration

INTERSECT(EXTEND(A, 0, START + 24 HOURS), EXTEND(B, 0, START + 24 HOURS), EXTEND(C, 0, START + 24 HOURS))

Patients that had at least 10 labs A within 30 days and a lab B and C done at the same time-line

INTERSECT(B, C, COUNT(A, EXTEND(A, 0, START + 30 DAYS), 10, MAX))

24 hours during which patients had A, B, C

EXTEND(INTERSECT(EXTEND(C, 0, START + 1 DAY), EXTEND(B, 0, START + 1 DAY), EXTEND(A, 0, START + 1 DAY)), 0, 24 HOURS)

## Iterative evaluation

To evaluate each time interval in a query separately, iterative evaluation can be used. Examples where iterative evaluation allows operations that cannot be done in regular approach.

```
FOR EACH (COMMAND) AS (LABEL_MAIN) {  
}
```

Takes each time interval from the command COMMAND and makes it accessible in a for loop as LABEL\_MAIN

Example:

```
FOR EACH (INTERSECT(ICD9=250.50, CPT=25000)) AS (DIABETES) {  
  RETURN DIABETES AS VARIABLE_1;  
}  
INTERSECT(VARIABLE_1, GENDER="MALE")
```

Takes each interval from the intersection of both codes, makes it available in a for loop under the name DIABETES. There is no algorithm in place so it just returns all the results as VARIABLE\_1. VARIABLE\_1 is then accessible in the global context and can be evaluated.

Commands available in the loop:

```
LABEL = COMMAND;  
CONTINUE;  
EXIT;  
FAIL PATIENT;  
CLEAR GLOBAL_LABEL;  
RETURN LABEL AS GLOBAL_LABEL;  
IF EMPTY (COMMAND) { ... }  
IF !EMPTY (COMMAND) { ... }  
IF (COMMAND1) == (COMMAND2) { ... }  
IF (COMMAND1) != (COMMAND2) { ... }  
IF (COMMAND1) IN (COMMAND2) { ... }  
IF (COMMAND1) !IN (COMMAND2) { ... }
```

## LABEL=COMMAND;

Stores the result of the command into a variable accessible by the label name in the current for each loop context. Variable is not accessible in a nested for each context and is not accessible in global context. This command can be used to reassign previously assigned variable as well as global variable.

At the end of each loop, all local variables are deleted. The persistence of local variables is only within a single loop cycle.

To store value persistently use the RETURN command.

Example:

```
FOR EACH (ICD9=250.50) AS (DIABETES) {  
  LONGER_THAN_3_YEARS = DURATION(DIABETES, SINGLE, 3 YEARS, MAX);  
  RETURN LONGER_THAN_3_YEARS AS RESULT_1;  
}  
RESULT_1
```

## CONTINUE;

Skips the rest of the FOR EACH loop.

Example:

```

FOR EACH (ICD9=250.50) AS (DIABETES) {
  LONGER_THAN_3_YEARS = DURATION(DIABETES, SINGLE, 3 YEARS, MAX);
  // skips intervals shorter than 3 years
  IF EMPTY(LONGER_THAN_3_YEARS) {
    CONTINUE;
  }
  // return command is skipped for shorter ones
  RETURN DIABETES AS RESULT_1;
}
RESULT_1

```

## EXIT;

Stops evaluation of the FOR EACH loop and exits. All the results stored in the global variable before the exit will be returned.

Example:

```

FOR EACH (ICD9=250.50) AS (DIABETES) {
  LONGER_THAN_3_YEARS = DURATION(DIABETES, SINGLE, 3 YEARS, MAX);
  // returns only those instances that are longer than 3 years and followed by the first shorter instance
  IF EMPTY(LONGER_THAN_3_YEARS) {
    EXIT;
  }
  RETURN DIABETES AS RESULT_1;
}
RESULT_1

```

## FAIL PATIENT;

Stops evaluation of the FOR EACH loop and causes the patient's evaluation to be FALSE (empty interval).

Example:

```

FOR EACH (ICD9=250.50) AS (DIABETES) {
  LONGER_THAN_3_YEARS = DURATION(DIABETES, SINGLE, 3 YEARS, MAX);
  // returns only those patients which had all their durations longer than 3 years
  IF EMPTY(LONGER_THAN_3_YEARS) {
    FAIL PATIENT;
  }
  RETURN DIABETES AS RESULT_1;
}
RESULT_1

```

Returns only the patients which did not have any ICD9=250.50 codes shorter than 3 years.

## RETURN LABEL AS GLOBAL\_LABEL;

Takes a variable or a command and stores the resulting value into the global context (if FOR EACH is nested in another FOR EACH command, returned nested variable will be stored in the parent context).

Example:

```

FOR EACH (ICD9=250.50) AS (DIABETES) {
  A = INTERSECT(CPT=250000, DIABETES);
  FOR EACH (A) AS (NESTED_A) {
    B = DURATION(NESTED_A, SINGLE, 3 YEARS, MAX);
    IF NOT EMPTY(B) {
      RETURN B AS GLOBAL_B;
    }
  }
}

```

```

}
}
// GLOBAL_B is accessible within this context since it was returned by the nested FOR EACH loop
RETURN GLOBAL_B AS RESULT_1;
}
// RESULT_1 is available in global context since it was returned by the parent FOR EACH
RESULT_1

```

## CLEAR GLOBAL\_LABEL;

Clears the contents of the global variable.

Example:

```

FOR EACH (ICD9=250.50) AS (DIABETES) {
  A = INTERSECT(CPT=250000, DIABETES);
  FOR EACH (A) AS (NESTED_A) {
    B = DURATION(NESTED_A, SINGLE, 3 YEARS, MAX);
    IF NOT EMPTY(B) {
      RETURN B AS GLOBAL_B;
    }
    IF EMPTY(B) {
      CLEAR GLOBAL_B;
      EXIT;
    }
  }
}
}
// GLOBAL_B is incrementally receiving values from each iteration unless B is empty which clears all the
// previous results and exits the loop
GLOBAL_B

```

## IF EMPTY (COMMAND) { ... }

Executes commands in curly braces if the command evaluates to an empty interval.

## IF !EMPTY (COMMAND) { ... }

Executes commands in curly braces if the command evaluates to a non empty interval.

## IF (COMMAND1) == (COMMAND2) { ... }

Executes commands in curly braces if the command1 time intervals equal command2 time intervals

## IF (COMMAND1) != (COMMAND2) { ... }

Executes commands in curly braces if the command1 time intervals are not equal to command2 time intervals

## Using multiple iterative loops

Iterative loops are evaluated sequentially.

In this example, first the ICD9=222.22 is pushed to a global variable DEF, then global variable DEF is used in the second loop and returned as DEF.

```

FOR EACH(ICD9=222.22) AS (ABC) {RETURN ABC AS DEF;}
FOR EACH(DEF) AS (DEF) {RETURN DEF AS DEF;}
DEF

```

This for loop will throw an error, since it relies on a variable DEF that has is unknown  
FOR EACH(DEF) AS (DEF) {RETURN DEF AS DEF;}  
FOR EACH(ICD9=222.22) AS (ABC) {RETURN ABC AS DEF;}  
DEF

Each FOR EACH loop is evaluated, even if it does not return any results, so just the presence of a FOR EACH loop modifies the query.

FOR EACH loop is ignored if it does not contain FAIL PATIENT OR does not RETURN a variable used elsewhere.