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Departamento de Lenguajes y Ciencias de la Computación

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TESIS DOCTORAL

De la Información al Conocimiento

**Aplicaciones basadas en implicaciones y
computación paralela**

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HACEN CONSTAR QUE:

D. Fernando Benito Picazo, Ingeniero en Informática, ha realizado en el Departamento de Lenguajes y Ciencias de la Computación de la Universidad de Málaga, bajo nuestra dirección, el trabajo de investigación correspondiente a su Tesis Doctoral titulado:

De la Información al Conocimiento
Aplicaciones basadas en implicaciones y computación paralela

Revisado el presente trabajo, estimamos que puede ser presentado al tribunal que ha de juzgarlo, y autorizamos la presentación de esta Tesis Doctoral en la Universidad de Málaga.

Málaga, Julio de 2018

Dr. Manuel Nicolás García-Oliveros Dr. Carlos Manuel Rossi Jiménez

A mi querida Aurora,

a mi padre y a mi madre,

a mi hermano,

*por el apoyo y la confianza que
siempre me habéis demostrado.*

*En todo objetivo conseguido hay siempre una especial tristeza,
en el conocimiento de que una meta largamente deseada
se ha logrado al fin, y que la vida tiene entonces que ser
moldeada y encaminada en busca de nuevos fines.*

La Ciudad y las Estrellas

A. C. Clarke

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Aportaciones

A continuación se expone una lista de los trabajos que han sido publicados como resultado de la investigación llevada a cabo a lo largo de esta tesis doctoral. Estas publicaciones avalan el trabajo realizado poniendo de manifiesto tanto su interés como su validez científica.

Revistas

- (i) Fernando Benito-Picazo, Pablo Cordero, Manuel Enciso, Ángel Mora. *Minimal generators, an affordable approach by means of massive computation.* The Journal of Supercomputing, Springer, 2018.
DOI: 10.1007/s11227-018-2453-z.
Factor de impacto en J.C.R. 2016: 1,349. Posición 52 de 104 (Q2) en la categoría: ‘Computer Science, Theory & Methods’.
- (ii) Fernando Benito-Picazo, Manuel Enciso, Carlos Rossi, Antonio Guevara. *Enhancing the conversational process by using a logical closure operator in phenotypes implications.* Mathematical Methods in the Applied Sciences, John Wiley & Sons Ltd, 2017.
DOI: 10.1002/mma.4338.
Factor de impacto en J.C.R. 2016: 1,017. Posición 108 de 255 (Q2) en la categoría: ‘Mathematics, Applied’.
- (iii) Fernando Benito-Picazo, Pablo Cordero, Manuel Enciso, Ángel Mora. *Reducing the search space by closure and simplification paradigms. A*

parallel key finding method. The Journal of Supercomputing, Springer, 2016.

DOI: 10.1007/s11227-016-1622-1.

Factor de impacto en J.C.R. 2016: 1,349. Posición 52 de 104 (Q2) en la categoría: ‘Computer Science, Theory & Methods’.

Congresos

- Fernando Benito-Picazo, Pablo Cordero, Manuel Enciso, Ángel Mora. *Closed sets enumeration: a logical approach.* Proceedings of the Seventeenth International Conference on Computational and Mathematical Methods in Science and Engineering, CMMSE, 2017. Cádiz, Spain, July 4-8, pp. 287-292, ISBN: 978-84-617-8694-7.
- Fernando Benito-Picazo, Manuel Enciso, Carlos Rossi, Antonio Guevara. *Conversational recommendation to avoid the cold-start problem.* Proceedings of the Sixteenth International Conference on Computational and Mathematical Methods in Science and Engineering, CMMSE, 2016. Cádiz, Spain, July 4-8, pp. 184-190, ISBN: 978-84-608-6082-2.
- Fernando Benito-Picazo, Pablo Cordero, Manuel Enciso, Ángel Mora. *Keys for the fusion of heterogeneous information.* Proceedings of the Fifteenth International Conference on Computational and Mathematical Methods in Science and Engineering, CMMSE, 2015. Cádiz, Spain, July 6-10, pp. 201-211, ISBN: 978-84-617-2230-3.
- Fernando Benito-Picazo, Pablo Cordero, Manuel Enciso, Ángel Mora. *Increasing the Efficiency of Minimal Key Enumeration Methods by Means of Parallelism.* Proceedings of the 9th International Conference on Software Engineering and Applications, ICSOFT-EA, Vienna, Austria, August 29-31, 2014, pp. 512-517.

DOI: 10.5220/0005108205120517.

Capítulo 1

Introducción

La gestión de la información es uno de los pilares esenciales de la Ingeniería Informática. No es de extrañar, por tanto, que conforme un amplio campo de investigación y conocimiento donde diversas disciplinas como las Matemáticas, la Lógica y la Ingeniería actúen conjuntamente para alcanzar mejores sinergias.

Dentro de esta filosofía y en aras de mejorar el funcionamiento de dos tipos de sistemas de información como son las bases de datos y los sistemas de recomendación, en esta tesis doctoral nos vamos a centrar en el Análisis Formal de Conceptos (FCA, por sus siglas en inglés: *Formal Concept Analysis*); concretamente, en la utilización de dos de sus conceptos fundamentales: los retículos de conceptos y los conjuntos de implicaciones. La gestión inteligente de estos elementos mediante técnicas lógicas y computacionales confieren una alternativa para superar obstáculos que podemos encontrar a la hora de trabajar con los sistemas de información mencionados. Comenzamos pues introduciendo brevemente FCA.

Podemos considerar FCA como una teoría matemática y una metodología para derivar una jerarquía de conceptos a partir de una colección de objetos y las relaciones que verifican. De esta forma, el propósito es poder representar y organizar la información de manera más cercana al pensamiento humano sin perder rigor científico. En este sentido se enmarca la cita de

Rudolf Wille: “*El objetivo y el significado del FCA como teoría matemática sobre conceptos y sus jerarquías es apoyar la comunicación racional entre seres humanos mediante el desarrollo matemático de estructuras conceptuales apropiadas que se puedan manipular con la lógica.*”

El término FCA fue acuñado por Wille en 1984 culminando años más tarde con la publicación más citada al respecto en colaboración con Bernhard Ganter [38]. Desde entonces FCA se ha aplicado con éxito en diferentes disciplinas de la Ciencia, como por ejemplo: minería de datos, biología celular [33], genética [54], economía, ingeniería del software [99], medicina [76], derecho [70], gestión de la información [85], etc.

La motivación principal de FCA aboga por representar conjuntos de objetos y atributos por medio de tablas de datos. Estas tablas se denominan contextos formales y representan las relaciones binarias entre esos objetos y atributos. Principalmente, existen dos formas básicas para representar el conocimiento: los retículos de conceptos y los conjuntos de implicaciones.

Desde hace años, existen en la literatura estudios [57] donde se han investigado y comparado diferentes algoritmos para obtener el retículo de conceptos a partir del conjunto de datos (en adelante *dataset* por su nomenclatura habitual en el campo). Sin embargo, debido a que el tamaño del retículo de conceptos es, en el peor de los casos, $2^{\min(|G|, |M|)}$, siendo G el conjunto de objetos y M el conjunto de atributos, el coste computacional de los métodos encargados de su construcción constituye una limitación de la aplicación de FCA.

Por otro lado tenemos el conjunto de implicaciones. Las implicaciones pueden considerarse *grosso modo* como reglas del tipo *si-entonces*, o en inglés: *if-then rules*, cuya noción principal es un concepto muy intuitivo: cuando se verifica una premisa, entonces se cumple una conclusión. Esta idea básica se refleja en numerosos campos de conocimiento bajo diferentes nombres. Así, en base de datos relationales se denominan dependencias funcionales [20], en FCA son implicaciones [38], en programación lógica son reglas lógicas de programación [78]. No obstante, al igual que en el caso del retículo de conceptos, también existen ciertas desventajas a la hora de trabajar con implicaciones, de hecho, la propia extracción del conjunto

completo de implicaciones de un dataset es una tarea que presenta una complejidad exponencial [22].

Trabajar con conjuntos de implicaciones nos permite utilizar técnicas automáticas basadas en la lógica, lo que nos conduce al propósito principal de esta tesis doctoral, que principalmente consiste en, utilizando los conjuntos de implicaciones, aplicar mecanismos lógicos para realizar un tratamiento eficiente de la información.

La aproximación a través de la lógica es posible gracias a sistemas axiomáticos válidos y completos como los Axiomas de Armstrong [3] y la Lógica de Simplificación [24] (SL por sus siglas en inglés: *Simplification Logic*). Estos métodos aplicados sobre conjuntos de implicaciones nos proporcionan la base para trabajar en esta tesis doctoral fundamentalmente sobre los siguientes tres campos de conocimiento: claves minimales, generadores minimales y sistemas de recomendación conversacionales.

En cada uno de los casos, vamos a aprovechar la información subyacente al conjunto de implicaciones, que son el núcleo principal de conocimiento en esta tesis, para realizar novedosas aproximaciones que nos permitan abordar problemas presentes en esos ámbitos. De esta forma, con los métodos desarrollados basados en implicaciones, hemos conseguido obtener resultados favorables para todos esos campos. Tales resultados se sustentan por una amplia gama de experimentos, en los cuales se ha utilizado tanto información real como conjuntos autogenerados y donde la computación paralela en entornos de supercomputación ha desempeñado un papel crucial.

Antes de entrar con mayor detalle en los anteriores tres campos, es necesario hacer una importante declaración previa. Tal y como se ha mencionado anteriormente, vamos a trabajar con el conjunto de implicaciones que se verifican en un dataset. Sin embargo, hay que dejar claro que no es competencia de este trabajo el estudiar técnicas para la extracción de estas implicaciones (lo cual es más una tarea de minería de datos), sino que la intención es partir del punto en el que ya contamos con el conjunto de implicaciones para trabajar con él. A este respecto, podemos mencionar los trabajos más citados en la literatura en relación a la extracción del conjunto de implicaciones a partir de datasets [48, 117, 118]. Son trabajos de suma

importancia desde el punto de vista teórico pero también desde el punto de visto práctico, pues incluyen las implementaciones de las aplicaciones que realizan la extracción de las implicaciones.

1.1 Claves Minimales

El concepto de clave es fundamental en cualquier modelo de datos, incluyendo el modelo de datos relacional de Codd [20]. Una clave de un esquema relacional está compuesta por un subconjunto de atributos que representan el *dominio* de una determinada función cuya *imagen* es la totalidad del conjunto de atributos. Estas funciones se pueden representar por medio de Dependencias Funcionales (FD, por sus siglas en inglés: *Functional Dependencies*) que especifican una relación entre dos subconjuntos de atributos, e.g. A y B , asegurando que para cualesquiera dos tuplas de una tabla de datos, si verifican A , entonces también verifican B . Nótese el cambio de denominación de implicación a dependencia funcional por estar en el entorno de los sistemas de base de datos relacionales.

La identificación de las claves minimales de una determinada relación es una tarea crucial para muchas áreas de tratamiento de la información: modelos de datos [97], optimización de consultas [55], indexado [68], etc. El problema consiste en el descubrimiento de todos los subconjuntos de atributos que componen una clave minimal a partir de un conjunto de FD que se cumplen en un esquema de una tabla del modelo relacional.

Las claves no sólo son parte fundamental a considerar durante la fase de diseño de sistemas de base de datos relacionales, sino que también se consideran una poderosa herramienta para resolver multitud de problemas referentes a diversos aspectos del tratamiento de la información. Como muestras de la relevancia de este problema, encontramos numerosas citas en la literatura, entre las que podemos destacar las siguientes. En [98], los autores afirman que: “*identification of keys is a crucially important task in many areas of modern data management, including data modeling, query optimization (provide a query optimizer with new access paths that can lead to substantial speedups in query processing), indexing (allow the database*

administrator to improve the efficiency of data access via physical design techniques such as data partitioning or the creation of indexes and materialized views), anomaly detection, and data integration”. Más recientemente, en referencia a áreas emergentes como el *linked-data*, en [82], los autores delimitan el problema manifiestando: “*establishing semantic links between data items can be really useful, since it allows crawlers, browsers and applications to combine information from different sources.*”

Para ilustrar de forma básica el concepto de clave, vamos a utilizar el siguiente Ejemplo 1.1.1.

Ejemplo 1.1.1. *Supongamos que disponemos de la Tabla 1.1. Es una pequeña tabla donde se refleja información que relaciona títulos de películas, actores, países, directores, nacionalidad y años de estreno.*

De esta información, utilizando los métodos comentados anteriormente, podemos extraer el siguiente conjunto de FDs:

$\Gamma = \{Título, Año \rightarrow País; Título, Año \rightarrow Director; Director \rightarrow Nacionalidad\}.$

Cuadro 1.1: Tabla de películas

Título	Año	País	Director	Nacionalidad	Actor
Pulp Fiction	1994	USA	Quentin Tarantino	USA	John Travolta
Pulp Fiction	1994	USA	Quentin Tarantino	USA	Uma Thurman
Pulp Fiction	1994	USA	Quentin Tarantino	USA	Samuel Jackson
King Kong	2005	NZ	Peter Jackson	NZ	Naomi Watts
King Kong	2005	NZ	Peter Jackson	NZ	Jack Black
King Kong	1976	USA	De Laurentiis	IT	Jessica Lange
King Kong	1976	USA	De Laurentiis	IT	Jeff Bridges
Django Unchained	2012	USA	Quentin Tarantino	USA	Jamie Foxx
Django Unchained	2012	USA	Quentin Tarantino	USA	Samuel Jackson

Esta tabla tiene una única clave minimal: {Título, Año, Actor} que corresponde con el conjunto de atributos necesario para identificar cualquier tupla de la relación.

Una característica muy importante de las claves es su minimalidad. Una clave se considerará minimal cuando todos y cada uno de los atributos que

la forman son imprescindibles para mantener su naturaleza de clave, es decir, no contiene ningún atributo superfluo.

El problema de la búsqueda de claves

El problema de la búsqueda de claves consiste en encontrar todos los subconjuntos de atributos que componen una clave minimal a partir de un conjunto de FD que se verifican en un esquema de una tabla de datos relacional. Es un campo de estudio con décadas de antigüedad como vemos en [34], donde las claves se estudiaron dentro del ámbito de la matriz de implicaciones u otros tantos trabajos como [39, 91] que se centran en averiguar estas claves minimales.

La dificultad al enfrentarnos con el problema de la búsqueda de claves surge debido a que, dado un conjunto de atributos A , la cardinalidad del conjunto 2^A hace que haya que abordar el problema aplicando técnicas que guíen la búsqueda de los conjuntos candidatos a ser claves minimales.

El cálculo de todas las claves minimales representa un problema complejo. En [65, 119] se incluyen resultados interesantes acerca de la complejidad del problema; los autores demuestran que el número de claves minimales para un sistema relacional puede ser exponencial respecto al número de atributos, o factorial respecto al número de dependencias. Además, establecieron que el número de claves está limitado por el factorial del número de dependencias, por tanto, no existe un algoritmo que resuelva el problema en tiempo polinómico. En definitiva, es un problema NP-completo decidir si existe una clave de tamaño a lo sumo k dado un conjunto de FDs [65].

Es significativo como el problema de la búsqueda de claves aparece en diversos campos de conocimiento. Por ejemplo, en [9] se hace mención a la importancia de conocer las claves en áreas emergentes como el *linked-data*. Por otro lado, en [25], muestran cómo el problema de las claves minimales en las bases de datos tiene su análogo en FCA, donde el papel de las FDs se manifiesta como implicaciones de atributos. En ese artículo, el problema de las claves mínimas se presentó desde un punto de vista lógico y para ello se empleó un sistema axiomático para gestionar las FDs y las implicaciones

que los autores denominaron *SLFD* [24].

Las principales referencias sobre este problema apuntan a los trabajos de Lucchesi y Osborn en [65] que muestran un algoritmo para calcular todas las claves candidatas. Por otro lado, Saiedian y Spencer [90] presentaron un algoritmo usando grafos con atributos para encontrar todas las claves posibles de un esquema de base de datos relacional. No obstante, demostraron que sólo podía aplicarse cuando el grafo de FDs no estuviera fuertemente conectado. Otro ejemplo lo encontramos en el trabajo de Zhang [122] en el cual se utilizan mapas de Karnaugh [53] para calcular todas las claves. Existen más trabajos sobre el problema del cálculo de las claves minimales como son [98, 115] y otra contribución actual que aborda el problema en un estilo lógico [25]. Asimismo, en [62, 108, 109] los autores propusieron el uso de FCA [38] para abordar problemas relacionados con la búsqueda y la gestión de las implicaciones, que pueden considerarse complementarios a nuestro trabajo.

Algoritmos para el cálculo de claves

Por nuestra parte, como objetivo, nos hemos centrado en los algoritmos de búsqueda de claves basados en la lógica, y más específicamente, en aquellos que utilizan el paradigma de Tableaux [75, 89] utilizando un sistema de inferencia.

De forma muy general, podemos decir que los métodos tipo Tableaux representan el espacio de búsqueda como un árbol, donde sus hojas contienen las soluciones (claves). El proceso de construcción del árbol comienza con una raíz inicial y desde allí, las reglas de inferencia generan nuevas ramas etiquetadas con nodos que representan instancias más simples del nodo padre. La mayor ventaja de este proceso es su versatilidad, ya que el desarrollo de nuevos sistemas de inferencia nos permiten diseñar un nuevo método. Las comparaciones entre estos métodos se pueden realizar fácilmente ya que su eficiencia va de la mano con el tamaño del árbol generado.

Esto nos lleva a un punto de partida fundamental, los estudios de R. Wastl (Universidad de Wurzburg, Alemania) [110, 111] donde se introduce

por primera vez un sistema de inferencia de tipo Hilbert para averiguar todas las claves de un esquema relacional. A modo de ejemplo básico, en la Figura 1.1 podemos ver un ejemplo de árbol de búsqueda según el paradigma de Tableaux desarrollado según las reglas de inferencia del sistema de inferencia \mathbb{K} de Wastl.

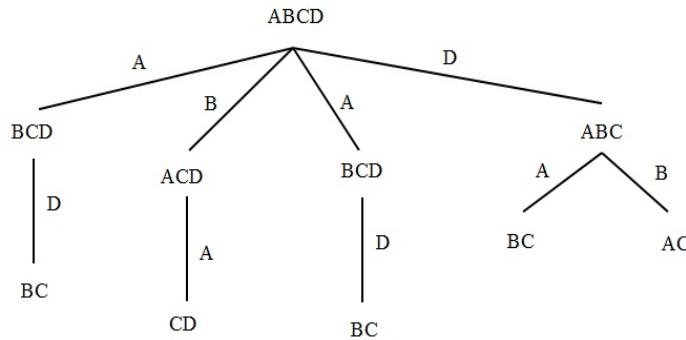


Figura 1.1: Ejemplo de Tableaux utilizando el sistema de inferencia \mathbb{K} de Wastl.

Siguiendo esta línea, en [23] los autores abordan el problema de la búsqueda de claves utilizando un sistema de inferencia basado en la lógica de simplificación para dependencias funcionales [24] demostrando como el árbol del espacio de búsqueda que se genera nos lleva a sobrepassar las capacidades de la máquina, incluso para problemas pequeños. En [73] los autores muestran la equivalencia entre SL_{FD} y los axiomas de Armstrong [3] junto con un algoritmo para calcular el cierre de un conjunto de atributos. Más tarde, en [25], los autores introdujeron el método SST, basado en la introducción del test de minimalidad que evita la apertura de ramas adicionales del árbol, por lo que el espacio de búsqueda se vuelve más reducido, logrando un gran rendimiento en comparación con sus predecesores.

Una propiedad muy interesante de los métodos basados en Tableaux es su generación de subproblemas independientes los unos de los otros a partir del problema original. De esta forma llegamos a nuestro objetivo

fundamental de utilizar las técnicas lógicas sobre una implementación paralela de los métodos que, mediante el uso de recursos de supercomputación, nos permitan alcanzar resultados en un tiempo razonable.

En esta línea, son varios los trabajos que han utilizado la paralelización para afrontar problemas relacionados con implicaciones o FCA. Un algoritmo paralelo para el tratamiento de implicaciones enmarcado en el campo de los hipergrafos lo podemos encontrar en [102]. A su vez, Krajca et al. [56] presentan un algoritmo paralelo para el cálculo de conceptos formales. Por nuestra parte, una primera aproximación a la paralelización del método de Wastl [110, 111] y el algoritmo de claves [23] fue presentado en [8], donde se muestra cómo el paralelismo puede integrarse de forma natural en los métodos basados en Tableaux.

Para la labor de computación de alto rendimiento, a lo largo de este trabajo se han adquirido y aplicado las competencias necesarias para poder trabajar en entornos de supercomputación; en concreto, durante los últimos años, se ha trabajado fervientemente con el Centro de Supercomputación y Bioinnovación de la Universidad de Málaga¹. La posibilidad de tratar con este Centro nos ha proporcionado dos beneficios fundamentales: por un lado, se ha alcanzado una elevada pericia para trabajar en entornos de HPC y para realizar implementaciones que aprovechen una alta cantidad de recursos, y por otro lado, se han podido obtener resultados empíricos sobre experimentos utilizando estrategias de paralelismo, que han desembocado en contribuciones científicas [8, 11] y que habría sido imposible conseguir en la actualidad sin contar con tales recursos computacionales.

Métodos *SST* y *CK*

En [25] se presentó un nuevo algoritmo, denominado SST, para calcular todas las claves minimales usando una estrategia de estilo Tableaux, abriendo la puerta a incorporar el paralelismo en su implementación. SST se basa en la noción de cierre de conjunto, una noción básica en la teoría de base

¹<http://www.scbi.uma.es/>

de datos que permite caracterizar el conjunto máximo de atributos que se puede alcanzar, desde un determinado conjunto de atributos A con respecto a un conjunto de FD, utilizando el sistema axiomático. Por lo tanto, si el cierre de A se denota como A_{Γ}^+ , el sistema de inferencia para FD nos permite inferir la FD $A \rightarrow A_{\Gamma}^+$. El enfoque con estilo lógico para el problema de las claves minimales consiste en la enumeración de todos los conjuntos de atributos A tales que se verifique la FD: $A \rightarrow \Omega$.

SST muestra un gran rendimiento en comparación con sus predecesores como hemos visto hasta ahora y como puede comprobarse en el amplio estudio realizado sobre el método en [7]. El beneficio principal en la reducción del espacio de búsqueda de debe a la introducción del test de inclusión para evitar la apertura de ramas extra. Gracias a ello, SST no abre algunas ramas que sabemos que van a producir las mismas claves que se calculan en otra rama.

Basándonos en el sistema axiomático de la lógica SL_{FD} 2.2, proponemos un nuevo método llamado *Closure Keys (CK)* que incorpora un mecanismo eficiente de poda que utiliza el método de cierre basado en SL_{FD} para mejorar el método SST. El nuevo operador de cierre definido en [73] permite reducir el espacio de búsqueda realizando reducciones en el camino hacia las hojas, donde finalmente se obtienen las claves. El método CK tiene una característica fundamental que lo convierte en un novedoso enfoque a los métodos clásicos de cierre, ya que proporciona una nueva funcionalidad: además del conjunto de atributos que constituye la salida del operador de cierre, el método proporciona un subconjunto de implicaciones del conjunto Γ original.

Por tanto, el método CK recibe un conjunto de implicaciones Γ y un subconjunto de atributos $X \subseteq \Omega$, calcula el conjunto cierre X^+ respecto a Γ , y además, un nuevo conjunto Γ' que contiene el conjunto de implicaciones que guarda la semántica que queda fuera del cierre X^+ . Si $\Gamma' = \emptyset$, entonces $X^+ = \Omega$ (véase [73] para más detalles).

Finalmente, la principal contribución de esta parte de la tesis se produce de la siguiente forma. Se ha desarrollado una implementación paralela tanto del método SST como del método CK basándose en el paradigma

MapReduce [30]. Básicamente, el algoritmo paralelo de búsqueda de claves se divide en dos partes principales. Utiliza una primera fase en la que se realiza una expansión del árbol de búsqueda trabajando sobre el problema original pero llegando únicamente hasta un cierto nivel de árbol, es decir sin alcanzar todavía las claves en las hojas del árbol. En ese momento interviene la segunda etapa en la que cada nodo de ese nivel del árbol, que constituye un problema equivalente al original pero simplificado como resultado de la aplicación de las reglas de inferencia hasta ese momento, se resuelve en paralelo mediante el uso de un elevado número de cores, es decir, aplica el algoritmo de búsqueda de claves, pero ahora sí, hasta alcanzar las hojas del árbol.

De esta forma, se han realizado multitud de experimentos para confirmar las mejoras que se han alcanzado, los cuales necesitan llevarse a cabo en entornos de supercomputación y cuyos resultados pueden consultarse en [11]. Principalmente, se ha demostrado como estos métodos son claramente adecuados para ejecutarse utilizando una implementación paralela. De esta forma, se consiguen resultados en tiempos razonables incluso cuando la cantidad de información de entrada es considerable y donde los métodos secuenciales no son capaces de finalizar.

1.2 Generadores Minimales

Como se ha mencionado anteriormente, una forma de representar en FCA el conocimiento es el retículo de conceptos. Esta representación otorga una visión global de la información con un formalismo muy fuerte, abriendo el puerta para utilizar la teoría de retículos como una metateoría para gestionar la información [13].

Los conjuntos cerrados son la base para la generación del retículo de conceptos ya que éste puede ser construido a partir de los conjuntos cerrados, considerando la relación de subconjuntos como la relación de orden. En este punto nace el concepto de generadores minimales como aquellas representaciones de los conjuntos cerrados que no contengan información superflua, es decir, representaciones canónicas de cada conjunto cerrado [38].

Los generadores minimales junto con los conjuntos cerrados son esenciales para obtener una representación completa del conocimiento en FCA, pero no sólo son interesantes desde un punto de visto teórico. La importancia de los generadores minimales puede apreciarse claramente a través de citas tales como: “*Minimal generators have some nice properties: they have relatively small size and they form an order ideal*”, existente en importantes estudios como el de Poelmans et al. [83] o [86]. Además, los generadores minimales se han usado como punto clave para generar bases, las cuales constituyen una representación compacta del conocimiento que facilita un mejor rendimiento de los métodos de razonamiento basados en reglas. Mis-saoui et al. [71, 72] presentan el uso de generadores minimales para calcular bases que impliquen atributos positivos y negativos cuyas premisas son generadores minimales.

Cualquier operador de cierre c en M puede asociarse con un sistema de implicaciones. Esta conexión establece una forma de gestionar el trabajo del operador de cierre c por medio de su derivación sintáctica y, como consecuencia, se puede elaborar un método para realizar esta gestión. Pero, ¿qué pasa con la conexión inversa? Es decir, dado un conjunto de implicaciones, ¿es posible generar el operador de cierre c asociado a él? Tal pregunta es el núcleo de esta parte de la tesis y su solución implica enumerar todos los conjuntos cerrados.

Si tenemos que $X, Y \subseteq M$ satisfacen que $X = Y_{\Sigma}^+$, es habitual decir que Y es un generador del conjunto cerrado X . Obsérvese que cualquier subconjunto de X que contiene Y es también un generador de X . Dado que trabajamos con conjuntos finitos de atributos, el conjunto de los generadores de un conjunto cerrado se pueden caracterizar por sus generadores minimales.

Métodos para el cálculo de generadores minimales

En nuestro caso, centrándonos una vez más en el tratamiento inteligente de los conjuntos de implicaciones, vamos a utilizarlos como los elementos para describir la información y además, como base del diseño de métodos

para enumerar todos los conjuntos cerrados y sus generadores minimales a partir de esta información, y no del dataset original (como en los trabajos mencionados), lo cual, hasta donde sabemos, no existe trabajo previo al respecto.

El nuevo método propuesto es una evolución del método presentado en [26] donde se utilizó la SL_{FD} como herramienta para encontrar todos los generadores minimales a partir de un conjunto de implicaciones. Este método trabaja sobre el conjunto de implicaciones aplicando unas reglas de inferencia y construyendo un espacio de búsqueda de árbol muy parecido a los árboles del caso de las claves minimales.

De manera específica, dado un conjunto de atributos M y un sistema de implicaciones Σ , el método realiza un mapeo $mg_\Sigma: 2^M \rightarrow 2^{2^M}$ que satisface la siguiente condición.

$$\forall X, Y \subseteq M$$

$$X \in mg_\Sigma(C) \text{ si y sólo si } C \text{ es cerrado para } (\)_\Sigma^+ \text{ y } X \text{ es un generador minimal para } C.$$

Ejemplo 1.2.1. Sea $\Sigma = \{a \rightarrow c, bc \rightarrow d, c \rightarrow ae, d \rightarrow e\}$, el mapeo mg_Σ se describe como:

X	\emptyset	b	e	be	de	ace	bde	$acde$	$abcde$
$mg_\Sigma(X)$	\emptyset	b	e	be	d	a	bd	ad	ab
						c		cd	bc

En otro caso, X no es cerrado y $mg_\Sigma(X) = \emptyset$. Nótese que \emptyset es cerrado y $mg_\Sigma(\emptyset) = \{\emptyset\}$, i.e. \emptyset es un generador minimal del conjunto cerrado \emptyset .

Tras el método presentado en [26] (que denominaron MinGen) se presenta ahora un nuevo método (MinGenPr) que aplica una importante mejora con respecto al anterior. Básicamente consiste en incorporar un mecanismo de poda, basada en el test de inclusión de conjuntos que involucra a todos los nodos del mismo nivel, para evitar la generación de generadores minimales y cierres redundantes. El propósito de esta poda es verificar la

información de cada nodo en el espacio de búsqueda, evitando la apertura de una rama completa. Finalmente, se propone un último método (Gen-MinGen) que generaliza la estrategia de poda anterior al considerar el test de inclusión del subconjunto no sólo con la información de los nodos del mismo nivel, sino también con todos los generadores minimales calculados antes de la apertura de cada rama.

En definitiva, se han estudiado e implementado cada uno de estos métodos en su versión secuencial, y para evaluar el rendimiento e ilustrar las mejoras obtenidas al pasar de un método a otro, se han realizado un gran número de pruebas utilizando información autogenerada e información real.

Generadores minimales y paralelismo

La contrapartida es que la obtención de todos los conjuntos cerrados y sus respectivos generadores minimales es un problema con complejidad exponencial. Sin embargo, dado que nuestro objetivo es explotar las posibilidades de operar con conjuntos de implicaciones, en este trabajo vamos a combinar ambos aspectos enumerando todos los conjuntos cerrados a partir de un conjunto dado de implicaciones. De hecho, iremos un paso más allá, calculando no sólo todos los conjuntos cerrados, sino que para cada uno de ellos produciremos sus respectivos generadores minimales.

No obstante, vamos a encontrarnos con un problema similar al que sucedía con las claves minimales, y es que, al tratar con grandes cantidades de información, los métodos realizados para producir los generadores minimales, conllevan unas necesidades de cómputo que sobrepasan los límites de la máquina. Por tanto, una vez más, tenemos que trasladar las implementaciones de los métodos a versiones paralelas que nos permitan funcionar bajo arquitecturas de supercomputación.

En este sentido, se han realizado tanto las implementaciones secuenciales como paralelas de los métodos de producción de generadores minimales (MinGen, MinGenPr, GenMinGen, MinGenPar). Para las versiones paralelas vamos a utilizar la misma filosofía de implementación que en el caso de las claves minimales, es decir, hemos desarrollado unos códigos,

basándonos en el esquema *MapReduce* [30], que se ejecutarán en dos etapas. Al igual que en las claves minimales, la primera etapa divide el problema de entrada en varios subproblemas equivalentes pero reducidos de forma que puedan ser tratados de forma independiente. En la segunda etapa, cada uno de estos subproblemas se resuelve en paralelo usando múltiples núcleos.

Una vez más, para verificar el rendimiento y la idoneidad de los métodos para aplicar estrategias paralelas, se ha realizado una amplia batería de pruebas tanto sobre información autogenerada como información real. Además, las pruebas han incluido tareas de estimación del número óptimo de cores a utilizar así como del valor de corte más apropiado en la etapa primera de los métodos paralelos. Los resultados obtenidos respecto a esta parte de la investigación pueden consultarse en [10] y, especialmente en [12], que constituye uno de los trabajos que avalan esta tesis doctoral.

1.3 Sistemas de Recomendación Conversacionales

La última aplicación que se ha llevado a cabo en esta tesis doctoral haciendo uso de los conjuntos de implicaciones y los conjuntos cerrados se produce en el campo de los sistemas de recomendación (SR).

De forma muy básica, podríamos considerar que un SR es un sistema inteligente que proporciona a los usuarios una serie de sugerencias personalizadas (recomendaciones) sobre un determinado tipo de elementos (ítems). De forma general, los SR estudian las características de cada usuario e ítem del sistema, y mediante un procesamiento de los datos, encuentra un subconjunto de ítems que pueden resultar de interés para el usuario. Una de las referencias más notables en el campo de los SR la encontramos en el libro de Adomavicius y otros [2].

Desde que el primer SR hizo su aparición en el mundo de las tecnologías de la información [47, 88], los SR han estado en continua evolución durante los últimos años [1]. Sin embargo, es con la expansión de las nuevas tecnologías cuando han tenido un acercamiento más directo a la mayor parte de la sociedad debido a su capacidad para realizar todo tipo de recomendaciones sobre diversos elementos al alcance de todos (libros [28], documen-

tos [84], música [58], turismo [16], películas², etc.).

En la actualidad, los SR constituyen un claro campo de investigación y estudio como demuestran el gran número de trabajos que se están realizando [32, 100] y cuya cantidad continúa aumentando día a día. Además, la relevancia de estos sistemas no se limita al ámbito investigador. Actualmente, muchos SR ya han sido implantados con éxito en fuertes entornos comerciales a nivel mundial. Este es el caso de empresas líderes en el sector como pueden ser Amazon [63], LinkedIn [87] o Facebook [106], que han realizado fuertes inversiones con el fin de generar mejores SR. Estas situaciones ponen de manifiesto la gran importancia de estos sistemas en ambas vertientes de la sociedad actual.

Abordar la generación de recomendaciones haciendo uso de FCA es una aproximación existente en la literatura desde hace años. En [31], los autores utilizan FCA para agrupar elementos y usuarios en conceptos para posteriormente, realizar recomendaciones colaborativas según la afinidad con los elementos vecinos. Más tarde, en [94], los autores introducen un modelo para el filtrado colaborativo basado en FCA para generar correlaciones entre datos a través de un diseño del retículo. Zhang et al. [121] propusieron un sistema basado en similitud agrupando la información contextual en grafos mediante el cual llevar a cabo recomendaciones sobre las interacciones sociales entre usuarios. En [60, 61], se utilizan relaciones difusas e implicaciones ponderadas para especificar el contexto y SL_{FD} para desarrollar un proceso lineal de filtrado que permite a los SR podar el conjunto original de elementos y así mejorar su eficiencia. Recientemente, en [124] se propone y utiliza un novedoso SR personalizado basado en el retículo de conceptos para descubrir información valiosa de acuerdo con los requisitos e intereses de los usuarios de forma rápida y eficiente. Todos estos trabajos subrayan claramente cómo FCA puede aplicarse con éxito en el campo de los SR.

²<https://www.movielens.org>

Técnicas de recomendación

Existen numerosos tipos diferentes de SR que normalmente se clasifican atendiendo a cómo se llevan a cabo las recomendaciones. Los más conocidos y extendidos son los sistemas de filtrado colaborativo (denominados CF, por sus siglas en inglés: *Collaborative Filtering*) basan su funcionamiento fundamentalmente en las valoraciones que otros usuarios han otorgado a los elementos disponibles; y los sistemas basados en contenido (denominados CB, por sus siglas en inglés: *Content-Based*) que se basan en categorizar los ítems a recomendar, proporcionando resultados que tengan características similares a otros que han sido bien valorados anteriormente por el usuario.

En los últimos años ha habido un gran crecimiento de los SR contextuales [93], capaces de tener en cuenta información relevante para la recomendación como puede ser la hora, el lugar, la compañía, la ubicación, etc. Los SR demográficos [5] que clasifican a los usuarios según diferentes parámetros personales (edad, localización, etc.). Por otro lado encontramos los denominados SR basados en conocimiento (KB, por sus siglas en inglés: *Knowledge-Based*) [67]. Estos sistemas gestionan el conocimiento inherente a los datos y revelan cómo un ítem puede satisfacer la necesidad del usuario, es decir, utilizan un método de razonamiento para inferir la relación entre una necesidad y una posible recomendación. Finalmente, llegamos a los SR más importantes desde el punto de vista de este trabajo, los denominados SR conversacionales [41, 59]. Estos SR están estrechamente relacionados con los conceptos de recomendador basado en críticas [19] y recomendaciones de información [107]. Resaltamos este tipo de SR porque será la estrategia principal sobre el que se sustenta el desarrollo de SR realizado y que ha dado lugar a una de las contribuciones que avalan esta tesis [36]. Se puede consultar una clasificación más detallada en el libro de Adomavicius y Tuzhilin [2] que, junto con la contribución de Bobadilla et al. [15], constituyen las referencias más citadas en el campo.

Podemos apreciar que existen diferentes tipos de estrategias para los SR, sin embargo, la historia ha demostrado ampliamente que la mejor alternativa consiste en combinar características de diferentes tipos de SR

para generar híbridos que se beneficien de las ventajas de cada uno de ellos [29]. Tal es nuestro caso, en el que nuestro trabajo ha culminado en un SR híbrido que combina las siguientes técnicas de recomendación:

- **SR basados en conocimiento.** En virtud de nuestro estudio de extracción de conocimiento de los datos utilizando FCA, los conjuntos de implicaciones y los operadores de cierre.
- **SR basados en contenido.** Necesario ya que para aplicar las técnicas lógicas empleadas necesitamos en primera instancia información sobre la que trabajar.
- **SR conversacionales.** Se presenta como estrategia central sobre la que aplicar y utilizar las dos anteriores respectivamente.

Pero no nos ocuparemos únicamente de la estrategia de recomendación sino también del proceso de cómo obtener una recomendación. En este sentido, se introduce el concepto de Recuperación de Información (IR por sus siglas en inglés, *Information Retrieval*), que básicamente se encarga de realizar búsquedas para determinar cuán bien responde cada objeto a una consulta. Numerosos trabajos en la literatura relacionan el uso de FCA en modelos basados en IR [21, 50].

Problemas comunes

Si bien es cierto que los SR están alcanzando una enorme importancia existen numerosas dificultades que han de afrontarse a la hora de diseñar e implementar un SR. En la lista de problemas relacionados con los SR [95] podemos encontrar: el arranque en frío [35, 101], privacidad [37], oveja-negra [40], escasez [44], ataques maliciosos [116, 123], sobreespecialización [64], escalabilidad [51], postergación [105], dimensionalidad [92], etc.

En concreto, nuestro trabajo ha estado orientado a abordar este último problema de la dimensionalidad en los SR. Este problema, también conocido como *the curse of dimensionality phenomenon* [77, 92] aparece cuando es

necesario trabajar sobre datasets con un alto número de características (variables o atributos). De forma intuitiva, podríamos introducirlo de la siguiente manera. Cuando hay pocas columnas de datos, es relativamente fácil para los algoritmos realizar tareas de tratamiento inteligente de la información como: aprendizaje automático, *clustering*, clasificación, etc. Sin embargo, a medida que aumentan las columnas o características de nuestros ítems, se vuelve exponencialmente más difícil hacer labores predictivas con un buen nivel de precisión. El número de filas de datos necesarias para realizar cualquier modelado útil aumenta exponencialmente a medida que agregamos más columnas a una tabla.

Para abordar este problema, podemos encontrar muchos trabajos en la literatura sobre la reducción de la dimensión de la información, especialmente mediante selección de características (*feature selection*), que pueden ayudarnos a descartar aquellas características que no son merecedoras de ser considerados según diferentes criterios. De hecho, estas técnicas ya se aplican en otras áreas como son: algoritmos genéticos o redes neuronales, normalmente centrándose en la aplicación de un proceso automatizado que se aplique de una vez (*batch mode*) mediante selección de características.

En definitiva, nuestro objetivo ha sido abordar el problema de la alta dimensionalidad en los SR haciendo uso de los conjuntos de implicaciones a través de un proceso de selección de atributos por parte del usuario mediante un SR conversacional que utilice características de los SR basados en contenido y en conocimiento.

Un trabajo interesante en esta área es [52], que establece la idoneidad de los enfoques basados en el conocimiento para los procesos conversacionales. En particular, estos autores utilizan el razonamiento basado en restricciones, en lugar de nuestro enfoque basado en la lógica. Además, este trabajo trata sobre concepto de optimización de consultas, análogo al aplicado en nuestra propuesta. Otro trabajo notable es [107], que comparte nuestro objetivo de disminuir el número de pasos de la conversación. Los autores proponen métricas acerca del número de pasos de la conversación y tasas de poda, ambos muy similares a los utilizados en nuestro trabajo. Por otro lado, en [18], los autores demuestran cómo la posibilidad de que

sea el usuario el encargado de la selección de atributos supera al hecho de que sea el sistema mismo el encargado de dicha selección. Este hecho respalda nuestro enfoque en el cual el experto humano guía la conversación y el proceso de selección de características.

Evaluación de los sistemas de recomendación

La evaluación de las predicciones y recomendaciones se ha convertido en un aspecto muy importante [17, 45]. Los SR requieren medidas de calidad y métricas de evaluación [42] para conocer la calidad de las técnicas, métodos y algoritmos para las predicciones y recomendaciones. Las métricas de evaluación [46] y los *frameworks* de evaluación [14] facilitan la comparación de varias soluciones para el mismo problema.

No obstante, hay que tener en cuenta que dependiendo del SR con el que estemos trabajando, la evaluación habrá que llevarla a cabo utilizando aquellas métricas, que por su naturaleza y significado, tengan cabida en relación al SR que se desea evaluar. En nuestro caso, dado que hemos desarrollado un SR conversacional, puede ser evidente que la primera medida que podemos aplicar es calcular el número de pasos que se producen en la conversación [69]. Por contra, otras métricas tan populares como son *Precision* y *Recall* [43] no son adecuadas de aplicar.

Aplicación desarrollada

La solución que hemos realizado consiste en gestionar el problema de la alta dimensionalidad mediante un proceso de selección de características guiado por el usuario (experto humano) dentro de un sistema conversacional [120]. Para ello, una vez más haremos uso de una gestión inteligente de las implicaciones y de los conjuntos cerrados que nos va a permitir reducir sustancialmente el número de pasos necesarios en el diálogo entre el usuario y la aplicación para conseguir una recomendación adecuada en tiempo y forma. Para ello contaremos con el apoyo de la *SLFD* introducida por los autores en [24] y en particular, se utilizará el algoritmo del cierre

de atributos SL_{FD} [73] como núcleo de un marco de selección de atributos que será el que nos permita reducir el número de etapas del diálogo.

Además, se han realizado numerosas pruebas de aplicación sobre la propuesta que se ha desarrollado. En concreto, se han realizado pruebas utilizando información real sobre enfermedades y fenotipos como podemos apreciar en una de las contribuciones que avalan este trabajo de investigación [36]. Además, nuestra propuesta, al igual que la gran mayoría de los SR, permite actuar sobre diferentes datasets de forma que podemos utilizar el mismo procedimiento para mejorar las recomendaciones conversacionales en diversos entornos. Esta versatilidad es una característica notoria, ya que permite libertad de maniobra en el caso de que se quieran introducir ciertos cambios, o simplemente que los datos sean diferentes. En ese sentido, la propuesta de este trabajo casa con los conceptos de adaptabilidad y longevidad de los SR ya que el funcionamiento es independiente de la información de base con la que trabaje, sólo necesitamos conocer el conjunto de atributos e implicaciones subyacente a los datos.

Finalmente, antes de llegar a las últimas líneas del capítulo que dedicaremos a establecer el contenido del documento y las contribuciones producidas, cabe resaltar ya en este punto la naturaleza dual de la tesis, en el sentido de que mantendremos una línea de investigación en fundamentos teóricos complementada con la aplicación de dichos resultados en los tres campos de conocimiento mencionados anteriormente, claves minimales, generadores minimales y sistemas de recomendación. Haremos especial hincapié en la parte aplicada del estudio con la intención de facilitar la transferencia de conocimiento a entornos diferentes del ámbito académico, como el mercado empresarial.

A modo de resumen gráfico, la Figura 1.2 muestra un esquema del camino que investigación que se ha seguido en el desarrollo de esta tesis doctoral, apoyado por los principales conceptos y referencias.

Pasamos entonces ahora a desglosar la estructura del documento y la producción conseguida.

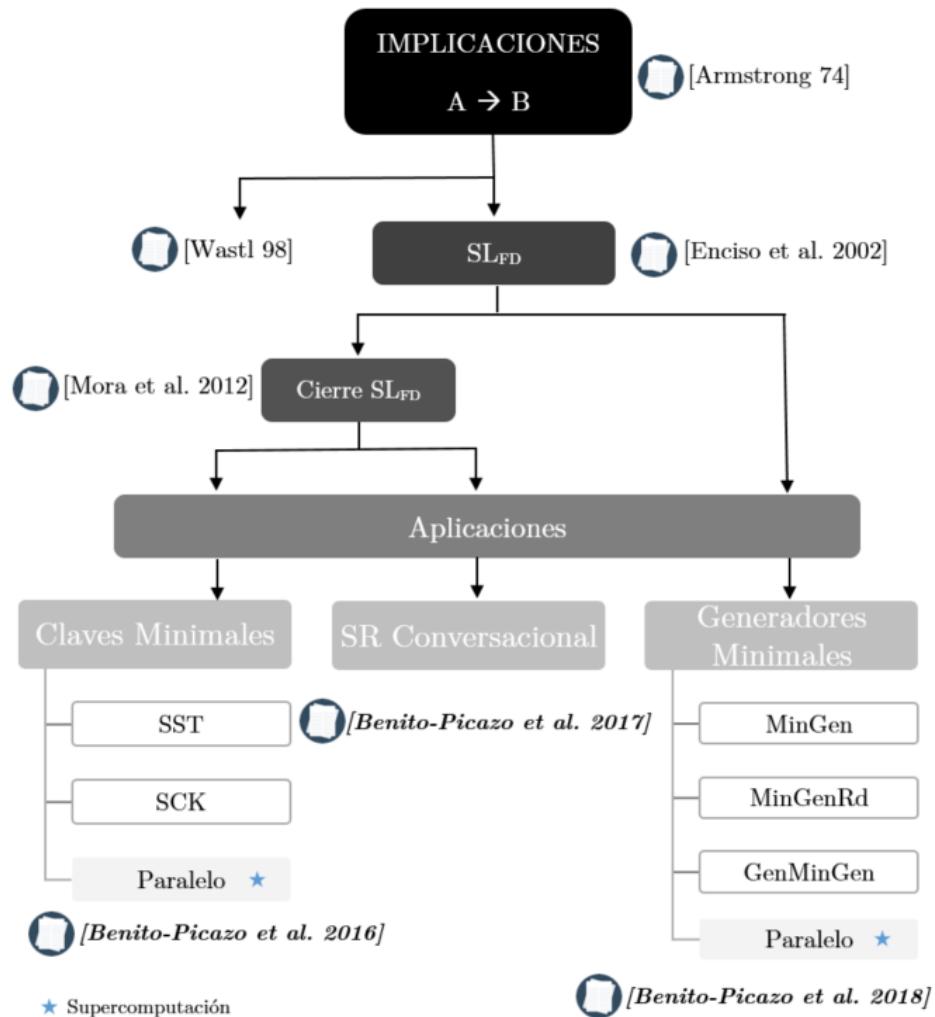


Figura 1.2: Esquema del estado del arte y las contribuciones generadas.

Estructura de la Tesis

En este primer capítulo de introducción, hemos fijado los puntos fundamentales de la tesis, como son: el marco de trabajo sobre el que vamos a actuar, las técnicas que utilizaremos y los principales objetivos que se pretenden alcanzar. Concretamente, hemos estipulado la utilización de FCA y los conjuntos de implicaciones como base del estudio sobre la que aplicar técnicas basadas en la lógica para mejorar el tratamiento de la información.

A continuación, entraremos en el Capítulo 2, en el que hemos recopilado un conjunto de conceptos previos necesarios relacionados con: FCA, la lógica de simplificación, los sistemas de implicaciones y los operadores de cierre. Tras estos dos primeros capítulos, ya contaremos con el conocimiento previo necesario para abordar el resto de capítulos de la tesis.

Tras la introducción y los preliminares, pasamos a un tercer capítulo 3 en el que se presenta la primera contribución que avala este trabajo de investigación y que corresponde con el trabajo realizado en el campo de la búsqueda de las claves minimales. De forma general, este artículo presenta nuevos métodos para resolver el problema de la inferencia de claves minimales en esquema de datos basándose en la SL_{FD} y el uso de implicaciones. Además, refleja las implementaciones y las ventajas obtenidas al aplicar técnicas de computación paralela para poder aplicar los métodos sobre conjuntos de datos de una entidad tal que técnicas secuenciales no son capaces de gestionar en cuanto a tiempo y recursos necesarios. Para ello se presentan los resultados obtenidos para los experimentos en entornos de supercomputación.

A continuación, llegamos a un capítulo análogo al anterior pero ahora para el tema de estudio referente a los generadores minimales 3. Este capítulo presenta un nuevo artículo en el cual se lleva a cabo un estudio de los métodos de producción de generadores minimales basados en la lógica y el tratamiento de implicaciones. Se comprueba las mejoras de rendimiento de los métodos al aplicar reducciones en el espacio de búsqueda basadas en estrategias de poda. Al igual que en el caso de las claves minimales, se presentan las implementaciones paralelas de los métodos para poder tratar

con conjuntos de datos de tamaño considerable y se incluyen las pruebas realizadas en entornos de supercomputación.

Como último capítulo dedicado a las aplicaciones desarrolladas mediante la gestión de implicaciones se presenta el capítulo 5. En este capítulo se incluye un novedoso artículo en el que se desarrolla una aproximación al tratamiento del problema de la dimensionalidad que aparece en el campo de los sistemas de recomendación. En él, mediante el uso eficiente de la *SLFD*, las implicaciones y los operadores de cierre, se consigue un modelo de sistema de recomendación conversacional que es capaz de gestionar el problema de la dimensionalidad aliviando la sobrecarga de información con la que el usuario debe lidiar a la hora de obtener una recomendación por medio de un sistema conversacional. Asimismo, se demuestra su utilidad en entornos reales mediante la realización de un experimento sobre información real.

Finalmente, cerraremos la tesis con un último capítulo 6 dedicado a recopilar las principales conclusiones obtenidas y a proponer caminos por los que seguir ahondando en la investigación en esta materia. Además, se incluye una relación de las referencias consultadas y los respectivos índices de términos, figuras y tablas.

En aras de la completitud, se incluyen como anexos finales aquellos artículos que han sido publicados a lo largo de este periodo de investigación, que si bien no se utilizan como respaldo para esta tesis doctoral por no corresponder con los criterios de calidad exigidos a tal efecto, han sido la semilla y experiencia inicial para conseguir el resto de trabajos que sí actúan como aval.

En la Figura 1.3 se muestra de forma gráfica el contenido de la tesis y las contribuciones publicadas.

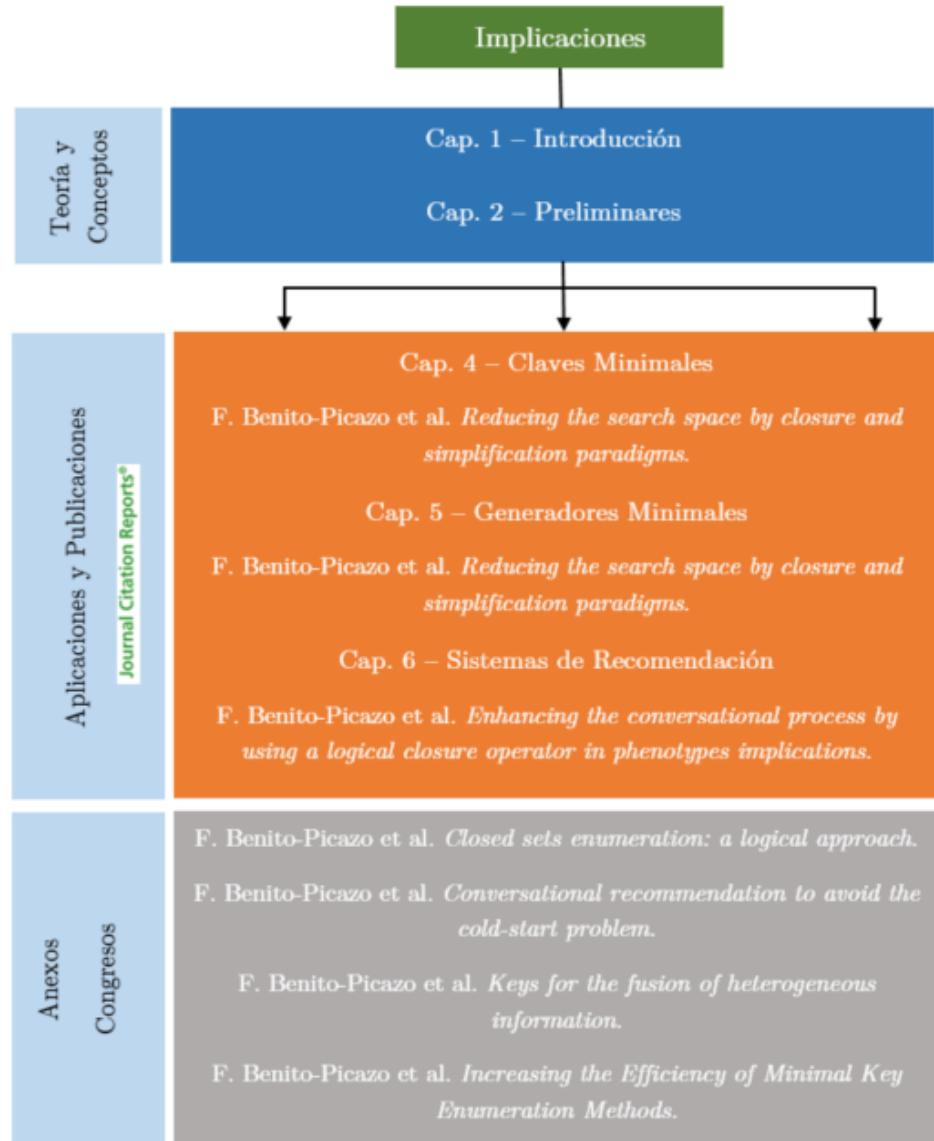


Figura 1.3: Esquema de la estructura de la tesis y las publicaciones.

Capítulo 2

Preliminares

A lo largo de este capítulo vamos a introducir las principales ideas, definiciones y propiedades de FCA como marco de investigación principal sobre el que se sustenta el trabajo de investigación desarrollado. Procederemos de forma que consigamos que el texto sea autocontenido en la medida de lo posible. Más concretamente, se introducen los principales conceptos sobre conjuntos de implicaciones y conjuntos cerrados que serán necesarios a lo largo del documento para poder plasmar en detalle las contribuciones realizadas. La referencia principal de este campo de conocimiento viene de la mano de Wille y Ganter en [38].

2.1 Análisis de conceptos formales

De forma general, podemos considerar FCA como un método de análisis de datos que facilita la extracción de conocimiento y la posibilidad de poder razonar al respecto. Es un marco conceptual con el que: analizar, estructurar, visualizar y sobre todo, revelar el conocimiento subyacente a los datos utilizando, generalmente, técnicas de minería de datos. Existen varias de estas técnicas para actuar sobre conjuntos de objetos, conjuntos de atributos y las relaciones que se establecen entre ellos [103, 104].

La característica principal de FCA es que, basado en sólidos funda-

mentos matemáticos, cubre una amplia gama de áreas de aplicación. Su motivación original fue la representación de retículos completos de objetos y sus propiedades por medio de contextos formales, que son tablas de datos que representan relaciones binarias entre objetos y atributos. Por otro lado, FCA puede verse como una alternativa razonable a la Teoría de Retículos en términos epistemológicos [114].

El punto de partida de FCA es un **contexto formal**, lo cual nos conduce necesariamente a las dos siguientes definiciones.

Definición 2.1.1 (Atributo). *Un atributo A es un identificador para un elemento en un dominio D .*

Definición 2.1.2 (Contexto formal). *Un contexto formal es una tripleta $\mathbf{K} := (G, M, I)$ que consiste en dos conjuntos no vacíos, G y M , y una relación binaria I entre ellos. Los elementos de G se llaman objetos del contexto, y los elementos de M se llaman atributos del contexto. Para $g \in G$ y $m \in M$, escribimos $< g, m > \in I$ o gIm si el objeto g posee el atributo m .*

De forma más específica, un contexto formal se puede considerar como un grafo bipartito que refleje las relaciones entre los conjuntos G y M , o también, como una tabla donde los objetos se sitúan en las filas de la tabla y los atributos en las columnas, de forma que un valor lógico en cada celda nos indica si un objeto $g \in G$ posee un atributo $m \in M$.

Ejemplo 2.1.3. Consideremos un ejemplo de contexto formal en el cual los objetos (G) son hoteles, los atributos (M) son diferentes servicios que ofrece el establecimiento y en cada celda de la tabla encontraremos una equis siempre y cuando el correspondiente hotel ofrezca el correspondiente servicio. Por tanto, tendremos el contexto formal $\mathbf{K} := (G, M, I)$ en el cual:

$G = \{\text{Fuerte Estepona Suites, Hotel Buenavista, Hotel Paraiso, Apts Marriot Playa, Hotel Piedra Paloma, Hostal Hospederia V Cent}\}$

$M = \{\text{AC, Bar, Gym, Internet, Masajes, Parking}\}$

La Tabla 2.1 muestra la relación binaria I del contexto formal K utilizando información real de varios hoteles de la Costa del Sol.

Cuadro 2.1: Ejemplo de contexto formal utilizando datos reales del sector hotelero

Nombre del Hotel	AC	Bar	Gym	Internet	Masajes	Parking
Fuerte Estepona Suites	✓	✓	✓	✓	✓	✓
Hotel Buenavista			✓			✓
Hotel Paraiso	✓	✓				✓
Apts Marriot Playa				✓		
Hotel Piedra Paloma				✓		✓
Hostal Hospederia V Cent	✓	✓			✓	✓
...						

Por tanto, vemos como el hotel Fuerte Estepona Suites ofrece todos los servicios disponibles en el contexto, mientras que los Apts Marriot Playa únicamente ofrecen el servicio de Internet.

A partir del contexto formal, se definen dos operadores llamados operadores de derivación.

Definición 2.1.4 (Operadores de derivación). *Dado un contexto formal $\mathbf{K} := (G, M, I)$, a continuación se definen dos funciones, denominadas operadores de derivación:*

$$(\)' : 2^G \rightarrow 2^M \quad (\)' : 2^M \rightarrow 2^G$$

$$A' = \{m \in M \mid g \ I \ m \quad \forall g \in A\} \quad B' = \{g \in G \mid g \ I \ m \quad \forall m \in B\}$$

Ambas funciones se denotan con el mismo símbolo porque no hay lugar a confusión. El primero asigna a cada conjunto de objetos, el conjunto de atributos comunes a todos los objetos, y el segundo asigna, a cada conjunto de atributos, el conjunto de objetos que tienen todos estos atributos.

El par de operadores de derivación constituye una conexión anti-monótona de Galois [79] con las consecuencias correspondientes, como que ambas composiciones son operadores de cierre, una de ellas en $(2^G, \subseteq)$ y la otra en $(2^M, \subseteq)$.

Definición 2.1.5 (Mapeo anti-monótono). *Sean G y M dos conjuntos no vacíos. Un mapeo de la forma $f : 2^G \rightarrow 2^M$ se dice anti-monótono si $A_1 \subseteq A_2$ implica que $f(A_2) \subseteq f(A_1) \quad \forall A_1, A_2 \subseteq G$.*

Definición 2.1.6 (Conexión de Galois). *Sean G y M dos conjuntos no vacíos. Un par (f,g) de dos mapeos $f : 2^G \rightarrow 2^M$ y $g : 2^G \rightarrow 2^M$ es una conexión de Galois entre G y M si, para cada $A \subseteq G$ y $B \subseteq M$ se cumple que: $A \subseteq g(B)$ si y sólo si $B \subseteq f(A)$.*

El operador de cierre " (i.e. aplicar el operador de derivación ' dos veces) verifica algunas propiedades interesantes que serán fundamentales para poder desarrollar la teoría formal.

Definición 2.1.7 (Operador de cierre). *Dado un conjunto no vacío M , un operador de cierre sobre M es una función $(\cdot)'' : 2^M \rightarrow 2^M$ que satisface las siguientes propiedades:*

- *Idempotente (ítems):* $X''' = X'' \quad \forall X \in 2^M$
- *Monótona (atributos):* $X \subseteq Y \rightarrow X'' \subseteq Y'' \quad \forall X, Y \in 2^M$
- *Extensa:* $X \subseteq X'' \quad \forall X \in 2^M$

En general no se verifica que $X'' = X$, no obstante, cuando un conjunto de objetos, $X \subseteq G$ verifica $X'' = X$, se llama conjunto cerrado. De manera análoga a los conjuntos de objetos, podemos hablar de conjuntos de atributos cerrados. El par $(M, '')$ se denomina sistema cierre y un conjunto $A \subseteq M$ se denomina conjunto cerrado para " si es un punto fijo para ", es decir, $"(A) = A$. La idempotencia de los operadores de cierre nos lleva al hecho de que, para todo sistema de cierre $(M, '')$ y todo $A \subseteq M$, el conjunto $"(A)$ es cerrado para ".

Además, los conjuntos cerrados definen lo que se denominan *conceptos formales*. De forma general, un concepto formal, que constituye un punto clave en FCA, permite describir formalmente un hecho del modelo y caracterizar un conjunto de objetos por medio de los atributos que comparten y viceversa.

De forma más intuitiva, un par (X, Y) es un concepto si se verifica que:

- Cada objeto en X tiene todos los atributos de Y .

- Para cada objeto en G que no está en X , existe un atributo en Y que el objeto no tiene.
- Para cada atributo en M que no está en Y , hay un objeto en X que no tiene ese atributo.

En otras palabras, (X, Y) es un concepto formal si X contiene todos los objetos cuyos atributos están en Y y, análogamente, Y contiene todos los atributos verificados por los objetos en X .

De esta forma, conseguimos introducir en la definición las dos partes esenciales: en primer lugar, el conjunto de objetos con propiedades comunes, y en segundo lugar, el conjunto de atributos que caracterizan a dichos objetos. Únicamente aquellos pares de conjuntos (X, Y) que tienen un cierre completo, establecen un concepto por sí mismos. En esta situación, los conjuntos X e Y son cerrados y se llaman, respectivamente, la *extensión* y la *intensión* del concepto. Para un conjunto de objetos X , el conjunto de sus atributos comunes X' , describe la similitud de los objetos de X , mientras que el conjunto X'' es la agrupación de aquellos objetos que tienen como atributos comunes a X' , en particular, todos los objetos de X , es decir, $X \subseteq X''$.

Más formalmente:

Definición 2.1.8 (Concepto formal). *Sea $\mathbf{K} := (G, M, I)$ un contexto formal y $X \subseteq G$, $Y \subseteq M$. El par (X, Y) se denomina concepto formal si $X' = Y$ y $Y' = X$. El conjunto de objetos X se denomina extensión del concepto (X, Y) mientras que el conjunto de atributos Y será la intensión del concepto.*

El conjunto de todos los conceptos formales de un contexto formal K constituye el denominado *retículo de conceptos* con la relación de inclusión que se muestra a continuación.

Si (X_1, Y_1) y (X_2, Y_2) son conceptos, se define un orden parcial, \leq , de forma que $(X_1, Y_1) \leq (X_2, Y_2)$ si $X_1 \subseteq X_2$, o equivalentemente, si $Y_2 \subseteq Y_1$.

Los pares de conceptos en este orden parcial tienen una única máxima cota inferior, que es el concepto generado por $X_1 \cap X_2$. De forma análoga,

poseen una única mínima cota superior, el concepto generado por los atributos $Y_1 \cap Y_2$. Por medio de estas operaciones de cálculo del máximo y del mínimo de dos conceptos, se satisfacen los axiomas que definen un retículo. La Figura 2.1 muestra un ejemplo básico de contexto formal y retículo de conceptos asociado.

Cuadro 2.2: Ejemplo de contexto formal básico

	adulto	joven	femenino	masculino
niño		✓		✓
niña		✓	✓	
hombre	✓			✓
mujer	✓		✓	

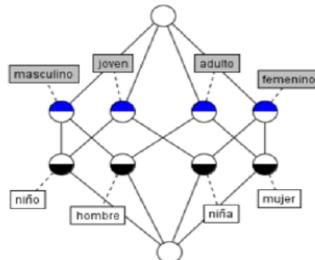


Figura 2.1: Retículo de conceptos

A partir del retículo de conceptos podemos obtener el contexto formal que representa. Los nodos del diagrama representan los objetos y atributos del contexto formal. Cada atributo del contexto corresponde a un elemento del retículo que tiene el conjunto de atributos minimal que contiene ese atributo, y un conjunto de objetos formado por todos los objetos con ese atributo. Por tanto, en la Figura 2.1 estaríamos hablando de: masculino, joven, adulto y femenino. Análogamente, cada objeto del contexto corresponde a un elemento del retículo que está formado por el conjunto de objetos minimal que contiene ese objeto, y un conjunto de atributos que consta de todos los atributos de dicho objeto. Esta vez, en la Figura 2.1 nos referimos a: niño, hombre, niña y mujer.

Una alternativa gráfica de definir los conceptos formales es la siguiente. Desde la representación del contexto formal, un concepto formal se puede reconocer por medio de una submatriz de tal manera que todas las celdas de la submatriz son verdaderas (no es necesario que la submatriz esté formada por celdas contiguas). De esta forma, en la Tabla 2.3, vemos como podemos extraer un concepto formal como el par ($\{ Plymouth\ Satellite, Chevrolet\ Malibu \}, \{ Consumo, Cilindrada, Movilidad, Aceleración \}$) a partir del contexto formal.

Cuadro 2.3: Conceptos formales a partir de la representación del contexto formal.

Modelo de coche	Consumo	Cilindrada	Movilidad	Potencia	Peso	Aceleración
Chevrolet Vega				✓		✓
Toyota Corolla				✓		
Ford Pinto			✓		✓	
Volkswagen Beetle	✓		✓	✓	✓	✓
Plymouth Satellite	✓	✓	✓			✓
Chevrolet Malibu	✓	✓	✓			✓

Otra alternativa aparece en la representación como grafo bipartito, donde un concepto formal aparece como un subgrafo bipartito completo, es decir, aquel que tiene todas las aristas posibles. Las conexiones que se establecen en el retículo nos indican que el objeto X tiene un atributo Y si y sólo si existe un camino estrictamente creciente o estrictamente decreciente que permite ir de X a Y en el retículo.

Por otro lado, otro concepto equivalente a las conexiones de Galois y los operadores de cierre en el marco de FCA es el denominado sistema de implicaciones [38]. Como ya se adelantó en el Capítulo 1, las implicaciones constituyen el eje fundamental de trabajo en esta Tesis doctoral pues serán el mecanismo que nos permita abordar el problema del tratamiento eficiente de la información por medio de la lógica. Reflejan una forma relativamente sencilla de trasladar las relaciones lógicas que encontramos entre los atributos que poseen los objetos de nuestros datasets ya que combinan una forma muy simple y natural de escribir reglas del tipo *si-entonces* con una gestión muy eficiente y automatizada de la información.

Un ejemplo básico del uso de implicaciones para reflejar información contenida en los datos podemos verlo utilizando la información de la Tabla 2.1. En ese ejemplo vemos que en ese dataset hay una relación que siempre se cumple y es la siguiente. Si el hotel provee servicio de *aire acondicionado* (*AC*) entonces tiene servicio de *Parking*. He ahí un claro ejemplo de regla *si-entonces*, o en nuestro caso, de implicación, que se verifica en el contexto.

Este concepto de implicación es el que guía este trabajo otorgándonos un potencial para automatizar técnicas de razonamiento, con el beneficio que eso conlleva. Además, las implicaciones son elemento fundamental para

el desarrollo de formas más eficientes que el retículo de conceptos de representación del conocimiento.

Por tanto, a partir de las implicaciones y de las propiedades básicas que verifican, podemos definir un cálculo lógico que nos permita realizar sistemas de deducción completos sobre el contexto actual. En cierto sentido, podríamos decir que hemos pasado de tener un conocimiento basado en ejemplos a disponer de un conocimiento abstracto que introduce sistemas de razonamiento más elaborados, partiendo únicamente de las observaciones concretas que hemos realizado, es decir, hemos aprendido reglas generales a partir de ejemplos.

Para finalizar, podemos considerar a las implicaciones, las conexiones de Galois y los operadores de cierre, como distintas alternativas de tratar la información que se puede extraer de un contexto formal. No obstante, las conexiones de Galois y los retículos de conceptos permiten una representación gráfica, mientras que los operadores de cierre y los sistemas de implicaciones nos facilitan el razonamiento por medio de la lógica.

En definitiva, todas estas nociones y propiedades son los fundamentos sobre los cuales se consolida FCA. Recordamos que la motivación original de FCA es la reestructuración de esta teoría utilizando la teoría matemática de conceptos y jerarquías con la intención de acercarse al raciocinio humano y facilitar las conexiones con las Lógicas Filosóficas del pensamiento [112,113].

2.2 Lógica de Simplificación

Introducimos la lógica SL_{FD} [24] describiendo sus cuatro pilares fundamentales: su lenguaje, su semántica, su sistema axiomático y su método de razonamiento automático.

Lenguaje

Definición 2.2.1. *Dado un conjunto M finito de símbolos (denominados atributos) no vacío, el lenguaje sobre M se define como:*

$$\mathcal{L}_M := \{A \rightarrow B \mid A, B \subseteq M\}$$

Las fórmulas $A \rightarrow B$ se denominan *implicaciones* y los conjuntos A y B reciben el nombre de *premisa* y *conclusión* de la implicación respectivamente. Los conjuntos $\sum \subseteq \mathcal{L}_M$ se denominan *sistemas de implicaciones* sobre M .

Para simplificar la notación:

- Usaremos letras minúsculas para denotar los elementos en M , mientras que las mayúsculas denotan sus subconjuntos.
- Omitimos las llaves en premisas y conclusiones, es decir, $abcde$ denota el conjunto $\{a, b, c, d, e\}$.
- Escribimos sus elementos por yuxtaposición, es decir, para $X \cup Y$ escribiremos XY .
- Para la diferencia, cambiamos $X - Y$ por $X \setminus Y$.

Definición 2.2.2 (Implicación). *Una implicación sobre un conjuntos de atributos M es una expresión de la forma $X \rightarrow Y$, donde X e Y son conjuntos de atributos, i.e. $X \subseteq M$ y $Y \subseteq M$.*

Definición 2.2.3 (Implicación trivial). *Una implicación $X \rightarrow Y$ es trivial si $Y \subseteq X$.*

Definición 2.2.4 (Implicación unitaria). *Sea X un conjunto de atributos y a un atributo del conjunto, $a \in X$, una implicación se dice unitaria si su conclusión es un conjunto unitario, i.e. $X \rightarrow a$.*

Ejemplo 2.2.5. *Sea $M = \{a, b, c, d\}$ un conjunto de atributos. Se cumple que $\{a \rightarrow b, a \rightarrow c\} \equiv \{a \rightarrow bc\}$ porque para cada contexto formal $\mathbf{K} := (G, M, I)$, se verifica que $\{b, c\}' = \{b\}' \cup \{c\}'$. Por tanto, $\{a\}' \subseteq \{b\}'$ y $\{a\}' \subseteq \{c\}'$ si y sólo si $\{a\}' \subseteq \{b, c\}'$. Este ejemplo nos permite asegurar que cualquier sistema de implicaciones es equivalente a un sistema de implicaciones unitarias.*

Proposición 2.2.6. *Sea M un conjunto de atributos y $\Sigma \subseteq \mathcal{L}_M$. Se verifica que:*

$$\Sigma \equiv \bigcup_{A \rightarrow B \in \Sigma} \{A \rightarrow b \mid b \in B\}$$

Semántica

Tras la definición del lenguaje pasamos ahora a introducir la semántica utilizada, para lo cual utilizamos el concepto de *operador de cierre* que reescribimos a continuación.

Un operador de cierre en M es una función $c: 2^M \rightarrow 2^M$ que es *extensivo* ($X \subseteq c(X) \forall X \subseteq M$), *monótono* ($X_1 \subseteq X_2$ implica que $c(X_1) \subseteq c(X_2) \forall X_1, X_2 \subseteq M$), e *idempotente* ($c(c(X)) = c(X) \quad \forall X \subseteq M$). Los puntos fijos para c (i.e. $X \subseteq M$ tales que $c(X) = X$) se denominan *conjuntos cerrados*.

Un operador de cierre c en M recibe el nombre de *modelo* para una implicación $A \rightarrow B \in \mathcal{L}_M$ si $B \subseteq c(A)$; se denota $c \models A \rightarrow B$. Además, dado un sistema de implicaciones $\Sigma \subseteq \mathcal{L}_M$ y un operador de cierre c en M , $c \models \Sigma$ denota que $c \models A \rightarrow B$ para todo $A \rightarrow B \in \Sigma$.

Decimos que $A \rightarrow B$ se *deriva semánticamente* de Σ , denotado $\Sigma \models A \rightarrow B$, si cualquier modelo para Σ también es un modelo para $A \rightarrow B$, i.e. $c \models \Sigma$ implica que $c \models A \rightarrow B$ para todo operador de cierre c en M .

Sistema Axiomático

Los conocidos Axiomas de Armstrong es el primer sistema descrito para tratar sistemas de implicaciones utilizando la lógica. Tiene su origen en [3] donde se utilizó para estudiar las propiedades de las dependencias funcionales en el modelo relacional de Codd [20]. Este sistema ha tenido una clara influencia en el diseño de varias lógicas sobre implicaciones, todas ellas construidas alrededor del paradigma de la transitividad. Con el paso de los años, otros trabajos han propuesto sistemas axiomáticos equivalentes [4, 49, 80].

Estos axiomas son válidos al generar sólo implicaciones dentro del cierre del conjunto de implicaciones (denotado como F^+) cuando se actúa sobre el conjunto F . También son completos ya que la aplicación repetitiva de las reglas generarán todas las implicaciones en el conjunto cerrado F^+ .

Este sistema axiomático está formado por un axioma, denominado *Reflexivo*, y dos reglas de inferencia, *Aumentativa* y *Transitiva* que se definen

a continuación

Sean A, B, C subconjuntos de atributos de M . Entonces, tenemos:

$$\text{[Ref]} \quad \frac{}{AB \rightarrow A} \quad \text{[Aug]} \quad \frac{A \rightarrow B \cup C}{AC \rightarrow BC} \quad \text{[Tran]} \quad \frac{A \rightarrow B, B \rightarrow C}{A \rightarrow C}$$

Si bien es cierto que el sistema de Armstrong es un trabajo que acumula innumerables citas en diferentes trabajos a lo largo de los años, su utilidad práctica se ve mermada debido a la dificultad que conlleva a la hora de plasmar las demostraciones. Es por ello que su uso está enfocado al estudio teórico de las implicaciones en vez de al desarrollo de nuevos algoritmos. No obstante, como se ha mencionado, es un buen punto de partida para el desarrollo de nuevas lógicas para el tratamiento de las implicaciones.

En este punto es donde aparece la $SLFD$ [74], que constituye un marco mucho más adecuado en términos de la automatización del razonamiento utilizando implicaciones. De hecho, $SLFD$ será el núcleo fundamental de los métodos y las aplicaciones que se han llevado a cabo a lo largo de esta tesis doctoral.

$SLFD$ constituye una lógica válida y completa [24] en la que el principal elemento del lenguaje son las implicaciones. Se define formalmente como:

Definición 2.2.7. *Sea M un conjunto finito, la formulación de $SLFD$ son expresiones, denominadas implicaciones, de la forma $X \rightarrow Y$, donde X e Y son subconjuntos de M .*

Las implicaciones se interpretan de forma conjuntiva, es decir, corresponden a las fórmulas $a_1 \wedge \dots \wedge a_n \rightarrow b_1 \wedge \dots \wedge b_m$ donde las proposiciones $a_1, \dots, a_n, b_1, \dots, b_m$ son elementos del conjunto M . La interpretación es la siguiente:

Definición 2.2.8. *Sea O y M dos conjuntos finitos, que representan objetos y atributos respectivamente, y I una relación en $O \times M$. Una implicación de $SLFD$, $X \rightarrow Y$, donde X y Y son subconjuntos de M , es válida en I si y sólo si*

$$\{o \in O \mid (o, x_i) \in I \ \forall x_i \in X\} \subseteq \{o \in O \mid (o, y_j) \in I \ \forall j_i \in Y\}$$

Además de su forma natural de expresar el conocimiento como regla, las implicaciones proporcionan un sistema de razonamiento lógico y de inferencia. Su tratamiento simbólico se propuso como hemos mencionado anteriormente en [3], sin embargo, debido al rol central que desempeña la transitividad en ese sistema axiomático, el desarrollo de métodos ejecutables para resolver problemas de implicaciones se basa en métodos indirectos. SL_{FD} evita el uso de la transitividad y se guía por la idea de simplificar el conjunto de implicaciones mediante la eliminación atributos redundantes de manera eficiente [73]. Por consiguiente, la introducción de SL_{FD} abrió la puerta al desarrollo de métodos de razonamiento automatizados directamente basados en su novedoso sistema axiomático [25, 27] que resumimos a continuación.

SL_{FD} se define como el par (L_{FD}, S_{FD}) donde S_{FD} tiene el siguiente esquema axiomático:

$$\text{[Ref]} \quad \overline{A \cup B \rightarrow A}$$

junto con las siguientes reglas de inferencia, denominadas *fragmentación*, *composición* y *simplificación* respectivamente.

$$\text{[Frag]} \quad \frac{A \rightarrow B \cup C}{A \rightarrow B} \quad \text{[Comp]} \quad \frac{A \rightarrow B, C \rightarrow D}{A \cup C \rightarrow B \cup D} \quad \text{[Simp]} \quad \frac{A \rightarrow B, C \rightarrow D}{A \cup (C \setminus B) \rightarrow D}$$

Remarcamos que el lenguaje SL_{FD} considera como fórmulas válidas aquellas en donde cualquiera de sus dos partes puede ser el conjunto vacío, denotado $A \rightarrow \emptyset$ y $\emptyset \rightarrow A$. Sus significados fueron discutidos en [24]. En ese trabajo, también presentamos el siguiente resultado donde la derivación de una implicación $A \rightarrow B$ se reduce a la derivación de la fórmula $\emptyset \rightarrow B$ teniendo $\emptyset \rightarrow A$. Este resultado se usará más adelante en el diseño de nuestro novedoso método de cierre.

Proposición 2.2.9. *Para cualquier Γ y $\forall X, Y \subseteq M$, $\Gamma \vdash X \rightarrow Y$ si y sólo si $\Gamma \cup \{\top \rightarrow X\} \vdash \top \rightarrow Y$*

Definición 2.2.10. *Se dice que una implicación $A \rightarrow B$ se deriva sintácticamente de un sistema de implicaciones Σ , y se denota por $\Sigma \vdash A \rightarrow B$, si*

existe una secuencia de implicaciones $\sigma_1, \dots, \sigma_n \in \mathcal{L}_M$ tal que $\sigma_n = (A \rightarrow B)$ y, para todo $1 \leq i \leq n$, la implicación σ_i satisface una de las siguientes condiciones:

- σ_i es un axioma, es decir, verifica el esquema [Ref].
- $\sigma_i \in \Sigma$.
- σ_i se obtiene a partir de implicaciones pertenecientes a $\{\sigma_j \mid 1 \leq j < i\}$ aplicando las reglas de inferencia [Frag], [Comp] o [Simp].

La secuencia $\sigma_1, \dots, \sigma_n$ constituye una *demonstración* para $\Sigma \vdash A \rightarrow B$.

La derivación sintáctica proporciona una gestión automatizada de las implicaciones. En particular, se puede usar para resolver el denominado problema de las implicaciones: dado un conjunto de implicaciones Γ y una implicación $A \rightarrow B$, queremos responder si $A \rightarrow B$ se deduce de Γ . Este problema se puede abordar utilizando el operador de cierre.

El sistema axiomático es válido y completo, es decir, se cumple que la derivación sintáctica y semántica coinciden. Esto significa que cada regla que se puede deducir con este sistema puede derivarse semánticamente (el sistema axiomático es válido) y viceversa (el sistema axiomático es completo).

Teorema 2.2.11. *Sea M un conjunto finito no vacío de atributos, $\Sigma \subseteq \mathcal{L}_M$ y $A \rightarrow B \in \mathcal{L}_M$. Entonces, $\Sigma \models A \rightarrow B$ si y sólo si $\Sigma \vdash A \rightarrow B$.*

La derivación sintáctica nos conduce a un nuevo operador de cierre denominado *cierre sintáctico*.

Definición 2.2.12. *Dado un conjunto de implicaciones $\Sigma \subseteq \mathcal{L}_M$, un conjunto $X \subseteq M$ se dice cerrado respecto de Σ si $A \subseteq X$ implica $B \subseteq X$ para toda $A \rightarrow B \in \Sigma$. Debido a que M es cerrado con respecto a Σ y cualquier intersección de conjuntos cerrados es cerrada, podemos definir el siguiente operador de cierre:*

$$(\)_\Sigma^+ : 2^M \rightarrow 2^M \quad X_\Sigma^+ = \bigcap \{Y \subseteq M \mid Y \text{ es cerrado con respecto a } \Sigma \text{ y } X \subseteq Y\}$$

El siguiente teorema es esencial para calcular cierres y a su vez también para introducir un método de razonamiento automático.

Teorema 2.2.13 (Teorema de deducción). *Sea $A \rightarrow B \in \mathcal{L}_M$ y $\Sigma \subseteq \mathcal{L}_M$. Entonces,*

$$\Sigma \vdash A \rightarrow B \quad \text{si} \quad B \subseteq A_\Sigma^+ \quad \text{si} \quad \{\emptyset \rightarrow A\} \cup \Sigma \vdash \{\emptyset \rightarrow B\}$$

Corolario 2.2.14. *Sea $\Sigma \subseteq \mathcal{L}_M$. $\forall X \subseteq M$, se tiene que*

$$X_\Sigma^+ = \max\{Y \subseteq M \mid \Sigma \vdash X \rightarrow Y\}.$$

La principal ventaja de SL_{FD} radica en que las reglas de inferencia pueden considerarse reglas de equivalencia. Como consecuencia, SL_{FD} se ha podido utilizar como núcleo principal para el desarrollo de métodos automáticos para diversas aplicaciones (e.g. obtener claves minimales, cálculos del cierre) como veremos más adelante. Un estudio más detallado al respecto, incluyendo teoremas y demostraciones, puede verse en [73].

Método de Razonamiento Automático

El problema de las implicaciones se ha abordado tradicionalmente utilizando un método básico que recibe $A \subseteq M$ como entrada y utiliza de forma exhaustiva la relación de subconjuntos recorriendo iterativamente Γ y agregando nuevos elementos al cierre. Este método, propuesto en la década de 1970 [66] y puede considerarse la base principal donde se han sustentado tantos otros. Podemos verlo detallado en el Algoritmo 2.1.

Más tarde, varios autores han desarrollado otros métodos mediante el uso de diferentes técnicas, resolviendo eficientemente este problema en tiempo lineal. En [81], los autores muestran que la complejidad del problema de cierre es $\Theta(|\mathcal{A}| |\Gamma|)$. En [73], presentamos un método de cierre de atributos estrechamente relacionado con el sistema axiomático SL_{FD} . También demostramos que nuestro método tiene un mejor rendimiento que aquellos basados en el cierre clásico.

Por nuestra parte, el método de razonamiento automático en SL_{FD} se basa en el Teorema de Deducción y un conjunto de equivalencias. Siguiendo

Algorithm 2.1: Cierre clásico

```

Data:  $\Gamma, A$ 
Result:  $A_\Gamma^+$ 
begin
   $A_\Gamma^+ := A$ 
  repeat
     $A' := A_\Gamma^+$ 
    foreach  $X \rightarrow Y \in \Gamma$  do
      if  $X \subseteq A_\Gamma^+$  and  $Y \not\subseteq A_\Gamma^+$  then
         $A_\Gamma^+ := A_\Gamma^+ \cup \{Y\}$ 
    until  $A_\Gamma^+ = A'$ ;
  return  $A_\Gamma^+$ 

```

la forma habitual, dos sistemas de implicaciones $\Sigma_1, \Sigma_2 \subseteq \mathcal{L}_M$ se dicen equivalentes si sus modelos coinciden, es decir, se cumple que para todo operador de cierre en M , $c \models \Sigma_1$ si y sólo si $c \models \Sigma_2$.

La siguiente proposición proporciona las tres equivalencias, denominadas también: *Fragmentación, Composición y Simplificación*, que se aplican y justifican el nombre de la lógica SL_{FD} porque, si las leemos de izquierda a derecha, eliminan la información redundante en el sistema de implicaciones (cf. [73]).

Proposición 2.2.15. *Sean $A, B, C, D \subseteq M$. Se verifican las siguientes equivalencias:*

- (i) $\{A \rightarrow B\} \equiv \{A \rightarrow B \setminus A\}$
- (ii) $\{A \rightarrow B, A \rightarrow C\} \equiv \{A \rightarrow B \cup C\}$
- (iii) $A \cap B = \emptyset$ y $A \subseteq C$ implica $\{A \rightarrow B, C \rightarrow D\} \equiv \{A \rightarrow B, C \setminus B \rightarrow D \setminus B\}$

Basándonos en el Teorema de Deducción y las equivalencias anteriores, en [73], los autores presentan un novedoso algoritmo para calcular cierres

utilizando $SLFD$, denominado **Cl_s** que actúa según el siguiente procedimiento.

Dado un conjunto $A \subseteq M$ y un conjunto de implicaciones Σ , **Cl_s** calcula el par (A_Σ^+, Γ) , siendo A_Σ^+ el cierre sintáctico de A con respecto a Σ y Γ el conjunto residual de implicaciones respecto a $\emptyset \rightarrow A_\Sigma^+$. Esto permite determinar si una implicación $A \rightarrow B$ puede deducirse de Σ .

Los pasos del algoritmo desglosados en lenguaje natural y de forma más detallada son:

- En primer lugar, se incluye la fórmula $\emptyset \rightarrow A$ en Σ y se usa como semilla por el método de razonamiento mediante las equivalencias mencionadas en la proposición anterior.
- A continuación, el algoritmo entra en un proceso iterativo en el cual se irán aplicando las siguientes equivalencias mientras no se alcance la condición de parada.
 - **Eq. I:** Si $B \subseteq A$ entonces $\{\emptyset \rightarrow A, B \rightarrow C\} \equiv \{\emptyset \rightarrow A \cup C\}$.
 - **Eq. II:** Si $C \subseteq A$ entonces $\{\emptyset \rightarrow A, B \rightarrow C\} \equiv \{\emptyset \rightarrow A\}$.
 - **Eq. III:** En otro caso $\{\emptyset \rightarrow A, B \rightarrow C\} \equiv \{\emptyset \rightarrow A, B \setminus A \rightarrow C \setminus A\}$.
- En el momento en el que no sea posible aplicar ninguna de las equivalencias anteriores, el algoritmo termina dando como resultado el conjunto cierre A_Σ^+ , es decir, el cierre sintáctico de A con respecto a Σ , y además, el conjunto residual de implicaciones Γ .

Formalmente, el algoritmo **Cl_s** puede verse en la Función Cls.

Aunque es cierto que existen en la literatura numerosas propuestas de algoritmos para calcular el cierre (la mayoría de ellas como modificaciones del cierre clásico de Maier [66]), la principal novedad y ventaja que aporta **Cl_s** es que, de manera simultánea, también reducimos el número de implicaciones, guardando el conocimiento complementario que describe la información que no pertenece al cierre. Este hecho nos coloca sin lugar a dudas en una posición privilegiada, ya que nos evita el elevado coste de

Function $\text{Cls}(A, \Sigma)$

input : A , conjunto de atributos sobre los que se quiere calcular el cierre;
 Σ , un sistema de implicaciones;
output: A_Σ^+ , cierre sintáctico de A con respecto a Σ ;
 Γ , el sistema de implicaciones residual;

repeat

$\Gamma := \Sigma$	
$\Sigma := \emptyset$ foreach $B \rightarrow C \in \Gamma$ do	
if $B \subseteq A$ then	
$A := A \cup B$	
else if $C \not\subseteq A$ then	
$\Sigma := \Sigma \cup \{B \setminus A \rightarrow C \setminus A\}$	

until $\Gamma = \Sigma$

return (A, Γ)

minería de datos para extraer el nuevo conjunto de implicaciones para el conjunto de datos reducido después de cada aplicación del método, algo imprescindible con las implementaciones clásicas. En consecuencia, el proceso supera los costos de minería de datos preservando una complejidad lineal a lo largo del proceso.

Finalizamos este apartado mostrando un ejemplo de aplicación del algoritmo del cierre propuesto.

Ejemplo 2.2.16. Sean $\Sigma = \{a \rightarrow c, bc \rightarrow d, c \rightarrow ae, d \rightarrow e\}$ y $A = \{c, e\}$. El algoritmo **Cls** devuelve $A^+ = \{a, c, e\}$. La siguiente tabla muestra la traza de ejecución paso a paso del algoritmo.

<i>Guide</i>	Σ			
$\emptyset \rightarrow ce$	$a \rightarrow c$	$bc \rightarrow d$	$c \rightarrow ae$	$d \rightarrow e$
$\emptyset \rightarrow ce$	$a \dashv$	$b \not\rightarrow d$	$\not\rightarrow a \not\rightarrow$	$d \dashv$
$\emptyset \rightarrow ace$		$b \rightarrow d$		

Una vez finalizado la ejecución del algoritmo, obtenemos: $\text{Cls}(\{c, e\}, \{a \rightarrow c, bc \rightarrow d, c \rightarrow ae, d \rightarrow e\}) = (\{a, c, e\}, \{b \rightarrow d\})$, donde el cierre es

$\{c, e\}_{\Sigma}^+ = \{a, c, e\}$ y el conjunto residual de implicaciones queda como:
 $\Gamma = \{b \rightarrow d\}$.

Capítulo 3

Claves Minimales

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Reducing the search space by closure and simplification paradigms

A parallel key finding method

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Abstract In this paper, we present an innovative method to solve the minimal keys problem strongly based on the Simplification Logic for Functional Dependencies. This novel method improves previous logic-based methods by reducing, in a significant degree, the size of the search space this problem deals with. Furthermore, the new method has been designed to easily fit within a parallel implementation, thereby increasing the boundaries current methods can reach.

Keywords Minimal keys · Parallelism · Logic

1 Introduction

Finding a set of items/attributes characterizing a greater set of data is a significant problem in several areas: data modeling, data integration, integration of heterogeneous databases, knowledge discovery, anomaly detection, query formulation, query optimization, and indexing. This problem is also interesting in emergent areas as

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linked data. In [14] authors delimit the problem: “establishing semantic links between data items can be really useful, since it allows crawlers, browsers and applications to combine information from different sources”. The proposal is the use of keys to infer identity links among the data distributed in the web.

The notion of key is fundamental in Codd’s relational model of data [2]. Calculating all minimal keys constitutes a tough problem; in [10, 22], authors prove that the number of minimal keys for a relational system can be exponential in the number of attributes, or factorial in the number of dependencies. The main approaches to this problem point to the works of Lucchesi and Osborn in [10] that show an algorithm to calculate all candidate keys. Saiedian and Spencer [15] presented an algorithm using attribute graphs to find all possible keys of a relational database schema, and Wastl [20] proposed an inference system for deriving all keys which can be inferred from a set of functional dependencies. Recent works about calculating minimal keys are [16, 21] and a very modern paper approaching the problem in a logic style [5]. In addition, in [9, 18, 19] the authors proposed the use of formal concept analysis [6] to face problems related with the discovery and management of implications, which can be considered complementary to our work.

We highlight [20] to be the one closest to our framework. In that work, a tableaux method to calculate all minimal keys is presented. In [3] we introduce the Key Algorithm, a method inspired in Wastl’s tableaux method where the Simplification Logic for Functional Dependencies [12] is used to find all minimal keys. Later, in [5] we introduce the SST method, achieving a great performance in comparison to its predecessors based on the introduction of the inclusion-minimality test which avoids the opening of extra-branches of the tree, and the search space becomes narrower.

A parallel algorithm for the manipulation of implications is described in [17], enclosed in the field of hyper-graphs. Parallel issues in similar problems have been addressed by several authors. For instance, Krajca et al. [8] present a parallel algorithm for computing formal concepts. A first view for the parallelization of Wastl’s method and the Key Algorithm was presented in [1], showing how parallelism could naturally be integrated in tableaux methods.

This paper makes the following contributions. We propose a new method named Closure Keys incorporating an efficient prune mechanism that use the closure method based on \mathbf{SL}_{FD} to improve SST method. The new method is based on the strong relation between the notion of key and the closure operator, not only in the definition issue but also in its construction. The closure operator defined in [12] allows us to highly reduce the search space by introducing shortcuts in the way to the leaves, where keys are finally produced. Moreover, a parallel implementation of SST method is presented along with several experiments to confirm the improvements that have been accomplished.

The remainder of the paper is organized as follows: Sect. 2 presents the key finding problem and the state of the art. In Sect. 3, we explain how a closure operator could help us improving the efficiency over minimal keys methods. In Sect. 4, the main aspects of the implementation of these methods will be exposed. Several results are shown in Sect. 5 with a collection of tables and charts. Finally, conclusions will end up the document.

2 Background

In this section we introduce the minimal key problem, the basic notions related with this issue and the last logic-based algorithm in the literature to tackle this problem.

Definition 1 (*Functional dependency*) Let Ω be a set of attributes. A functional dependency (FD) over Ω is an expression of the form $X \rightarrow Y$, where $X, Y \subseteq \Omega$. It is satisfied in a table R if for every two tuples of R , if they agree on X , then they agree on Y .

A key of a relational table is an attribute subset that allows us to uniquely characterize each row, and it is defined by means of FDs as follows:

Definition 2 (*Key*) Given a table R over the set of attributes Ω , we say that K is a key in R if the functional dependency $K \rightarrow \Omega$ holds in R .

We would like to manage keys with no superfluous attributes, named minimal keys. Due to space limitation, we refer those readers non-familiar with the formal notions of FDs, keys and relational tables to [11]. In addition, it is remarkable that this classical problem appears in several areas. For instance, in [5], we show how the minimal key problem in databases has an analogous one in formal concept analysis, where the role of FDs is played by attribute implications. In that paper, minimal key problem was presented from a logical point of view. An axiomatic system to manage both FDs and implications, named Simplification Logic and denoted \mathbf{SL}_{FD} , was introduced in [4]. The inference system of such logic is presented as follows:

Definition 3 (*Axiomatic system*) \mathbf{SL}_{FD} has a reflexivity axiom $\frac{B \subseteq A}{A \rightarrow B}$; and inference rules named fragmentation, composition and simplification.

$$\text{[Frag]} \quad \frac{A \rightarrow BC}{A \rightarrow B}; \quad \text{[Comp]} \quad \frac{A \rightarrow B, C \rightarrow D}{AC \rightarrow BD}; \quad \text{[Simp]} \quad \text{If } A \subseteq C, A \cap B = \emptyset, \frac{A \rightarrow B, C \rightarrow D}{C - B \rightarrow D - B}$$

This logic allows the development of efficient methods to manage FDs and implications. In [5] a new algorithm, named SST, for computing all minimal keys using a tableaux-like strategy was introduced, opening the door to embed a massive parallelism into its execution.

SST relies on the notion of set closure, a basic notion in database theory which allows to characterize the maximum attribute set that can be reached from a given attribute set A with respect to a set of FDs using the axiomatic system. Thus, if the closure of A is denoted as A_{Γ}^+ , the inference system for FDs allows us to infer the FD $A \rightarrow A_{\Gamma}^+$. The logical-style approach to the minimal key problem consists in the enumeration of all attribute sets A such that the following FD holds $A \rightarrow \Omega$.

SST algorithm is strongly based on two proper extensions of the \mathbf{SL}_{FD} simplification rule, named [sSimp] and [lSimp]. These rules guide the construction of the search space to look for all minimal keys (see [5] for further details). The method works step by step by building a tree from the original problem to the solution, i.e. the set of minimal keys. They are applied to each pair (attribute set, implication set) labeling each node to produce new subnodes that will be the origin of new branches in the search space. Applying [lSimp] to the subset of attributes in each node and the implication

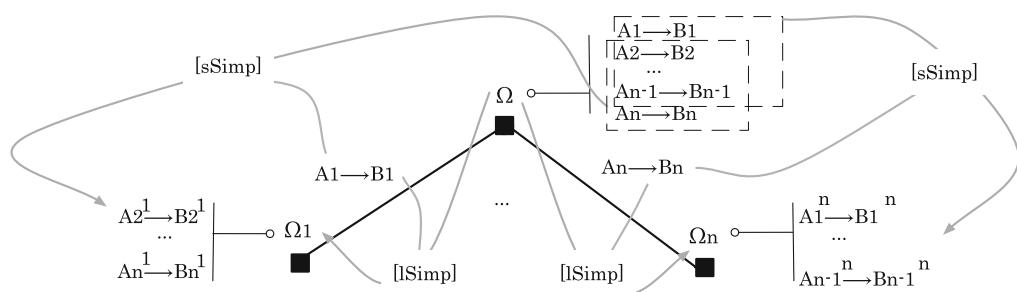


Fig. 1 [lSimp] generates new implication subsets and [sSimp] generates new roots

in the corresponding edge, we obtain the new root in the branch (e.g. in Fig. 1 from Ω and $A_1 \rightarrow B_1$, we obtain the new subset Ω_1). Moreover, the application of [sSimp] over the implication set in each node is done by taking every implication as a pivot and applying the rule to the rest of the implications with the corresponding pivot to generate new branches (e.g. in Fig. 1 the first branch is generated by taking $A_1 \rightarrow B_1$ as the pivot and facing it to the rest of the implication set $A_2 \rightarrow B_2, \dots, A_n \rightarrow B_n$).

When the set of implications in a node is the empty set, we have reached a leaf of the tree and a minimal key is added to the solution. The full situation is depicted in Fig. 1, where a schema of the application of both rules to a generic node is shown.

SST shows a great performance compared with its predecessors. The main benefit in the reduction of the search space was the introduction of the inclusion-minimality test to avoid the opening of extra-branches. Thus, SST does not open some branches which are going to produce the same keys that are calculated in another branch. The characterization of such branches rendering leaves with duplicate information is not a trivial issue. To approach it, we have defined the notion of minimal implication w.r.t. an implication set Γ : $A \rightarrow B \in \Gamma$ is minimal if for all $C \rightarrow D \in \Gamma$ we have that $C \not\subseteq A$.

3 Improving minimal keys methods by means of closure operator

As it has been established this far, the enumeration of all minimal keys is a non-trivial problem that can be approached using logic-based methods. The main goal to progress in this line is the reduction of the search space that can be measured by means of the number of nodes in the tree. In the previous section, we have summarized the main characteristics of the SST method and how it incorporates a strategy to shorten the width of the tree corresponding to the search space. Thus, it provides a narrower tree regarding previous methods. In this spirit, we have studied how to reduce even more the size of the tree by shorting its depth. The kernel of the method presented here is a logic-based closure algorithm presented in [12], strongly based on Simplification Logic \mathbf{SL}_{FD} . The closure of an attribute set can be solved in linear time w.r.t. the cardinality of the implication set. There exists in the literature several approaches to tackle this problem, but most of them may be considered a modification of classical Maier's method [11].

The method presented in [12], named \mathbf{SL}_{FD} Closure, can be considered a novel approach to classical closure methods, because it provides a new feature: In addition to the set of attributes that constitutes the output of the closure operator, \mathbf{SL}_{FD} Closure provides a subset of implications of the original Γ set. This new implication subset encloses the knowledge that can be considered as the complement of the closure and, as we shall see, may be used in a very smart way to find minimal keys by means of the closure operator.

As mentioned before, closure operator may be used to define the notion of key. In this section we propose it as the backbone of a new method to compute all minimal keys. In this approach, \mathbf{SL}_{FD} Closure plays an outstanding role. The main point of our method is that it receives a set of implications Γ and a subset of attributes $X \subseteq \Omega$. It renders the closure set X^+ w.r.t. Γ , and also, a novelty output Γ' consisting in a new implication set that covers the semantics that reaches beyond X^+ . If $\Gamma' = \emptyset$ then $X^+ = \Omega$ (see [12] for further details).

We propose to apply \mathbf{SL}_{FD} Closure to each minimal implication in the Γ set in each node to open new branches with the output produced by \mathbf{SL}_{FD} Closure. The definition of our new minimal key method is presented in Algorithm 1. An illustrative example showing the reduction in the tree provided by Closure Keys w.r.t. SST method is shown in Figs. 2 and 3 respectively.

The main call of Algorithm 1 must be $\text{Closure_Keys}(\Omega, \Gamma, \emptyset, \emptyset)$. Later, to get all minimal keys we pick up the minimal elements of the fourth parameter (which acts as an accumulator in a in/out mode) of this procedure call with regard to $(2^\Omega, \subseteq)$. The following theorem ensures that Algorithm 1 provides a sound and complete method.

Theorem 1 Let Ω be an attribute set and Γ be an implicational system over M . Then Algorithm 1 renders all minimal keys.

Proof Firstly, the termination of the algorithm is ensured because, in each recursive call of Closure_Keys , the cardinality of the implicational system is strictly decreased.

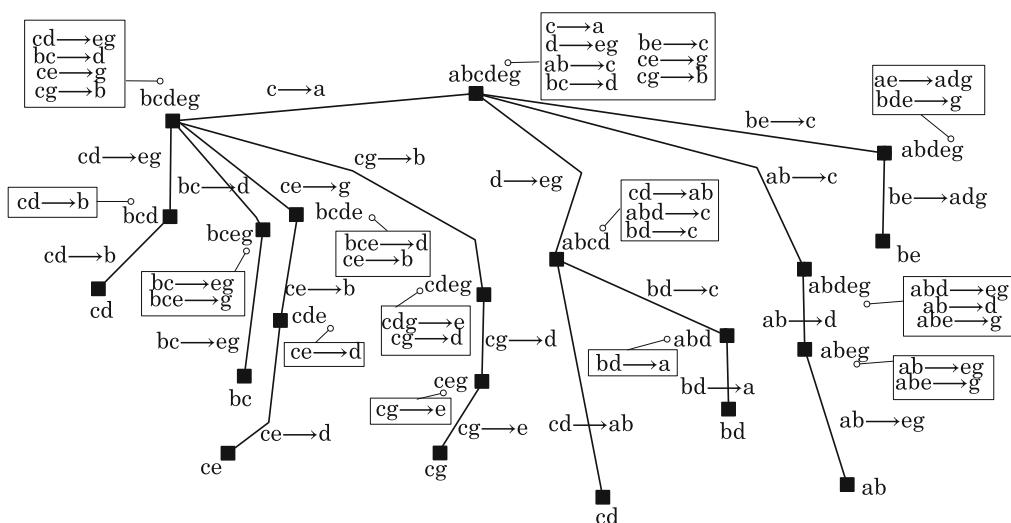


Fig. 2 A complete example using SST method

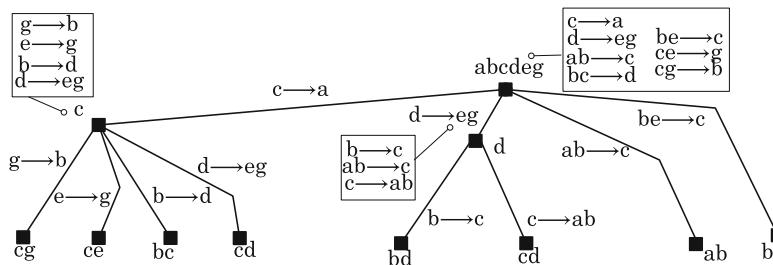


Fig. 3 SL_{FD} Closure Keys Method example. Notice how the treatment of the implication set in each node produces a big reduction of the tree of Fig. 2 from 21 to 11 nodes

Moreover, the number of recursive calls is limited by the cardinality of Γ . The kernel of the method is the SL_{FD} Closure presented in [12] providing a compound output, i.e., $\text{SL}_{\text{FD}} \text{Closure}(X, \Gamma) = \langle X^+, \Gamma' \rangle$.

This method produces all minimal keys because, in each step, it renders a closure X^+ . If it is not the entire Ω , we go ahead by selecting a new antecedent from the implication set Γ' . We remark that this implication set has been depurated by means of SL_{FD} Closure. By virtue of this, once we reached a leaf, the union of the antecedents in the way through conform a key and, therefore, the method is sound.

Concerning its completeness, any key in the system is produced since the search space induced by our method constitutes an exhaustive search. As a conclusion from the key study presented in [13], keys have to be composed with bricks being the antecedent of the implications. Those pruned branches do not fall down the completeness since the removed keys were redundant.

Algorithm 1: Closure_Keys

Data: Ω a set of attributes, Γ a set of implications, \mathcal{C} an accumulator variable of a subset set of attributes, \mathcal{K} a accumulator variable with sets of attributes.

```

begin
     $\mathcal{A} := \Omega$ 
    if  $\Gamma = \emptyset$  then Add( $\mathcal{K}$ ,  $\{\mathcal{A} \cup \mathcal{C}\}$ )
    else
        foreach  $X \rightarrow Y \in [\text{IM}](\Gamma)$  do
             $\langle X', \Gamma' \rangle := \text{SL}_{\text{FD}} \text{Closure}(X, \Gamma)$ 
             $\mathcal{C} := \mathcal{C} \cup X$ 
            Closure_Keys( $X'$ ,  $\Gamma'$ ,  $\mathcal{C}$ ,  $\mathcal{K}$ )
        end
    /* Where  $[\text{IM}](\Gamma)$  denotes the implications in  $\Gamma$  being minimal FD. */

```

4 Implementation

The approach presented determines that, due to the construction of the search tree, parallel strategies and big hardware resources are strongly needed when we try to resolve problems containing substantial size at the input. In order to test the improvement, we are going to proceed with a parallel implementation of SST and Closure_Keys algorithms that works in two steps, (1) a partial method splits the entry problem into several subproblems and (2) a parallel method resolves all these generated subproblems and composes the final result, the minimal key set (see Fig. 4).

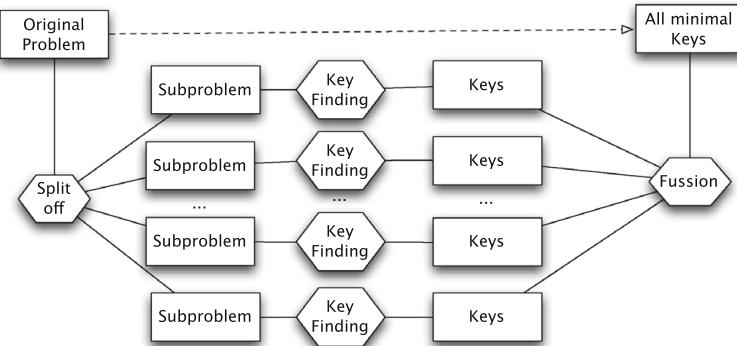


Fig. 4 Parallelism diagram

We can go along this kind of strategy over our parallel implementations because, due to the tableaux nature, each branch will be totally independent to others, so every node can be treated on its own. That is the great point of using parallelism within these techniques; we can deliver a huge number of problems to different cores so they can be solved simultaneously. By virtue of this, the size of the problems at the input could be greater without exceeding the machine limitations; at least, we have distanced the limit in a substantial way for the moment.

The application of the partial code moves us to split the entry problem into several subproblems. At this regard, we will stop at a certain level of the tree, and there, every node constitutes an independent problem by itself. Each of these problems represents a branch of the search tree and will be the input for the application of the parallel method, that will end up in a leaf of the tree. However, it is needed to go forward determining the value that will decide the splitting point, and this is, indeed, not an easy issue so far, as we discuss in the following paragraph.

As break-off value (BOV), determining the level in which the branch is considered an atomic to be treated by the parallel stage, we use the cardinal of the set of implications of the current node, since we have observed that, the bigger this cardinal is, the longer the branch *used to* reach. Once we have all these subproblems, and due to all of them conform an individual state of the algorithm, they are meant to be solved by the parallel code. However, if we are treating with big problems (those containing a substantial number of attributes and implications), the number of generated subproblems could be huge (we have run experiments with 50,000+ subproblems). Thus, handling them is a task only available to a significant amount of resources, as Supercomputing and Bioinnovation Center of the University of Málaga¹ supports us.

5 Experiments and results

We have randomly generated a set of experiments, varying the number of attributes and implications. We start with problems with 100 implications, each one built using 100 different attributes as well. Then, we move forward considering bigger problems

¹ <http://www.scbi.uma.es>.

Table 1 Attempts trying to improve execution times by increasing the number of cores

Problem and method	Implications 150 Partial _t (s)	Attrib 150 Total _t (s)	BOV 140 Nodes	Cores	Ratio
150150-4-SST	581	885	55.211	32	62
150150-4-CK	48	65	25.477	32	391
150150-4-SST	576	880	55.211	64	62
150150-4-CK	46	64	25.477	64	398

counting on 150 implications and 150 attributes. Notice that these numbers go far beyond machine capabilities, as it has been demonstrated in previous studies of this work [3]; they even substantially exceed the results given in [1], where parallelism techniques had already been applied.

Obviously, both methods obtain the same keys. Therefore, we have omitted this parameter in result tables since it only experimentally validates the method. Consequently, in order to compare SST method versus Closure Keys method, we focus on execution times and the number of nodes of the tree (so, we can figure out the size of the problem).

Number of nodes will always be the same for an individual experiment no matter the number of runs we do. However, execution times will not go along exactly with this statement. They could be slightly different due to its intrinsic nature. Differences concerning number of nodes will show how new methods have improved the algorithm as the depth of the tree has been sharply reduced, and consequently, execution times become better. Besides, we have included a last column in order to show the ratio of time and nodes between SST and CK. Regarding all these points, needless to say that every execution time shown here is the fruit of a post-statistical study [7, 23] behind the results obtained from every run, so we kept the most reliable ones.

Before we reach the results section, the hardware architecture that has been used for developing every test shown in the paper can be visited in <http://www.scbi.uma.es/site/scbi/hardware>.

Every experiment we show here have been carried out using a cluster of 32 cores from those available within the hardware architecture mentioned above. Few subsequent experiments increasing the number of cores available went not up to expectations results in terms of execution times, as we briefly show in Table 1. Increasing the number of cores to engage parallelism within these kind of problems is a matter where much still remains to be investigated.

The first experiment starts solving a battery of hard load problems counting on 100 attributes and 100 implications. Results are shown in Table 2.

Even with such a big number of attributes and dependencies, it's been just around 10 min long for the slowest algorithm to finish (problem 100100-3). Yet, less than 300k nodes have been the maximum size of the Tableaux of all of them (problem 100100-7). It is absolutely out of mind thinking about solving these problems using sequential versions of the algorithms, since execution times would go out of hands.

Table 2 Parallel methods applied to big-size problems (I)

Problem and method	Attrib 100 Subp	Implications 100 Partial _t (s)	BOV 90 Total _t (s)	Cores 32 Nodes	Ratio
100100-1-SST	14	0	1	33	33
100100-1-CK	0	0	0	15	15
100100-2-SST	1.354	36	105	25.621	244
100100-2-CK	212	4	15	12.715	847
100100-3-SST	8.602	183	644	192.574	299
100100-3-CK	1.286	37	99	94.255	952
100100-4-SST	400	7	26	1.704	65
100100-4-CK	15	1	2	751	375
100100-5-SST	39	0	2	119	59
100100-5-CK	0	0	1	42	42
100100-6-SST	1.808	37	123	7.856	63
100100-6-CK	115	4	9	3.698	410
100100-7-SST	6.167	182	489	275.429	563
100100-7-CK	1.378	24	90	118.884	1.320
100100-8-SST	5.104	146	415	182.167	438
100100-8-CK	1.014	19	68	81.632	1.200
100100-9-SST	314	11	25	868	34
100100-9-CK	0	1	1	341	341
100100-10-SST	1.130	27	84	12.541	149
100100-10-CK	136	4	10	6.128	612

In fact, these are encouraging results since, earlier experiments developed using older methods [1] couldn't even have imagined to deal with such a huge problems.

This first experiment confirms which it was mentioned above: subproblems and nodes show a better performance by Closure Keys method.

A very remarkable outcome can be appreciated in problems 100100-{1,5,9}. Notice that Closure Keys does not create any subproblems. As this regard, the improvement is twofold: (1) in some cases, with the same BOV, the new method doesn't even need to move on the parallel implementation, partial one is enough to resolve these problems, and (2) to exploit this discovery, at the moment we need to deal with even more complex problems, we can establish a BOV nearer to the root of the tree and then, partial time will be significantly reduced.

Second experiment goes ahead giving another turn of the screw increasing the input of our experiments. We develop a new battery of 10 problems, increasing the number of implications and available attributes up to 150.

As far as these problems become more complex, the better is the improvement achieved by the new method. Executions times and number of nodes have been drastically reduced as shown in Table 3. Actually, we decided to add one more experiment (problem 150150-EXTRA) due to the specific remarkable results it reached.

Table 3 Parallel methods applied to big-size problems (II)

Problem and method	Attrib 150 Subp	Implications 150 Partial _t (s)	BOV 140 Total _t (s)	Cores 32 Nodes	Ratio
150150-1-SST	165	6	14	911	65
150150-1-CK	11	2	3	374	124
150150-2-SST	2.949	229	394	116,517	295
150150-2-CK	347	25	44	54,375	1.235
150150-3-SST	12.968	1.049	1.716	157,947	92
150150-3-CK	822	125	165	68,531	415
150150-4-SST	5.352	581	885	55,211	62
150150-4-CK	344	48	65	25,477	391
150150-5-SST	5.361	211	484	32,377	66
150150-5-CK	168	27	36	12,522	347
150150-6-SST	771	72	155	17,298	111
150150-6-CK	79	7	11	8,110	737
150150-7-SST	9.473	638	1.252	576,912	460
150150-7-CK	1.754	97	187	262,621	1.404
150150-8-SST	5.466	424	857	510,627	595
150150-8-CK	966	57	104	257,267	2.473
150150-9-SST	235	25	45	3,632	80
150150-9-CK	24	3	4	1,726	431
150150-10-SST	3.403	348	555	102,537	184
150150-10-CK	277	31	46	45,962	999
150150-EXTRA-SST	31.401	2,950	30,983	21,404,732	690
150150-EXTRA-CK	8.049	354	1,320	10,614,386	8.041

Several experiments are worth to be discussed separately. As a matter of fact, if we set our sight on problems 150150-{3,7}, number of subproblems generated are much less for Closure Keys method than SST. Difference is not trivial so far, it also guides us directly to a great reduction of partial execution times. Total execution times go along in the same direction. Regarding the size of the tree, the benefits by introducing the closure are pretty striking. Most of cases, the number of nodes becomes near to 50 % lower. Figure 5 would help us appreciating the differences more clearly.

6 Conclusions

We presented here a method to solve the key finding problem. Among the different strategies introduced in the literature, we follow the logic-based line pioneered in [20]. The new method presented here uses the Simplification Logic and introduces an alternative to the SST method presented in [5]. Here, we provide a further reduction of the tree size by shortening its depth, empirically confirmed by means of the number of

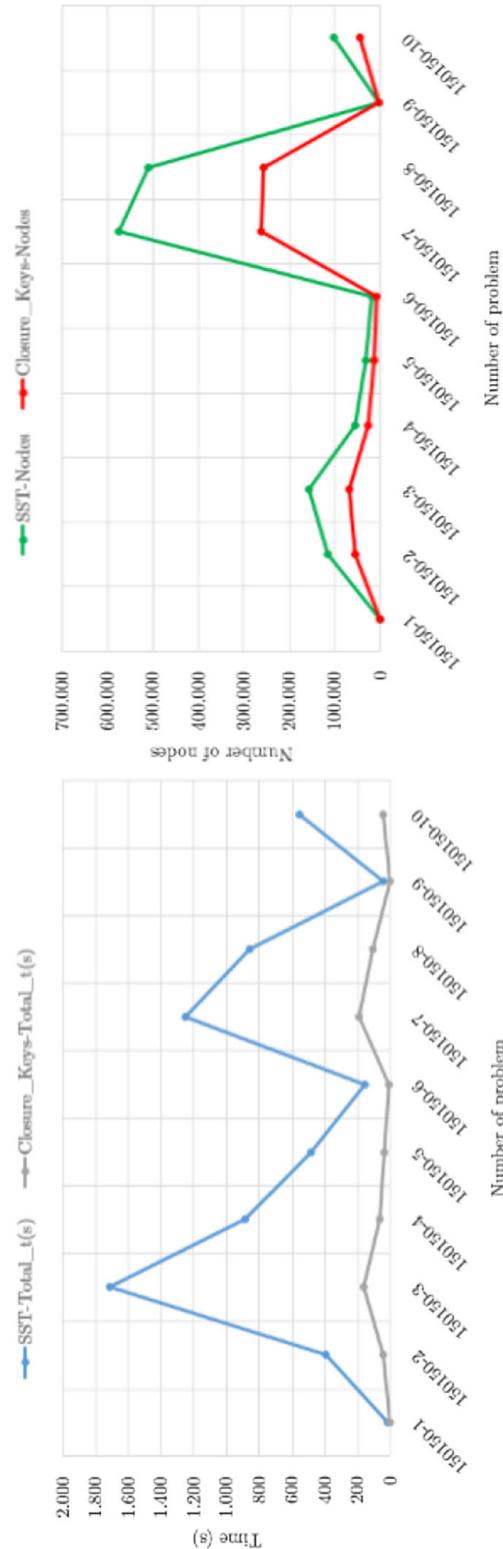


Fig. 5 Execution times and number of nodes for Big-size problems (II)

generated nodes. In addition, a parallel implementation has been developed following a map-reduce architecture. The inherent parallel design of the new method allows us to face problems containing a significant size at the input.

We have set two priorities for future works:

- Applying these mechanisms over real data stores, so we can figure out how they really could take advantage of parallelism working on these kind of datasets.
- The study of the impact concerning the number of cores involved in the experiments.

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Capítulo 4

Generadores Minimales

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Minimal generators, an affordable approach by means of massive computation

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Abstract Closed sets and minimal generators are fundamental elements to build a complete knowledge representation in formal concept analysis. The enumeration of all the closed sets and their minimal generators from a set of rules or implications constitutes a complex problem, drawing an exponential cost. Even for small datasets, such representation can demand an exhaustive management of the information stored as attribute implications. In this work, we tackle this problem by merging two strategies. On the one hand, we design a pruning, strongly based on logic properties, to drastically reduce the search space of the method. On the other hand, we consider a parallelization of the problem leading to a massive computation by means of a map-reduce like paradigm. In this study we have characterized the type of search space reductions suitable for parallelization. Also, we have analyzed different situations to provide an orientation of the resources (number of cores) needed for both the parallel architecture and the size of the problem in the splitting stage to take advantage in the map stage.

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1 Introduction

Knowledge representation and reasoning are the two main pillars of artificial intelligence (AI). The growing interest in AI arisen in several areas beyond computer science is firmly rooted in the recent techniques to extract knowledge from the data and to be efficiently managed. One outstanding framework, based on a strong theoretical basis and also providing executable methods, is the Formal Concept Analysis (FCA) introduced by Ganter and Wille [11]. In FCA, data are stored in a table, representing a binary relation between G (a set of objects) and M (a set of attributes). FCA is not an approximate approach since it pursues to capture, manage and analyze the complete knowledge from the information. This feature has a direct impact on the cost of the methods developed to extract and manage the information.

In this framework, two alternative knowledge representations are used: concept lattice and implications. The second one can be viewed as if-then rules already introduced in other areas, dressed with different clothes. Thus, in relational databases [4] they are named Functional Dependencies and in FCA they are named Implications [11]. All these notions capture the same idea (a pretty intuitive one): when some premise occurs, then a conclusion holds.

Since the size of the concept lattice (in the worst case) is $2^{\min(|G|, |M|)}$, the computational cost of the methods to build it has been considered as a limitation for the application of FCA. Algorithms to infer the concept lattice from the dataset were deeply studied and compared in [15]. Moreover, the extraction of the complete set of implications (basis) from a data set also presents an exponential behavior [5]. In both problems, the density of the dataset plays a major role in the performance of the execution of the method. To improve the computational behavior, in [14] the authors introduce the notion of redundant attributes to avoid the inclusion of such attributes in the extraction of the implications. Recently, in [8] a wide range of parallel methods to solve this second problem has been presented. These works motivate the need to combine some kind of reduction in the search space and a parallel execution to solve these complex problems.

In the two problems already mentioned, the input is the dataset and the output is its knowledge in terms of implications (basis) or attribute closed sets (concept lattice). However, here we deal with a complementary problem that allows to connect both knowledge representations: the enumeration of all the closed sets from a given set of implications. Moreover, we propose a method to produce not only all closed sets but also, for each of them, their canonical representations, named minimal generators.

Minimal generators are interesting not only from a theoretical point of view. In recent works, their computation is the core for solving other problems. The significance of minimal generators is well emphasized in the survey of Poelmans et.al. [21]: “Minimal generators have some nice properties: they have relatively small size and they form an order ideal”. Qu et al. [22] pointed out that decision implications involve to know all minimal generators.

Furthermore, they have been used as a key point to build basis, which constitutes a compact representation of the knowledge allowing a better performance of the reasoning methods based on rules. Missaoui et al. [17, 18] present the use of minimal generators to compute basis involving positive and negative attributes whose premises are minimal generators. All the mentioned authors have considered the dataset as the input of the problem, namely, the minimal generators and closed sets are inferred from the plain data. In [10, 20] some methods for solving this problem have been proposed.

In this work, we consider implications as elements to describe the information and we design a method to enumerate all closed sets and their minimal generators from this information, and not from the dataset. This is a complementary problem to the extraction of minimal generators from the dataset and, as far as we know, no previous work has been developed. In summary, we propose a method to transform the knowledge in terms of implications, into a more organized form so that a complete representation of the closed sets of attributes (and their minimal generators) is obtained. This complete and precise specification allows a further fast management of the semantics of the information contained in the dataset.

Our starting point is [6], where a logic-based method based on SL_{FD} , a sound and complete logic for implications was introduced. In that work, we present the MinGen algorithm which works by traversing the set of implications and applying a set of inference rules, building a search tree space. This shape of the search space limits its execution for medium-sized problems, because of the overwhelming resources of the sequential MinGen algorithm. Although the search space is suitable for a parallel extension of the method, some further research has to be carried out. On the one hand, we first consider the integration of some strategies to reduce the search space. In this work, we present two approaches, MinGenRd and GenMinGen, providing a single-level or a multi-level strategy, respectively. We also show how the MinGenRd is suitable for parallelization whereas GenMinGen is not. On the other hand, we approach a parallelization of the MinGenRd, rendering the MinGenRdPar algorithm. This new method is driven by a like-Map-Reduce strategy.

In addition to the MinGenRdPar method, we also provide a set of experiments in different situations to characterize the resources needed for the parallel execution and to establish a threshold in the splitting stage. In both issues, our goal is to look for an efficient execution of the method. We remark that we have used a random generation of synthetic problems and also real-world problems. Our intention is to show that our approach is valid in both situations.

The paper is organized as follows: after presenting the needed background in Sect. 2, we revisit in Sect. 3 our starting point: a previous logic-based method to enumerate all minimal generators and closed sets (MinGen). Then the reduction strategy by eliminating redundant branches is described in Sect. 4, introducing the MinGenRd and GenMinGen methods. Later, in Sect. 5, we design a parallel version of the MinGenRd algorithm, named MinGenRdPar, running on a massive computation architecture. We provide some stats of its performance together with an illustrative execution of the parallel algorithm over a real (and well-known) problem. We will end up with Conclusions and Future Works. Figure 1 illustrates the content of the paper.

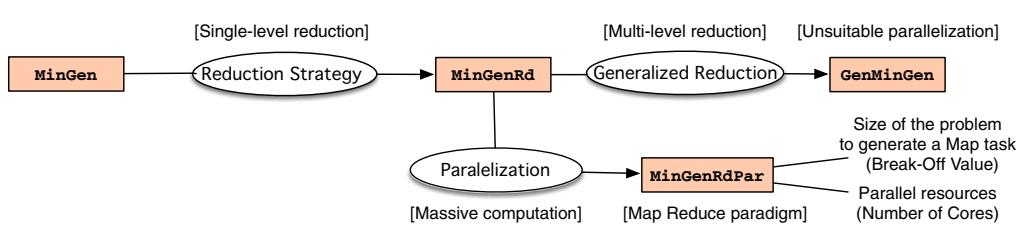


Fig. 1 Illustration of the content of the work

2 Preliminaries

Simplification Logic [7] is introduced by describing its four pillars: its *language*, its *semantics*, its *axiomatic system* and its *automated reasoning method*. Let M be a finite set of symbols (called *attributes*). The language over M is defined as

$$\mathcal{L}_M := \{A \rightarrow B \mid A, B \subseteq M\}$$

Formulas $A \rightarrow B$ are called (attribute) *implications* and the sets A and B are called *premise* and *conclusion* of the implication, respectively. Sets of implications are called *implicational systems*. In order to simplify the notation we omit the brackets in premises and conclusions and write their elements by juxtaposition. Thus, for instance, $ab \rightarrow cde$ denotes $\{a, b\} \rightarrow \{c, d, e\}$.

For introducing the semantics we use the notion of *closure operator*. For a more detailed description of closure operator see [23]. A closure operator on M is a mapping $c: 2^M \rightarrow 2^M$ that is *extensive*, *isotone* and *idempotent*. The fix-points for c are called *closed sets*. Closure operators are strongly connected to knowledge representation in different areas [1, 3, 9, 12, 16].

Since we use the logic as an executable tool, we have to substitute the semantic interpretation and inference for an efficient symbolic management. Thus, we introduce an axiomatic system considering reflexivity as axiom scheme

$$\text{[Ref]} \quad \overline{A \cup B \rightarrow A}$$

together with the following inference rules, called *fragmentation*, *composition* and *simplification*, respectively:

$$\text{[Frag]} \quad \frac{A \rightarrow B \cup C}{A \rightarrow B} \quad \text{[Comp]} \quad \frac{A \rightarrow B, C \rightarrow D}{A \cup C \rightarrow B \cup D} \quad \text{[Simp]} \quad \frac{A \rightarrow B, C \rightarrow D}{A \cup (C \setminus B) \rightarrow D}$$

This axiomatic system is *sound* and *complete* (both semantic and syntactic derivations coincide). This result allows us to design automated reasoning methods. These methods can be approached by using a new closure operator called *syntactic closure*: give an implicational system $\Sigma \subseteq \mathcal{L}_M$, a set $X \subseteq M$ is said to be closed w.r.t Σ if $A \subseteq X$ implies $B \subseteq X$ for all $A \rightarrow B \in \Sigma$. Since M is closed w.r.t Σ and any intersection of closed sets is closed, we can define the following closure operator:

$$(\cdot)_\Sigma^+: 2^M \rightarrow 2^M \quad X_\Sigma^+ = \bigcap \{Y \subseteq M \mid Y \text{ is closed w.r.t } \Sigma \text{ and } X \subseteq Y\}$$

The automated reasoning method in Simplification Logic is based on the Deduction Theorem and three syntactic equivalences which allow us to transform the set of

implications with no semantic loss, i.e., the meaning of the knowledge representation is preserved.

Theorem 1 (Deduction Theorem) *Let $A \rightarrow B \in \mathcal{L}_M$ and $\Sigma \subseteq \mathcal{L}_M$. Then,*

$$\Sigma \vdash A \rightarrow B \text{ iff } B \subseteq A_{\Sigma}^{+} \text{ iff } \{\emptyset \rightarrow A\} \cup \Sigma \vdash \{\emptyset \rightarrow B\}$$

The following corollary characterizes the closed sets built with the syntactic closure as a maximum set enclosing the information of implications.

Corollary 1 *Let $\Sigma \subseteq \mathcal{L}_M$. For all $X \subseteq M$, one has $X_{\Sigma}^{+} = \max\{Y \subseteq M \mid \Sigma \vdash X \rightarrow Y\}$.*

In [19] we present a novel algorithm to compute closures using Simplification Logic. Given a set $A \subseteq M$, to compute A_{Σ}^{+} , the formula $\emptyset \rightarrow A$ is added to Σ and used as a seed by the reasoning method by using the equivalences provided by the previous proposition. Specifically, the algorithm uses the following equivalences:

- **Eq. I:** If $B \subseteq A$ then $\{\emptyset \rightarrow A, B \rightarrow C\} \equiv \{\emptyset \rightarrow A \cup C\}$.
- **Eq. II:** If $C \subseteq A$ then $\{\emptyset \rightarrow A, B \rightarrow C\} \equiv \{\emptyset \rightarrow A\}$.
- **Eq. III:** Otherwise $\{\emptyset \rightarrow A, B \rightarrow C\} \equiv \{\emptyset \rightarrow A, B \setminus A \rightarrow C \setminus A\}$.

One outstanding characteristic of our algorithm, named as Function `Cls`, is that, besides the attribute set corresponding to the closure, it also renders a set of implications corresponding to the complementary knowledge that describes the information which is not within the closure. This new set could be further treated in an iterative process, as our Minimal Generator algorithm will do. An illustrative execution of this function is shown in Example 1.

Example 1 Let $\Sigma = \{a \rightarrow c, bc \rightarrow d, c \rightarrow ae, d \rightarrow e\}$ and $A = \{c, e\}$. The algorithm `Cls` returns $A^{+} = \{a, c, e\}$ and the following table summarizes its trace:

Guide	Σ
$\emptyset \rightarrow ce$	$a \rightarrow c \quad bc \rightarrow d \quad c \rightarrow ae \quad d \rightarrow e$
$\emptyset \rightarrow ce$	$a \rightarrow \emptyset \quad b \not\rightarrow d \quad d \not\rightarrow a \not\rightarrow e \quad d \rightarrow \emptyset$
$\emptyset \rightarrow ace$	$b \rightarrow d$

Therefore, $\text{Cls}(\{c, e\}, \{a \rightarrow c, bc \rightarrow d, c \rightarrow ae, d \rightarrow e\}) = (\{a, c, e\}, \{b \rightarrow d\})$.

An implicational system is said to be *complete for a closure operator* if it captures all the knowledge relating to that operator, that is to say, the implicational system expresses in the language of logic all knowledge relative to it. Therefore, when an implicational system Σ is complete for a closure operator c , its syntactic closure coincides with c , i.e., $X_{\Sigma}^{+} = c(X)$ for all $X \subseteq M$.

To efficiently manage all this knowledge, we characterize it with all the closed sets and their minimal generators. That is, given a set of implications Σ , closed sets characterize attribute sets with maximal meaning and, looking for a better integration in applications, in addition to the closed sets it would be very appreciated to provide a

compact representation of these closed sets. In other words, enumerate for each closed set those subsets that generate them. The following definition comes to formalize these kind of subsets.

Definition 1 Let Σ be a set of implications, $(\cdot)_{\Sigma}^+ : 2^M \rightarrow 2^M$ be its closure operator and $C \subseteq M$ a closed set, i.e., $(C)_{\Sigma}^+ = C$. The set $A \subseteq M$ is said to be a minimal generator (mingen) for C if $(A)_{\Sigma}^+ = C$ and, for all $X \subseteq A$, if $(X)_{\Sigma}^+ = C$ then $X = A$.

As we mentioned before, the generation of the implicational system associated with a closure operator is a hard problem and the reverse one has also an exponential behavior. Thus, here we deal with the definition of a method to solve this problem (Sect. 3) and also with the design of its efficient implementation (Sect. 5), particularly approaching the problem by using parallelism.

3 Minimal generators method

In [6], the Simplification Logic was used as the tool to find all the minimal generators (mingens) from a set of implications. The method applies the Function `Cls` to guide the search of new minimal generator candidates. Specifically, given a set of attributes M and an implicational system Σ , the algorithm renders a mapping $mg_{\Sigma} : 2^M \rightarrow 2^{2^M}$ that satisfies the following:

$$\forall X, Y \subseteq M, X \in mg_{\Sigma}(C) \text{ iff } C \text{ is closed for } (\cdot)_{\Sigma}^+ \text{ and } X \text{ is a mingen for } C.$$

Example 2 For the implicational system introduced in Example 1, the mapping mg_{Σ} is described as follows:

X	\emptyset	b	e	be	de	ace	bde	$acde$	$abcde$
$mg_{\Sigma}(X)$	\emptyset	b	e	be	d	a	bd	ad	ab
						c		cd	bc

Otherwise, X is not closed and $mg_{\Sigma}(X) = \emptyset$. Notice that \emptyset is closed and $mg_{\Sigma}(\emptyset) = \{\emptyset\}$, i.e., \emptyset is a minimal generator of the closed set \emptyset .

The algorithms we introduce in this section need to use the following operation for this kind of mappings. Given two mappings $mg_1, mg_2 : 2^M \rightarrow 2^{2^M}$, the mapping $mg_1 \sqcup mg_2 : 2^M \rightarrow 2^{2^M}$ is defined as

$$(mg_1 \sqcup mg_2)(X) = \text{minimals}(mg_1(X) \cup mg_2(X)) \quad \forall X \subseteq M$$

Thus, $Y \in (mg_1 \sqcup mg_2)(X)$ if and only if Y is a minimal set of $mg_1(X) \cup mg_2(X)$ in $(2^M, \subseteq)$, i.e., $Y \in mg_1(X) \cup mg_2(X)$ and there does not exist another attribute set $Z \in mg_1(X) \cup mg_2(X)$ such that $Z \subsetneq Y$.

We already have all the tools needed to define the Minimal Generator method. It was introduced in [6] and here we describe it as Function `MinGen` depicted below, together with an illustrative example of its application (see Example 3).

Minimal generators, an affordable approach by means of...

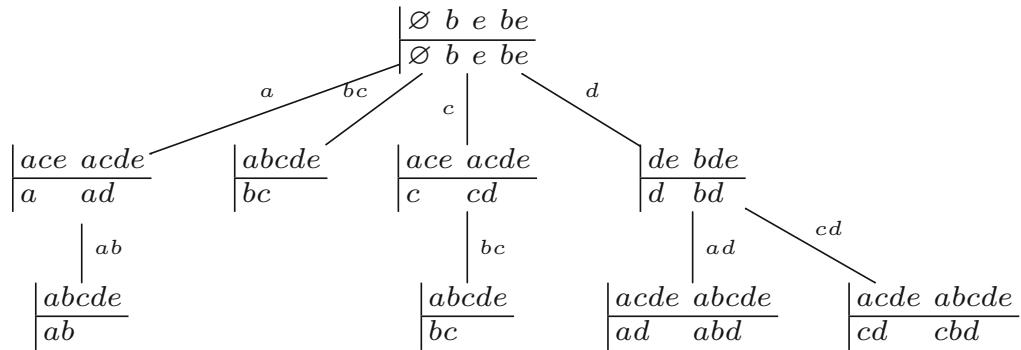


Fig. 2 Search tree for MinGen in Example 3

Function MinGen(M ,Label,Guide, Σ)

input : M , the set of all attributes;
 Label, an auxiliar set to build a minimal generator;
 Guide, an auxiliar set to build a closed set;
 Σ , an implicational system on M ;
output: The mapping mg_{Σ}
begin
foreach $X \subseteq M$ **do**
 $mg_{\Sigma}(X) := \emptyset$ ($Guide, \Sigma$):=cls($Guide, \Sigma$);
 $M := M \setminus Guide$;
 Premises := $\{A \subseteq M \mid A \rightarrow B \in \Sigma \text{ for some } B \subseteq M\}$;
 ClosedSets:= $\{X \subseteq M \mid A \not\subseteq X \text{ for all } A \in \text{Premises}\}$;
foreach $X \in \text{ClosedSets}$ **do**
 $mg_{\Sigma}(Guide \cup X) := \{\text{Label} \cup X\}$
foreach $A \in \text{Premises}$ **do**
 $mg_{\Sigma} := mg_{\Sigma} \sqcup \text{MinGen}(M, \text{Label} \cup A, \text{Guide} \cup A, \Sigma)$
return mg_{Σ}

Example 3 For the implicational system introduced in Example 1, the Function **MinGen** ($abcde$, \emptyset , \emptyset , $\{a \rightarrow c, bc \rightarrow d, c \rightarrow ae, d \rightarrow e\}$) returns

X	\emptyset	b	e	be	ace	$acde$	$abcde$	de	bde
$mg_{\Sigma}(X)$	\emptyset	b	e	be	a	ad	ab	d	bd

The search tree is shown in Fig. 2.

4 Reducing the search space in the MinGen method

In this subsection, we present a further MinGen method corresponding to a significantly enriched version of the original one. We have integrated a reduction to avoid the generation of redundant minimal generators and closed sets. The aim of this pruning is to identify redundant branches in the search space to avoid their exploration. This reduction must only be carried out if we can ensure that all the information regarding minimal generators has been collected in other branches. In Function **MinGenRd** this reduction strategy is implemented in line #1. Thus, to ensure that the information generated in one branch is superfluous, we design a prune based on set inclusion involving

all the nodes at the same level. We illustrate how this technique works in the following example:

Example 4 Given the search tree previously introduced in Example 3, the Function **MinGenRd** ($abcde, \emptyset, \emptyset, \{a \rightarrow c, bc \rightarrow d, c \rightarrow ae, d \rightarrow e\}$) applies a reduction strategy avoiding to open the branch whose label is a superset of another edge at the same level. Particularly, the branch labeled bc in Fig. 2 will not be opened because it is not minimal in the set $\{a, bc, c, d\}$.

Function MinGenRd($M, \text{Label}, \text{Guide}, \Sigma$)

```

output: The mapping  $mg_{\Sigma}$ 
begin
  foreach  $X \subseteq M$  do
     $mg_{\Sigma}(X) := \emptyset$  ( $\text{Guide}, \Sigma$ ):=Cls( $\text{Guide}, \Sigma$ );
     $M := M \setminus \text{Guide};$ 
  [<#1] Premises := Minimals{ $A \subseteq M \mid A \rightarrow B \in \Sigma$  for some  $B \subseteq M$ };
  ClosedSets := { $X \subseteq M \mid A \not\subseteq X$  for all  $A \in \text{Premises}$ };
  foreach  $X \in \text{ClosedSets}$  do
     $mg_{\Sigma}(\text{Guide} \cup X) := [\text{Label} \cup X]$ 
  foreach  $A \in \text{Premises}$  do
    [ $mg_{\Sigma} := mg_{\Sigma} \sqcup \text{MinGenRd}(M, \text{Label} \cup A, \text{Guide} \cup A, \Sigma)$ 
  return  $mg_{\Sigma}$ 
```

4.1 A generalization of the reduction strategy

In this section, we propose a generalization of the reduction strategy by considering the subset inclusion test not only with the nodes at the same level, but with all the minimal generators computed before the opening of each branch. One step further is to take advantage of the minimal generators already computed to increase the number of branches not needed to be opened. As Function **GenMinGen** shows, we consider this generalized pruned in #1, and then, the list of minimal premises to be considered in each stage is built in #2.

Function GenMinGen($M, \text{Label}, \text{Guide}, \Sigma, \text{MinGenList}$)

```

output: The mapping  $mg_{\Sigma}$ 
begin
  foreach  $X \subseteq M$  do
     $mg_{\Sigma}(X) := \emptyset$  ( $\text{Guide}, \Sigma$ ):=Cls( $\text{Guide}, \Sigma$ );
     $M := M \setminus \text{Guide};$ 
    Premises := Minimals{ $A \subseteq M \mid A \rightarrow B \in \Sigma$  for some  $B \subseteq M$ };
    ClosedSets := { $X \subseteq M \mid A \not\subseteq X$  for all  $A \in \text{Premises}$ };
    foreach  $X \in \text{ClosedSets}$  do
       $mg_{\Sigma}(\text{Guide} \cup X) := [\text{Label} \cup X]$ 
    foreach  $A \in \text{Premises}$  do
      [ $\text{if there no exists } Y \in \text{MinGenList such that } Y \subseteq A \text{ then}$ 
       [ $mg_{\Sigma} := mg_{\Sigma} \sqcup \text{GenMinGen}(M, \text{Label} \cup A, \text{Guide} \cup A, \Sigma, \text{MinGenList})$ 
  [<#1] [<#2] add  $A$  to  $\text{MinGenList}$ ;
  return  $(mg_{\Sigma})$ 
```

Notice that in Fig. 2 the branches labeled with ad and cd will now not be opened because in the previous level labels a and c were (respectively) opened. In the output of Function `GenMinGen` the minimal generators ad and cd appears in previous branches whereas abd and cbd are not computed because they are not really minimal generators (see the enumeration closed sets and minimal generators for this problem in Example 3).

4.2 Testing the performance of the reducing techniques

Once we have presented the original Minimal Generator method (MinGen) along with the two approaches (MinGenRd and GenMinGen) and an illustration of their search spaces has been shown, we present now a global comparison of the performance achieved by each of them.

For that matter, we have developed the corresponding implementations to apply the methods to a battery of sets of implications randomly generated. To evaluate this comparison, two different metrics are applied: (1) the execution time of the algorithm and (2) the number of nodes in the search tree built by the method. The reason for this selection becomes reasonable as follows. The execution time arises as the classical measure to test the performance, but it is always hardly linked to the resources we are working with. So, we add the number of nodes of the tree as a metric to compare these algorithms since it could better guide us to put forward an argument to decide which algorithm is the best as it is an architecture-independent value.

In addition, due to the intrinsic nature of execution time, it is also imperative to state that every experiment shown throughout this document has been repeated several times, so we can now write down the most reliable average values w.r.t. the execution time metric.

In the experiments, the hardware configuration used is: Intel(R) Core(TM) i7-6700HQ CPU 2.60 GHz, 8 Gb RAM memory, running over Windows 10. We have generated a battery of different inputs to be used in the sequential implementation to show the improvements of the methods, that is:

- A synthetic test. Contains 5 testing files with 50 implications built using 50 possible different attributes. The implications are randomly generated.
- A real dataset. We focused on MovieLens datasets¹ which have been widely used in education, research, and industry [13]. In this spirit, we chose the MovieLens10M dataset which is a dataset totally accessible through MovieLens web page and contains huge information about movies, genres, ratings, users, etc. From all this information, we are going to generate a table with 10.681 rows, where we face every movie with all the possible genres included in the dataset. The extraction of implications from the dataset renders a set of 19 attributes (genres) and a set of 245 implications.

Having said that, the results of the experiments are shown in Fig. 3. Giving an overall view of the results, it can be clearly seen how the reduction strategies significantly reduce both the number of nodes and the execution times. Indeed, experiment

¹ <https://grouplens.org/datasets/movielens/>.

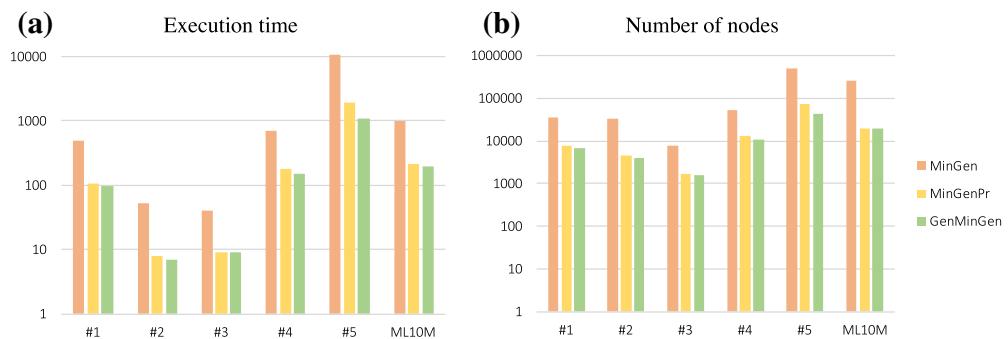


Fig. 3 Execution times (a) and number of nodes (b) results for the sequential experiments. Notice the logarithmic scale applied to the axes to better view the results

over ‘sequential-5’ file shows how both metrics have been drastically reduced. The execution time has been reduced from more than 10.000 s to less than 2.000 s. Besides, it is also worth to mention the reduction of the number of nodes which implies less resources needed in terms of memory and storage. Overall, like it was meant to happen, MinGen is surpassed by MinGenRd, and GenMinGen improves the latter as well.

Although GenMinGen has stated to have a better performance than MinGenRd (and both of them better than MinGen), the former is not suitable for parallelization. This is because it demands communication among all the subproblems previously obtained. This breaks our parallel philosophy, avoiding the use of massive computation. For this reason, we have studied a parallelization of the MinGenRd method.

5 Parallel computation of minimal generators

We have already shown the improvements reached by the reduction strategies within the minimal generators methods. However, once it comes the case that we want to use these methods over larger inputs, we can figure out in the light of the results obtained, that execution times of the sequential methods would go too far to be easily handled. Nonetheless, taking into account that every branch of the tree created by the methods constitutes a problem on its own, we can think about resolving them simultaneously and, later, combining the partial solutions to get the final output. This natural view of the problem as a parallel execution, provided by our logic-based methods, leads us to develop a new and parallel version of the minimal generators method.

Before going further, we introduce that the supercomputing resources and architecture that have been used to run the following parallel experiments are those provided by the Supercomputing and Bioinnovation Center of the University of Málaga.² In particular, we have developed each experiment using 32 nodes cluster SL230, counting on 16 cores and 64GB RAM memory and 7 nodes cluster DL980, counting on 80 cores and 2 TB RAM memory. Communications are carried out on Infiniband Net FDR and QDR. These cores are reserved just for our use so we can assess reliable results regarding execution times.

² <https://www.scbi.uma.es/>.

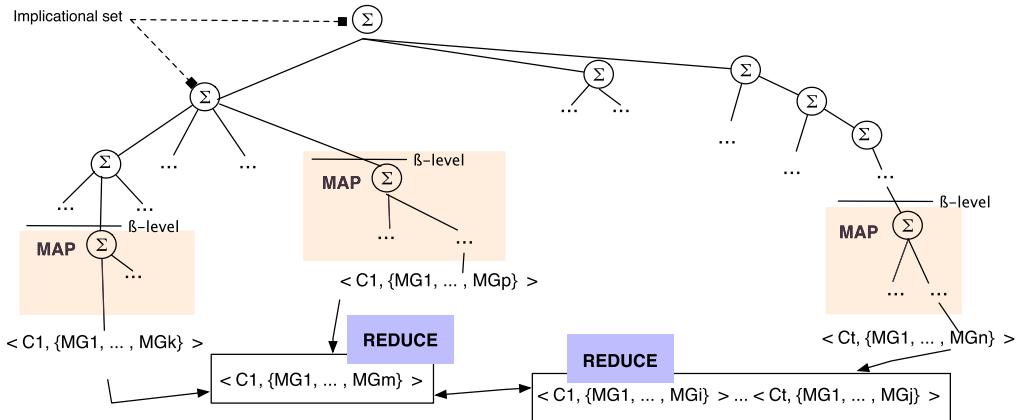


Fig. 4 Illustration of the map-reduce paradigm used in the design of MinGenRdPar

5.1 Parallel algorithm

We have developed a parallel implementation of the methods following the framework shown by the authors in [2]. In summary, it takes advantage of massive computation following the map-reduce paradigm. This parallel implementation of the methods performs in two stages. First one is the splitting stage. There, a single core builds the search space until it reaches a certain depth level of the tree. The goal is to split the original problem into several problems that can be treated by multiples cores, playing the role of a “Map” procedure. It goes this way until the size of the current node becomes lower than a given size (β) w.r.t the number of implications. From that moment on, each of these subproblems is resolved in parallel using multiple cores until we reach the leaves of the tree. Finally, a single core is used to compose the global result by merging the different outputs of each subproblems into a final one. This becomes necessary in order to eliminate redundancies to obtain the *minimal* generators. It is in this stage where the Reduce procedure acts as a filter to avoid redundant information in the global output. The architecture of our approach is depicted in Fig. 4.

In addition, we show the pseudocode of our parallel *MinGenRd* algorithm in Function [MinGenRdPar](#). We have emphasized the main stages of our implementation that likely correspond to the Map-Reduce paradigm.

The β level, named *BOV* (Break-Off Value) in the algorithm, is a critical parameter since it is in charge of deciding when to split the original problem into subproblems to manage them in parallel, generating a new Map task. As we will broadly explain later in Sect. 5.3, a hard experimental study has been needed to determinate the most suitable value for *BOV*.

Function MinGenRdPar(M ,Label,Guide, Σ ,BOV)

```

output: The mapping  $mg_{\Sigma}$ 
begin
  foreach  $X \subseteq M$  do  $mg_{\Sigma}(X) := \emptyset$  (Guide, $\Sigma$ ):=Cls(Guide, $\Sigma$ );
   $M := M \setminus$ Guide;
  Premises := Minimals{ $A \subseteq M$  |  $A \rightarrow B \in \Sigma$  for some  $B \subseteq M$ };
  ClosedSets:=  $\{X \subseteq M$  |  $A \not\subseteq X$  for all  $A \in$  Premises};
  foreach  $X \in$  ClosedSets do  $mg_{\Sigma}(\text{Guide} \cup X) := \{\text{Label} \cup X\}$ 
  If  $|\Sigma| \leq BOV$  then do in parallel
    // [MAP]
    foreach  $A \in$  Premises do
      // [REDUCE]
       $mg_{\Sigma} := mg_{\Sigma} \sqcup$  MinGenRd( $M$ , Label  $\cup A$ , Guide  $\cup A$ ,  $\Sigma$ )
    return  $mg_{\Sigma}$ 
  else
    // [Splitting stage]
    foreach  $A_k \in$  Premises do
       $mg_{\Sigma}^k =$  MinGenRdPar( $M$ , Label  $\cup A_k$ , Guide  $\cup A_k$ ,  $\Sigma$ , BOV)
    forall the  $mg_{\Sigma}^k$  do
      // [REDUCE]
       $mg_{\Sigma} := mg_{\Sigma} \sqcup mg_{\Sigma}^k$ ;
    return  $mg_{\Sigma}$ 

```

5.2 Sequential versus parallel experiments

This time, we have carried out an experiment to test the performance of the parallel implementation `MinGenRdPar` versus the sequential implementation `MinGenRd`. The input files we are going to use will count on numbers raised up to 150 attributes and 150 implications. Even with these numbers, three times higher than the sequential experiment in Sect. 4.2, parallel version obtains results in an admissible time as we can notice on the results given in Table 1. Looking closely, Table 1 collects the following information from left to right: (1) identifier of the input file and the method used to resolve, (2) execution time of the sequential version, (3) number of subproblems generated by the splitting stage of the parallel implementation, (4) execution time of the splitting stage, (5) execution time of parallel resolution, (6) execution time of the whole process, (7) number of nodes of the generated tree and (8) number of minimal generators obtained.

In conclusion, thanks to the application of the parallel implementation, we have been able to reduce the execution times from hours and days (experiments over #-{2,3,7,8,10}-sequential) down to just a few minutes. In other words, parallel computation improves the execution times even when dealing with big-sized problems.

5.3 Estimation of the BOV

There is a crucial aspect we have to keep in mind in the first stage of the parallel implementation, i.e., the splitting. We need to decide when a subproblem has to be generated (Map procedure) to be simultaneously executed. To this end, we are going to

Table 1 Comparison between both sequential and parallel versions of MinGenRd algorithm applied to big-sized problems

Problem	Seq.time (s)	Subp	Split _t (s)	Parallel _t (s)	Total _t (s)	Nodes	MinGens
#1	43	11	3	1	4	374	216
#2	17.352	347	220	55	275	54.375	6.273
#3	33.338	822	2.338	251	2.589	68.531	6.529
#4	4.612	344	350	97	447	25.477	2.478
#5	1.585	168	432	30	462	12.522	1.159
#6	1.653	79	35	7	42	8.110	1.436
#7	107.238	1.754	958	242	1.200	262.621	9.113
#8	61.381	966	253	188	441	257.267	5.538
#9	372	24	7	2	9	1.726	683
#10	7.484	277	186	65	251	45.962	2.969

use a value, denominated *BOV*, which represents the cardinal of the set of implications of the current node, since we have empirically observed that as long as this cardinal grows, the longer the branch of the tree uses to reach. The *BOV* is computed as a percentage of the size of the input that has to be estimated to balance the work done by each core.

Yet, deciding the *BOV* is a decisive task of the current investigation due to the following difficulties. On the one hand, if we decide to stop at a level near to the root of the tree by selecting a low *BOV* (i.e., a high number of implications), we are certainly reducing the execution time of the splitting stage and only a few subproblems would be created. Accordingly, since the tree would not have been able to spread out yet, then we will not have enough material to be managed in parallel using different cores. On the other hand, if we stop the split in a depth level far from the root, the splitting process will surely create a large amount of subproblems but its execution time would surely grow up. This is why we have selected a *BOV* empirically, as it is really difficult to get the right value by just analyzing the input theoretically. However, after making a lot of experiments, several aspects worth to be mentioned and this section stands to this effect.

We have hitherto established a *BOV* of 140 (recall that our number of implications is 150) within the experiments as it has shown to be the best one to leverage parallelism based on our experience. That is to say, we are choosing a *BOV* that corresponds to the $\approx 93.33\%$ of the original set of implications. However, in order to explore this fact, we have repeated our parallel experiments using different *BOVs*. We take lower *BOVs* that makes the splitting stage to go deeper into the tree, specifically, we will use both 130 and 100 *BOVs*, i.e., the ≈ 86 and $\approx 66\%$ of the whole original set of implications, respectively. This particular selection of values is not made by chance but with the aim of conveying several situations we have found after carrying out plenty of experiments which we come to analyze next. Results concerning execution times are gathered in Table 2.

Table 2 Experiments using different BOV within the parallel version of MinGenRd algorithm applied to big-sized problems

Problem	Subp	Split _t (s)	Parallel _t (s)	Total _t (s)	BOV (%)
#1	0	44	–	44	66.67
	0	41	–	41	86.67
	11	3	1	4	93.33
#2	0	18.533	–	18.533	66.67
	885	8.532	58	8.590	86.67
	347	220	55	275	93.33
#3	0	33.377	–	33.377	66.67
	0	33.285	–	33.285	86.67
	822	2.338	251	2.589	93.33
#4	0	4.858	–	4.858	66.67
	308	3.703	22	3.725	86.67
	344	350	97	447	93.33
#5	0	1.601	–	1.601	66.67
	0	1.547	–	1.547	86.67
	168	432	30	462	93.33
#6	0	772	–	772	66.67
	144	492	11	503	86.67
	79	35	7	42	93.33
#7	0	167.451	–	167.451	66.67
	5.412	96.433	295	96.728	86.67
	1.754	958	242	1.200	93.33
#8	0	75.060	–	75.060	66.67
	5.344	41.404	375	41.779	86.67
	966	253	188	441	93.33
#9	0	82	–	82	66.67
	24	42	2	44	86.67
	24	7	2	9	93.33
#10	0	9.438	–	9.438	66.67
	697	6.569	50	6.619	86.67
	277	186	65	251	93.33

There are several outcomes within that table. First and foremost, except for problem #1 where there exist a subtle difference, all the problems behave worse regarding the execution time when we delay the splitting point. To find out an explanation, we put our attention in the number of generated subproblems where four different behaviors arise when we vary the BOV . We proceed to enumerate each of them with the support of the results obtained in Table 2.

Let $BOV_1 > BOV_2$, and let Sp_1, Sp_2 the number of generated subproblems associated with each of them. So, there might be situations where:

- $Sp_1 < Sp_2$. The algorithm has more scope to expand the tree and generate more subproblems. In this way, parallelism would take advantage but the time needed to split scupper the global execution time. Problems #\{2,6,7,8,10\} reflect this situation.
- $Sp_1 > Sp_2$. It can happen that even going deeper into the tree we obtained less nodes, i.e., less subproblems. This is because there are branches that end before reaching the BOV and they are resolved within the split stage, so the execution time grows higher. We can see this case in problem #4.
- $Sp_1 \neq 0 \wedge Sp_2 = 0$. Following on from the previous point, we can fall in an extreme situation where no subproblems are generated within the split stage because every branch of the tree ends up before the splitting point. This situation may be considered as the worst case since the parallel implementation performs the same as the sequential one. This happens for every problem when using a $BOV \approx 66\%$ and also in problems #\{1,3,5\} even with a $BOV \approx 86\%$.
- $Sp_1 = Sp_2$. This reflects an entangled situation in which lower levels of the tree may count on the same number of nodes than the higher ones. The point is that when we go deeper in the tree we may be generating more nodes as the tree expands and so the number of subproblems would grow up, but also, it may be several branches that end before going deeper and so it reduces the number of nodes. Therefore, with this additions and subtractions, the global calculation of subproblems can end up in a draw with different $BOVs$ as shown in problem #9. But that's not all. Problem #9 indeed shows the same number of subproblems for $BOV \approx 93.33\%$ and $BOV \approx 86\%$, however, execution time is not the same, actually, it is worse for a $BOV \approx 86\%$ since we have stretched on the split stage.

As we have already mentioned before and after the analysis shown in this part, it becomes clear that trying to infer the best BOV for an experiment is not an easy issue so far as it depends on many possible situations we can face when dealing with the minimal generators enumeration problem.

5.4 Estimation of the suitable number of cores

Up to now, we can think it is just a matter of resources that we can increase the input size and still get results within a reasonable time. However, it is not so obvious and this subsection is dedicated to analyze this fact. For this purpose, the big-sized problems used before in Sect. 5.2 will be tested again by using 16, 32, 48, 64 and 80 cores each time. Results are shown in Table 3.

This is not up to expectations results. Although resources are better now, results do not come along. It can be seen that, we do improve the performance by using more cores (e.g., from 16 to 64 cores), however, there comes a moment where no profit is obtained; results are pretty similar for 48, 64 and 80 cores. This is due to the fact that there are several branches in the tree (maybe just one) that take so long to finish and then, no matter how much we improve the resources, the global time remains almost the same. The problem is that we cannot predict for the time being whether a branch will result in a long or a short one. This situation stalls our first intentions of blindly growing the amount of resources and shows up that, the number of cores for

Table 3 Execution times (in seconds) results obtained when increasing the number of cores applied

Problem	Number of cores				
	16	32	48	64	80
#1	5	4	3	3	3
#2	310	275	199	190	197
#3	2.742	2.589	2.122	2.078	2.075
#4	512	447	444	440	446
#5	598	462	416	445	442
#6	50	42	34	35	34
#7	1.457	1.200	1.078	1.010	1.012
#8	499	441	432	430	425
#9	9	9	7	7	7
#10	267	251	196	194	195

Table 4 Parallel minimal generator algorithm applied to a real-world dataset

Problem and Method	Subp	Split _t (s)	Parallel _t (s)	Total _t (s)	Nodes	MinGens
Mushrooms-parallel	224	152	9	161	81.363	17.127

this problem can be established to 64 as an optimal value, achieving a balance between resources needed and benefits obtained.

5.5 Real-world dataset experiment

To properly finalize this section, we bring now the results obtained when applying our algorithm to a real-world dataset. In particular, we put our focus in the Mushroom Data Set³ accessible from the website of the University of California, Irvine (UCI).⁴ Basically, this dataset includes descriptions of hypothetical samples corresponding to 8.124 species of gilled mushrooms using 22 attributes. On this dataset, an adaptation is made in order to convert multi-valued information into binary information. As a result, we will manage a dataset with 126 attributes. The number of implications inferred from this dataset is 1.587. We remark that with this number of implications, sequential version of the algorithm does not finish.

To carry out this experiment we use the conclusions reached by the previous sections. Therefore, we will use 64 different cores and a $BOV \approx 93.33\%$, i.e., $BOV = 1.481$, as they seem to be the best values to proceed. The results of this experiment are shown in Table 4 where it clearly arises how our algorithm fulfills when dealing with a big-sized and real-world dataset.

³ <https://archive.ics.uci.edu/ml/datasets/mushroom>.

⁴ <http://archive.ics.uci.edu/ml/>.

6 Conclusions and future works

Enumerating all the closed sets and their minimal generators is a hard problem, but essential in several areas and an opportunity to show the benefits of FCA for real applications. Such information can be obtained from a dataset or from a set of implications. In this work we focus on this second problem, which have been approached in a lesser extent. In order to face this task, this work presents an efficient reduction of the search space technique to improve the performance of minimal generator enumeration. The new method has been designed to fit the Map-Reduce architecture. Here is where parallel computation comes to make it possible for us to deal with such amount of information.

In particular, we have designed two new methods (MinGenRd and GenMinGen) implementing pruning strategies to reduce the search space. The empirical study proves how these methods surpass the previous one in the literature (MinGen). We also establish the adequacy of MinGenRd on being implemented following a parallel implementation by means of the Map-Reduce paradigm. Therefore, as the main contribution of this work, we propose the parallel implementation of MinGenRd method. The empirical study proves the very significative improvement achieved w.r.t the original sequential version. The parallel methods to compute minimal generators can make really usable these methods in practical applications.

To properly characterize the parallel approach, we have developed two battery of experiments: (1) we have established the *B OV* threshold, a parameter included in the algorithm to effectively apply the Map procedure taking the most of this paradigm and, (2) we have established the number of cores need to execute this method, thus delimiting the resources of the parallel architecture.

As future works, we plan to study how to use different software and hardware resources to achieve a parallel implementation of the best sequential method, that we have proposed in this paper, GenMinGen method. Such implementation requires a complex design to store the previously generated minimal generators without interfering the parallel execution.

A deeper study of the best values to be considered as *B OV* to maximize the results will be also developed, such as taking into account another parameters beyond the implication set cardinality.

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Capítulo 5

Sistemas de Recomendación Conversacionales

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Enhancing the conversational process by using a logical closure operator in phenotypes implications

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In this paper, we present a novel strategy to face the problem of dimensionality within datasets involved in conversational and feature selection systems. We base our work on a sound and complete logic along with an efficient attribute closure method to manage implications. All of them together allow us to reduce the overload of information we encounter when dealing with these kind of systems. An experiment carried out over a dataset containing real information comes to expose the benefits of our design. Copyright © 2017 John Wiley & Sons, Ltd.

Keywords: conversational systems; feature selection; implications; logic

1. Introduction

A common problem related to knowledge discovering within the clinical context appears when it is necessary to work over datasets with a high number of features (variables or attributes). This situation is known as the curse of dimensionality phenomenon. In these cases, trying to apply data mining techniques in its different approaches (classification, regression, clustering, association rule analysis, etc.) becomes a hard task.

To address this issue, we can find many works in the literature about data reduction, specially in feature selection, that can help us to discard those features not worthy to be considered by means of different criteria. As this respect, several techniques have already been applied such as genetic algorithms, regression, neural networks and many others. All of them guided towards the application of an automated process that is applied at once (batch mode) by feature selection.

The problem of managing large volumes of information is highly present on another hot topic field of knowledge: recommender systems. The major goal of these systems is to help the user when dealing with an extremely high number of alternatives. Recommender systems are present in many different areas of today's society (e-commerce, tourism, films, music, news, etc.) in which large amount of data are pretty much often. Most recommender systems base the retrieval of items in predictions about how suitable is an item to satisfy a user's need. These predictions could be performed from user's preferences, profile, context and so on. To achieve that, different strategies are applied to enclose a recommender system into different types. Best known are content-based, knowledge-based, collaborative filtering and context-based. From the point of view of this work, we centre our efforts not just in the recommendation strategy but also in the process of obtaining a recommendation.

As is usually the case, in order to properly make a selection of items within recommenders, the user needs to introduce information over and over. That used to be something of a chore because many items may be too much detailed drawing the high-dimensionality problem mentioned earlier. An example of this situation appears when medical professionals try to elaborate a diagnosis checking symptoms from a vast list of possible ones.

One solution to tackle this issue in the field of recommender systems is the trend of conversational systems (critiquing recommender systems overall). In these systems, an iterative process is applied in which the user sets one or more features for items to fit. From this input and using different techniques, the system progresses choosing (or even predicting) subsets of items that agree with the user's preferences, until it comes to a final output with a suitable size. The problem is that, if the dataset presents a high dimensionality, the number of steps until we acquire a fitting recommendation set could be huge.

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The solution we proposed in this work is to manage the problem of the high dimensionality by means of a feature selection process guided by the user (human expert) within a conversational system. To achieve that, we base the approach on a novel management of implications and closure implementation. Our proposal avoids the problem mentioned in [1] remarking that batch mode feature selection systems extract them randomly. Their only criterion in the selection is the feature predictive capacity, without considering the medical knowledge. For instance, the Computer Feature Selection over Cleveland dataset discards some key features such as cholesterol, age or ECG characteristics. On the contrary, we surpass this issue by making an expert-driven automated selection.

Our major goal is to enhance the diagnosis process preserving the accuracy of the results (that is granted when using implications) and accelerating the process by reducing the necessary steps in the dialogue. Besides, the reduction of the complexity within the process is also a must, so we can obtain results timely and in due form.

As already mentioned, this work is based on implication management. Nonetheless, it is necessary to clarify that it is out of the scope of this work to describe how to extract these implications, because there are already data mining techniques specialised on this task [2].

The set of implications is the heart of the knowledge in the process we propose. We support our approach on a sound and complete logic named simplification logic (SL_{FD}), presented in several previous works, which was designed to develop deduction methods. In particular, we propose to use SL_{FD} attribute closure algorithm as the core of a feature selection framework. This algorithm is used as a basis for reasoning over the features (symptoms, phenotypes, signs) of the items (diseases) of the dataset we are going to use, reducing the number of steps in the conversation to reach a suitable diagnosis.

The key point of our framework comes from the SL_{FD} closure nature that renders a set of features that corresponds with the closure and, in addition, a new implication set corresponding with the knowledge not already used in the conversation (selection o diagnosis) process. Such implication set can be obtained by using other methods but, with SL_{FD} closure method, we compute it in linear time. This low complexity along with the reduction of the number of steps provide an overall time reduction for the process.

Our framework has been tested on a dataset that puts together real information about diseases and phenotypes. Particularly, we have selected a set of haematological diseases with phenotypes related to them. In the experiment, several metrics to evaluate the system's performance have been applied, and results have come to demonstrate the highlights of our framework.

The remainder of the paper is organised as follows: Section 2 draws attention about the state-of-the-art references and their motivations along with a brief introduction of the main elements of our approach. Section 3 brings us a detailed explanation about how the information will be managed, the notions of the SL_{FD} , that apply in this approach and the specification and the usefulness of the closure algorithm. The conversational processes our approach carries out along with major benefits it reaches are presented in Section 4. Section 5 presents an empirical test considering a real case in a medical environment and the metrics used to evaluate the performance. Conclusions and future works close the paper.

2. Related works

Our approach tries to solve the problem of searching information in high-dimensionality datasets by combining techniques from diverse fields (recommender systems, feature selection, logic) in a knowledge-based framework. In this section, we analyse some outstanding related works in each area.

As mentioned, in general terms, the problem we aim to solve is to search a precise result (e.g. a diagnosis) from a high volume of data (e.g. a diseases dataset) without previous user information. By these means, our approach is inspired in one of the solutions proposed in this field, the so-called conversational recommender systems [3]. These are closely related with the concepts of critiquing recommender systems [4] and information recommendation [5]. In these systems, recommendation is generated by means of a dialogue with the user that allows an incremental elicitation of preferred item features, that is, user requirements are directly elicited within a recommendation session. An interesting work in this area is [6], which states the suitability of knowledge-based approaches for conversational processes. In particular, these authors use constraint-based reasoning, instead of our logic-based approach. Besides, this work deals with the concept of query tightening, analogous to the one applied in our proposal.

Another remarkable work is [5], which shares our aim to decrease the conversation length (number of steps). The theoretical framework they use is a model of attribute dominance with two versions, a qualitative and a quantitative one. These authors propose metrics about conversation number of steps and pruning rates, both of them very similar to the ones used in our work. Nevertheless, they present an experiment focused on tourism.

Regarding critiquing recommender systems, Chen and Pu [7] explain how user self-motivated attribute selection enable users to achieve a higher level of decision accuracy than system-proposed attribute selection. This fact supports our approach in which human expert guides the conversation and the feature selection process.

The main elements of our theoretical framework are SL_{FD} as well as the concept of attribute closure and the algorithm we have defined to compute it. One of the main advantages of this algorithm is the reduction of the time required by the selection process. For a better understanding, works related with the theoretical basis are exposed in the next section.

The starting point of our selection process is the set of implications (also named exact association rules) derived from the working dataset. To acquire these implications, association rule mining is needed [8]. It is easy to find in literature works evaluating association rule mining techniques both in the biological-medical [9] and recommender systems [10] fields.

Once association rules had been inferred from the dataset, they can be used in the main medical tasks, such as screening, diagnosis, treatment, prognosis, monitoring and management [11, 12]. For example, Nahar *et al.* [13] uses association rule mining to analyse factors related with heart diseases. In the same way, many other works use association rules to build a decision support system or prediction model [14–17].

Closer to our work, Mansing *et al.* [18] deals with the fact that the number of association rules used to be very high. This work proposes an association rule pruning mechanism, based on well-known concepts as support, confidence and reliability. A similar idea is proposed in [19] and [20], although the latter applies temporal abstraction in the decision support system. Our approach includes an association rule filtering process based on the closure concept. Hu *et al.* [21] proposes the use of association rules for predicting combination of alarms generated by bedside monitors. These authors identify frequent alarm combinations and then carry out a variance analysis to measure the data mining process performance. Regarding our work, it should be noticed that they use a closure-like concept to define a heuristic that controls combination size. In any case, we must remark that our work is not strictly comparable with the previously cited ones, because we do not aim to build a prediction model but to improve a conversational process.

In the introduction, we affirm that our approach may be described as an expert-driven feature selection process. As Fang *et al.* [22] exposes, feature selection is a problem profusely studied in the literature. So, there are a number of solutions based on statistical techniques such as principal component analysis, linear discriminant analysis and independent component analysis, all of them suitable to deal with high-dimensionality datasets. Besides, other authors use evolutionary computing techniques such as particle swarm optimisation [23, 24]. Nevertheless, it should be remarked that most of feature selection works are focused on batch (non-interactive) processes. This way, the system, without user intervention, determines and selects more relevant features according to their predictive significance.

Recently, some papers dealing with iterative feature selection processes have been published. For example, Fialho *et al.* [25] proposes a tree feature selection process based on fuzzy modelling (and fuzzy rules). They use two tree searching techniques (sequential forward selection and sequential backward elimination). This interesting work achieves good results in well-known metrics such as Area Under Curve (AUC) or accuracy, but requires many inputs when the dataset has a high number of features. Analogously, Shilaskar and Ghatol [26] apply the same techniques but using Support Vector Machine (SVM) as classifier.

Some authors analyse the relation between features as means to reduce the dataset, for example, using correlation feature selection as a basis. This is the case of [27], although they propose a batch, non-interactive, process.

Closer to an interactive feature selection, Li *et al.* [28] define a selection process in which the input is a feature stream. They use information theory techniques (mutual information, conditional mutual information, entropy) to build a prediction model and achieve good results about the number of selected features. Although they process the features sequentially, the work is not oriented to perform as a conversational selection process.

In this way, we must remark [29], which highlights the importance of an online feature selection, so that features are processed one-by-one. This work is aimed to an extremely high-dimensionality context (in the order of millions of features) and is based on information theory. They use pairwise correlation analysis as a mean to remove redundant features. The goal of this proposal is to build a prediction mode with scalability as its main advantage.

In [30], an interesting work is presented that, as ours, deals with association rules and feature selection. Nevertheless, they base the feature extraction on methods such as partial least squares or principal component analysis, and it is not directly comparable with our approach.

3. Knowledge representation and automated deduction

In this section, we address the issue of specifying and efficiently managing the information. Knowledge-based systems have to balance both commitments to obtain, at the same time, powerful and efficient techniques. In our opinion, one of the tools that has shown a better behaviour is implications. They combine a very simple and natural way to write if-then-rules with an efficient and automated management. One evidence supporting our choice is its widespread use in different areas: databases, logic programming, formal concept analysis, artificial intelligence and so on.

In this work, knowledge is stored by considering implications, following the interpretation adopted in formal concept analysis [31], because of their simplicity. In this area, implications are inferred from datasets that are considered as binary relations between a finite set of objects and their attributes (depicted by rows and columns, respectively). Such interpretation is the following: Given a formal context K , an implication is an expression $A \rightarrow B$, where A and B are subsets of attributes, and it is said to be valid in K if and only if every object that has all the attributes from A also has all the attributes from B . Example 1 illustrates the knowledge captured by using implications.

Example 1

Let K be the formal context described in Table I showing the 22 common viruses and their usual ways of transmission.

The implications that hold in this dataset are as follows:

$\text{Blood} \rightarrow \text{Sexual}$	$\text{Droplet} \rightarrow \text{Direct}$
$\text{Fluids} \rightarrow \text{Vertical}$	$\text{Respiratory} \rightarrow \text{Direct, Droplet}$
$\text{Direct, Sexual, Droplet} \rightarrow \text{Faecal}$	$\text{Direct, Vertical} \rightarrow \text{Sexual}$
$\text{Faecal, Sexual} \rightarrow \text{Direct, Droplet}$	$\text{Faecal, Direct} \rightarrow \text{Droplet}$

In this way, implications allow to express a strong relation between two subsets of attributes of our system. Moreover, such information can be interpreted in a natural way. For instance, the last implication tells us that if a virus is transmitted by a direct contact and with faecal transmission, we also have to be vigilant about the sneezes.

Our proposal to integrate implications into the conversational issue is based on the SL_{FD} [32], which constitutes a sound and complete logic. As we shall see, such a strong basis allows us to include a reasoning method in the dialogue process. As mentioned, we built our framework on implications, which constitutes the main element of SL_{FD} language. It is formally defined as follows:

Table I. Viruses and usual way of transmissions dataset.

	Faecal	Direct	Vertical	Sexual	Respiratory	Saliva	Fluids	Droplet	Blood
Adenovirus	×	×		×				×	
Coxackievirus	×	×			×			×	
Epstein-Barr						×			
Hepatitis A	×								
Hepatitis B			×	×			×		
Hepatitis C				×				×	
Herpes type 1		×				×			
Herpes type 2			×	×					
Cytomegalovirus			×				×		
Herpesvirus type 8				×		×			
HIV			×	×					
Influenza		×						×	×
Measles virus		×						×	
Mumps virus		×						×	
Papillomavirus		×	×	×					
Parainfluenza		×						×	
Poliovirus	×								
Rabies		×						×	
Respiratory syncytial		×						×	
Rubella		×			×			×	
Varicella zoster		×						×	

Definition 1

Let M be a finite set, the formulae of SL_{FD} are expressions, named implications, of the form $X \rightarrow Y$, where X and Y are subsets of M .

From now on, we use lower case letters to denote the elements in M while uppercase letters denote its subsets. We use the standard notation and symbols of set theory. For the sake of readability, inside of a formula, $X - Y$ denotes the set difference operator $X \setminus Y$, and XY denotes the union operator $X \cup Y$.

Implications are interpreted in a conjunctive way, that is, they correspond to formulas $a_1 \wedge \dots \wedge a_n \rightarrow b_1 \wedge \dots \wedge b_m$ where propositions $a_1, \dots, a_n, b_1, \dots, b_m$ are elements of the set M . The interpretation is the following:

Definition 2

Let O and M be two finite sets, named objects and attributes, respectively, and I a relation in $O \times M$. An implication of SL_{FD} $X \rightarrow Y$, where X and Y are subsets of M , is valid in I if and only if

$$\{o \in O \mid (o, x_i) \in I \forall x_i \in X\} \subseteq \{o \in O \mid (o, y_j) \in I \forall y_j \in Y\}$$

Apart from its natural way to express knowledge as a rule, implications provide a logic reasoning and inference. Their symbolic management was originally proposed in [33]. However, because of the central role that transitivity plays in that axiomatic system, the development of executable methods to solve implications problems has rest on indirect methods. The introduction of the SL_{FD} opened the door to the development of automated reasoning methods directly based on its novel axiomatic system [34, 35]. The axiomatic system of SL_{FD} is introduced as follows:

Definition 3

The axiomatic system of SL_{FD} considers reflexivity as axiom scheme

$$[\text{Ref}] \quad \overline{A \rightarrow A}$$

together with the following inference rules called fragmentation, composition and simplification, respectively.

$$[\text{Frag}] \quad \frac{A \rightarrow BC}{A \rightarrow B} \quad [\text{Comp}] \quad \frac{\begin{array}{c} A \rightarrow B, C \rightarrow D \\ AC \rightarrow BD \end{array}}{AC \rightarrow BD} \quad [\text{Simp}] \quad \frac{A \rightarrow B, C \rightarrow D}{A(C - B) \rightarrow D}$$

As we mentioned earlier, in [32], we defined, in the usual way, the semantic entailment ($\Gamma \models A \rightarrow B$) and syntactic derivation ($\Gamma \vdash A \rightarrow B$). Because we also proved the soundness and completeness of this logic, both notions can be equivalently used. We also introduced the notion of equivalence between sets of implications $\Gamma_1 \equiv \Gamma_2$ iff for all $A \rightarrow B \in \Gamma_1$, we have that $\Gamma_2 \vdash A \rightarrow B$ and vice versa.

We remark that SL_{FD} language considers as valid formulae those ones where any of their two parts can be the empty set, denoted $A \rightarrow \top$ and $\top \rightarrow A$. Their meanings were discussed in [32]. In that work, we also introduced the following result where the derivation of an implication $A \rightarrow B$ is reduced to the derivation of the formula $\top \rightarrow B$ having $\top \rightarrow A$. This result will be used later in the design of our novel closure method.

Proposition 3.1

For any Γ and for all $X, Y \subseteq M$, $\Gamma \vdash X \rightarrow Y$ if and only if $\Gamma \cup \{\top \rightarrow X\} \vdash \top \rightarrow Y$

The syntactic derivation provides an automated management of implications. In particular, it can be used to solve the so-called implication problem: Given a set of implications Γ and an implication $A \rightarrow B$, we want to answer whether $A \rightarrow B$ is deduced from Γ . This problem can be approached by using the closure operator.

Definition 4

Let $A \subseteq M$ be a set of attributes and Γ a set of implications; we define its closure with respect to Γ as the maximum subset $A_\Gamma^+ \subseteq M$ such that the $\Gamma \vdash A \rightarrow A_\Gamma^+$.

Moreover, the set A is named closed iff we have $A_\Gamma^+ = A$.

Implication problem has been traditionally tackled by using a basic method that receives $A \subseteq M$ as input and exhaustively uses the subset relation by iteratively traversing Γ and adding new elements to the closure. This method was proposed in the 1970s [36], and it is sketched in Algorithm 1.

Algorithm 1: Standard Closure

```

Data:  $\Gamma, A$ 
Result:  $A_\Gamma^+$ 
begin
1    $A_\Gamma^+ := A$ 
2   repeat
3      $A' := A_\Gamma^+$ 
4     foreach  $X \rightarrow Y \in \Gamma$  do
5       if  $X \subseteq A_\Gamma^+$  and  $Y \not\subseteq A_\Gamma^+$  then
6          $A_\Gamma^+ := A_\Gamma^+ \cup \{Y\}$ 
7     until  $A_\Gamma^+ = A'$ ;
8   return  $A_\Gamma^+$ 

```

Later, several authors have developed several methods by using different techniques, efficiently solving this problem in linear time. In [37], the authors show that the complexity of closure problem is $O(|A| |\Gamma|)$. They also mention that 'in the literature, $O(|A| |\Gamma|)$ is usually considered as the order of the input. From this point of view, this is a linear time complexity for the computation of the closure of a set of attributes'.

In [38], we presented an attribute closure method closely tied to the SL_{FD} axiomatic system. We also showed that our method has a better performance than those based on classical closure. In this paper, we are going to use this method taking advantage of its novel characteristic.

Apart from having a strong base and good performance, one innovative feature of our method is that its output is twofold: besides the A_Γ^+ set constituting the closure of the input attribute set A , it also renders a reduced set of implications that encloses the semantics that is outside the set A_Γ^+ . We would like to remark that these two inputs are computed in linear time, because the subset of reduced implications is computed by the algorithm at the same time it computes the attribute closure.

The kernel of this closure method is the existence of three equivalences that can be enunciated by using SL_{FD} :

- Equivalence I: If $U \subseteq W$, then $\{\top \rightarrow W, U \rightarrow V\} \equiv \{\top \rightarrow WV\}$
- Equivalence II: If $V \subseteq W$, then $\{\top \rightarrow W, U \rightarrow V\} \equiv \{\top \rightarrow W\}$
- Equivalence III: If $U \cap W \neq \emptyset$ or $V \cap W \neq \emptyset$, then $\{\top \rightarrow W, U \rightarrow V\} \equiv \{\top \rightarrow W, U - W \rightarrow V - W\}$

The closure method works as follows. To compute the attribute closure of A with respect to Γ , the method is triggered by the seed formula $\top \rightarrow A$ and, exhaustively executing these three equivalences, it modifies this implication rendering $\top \rightarrow A_\Gamma^+$ and, at the same time, a reduced set of implications Γ' . The method is described in Algorithm 2.

We end this section with an illustrative example.

Example 2

Let Γ be the set of implications from Example 1. We show how SL_{FD} closure computes the closure of the attribute `Droplet`. In Table II, we show the application of the equivalences to each implication in the Γ set and how the set of attributes grows. We would like to remark that, in this example, only one repeat loop is needed.

As a final conclusion of this example, our method received the set Γ and `Droplet` attribute and renders as output the pair: $\{\text{Droplet}, \text{Direct}\}, \{\text{Blood} \rightarrow \text{Sexual}; \text{Fluids} \rightarrow \text{Vertical}; \text{Sexual} \rightarrow \text{Faecal}; \text{Vertical} \rightarrow \text{Sexual}\}$. That is to say, Droplet^+ and the reduced set of implications that stores the knowledge complementing the closure set.

Algorithm 2: The SL_{FD} Closure

```

Data:  $\Gamma, X$ 
Output:  $(X^+, \Gamma')$ 
1   begin
2      $\Delta := \langle X, \Gamma \rangle$ 
3     repeat
4        $\Gamma' := \Gamma$ 
5       foreach  $Y \rightarrow Z \in \Gamma$  do
6         if  $Y \subseteq X$  then /* Equivalence I */
7            $\Delta := \langle XZ, \Gamma \setminus \{Y \rightarrow Z\} \rangle$ 
8         else if  $Z \subseteq X$  then /* Equivalence II */
9            $\Delta := \langle X, \Gamma \setminus \{Y \rightarrow Z\} \rangle$ 
10        else if  $Y \cap X \neq \emptyset$  or  $Z \cap X \neq \emptyset$  then /* Equivalence III */
11           $\Delta := \langle X, (\Gamma \setminus \{Y \rightarrow Z\}) \cup \{(Y - X) \rightarrow (Z - X)\} \rangle$ 
12      until  $\Gamma' = \Gamma$ ;
13      return  $\Delta$ 

```

Table II. An application example of SL_{FD} closure.

Closure	Implication	New implication	
Droplet			
Droplet	$Blood \rightarrow Sexual$	$Blood \rightarrow Sexual$	-
Droplet, Direct	$Droplet \rightarrow Direct$	x	Equiv. I
Droplet, Direct	$Fluids \rightarrow Vertical$	$Fluids \rightarrow Vertical$	-
Droplet, Direct	$Respiratory \rightarrow Direct, Droplet$	x	Equiv. II
Droplet, Direct	$Direct, Sexual, Droplet \rightarrow Faecal$	$Sexual \rightarrow Faecal$	Equiv. III
Droplet, Direct	$Direct, Vertical \rightarrow Sexual$	$Vertical \rightarrow Sexual$	Equiv. III
Droplet, Direct	$Faecal, Sexual \rightarrow Direct, Droplet$	x	Equiv. II
Droplet, Direct	$Faecal, Direct \rightarrow Droplet$	x	Equiv. III

SL_{FD} , simplification logic.

4. Conversational process

Once we have presented our basis, in this section, we depict a detailed explanation of the dialogue process system along with a basic schema to facilitate the comprehension.

First of all, we depart from the premise that we have a dataset containing diseases and phenotypes and the set of implications that holds on it. This is considered the starting point at which this work begins and, as mentioned before, it stays out of the scope of this work. From there on, the dialogue process will go along through the following points:

- (1) Once we count on this information, the user starts interacting with the system by selecting a phenotype she suspects a disease to be related to. In this paper and with the intention of illustration, we are limiting the number of phenotypes to be selected to one at a time in each step of the conversation. This way, we can easily appreciate how the system performs. Yet a generalisation system considering more than one phenotype per step showed a similar behaviour but going faster.
- (2) Then, the process flows into the closure algorithm calculating both the phenotype's closure set for the selected one and also the reduced set of implications that corresponds with the complement of this closure.
- (3) Once the closure algorithm has finished, a first possible diagnosis is shown. This diagnosis shows the list of diseases in the dataset agreeing with the selected phenotypes. In order to acquire this list, the system launches a query to the database requesting those diseases that verify the selected phenotypes.
- (4) At this point, the user can stop the dialogue in the case that she is already satisfied with the result (a list of diseases), or she can go ahead trying to acquire a more reduced diagnosis.
- (5) Finally, the user can return to step 2 selecting new phenotypes until she acquires a satisfying set of disease result (possible diagnosis) or the system runs out of phenotypes.

The main contribution of this paper is the benefits of reducing the number of available phenotypes (in general, the attributes space), clarifying in each step the user scene. For further steps in the dialogue, we reduce the number of available phenotypes deleting those included in the closure set. As a consequence, some closure's phenotypes remain hidden to the user, in an intelligent and consistent way, because they are already implicitly included. This saves the user unnecessary efforts about information overload. However, even that this could be accomplished by classic closure algorithms, the major novelty of our method is that, *at the same time*, we also reduce

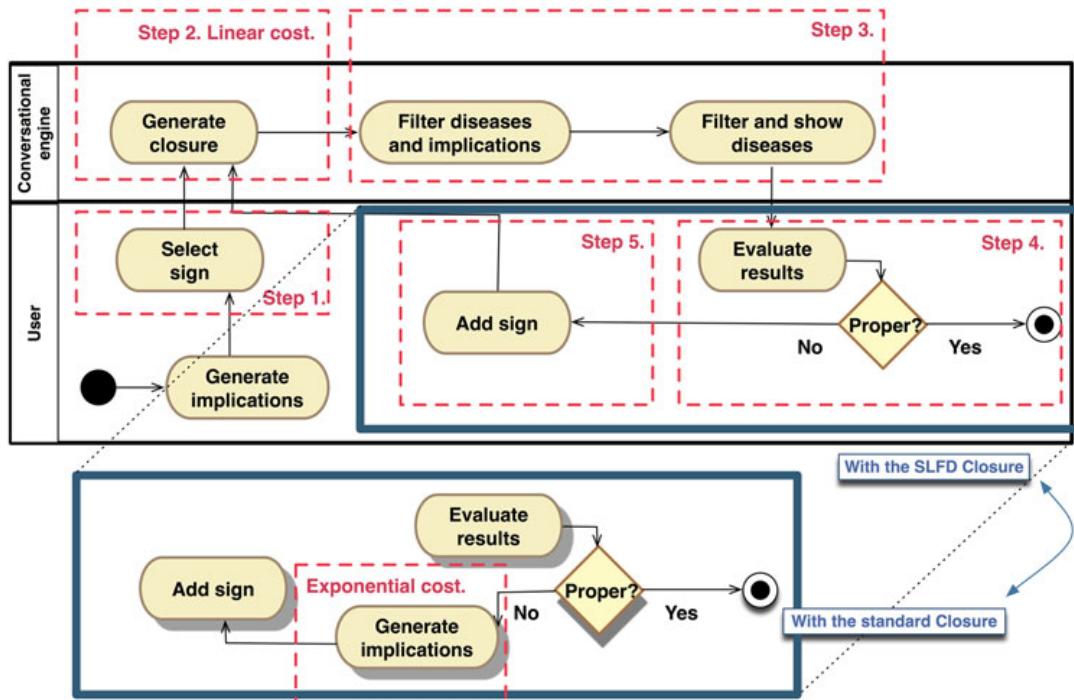


Figure 1. Dialogue process schema using simplification logic (SL_{FD}) closure versus standard closure.

the number of implications. This fact totally places us in a privileged position, overtaking the hard cost of a data mining process to extract the new set of implications for the reduced dataset after each searching step, as necessary when using classical implementations. On the contrary, in every refining attempt in the dialogue using our approach, we do not need to start the process from the beginning but continuing from there, where both symptoms and implications have been decreased. Consequently, the process overcomes the data mining costs preserving a linear complexity along the dialogue, and the interaction becomes truly faster. Figure 1 depicts the aforementioned steps by using the activity diagram of the UML language.

As mentioned earlier, in step 0, we need to count on the inferred set of implications within the dataset (generate implications activity). Although this is an exponential task, it is independent of the closure strategy we are willing to use. Thereafter, we execute step 1 and select the observed sign to continue with the execution of the closure method itself (generate closure activity). Observe that either our closure implementation or classical ones are able to perform in linear time. Now, however, it is remarkable that no matter how many steps we want the dialogue to go further, there is no need of mining any new set of implications; we already have it each time our closure implementation is applied because it is narrowing both attributes and implications at the same time thanks to the output of SL_{FD} closure (filter diseases and implications activity).

Nevertheless, in the case of classical closure implementations, the disadvantage arises as follows. Because only attributes are narrowed each time the closure applies, in order to proceed with the next step of the conversation, we need to generate a new set of implications accordingly to the new narrowed set of attributes. Therefore, this is an exponential task that is mandatory after every step of the dialogue. As a consequence, the overall complexity of the process becomes exponential.

Once the advantages of our method have been exposed, we proceed now with an example that illustrates the achievement of our approach about avoiding the exponential complexity of the conversational process.

Example 3

In the main experiment of this paper (which will be explained in Section 5.2), we have used a real dataset matching information about diseases and phenotypes. From the different runs of this experiment, five steps have been the maximal length the dialogue reached in a simulated conversation. Therefore, we are going to compare here this dialogue with such limit situation in two possible scenarios, the use of SL_{FD} closure and the use of any other classical closure implementations regarding the implication mining task. Here, we do not worry about the execution time of the closure method itself, because it is common to both approaches and not relevant compared with the implication mining cost.

This comparison is depicted in Table III. This table counts on five columns. First one indicates the number of steps along the conversation. Second one denotes the phenotype selected at each step of the dialogue. Columns 3 and 4 are the most important ones; they show the time needed to enumerate the new set of implications after each new step using classical closure implementations (column 3) and SL_{FD} closure (column 4). Last column shows the cardinal of the regenerated set of implications. At length, we show the time needed in the process using either a classical closure implementation or SL_{FD} .

Table III. Total time saved enumerating the set of implications by using SL_{FD} closure.

Step	Phenotype selected	Standard closure	SL_{FD} closure	Number of implications
1	HPO_7	22 s 181 ms	—	8.578
2	HPO_639	21 s 526 ms	—	8.270
3	HPO_2910	20 s 324 ms	—	7.988
4	HPO_1250	19 s 627 ms	—	7.534
5	Dialogue ends	—	—	—
	Total time saved	83 s 658 ms	—	—

SL_{FD} , simplification logic; HPO, Human Phenotype Ontology Consortium.

These measures have been obtained by virtue of the application of the specific package for R language named *Arules: Mining Association Rules and Frequent Itemsets*[‡]. Moreover, the hardware configuration used goes as follows: Intel Core 2 Duo 2.6 Ghz, 4 Gb RAM running over Windows 7.

The significant overall reduction of time obtained by using SL_{FD} closure is because of the linear calculation of the new set of implications to be used in the next step, whereas classical implementations suffer from regenerating implications again and again in each step.

5. Application of SL_{FD} conversational method to hematologic diseases selection

In this section, we describe the promising results obtained by our method in a real case. First, we establish the metrics defined to measure the benefits of our method and, later, we describe the selected dataset and the results of the experiment over it.

5.1. Evaluation metrics

When trying to enhance the interaction within a conversational system, evaluating the length of the dialogue for a typical query could be considered the most basic test [39]. Another popular measures to evaluate the effectiveness of recommender systems point to *precision* and *recall* measures [40]. Precision, defined as $P = TP / (TP + FP)$, where TP means the number of true positives, FP the number of false positives and FN the number of false negatives, determines the fraction of relevant items retrieved out of all items retrieved. On the other hand, recall $R = TP / (TP + FN)$, which determines the fraction of relevant items retrieved out of all relevant items. These two popular measures may not shed light on the matter of evaluating this approach, and the reason is twofold. First, every disease belonging to the list of resulting diagnoses agrees with the symptoms selected by the user as the database queries launched to retrieve diseases reflect these constraints. And second, after every loop of the process, we retrieve all the existing diseases in the dataset that hold with the symptoms selected, that is, all the relevant elements. Something similar occurs with other historical measures like mean absolute error or root mean square error. These two measures based on ratings have no place in this approach because no ratings are considered in order to conduct the conversation and the final items retrieval. In addition, there is no need to explicitly consider an accuracy measure because we do not build a prediction model, instead the use of implications ensures full accurate results. Fortunately, there are also others measures that could fairly reflect the statistics of the experiments. We continue describing those ones considered in this work.

5.1.1. Number of steps (N). This metric evaluates the actual length in steps of the conversation. It is interesting in the sense that it offers a rapid overview of the length of the interaction between the user and the conversational system. That gives an idea of whether the conversation has quickly satisfied the user or too many steps were needed. At this regard, within the following experiments, we will consider, without loss of generality, the user satisfied when a result of five (or less) diseases is returned. We set this limit because it deems advisable in order to constitute a proper diagnosis. Simultaneously, it represents the number of attributes (phenotypes) requested by the user because we are hitherto selecting one attribute at a time.

$$N = |\text{Selected attributes}|, \quad \text{where } |A| \text{ represents the cardinal of } A.$$

5.1.2. Speed of pruning at step i (S_i). This metric evaluates the percentage of the attributes the user is saving over the course of the conversation, accumulating from one step to another. When using this metric, we are willing to noticed whether the pruning rates have been better at the first steps of the conversation or at the last ones. Overall, it is a metric to measure how *fast* the system removes the overload of information. Notice that as mentioned before, we are taking one attribute at a time in this approach.

$$S_i = \frac{|\text{Attribute Closure}|_i - i}{|M|}, \quad i = 1, \dots, N, \text{ and } M \text{ represents the whole set of attributes as shown in Section 3.}$$

5.1.3. Attributes pruning (P). This last metric is equal to the speed of pruning but taking values at the end of the dialogue. It represents the percentage of the attributes that have been removed from the original set throughout the conversation. The pruning of these attributes is consequent because they are implicit by the user's selected ones. Formally,

$$P = S_N$$

Once all the basis have been defined and the metrics are explained and formulated, we are ready to begin the experiments.

[‡]<https://cran.r-project.org/web/packages/arules/index.html>.

Disease ID	HPO_1249	HPO_1250	HPO_1251	HPO_1252	HPO_1254	HPO_1257	...
274000		X					
275630	X		X				
277380				X		X	
300884		X			X		
300322	X				X		X
...							

HPO, Human Phenotype Ontology Consortium.

5.2. Experiments and results

In this section, we are going to perform experiments on a real-world dataset with a substantial amount of information. First of all, we are going to provide information about the dataset we are going to work with.

The source from which we have extracted the data is the Human Phenotype Ontology Consortium[§] (HPO). As can be read in their web page: '*HPO [41] aims to provide a standardized vocabulary of phenotypic abnormalities encountered in human disease. Each term in the HPO describes a phenotypic abnormality. The HPO is currently being developed using the medical literature, Orphanet[¶], DECIPHER^{||}, and OMIM^{**}. The HPO is developed within the context of the Monarch Initiative^{††}.*'

From this information, we have been able to generate a dataset on which we shall perform our experiments. Because the amount of information of HPO is huge, in this first approach, we are just going to use an extract of all the information available. In combination with Online Mendelian Inheritance in Man (OMIM) to distinguish amidst different types of diseases present in HPO databases, we have generated a table matching haematologic diseases and phenotypes, because the resulting information is substantial. Table IV shows an extract of the whole generated table. In the case we wish to obtain a detailed explanation of every disease shown forward, we commend the reader to visit OMIM web page and feed its search engine with the identifiers listed in Disease ID column. As an example, Disease ID 275630 corresponds to Chanarin-Dorfman syndrome.

Now, once we have presented our dataset, the next stage goes in using one of the previously mentioned techniques to retrieve all the implications that hold on it. Summing up, we are going to work over a dataset with 446 diseases, 100 different phenotypes and the set of implications that holds on it. Unfortunately, we cannot show all these implications here because of the obvious space limitations because the implications set goes beyond 6.000 implications.

In this point, we need to make an aside because from a purist view, if we calculate the so-called Duquenne-Guigues base [42] of implications that holds in context, there are 8.811 implications; so why have we pruned the set? Actually, the main feature of Duquenne-Guigues base of implications is that this base has a minimal possible number of implications among all possible bases of implications that hold in context. However, there will be several implications where there are no items that support (as stated in association rules theory) them, and usually such implications mean that set of items, contained in premise, does not occur together in context. Also, such implications include all attributes from context. Hence, the sense of this kind of implications is just theoretical and has nothing remarkable when dealing with real-world applications; that is why we ended up dismissing them. Nonetheless, the attributes present in these zero-supported implications are of course considered along the process and are fully accessible for the user to be selected.

That being said, the way on how we are going to proceed goes as follows. We are going to perform a test consisting of 1.000 simulations following to the letter the process depicted in Section 4. However, these runs will be carried out as random simulated dialogues. That is to say, we are going to conduct every dialogue by selecting phenotypes randomly from the set of available ones in each step. This strategy could give different readings. At a glance, it may deem advisable to go ahead with the random selection of phenotypes, so the reliability of the experiment is granted, and there is no possibility of inducing favourable situations for us to obtain better results. Yet the random strategy could also overshadow the benefits of our approach. On the one hand, imagine a situation where the random strategy falls into selecting phenotypes without any relation to each other. Then, the dialogue will end up quickly as there will be few diseases (or even no one) matching this phenotype's selection, and the process would finish with no possibility of applying any pruning at all. On the other hand, suppose a real dialogue where the phenotypes related to the patient share some sort of relation to each other. Then, during the dialogue, our process will be able to perform better pruning as the input phenotypes do have certain linkages; so, such a *fair-play* interaction would definitely highlight our approach. Needless to say that every result shown along with the experiments is the fruit of a statistical study [43] behind the results given from every run, so we kept the most reliable ones. Finally, in line with everything earlier, we are ready to launch the experiment.

At the end of the experiment, Figure 2 comes to clearly illustrate the results obtained for the number of steps metric after simulating 1.000 different conversations.

[§]<http://www.human-phenotype-ontology.org>.

[¶]<http://www.orpha.net/consor/cgi-bin/index.php>.

^{||}<https://decipher.sanger.ac.uk>.

^{**}<http://www.omim.org>.

^{††}<https://monarchinitiative.org>.

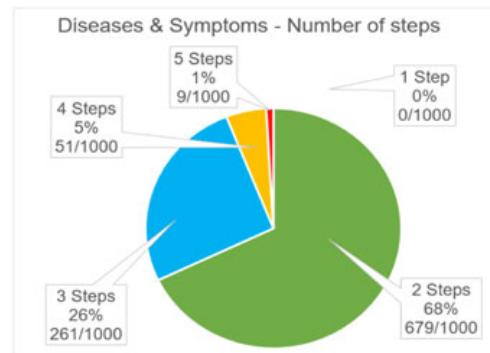


Figure 2. Number of steps metric results for the complete experiment.

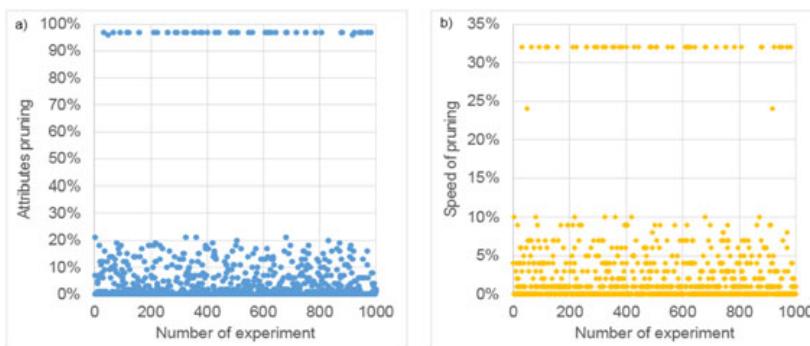


Figure 3. Attributes pruning (a) and speed of pruning (b) values for the complete experiment.

A glance is enough to easily realise that the conversational system is capable of guiding the dialogue to a suitable final diagnosis in two to three steps most of case. Keeping in mind the size of this dataset, these are encouraging results so far. There are other cases where the conversation has taken longer reaching four steps or even five, yet these cases are few and far between. Finally, only two experiments within the 1.000 runs have finished in just one step. These cases appear because the patient reveals a phenotype that appears in less than five of the diseases of the dataset and then the dialogue finishes at a glance (remember that for the random experiments, we established a limit of five or less diseases as the maximum to consider a diagnosis as a good option). Overall, we can consider that this number of step values fairly surpass previous studies [44] where the amount of input data provided is less than the one tested here.

Regarding the attributes pruning values, Figure 3(a) shows that in the vast majority of cases, the conversational system has freed us to worry about around 5–20% of attributes along the dialogue. In addition, there have been cases where the pruning did its utmost, reaching 97% of phenotypes, because the successive selected phenotypes conform a combination that brought the closure implementation to highly reduce the sets of both attributes and implications.

Speed of pruning values go hand-in-hand with the attributes pruning as it can be easily realised by matching Figure 3(a) and (b). Therefore, we can appreciate a general trend of experiments reducing attributes at an average speed of 5–10% per step and other ones achieving higher rates hovering 30–35% of attributes.

6. Conclusions and future works

We have presented here a novel application of our SL_{FD} closure algorithm in order to face the problem of the overwhelming dimensionality within the datasets. Our solution proposes a conversational process of user-driven feature selection. This work merges features of knowledge-based systems in combination with an appropriate management of implications through SL_{FD} closure. All these characteristics move us to favourably improve the diagnosis process with a high reduction in the length of the conversation and yet preserving the system's accuracy. That is not to forget the benefits we provide in terms of execution times reducing the conversational process from exponential to linear complexity.

Additionally, in the light of the results obtained over a dataset containing real information, we are of the opinion that this course of action heads the research to the right direction. The pruning rates reflect the good deeds of our approach improving the user–system interaction. Also, the number of step values encourage us to go straight on larger datasets because the actual numbers are admissible. Not forgetting, of course, the fact that results ensure 100% of accuracy.

Finally, our system constitutes a framework that can be integrated on diverse datasets and results stay admissible. This is certainly a major contribution of this work, because the possibilities it offers can reach many applications in different areas.

As future works, our results motivate a number of important directions for further research. Trying to discover which characteristics concerning the dataset (dimensionality, sparsity, etc.) are relevant to explain how the information extracted from the dataset behaves is an extended avenue for future researching tasks. In the same direction, identifying elements within the dataset, which play a more significant role amidst the others (either for being more frequent, unique, close-related to others, etc.), could help us to guide the conversation into the proper direction in a more comfortable way.

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Capítulo 6

Conclusiones y Trabajos Futuros

Fundamentalmente, podemos decir que esta Tesis ha englobado dos vertientes principales. Por un lado, se ha realizado un profundo estudio de los métodos basados en la lógica para el tratamiento eficiente de la información utilizando los conjuntos de implicaciones que se verifican en un determinado dataset. Y por otro lado, de forma más extensa, se han realizado las tareas de implementación necesarias para poder llevar estos métodos teóricos a la práctica. Hemos trabajado sobre tres campos diferentes: claves minimales, generadores minimales y sistemas de recomendación. Para cada uno de ellos se han realizado multitud de experimentos que demuestran la utilidad y la validez del trabajo realizado.

En esta tesis hemos podido apreciar el hecho de que contar con una sólida teoría basada en la lógica y las matemáticas nos concede la base para la creación de métodos automatizados con los que poder afrontar el desarrollo de aplicaciones de ingeniería. Como hemos podido comprobar, existe una gran cantidad de información implícita en los datos que solemos manejar. El descubrimiento de toda esta información y su gestión inteligente es sin duda un claro tema de investigación con fuerte actividad y repercusión en la actualidad. Esta ha sido nuestra intención a la hora de trabajar con FCA, los conjuntos de implicaciones y los operadores de cierre. Pasar de la teoría a la práctica y viceversa ha sido uno de los principales desafíos tanto

de esta tesis como lo es para el propio FCA si se pretende que se convierta en una herramienta fructífera para la representación, gestión y análisis del conocimiento en situaciones reales.

Antes de entrar plenamente en el apartado de conclusiones, queremos indicar la manera de certificar la validez de los resultados obtenidos a lo largo de la tesis. Como hemos podido advertir en los capítulos anteriores, nuestra labor se centra en actuar sobre conjuntos de implicaciones. En ese sentido, para los experimentos realizados hemos contado con unos ficheros de entrada que contenían la información necesaria, y sobre ellos hemos obtenido unos resultados. Ahora bien, la forma de verificar que esos resultados son correctos es la siguiente. En primer lugar y con respecto a los resultados de claves y generadores minimales, se han realizado y corregido numerosos ejercicios en papel intentando buscar casos límites donde la implementación pudiera no ser precisa y se ha comprobado que los resultados obtenidos en papel coincidían exactamente con los calculados por las implementaciones en la máquina. Además, dado que para muchos de los ejemplos probados en los que se llegaban a calcular millones de nodos de un árbol no era posible comprobar si cada uno de esos cálculos era correcto, para el caso concreto de los experimentos relacionados con claves minimales, la validez de los experimentos viene dada al haber cotejado los resultados con aquellos obtenidos sobre un amplio abanico de ficheros utilizados en trabajos anteriores [6, 7] donde su validez quedó demostrada. Básicamente, la validez de los ejercicios más grandes se ha extrapolado de los resultados correctos obtenidos para los ejercicios más pequeños. Asimismo, los resultados se corroboran igualmente al alcanzar las mismas soluciones para diferentes métodos cuando cada uno de ellos hace un tratamiento de la información diferente con respecto al otro. En relación a los experimentos con SR conversacionales, dado que los experimentos no alcanzan números tan altos, la validez puede demostrarse de forma más asequible siguiendo un desarrollo explícito en papel.

Y el otro punto que queremos remarcar es el siguiente. A lo largo de todo el proyecto siempre hemos hablado de trabajar sobre sistemas de implicaciones. No obstante, queda fuera del ámbito de esta tesis el procedimiento

mediante el cual se obtiene esos elementos para un sistema de datos. Para ello, encomendamos al avezado lector a visitar [48, 117, 118]. En nuestro caso, nuestro cometido comienza con el tratamiento de la información una vez de ha extraído el conjunto de implicaciones de un sistema concreto.

Aclarados estos puntos, nos encontramos ahora en el último capítulo de la Tesis, en el cual vamos a recopilar la conclusiones más importantes que hemos alcanzado como resultado del trabajo de investigación realizado. Seguidamente, cerrarán el capítulo una serie de tareas con las que continuar a partir de este punto y que se introducen como trabajos futuros.

Conclusiones

Como hemos mostrado anteriormente, conocer las claves es fundamental en cualquier modelo de datos. En este sentido, hemos presentado una serie de métodos que nos permiten averiguar el conjunto de claves a partir del conjunto de implicaciones y haciendo uso de métodos de razonamiento automatizado, en concreto, la *SLFD*. Se ha investigado, pasando desde la teoría a la práctica, los diferentes métodos implementados haciendo uso del paradigma de Tableaux y se ha comprobado como, hasta donde sabemos, los resultados obtenidos superan las aproximaciones anteriores.

Por otro lado, enumerar todos los conjuntos cerrados y sus generadores minimales es un problema muy complejo pero esencial en varias áreas de conocimiento y una oportunidad para mostrar los beneficios de FCA cuando trabajamos para aplicaciones reales. Para abordar esta tarea, se han presentado dos métodos de poda para mejorar el rendimiento de la enumeración de los generadores minimales. Para ello se ha hecho un uso intensivo de la *SLFD* sobre conjuntos de implicaciones. Finalmente, se han creado, analizado y probado enfoques diferentes (MinGen, MinGenPr, GenMinGen), mostrando claramente las mejoras alcanzadas por cada uno.

En ambas situaciones, es decir, tanto para claves minimales como para generadores minimales, se han desarrollado los códigos necesarios para poder actuar sobre grandes cantidades de información. No obstante, para resolver problemas reales donde la cantidad de información de entrada sea

considerable, es incuestionable la necesidad de unos recursos enormes tales como los que ha proporcionado el Centro de Supercomputación y Bioinnovación de la Universidad de Málaga; sin ellos habría sido inviable haber podido realizar gran parte de las pruebas. De acuerdo con esto, es absolutamente necesario que las implementaciones tengan en cuenta el correcto uso de recursos de memoria. Incluso para problemas pequeños, la cantidad de memoria que se puede necesitar puede dispararse escandalosamente.

Asimismo, el parecido y la cualidad de independencia que tiene cada nodo del árbol tanto para los métodos de claves minimales como para los de generadores minimales, ponen de manifiesto que la utilización de estrategias paralelas sea la mejor opción para abordar estos problemas. No obstante, el primer punto que queremos aclarar es que el concepto de paralelismo que hemos utilizado se refiere a un paralelismo de tipo *hardware*. Con esto nos referimos a que los beneficios que obtenemos del paralelismo se deben a la utilización de un conjunto de procesadores que se encargan de ir resolviendo cada uno de los subproblemas de forma simultánea. Por lo tanto, no estamos ante un caso de desarrollo de código paralelo desde una visión más purista, sino que es más acertado considerarlo como una aplicación basada en una estrategia *MapReduce* [30].

Para la mayoría de los casos, en relación a claves y generadores minimales, existe un serio inconveniente. En primera instancia es prácticamente imposible, por el momento, prever cuál va a ser la magnitud que va a alcanzar la resolución del problema a la vista de la información de entrada. Esto nos va a obligar a realizar una serie de experimentos previos de forma que podamos aproximar las necesidades que va a tener un determinado experimento.

Para cada una de las implementaciones realizadas, existía la necesidad de establecer criterios que nos permitieran evaluar el rendimiento de las pruebas de forma que pudiéramos comparar unos métodos con otros. En el caso de los experimentos relacionados con claves y generadores minimales, cuando nos planteamos la idea de la comparación de resultados, lo primero que se nos ocurrió fue la medición de los tiempos que necesitaba cada uno de los métodos para obtener los resultados. No obstante, advertimos como

este parámetro está íntimamente ligado a la arquitectura que estemos utilizando para ejecutar el experimento, lo cual hace que el resultado dependa en gran medida de los recursos que se están utilizando y no tanto de la calidad o eficiencia del propio algoritmo. En consecuencia, se oscurecía la utilidad teórica de los resultados obtenidos. Por tanto, se decidió contabilizar magnitud del árbol y la cantidad de resultados redundantes que se obtienen. De esta forma, si alguien hiciera otro método con un código en cualquier otro lenguaje o utilizase recursos *hardware* diferentes que desembocaran en una mejora del tiempo, siempre podríamos atenernos al tamaño del árbol pudiendo defender si realmente es una mejora en el método o en la ejecución debido a la arquitectura.

En cuanto a los SR conversacionales, también son varias las conclusiones a las que hemos llegado. La más importante es que, efectivamente, el tratamiento que realizamos de la información por medio de implicaciones y la *SLFD* puede aplicarse con éxito en este campo de conocimiento. Esto ya quedaba patente, como hemos mencionado con anterioridad, por la existencia de multitud de trabajos en la literatura de SR que utilizan conceptos de FCA; nuestro trabajo viene a reforzar este hecho.

Concretamente, presentamos una novedosa aplicación del algoritmo de cierre Cls para afrontar el problema de la sobrecarga de la información que a menudo presentan los SR. Nuestra solución propone un proceso conversacional de selección de atributos por parte del usuario. Este trabajo combina características de sistemas basados en contenido con sistemas basados en conocimiento mediante una gestión inteligente de las implicaciones a través del cierre Cls. Nuestro sistema constituye un marco que puede integrarse en diversos datasets y los resultados siguen siendo admisibles. Esto es, ciertamente, una gran contribución de este trabajo, pues ofrece la posibilidad de aplicarse a aplicaciones en diferentes áreas.

Otra conclusión importante es que existen numerosas técnicas de recomendación. De esta forma, una tarea fundamental en el desarrollo o en el análisis de un SR será identificar qué técnica es la más adecuada para su funcionamiento según el contexto de uso esperado. Del mismo modo, es excepcionalmente difícil abarcar todos los aspectos que involucran a un SR

para evitar los diferentes problemas que pueden aparecer, por tanto, hay que tener presente con qué expectativas queremos que actúe. Además, hay que tener en cuenta que son muy pocos los SR que basan su funcionamiento en una única técnica de recomendación; lo habitual es que sean sistemas híbridos con la intención de poder beneficiarse de las ventajas que ofrecen unas estrategias de recomendación y otras. Nuestro caso en un claro ejemplo; el tipo de SR desarrollado mezcla las estrategias de recomendación basada en contenido, basada en conocimiento y conversacional.

Por otro lado, existen numerosas opciones a la hora de evaluar el funcionamiento de un SR. Esta es una conclusión razonable en tanto en cuanto el número de técnicas diferentes con las que trabajan los SR es igualmente alto. Es fundamental decidir qué métricas son oportunas de aplicar dependiendo del tipo de SR que queramos evaluar, pues evidentemente habrá casos en los que una métrica no tenga cabida para un tipo de SR determinado. Asimismo, es recomendable no pretender obtener valores óptimos en la mayoría de las métricas. Hay que ser realista y darse cuenta de que es extremadamente difícil cumplir plenamente con unas y otras simultáneamente. La clave está en decidir cuál es el cometido principal de nuestro sistema, aplicarle las métricas de evaluación apropiadas y sacar las conclusiones pertinentes dentro de ese marco de actuación.

Finalmente, queremos remarcar que el sistema desarrollado es capaz de ir más allá del ámbito académico o de investigación. Las pruebas realizadas sobre *datasets* reales demuestran la viabilidad y la utilidad que el sistema puede aportar en entornos empresariales. De esta forma, apostamos de nuevo por una eficaz transmisión de conocimiento que acerque la empresa a la academia.

Trabajos Futuros

Finalmente, nuestros resultados motivan una serie de direcciones importantes para futuras investigaciones.

Respecto a las aportaciones referentes al tema de claves y generadores minimales existen varios aspectos con los que continuar a partir de este

trabajo de investigación. Los iremos introduciendo uno a uno sin perjuicio de que el orden de aparición denote una mayor o menos importancia.

En primera instancia es prácticamente imposible prever cuál va a ser la magnitud que va a alcanzar la resolución del problema a la vista de la información de entrada. Esto va a constituir en la mayoría de los casos un serio inconveniente.

Tanto en claves minimales como en generadores minimales hemos podido ver que existe un elemento fundamental en el diseño de las implementaciones paralelas: el valor de corte o BOV. Recordamos que el BOV es el valor a partir del cual la ejecución secuencial del código paralelo termina y se forman los diferentes subproblemas que serán resueltos en paralelo. Nos encontramos entonces con que establecer un valor de corte adecuado se convierte en una tarea realmente compleja. En primera instancia, hay que tener en cuenta la forma de expansión que tiene el Tableaux del experimento que estemos realizando y esto sólo lo podemos averiguar empíricamente. En definitiva, hay que buscar estrategias para encontrar un valor de corte tal que el aprovechamiento de los recursos computacionales sea óptimo.

Es necesario avanzar en el diseño de un *benchmark* que tenga en cuenta los diferentes aspectos y la naturaleza de estos problemas y algoritmos de manera que podamos dirigir las búsquedas de forma más eficiente.

Hay que profundizar en el hecho de que aumentar el número de cores para la resolución de un problema no siempre redunda en una mejora del rendimiento. Hay casos en los que aumentar los recursos utilizados puede ser incluso contraproducente como hemos comentado anteriormente.

Nuestra intención futura es aplicar el cálculo de claves y generadores minimales en más situaciones donde tratemos con datos reales, en especial a casos donde la información de entrada sea considerable, de forma que podamos valorar el rendimiento alcanzado gracias al paralelismo.

Otra tarea sobre la que continuar el trabajo consiste en investigar cómo realizar un nuevo diseño de los códigos paralelos que nos permita establecer comunicación entre las diferentes resoluciones paralelas de un mismo problema de forma que podamos mejorar en la reducción de los cálculos redundantes. Para el caso concreto del cálculo de los generadores minimales,

este mismo objetivo nos serviría para obtener una versión paralela del mejor método secuencial estudiado (GenMinGen).

En relación a los SR conversacionales, aparecen dos aspectos interesantes a tener en cuenta en el futuro próximo. En primer lugar, sería muy valioso poder identificar desde un primer momento qué elementos del dataset presentan una importancia superior entre los demás, ya sea por aparecer con mayor frecuencia, ser único, tener relación con un mayor número de elementos del conjunto, etc. En segundo lugar y siguiendo la misma idea, sería sin duda crucial saber qué características del dataset (tamaño, escasez, etc.) son las más influyentes en el rumbo que toman las conversaciones con el usuario. Tener un mayor conocimiento de estos dos factores nos permitiría poder influir de manera positiva en la conversación del sistema con el usuario y de esta forma, deparar en una mejor experiencia de usuario.

Asimismo, hay aspectos de los SR que sería recomendable investigar si sería acertado incluir en el sistema conversacional; tal puede ser el caso de proporcionar explicaciones que justifiquen las recomendaciones que el usuario ha recibido. Este es un aspecto importante en un SR, ya que ayuda a mantener un mayor grado de confianza del usuario en los resultados generados por el sistema [96].

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Anexo

Apéndice A

Closed sets enumeration: a logical approach

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Closed sets enumeration: a logical approach.

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Abstract

Closed sets are the basis for the development of the concept lattice, a key issue in formal concept analysis. The enumeration of all the closed sets is a complex problem, having an exponential cost. In addition to the closed set, it is very useful for applications to add the information of all the minimal generators for each closed set. In this work we explain how to approach this problem from a complete set of implication by means of a sound and complete logic.

Key words: Formal concept analysis, closed sets, minimal generator, logic.

1 Introduction

Formal concept analysis (FCA) is a theoretical and practical framework to store information and manage them [GW99]. Data is stored in a table, representing a binary relation between a set of objects and attributes. The success of FCA relies on its solid theoretical framework and a wide set of methods and techniques to extract the knowledge from this data and manipulate it. One outstanding representation of the knowledge is the concept lattice, built over the closed sets, considering the subset relation as the order relation. Such representation depict a overall view of the information with a very strong formalism, opening the door to use the lattice theory as a metatheory to manage the information [BDVG17].

If-then rules have been introduced in several areas, dressed with different clothes. Thus, in relational databases [Cod71] they are named Functional Dependencies, in FCA they are named Implications and in Logic Programming (fuzzy logic) [BV06a] they are named if-then rules. All this notions captures a very intuitive idea: when the premise occurs, then the conclusion holds. Nevertheless, their semantics are very different and they further use are

also distinct. In this work we consider implications as elements to describe the information and we design a method to enumerate all closed sets and their minimal generators.

The proposed method is an evolution of [CEMO12], where the authors introduce a logic-based method based on \mathbf{SL}_{FD} , a sound and complete logic for implications. That method works by traversing the set of implications and applying a set of inference rules, following a tree paradigm in its execution. In that method, an exhaustive search was developed, producing the intended result but with an improvable performance. Here, we propose the design of several pruning strategies to improve such performance. These strategies are motivated by the idea of avoiding the opening of full branches in the tree or reducing the size of the information in their nodes.

The rest of the work is organized as follows: in the following section we present \mathbf{SL}_{FD} and the axiomatic system which constitutes the basis of the MinGen algorithm. In Section 3 we present the algorithm to enumerate the closed sets and minimal generators and summarize the strategies to improve its execution in practice. The work ends with a brief conclusion.

2 Logic for implications

First, the main notions related formal concept analysis needed in this work are presented.

A **formal context** is a triple $\mathbf{K} := (G, M, I)$ where G is a set of objects, M is a set of attributes and $I \subseteq G \times M$ is a binary relation between G and M such that, for $o \in G$ and $a \in M$, $o \ I \ a$ means that the object o has the attribute a . Then, two mappings are defined $(\cdot)': 2^G \rightarrow 2^M$ defined for all $A \subseteq G$ as $A' = \{m \in M \mid g \ I \ m \text{ for all } g \in A\}$, and $(\cdot)'': 2^M \rightarrow 2^G$ defined for all $B \subseteq M$ as $B'' = \{g \in G \mid g \ I \ m \text{ for all } m \in B\}$. We use the same symbol since no confusion arises. This pair of mappings is a Galois connection.

The composition of the intent and the extent mappings, and vice versa, introduces two closure operators $(\cdot)'': 2^G \rightarrow 2^G$ and $(\cdot)': 2^M \rightarrow 2^M$. The notion of closed set (as a fixpoint of a closure operator) is defined as follows:

Definition 1 A **formal concept** is a pair (A, B) such that $A \subseteq G$, $B \subseteq M$, $A' = B$ and $B'' = A$. Consequently, A and B are closed sets of objects and attributes, respectively called *extent* and *intent*.

In this work we focus on the attributes closed sets. A key point in this work is the notion of the minimal generator (mingen) [GW99], which provides a minimal representation for each closed set, and is defined as follows:

Definition 2 Let $\mathbf{K} = (G, M, I)$ be a formal context and $A \subseteq M$. The set of attributes A is said to be a minimal generator (**mingen**) if, for all set of attributes $X \subseteq A$ if $X'' = A''$ then $X = A$.

Remark that the above definition allows to characterize each closed set by means of a minimal subset to provide a canonical representation of the closed sets. Moreover, we would like to remark that such representation is not unique, since a given closed sets can have several minimal generators.

The notion of minimal generator can also be defined from the point of view of implications. They are expressions $A \rightarrow B$ where A and B are attribute sets. A context satisfies the implication $A \rightarrow B$ if every object that has all the attributes from A also has all the attributes from B .

Definition 3 An (attribute) implication of a formal context $\mathbf{K} = (G, M, I)$ is defined as a pair (A, B) , written $A \rightarrow B$, where $A, B \subseteq M$ and $A \cap B = \emptyset$. Implication $A \rightarrow B$ holds (is valid) in \mathbf{K} if $A' \subseteq B'$.

The set of all valid implications in a context satisfies the well-known Armstrong's axioms [Arm74], which constitutes the pioneer logic to manage implications. The author introduces a sound a complete axiomatic system to infer new implications holding in a context from a given set of implications. Moreover, this logic constitutes a proposal to solve the attribute closure, i.e. to find the maximal set of attributes A^+ such that the implication $A \rightarrow A^+$ holds. As we mentioned, this maximal set is a closed set as defined before and this closure operator $(\cdot)^+$ allows us to guide the automatization the search for closed sets. Thus, we introduce a new logic suitable for this goal.

The introduction of the Simplification Logic [MECF12], named \mathbf{SL}_{FD} , opened the door to the development of automated reasoning methods directly based on its novel axiomatic system. \mathbf{SL}_{FD} considers reflexivity as axiom scheme

$$[\text{Ref}] \quad \overline{A \rightarrow A}$$

together with the following inference rules called Fragmentation, Composition and Simplification respectively.

$$[\text{Frag}] \quad \frac{A \rightarrow BC}{A \rightarrow B} \quad [\text{Comp}] \quad \frac{A \rightarrow B, C \rightarrow D}{AC \rightarrow BD} \quad [\text{Simp}] \quad \frac{A \rightarrow B, C \rightarrow D}{A(C \setminus B) \rightarrow D}$$

Similarly to the dual vision of closed set (in terms of Galois connection and implications), a dual definition of minimal generator can be done in terms of implications:

Definition 4 Let $\mathbf{K} = (G, M, I)$ be a formal context and $A \subseteq M$. The set of attributes A is said to be a minimal generator (**mingen**) if, for all set of attributes $X \subseteq A$ if $X \rightarrow A^+$ then $X = A$.

In the following section we introduce the algorithm to enumerate the minimal generator based on \mathbf{SL}_{FD} and describe the strategies to improve its performance.

3 An algorithm to enumerate all closed sets and their minimal generators

Simplification logic has allowed us to design several executable methods to manage implications. Thus in [MECF12] we developed a novel method to compute attribute closure strongly based on \mathbf{SL}_{FD} inference rules. This method has been showed to have a better performance than the classical methods based on indirect techniques. One outstanding characteristics of \mathbf{SL}_{FD} closure is the output it renders: given a set of attributes $X \subseteq M$ and a set of implications Γ , it renders its closure X^+ and a new set of implications Γ' which describes the remaining knowledge in the set $M \setminus X^+$.

This logic-based closure method is the basis of another method, named MinGen, to compute the set of all minimal generators from a set of implicant set presented in [CEMO12]. The algorithm works by applying the \mathbf{SL}_{FD} Closure algorithm to each implication in the set, opening a new branch. This application provides a new candidate to be added to mingens and a smaller implications set which guides us in the search of new sets of attributes to be added to mingens, producing a tree-like execution.

In summary, the input of this algorithm is a set of attributes M and a set of implications Γ over the attributes in M . The output is the set of closed sets endowed with all the minimal generators, i.e. $\{\langle C, mg(C) \rangle : C \text{ is a closed set of attributes}\}$ where $mg(C) = \{D : D \text{ is a mingens and } D^+ = C\}$. In this work we only consider non-trivial minimal generators, i.e. pairs of closed set and minimal generator $\langle X, Y \rangle$ where $Y \subsetneq X$.

For example, if $M = \{a, b, c, d, e, f\}$ and $\Gamma = \{a \rightarrow b, bc \rightarrow d, de \rightarrow f, ace \rightarrow f\}$ the output is the set $\{\langle abcdef, \{ace\} \rangle, \langle abdef, \{ade\} \rangle, \langle abcde, \{ac\} \rangle, \langle bcd, \{bc\} \rangle, \langle def, \{de\} \rangle, \langle ab, \{a\} \rangle, \langle c, \{c\} \rangle, \langle \emptyset, \{\emptyset\} \rangle\}$. The execution of the method is depicted in Figure 1. We refer the reader to [CEMO12] for a detailed description of the method and its theoretical results.

In this work, we propose two pruning strategies to improve MinGen method. We briefly describe them as follows:

- The first strategy characterizes the branches that can be considered a superfluous one because all their nodes explore closed sets and minimal generators already considered in another branches. To implement this strategy we will consider a subset test on the branches of the same level and in the same branch.
- The second strategy is to expedite the execution of the method by including the closure method in each node in two steps, so that in the first one the closure set is computed and, in the second one, the resulting set of implications is computed taking into account this closed set. In this way, the resulting set of implications will be a smaller one and the method will have a better performance.

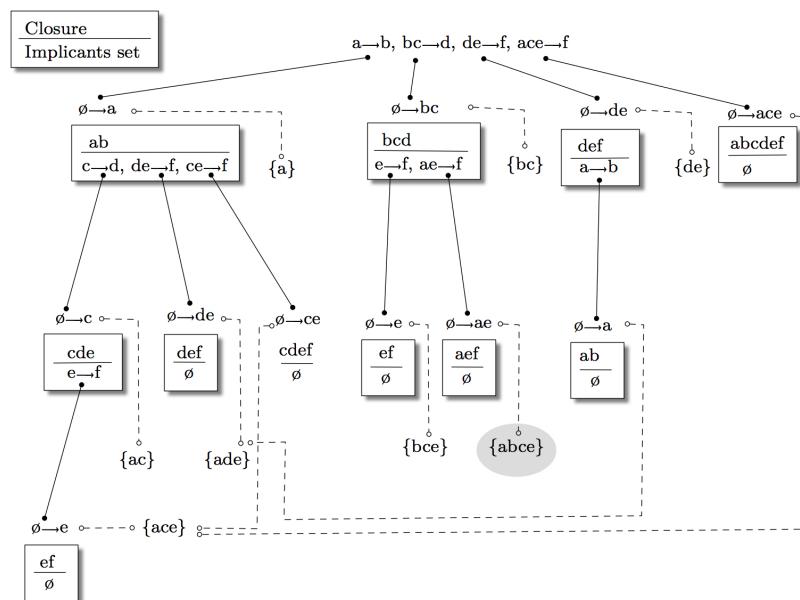


Figure 1: Exemplification of

4 Conclusion and future works

In this work we have studied the state of the art in the enumeration of closed sets and minimal generators based on logic. We have considered the MinGen method based on Simplification Logic as the target of our work and we propose to improve it by means of several prunes to improve its performance.

In a future work, we propose to establish the theoretical results to state these strategies and to develop an exhaustive practical experiment to show its benefits.

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Apéndice B

Conversational recommendation to avoid the cold-start problem

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Conversational recommendation to avoid the cold-start problem

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Abstract

Recommender systems has become a widespread topic, allowing to connect user demands to those products more suitable to their preferences. The more information we provide to the system, the better the system works. This is a weak point of recommenders: they need an initial information belonging to each new user. In this paper we propose to avoid the so-called cold-start problem by using a conversational recommendation approach. We consider products characteristics as attributes and deal with the attribute implications by means of the simplification logic to guide the user in the search.

Key words: *Recommendation systems, conversational recommendation, logic, implications*

1 Introduction

Nowadays recommender systems have established a solid field of knowledge within information technologies. They are a kind of software that group together a wide range of techniques and applications with the aim of providing the best user experience [18]. There has been much progress done towards recommender systems during last decade [1] but there is still so much work remained. Examples of the applications concerning recommender systems go over many different topics of today's society such as recommending books, music, documents, e-commerce, tourism, medical diagnosis, among others. Recommender Systems constitute a hot topic indeed, as we can notice by the way in which many top companies worldwide spend their efforts and resources developing more and better systems for them to

increase their benefits. By these means, companies with absolutely different market niches delegate their most important duties to recommender systems due to the wide range of possibilities they offer; and yet companies selling products are not all of them, global leaders in other fields as totally come aboard with recommender systems by recommending new friends, groups, followers, and other social connections.

When recommendations are based on the element evaluation made by other users or by similarity between the user preferences and these characteristics, recommender systems need to face many problems before they can flow into good recommendations. The first one we need to remark is the well-known cold-start problem [10], that appears when recommender systems try to elaborate reliable recommendations from the absence of initial information. Cold-start problem may be handled by requesting other agents to share what they have already learned from their respective users [11]. Also, new items (those which have not received any ratings from the community yet) would be assigned a rating automatically, based on those given by the community to other similar items [20] and so, we are at the mercy of similarity rules. In the same direction, until the new element has not been evaluated by a significant number of users, the system will not be able to recommend it. An item that is not recommended remain unnoticed by most of the user community, thus, we can enter into a vicious circle in which a set of elements of the recommender systems will be left out of the rating process and/or recommendations continuously [16]. In most of cases, users do not rate all the features we would desire for the optimum running of the recommender systems, this reveals scarcity problem.

In this work, we propose to deal with the cold-star problem by introducing an information flow based on the dialogue with the user. The lack of initial information is avoided with the design of a process with allows to collect this information of the user and storing them for further access to the system. This process, as we shall see, is a recommender-like system, to allow the user for getting some usefulness in its use.

2 Recommender systems and the conversational issue

There exists different kinds of recommender systems usually classified on how recommendations are made [1]. The most known and extended ones are collaborative filtering, content-based and demographic systems. Besides, in recent years there has been a great expansion of context-aware recommender systems [2] and knowledge-based recommender systems [14]. Other group of recommender systems that worths to be considered is that one focused on recommendations involving group of people [9]. Collaborative filtering systems [13], recommend items that other users have already rated before. Recommendations made by content-based systems present items similar to the ones the user preferred in the past [12]. Context-aware recommender systems try to adapt their recommendations to the world around the user. Finally, knowledge-based approaches are different; they manage functional

knowledge about how an item matches a particular need, and they can therefore reason about the relationship between a need and a possible recommendation. These characteristics make knowledge-base recommender systems not only valuable systems on their own, but also highly complementary to other types of recommender systems. However, the history of recommender systems has broadly demonstrated that best strategies are those who merge characteristics from different kinds of recommender systems in order to generate hybrids conforming best features of each one [6, 4].

In general, most of widely used recommendation techniques requires information to build a user profile before generating a result. In some cases, that information may be gathered explicitly: for example, requiring data about age, gender, etc. during a registration process, or by means of ratings and opinions about the recommended items. In other cases, the system may get implicit information from the browsing and/or purchase user history.

Nevertheless, there are contexts in which this previous information it is not available. This is the case of the well-known cold-start problem, when a new user asks for his first recommendation and obviously the system has not any information about him. This situation also occurs in systems where users make occasional use.

An interesting approach to solve this problem us the use of the so-called conversational recommender systems [7, 8]. These are closely related with critiquing recommender systems [17, 21]. In these works, recommendation is enriched by means of a dialog with the user that allows an incremental elicitation of his preferred item features. To promote an effective use of this approach, our proposal produces as an output a recommendation only based on the user dialogue information. In this way, the system is attractive for those user that are new in the system and can be used as a preliminary system to store user preferences for further accesses.

3 A logic approach to conversational recommendation

Our proposal to integrate recommender systems and the conversational issue is based on a sound and complete logic. As we shall see, such an strong basis allows us to include a reasoning method in the process and allows us to store the information in a natural way to be managed in the future by knowledge-base recommenders.

We built our framework on a basic elements, the implications. They correspond to formulas $a_1 \wedge \dots \wedge a_n \rightarrow b_1 \wedge \dots \wedge b_m$. The propositions $a_1, \dots, a_n, b_1, \dots, b_m$ are elements of a set Ω and they are interpreted as properties concerning attributes. For this reason, propositional symbols are named attributes. To compact notation it is usual to denote the above formulas as $A \rightarrow B$ being $A = \{a_1, \dots, a_n\}$ and $B = \{b_1, \dots, b_m\}$ i.e. sets of attributes are conjunctively interpreted.

The symbolic management of implications was originally proposed in [3]. However, due to the central role that transitivity plays in this axiomatic system, the development

of executable method to solve implications problems has rest on indirect methods. For instance, the proposal to solve the attribute closure, i.e. to find the maximal set of attributes A^+ such that the implication $A \rightarrow A^+$ holds has been traditionally tackle by using a basic method which exhaustively uses the subset relation to add new elements in the conclusion.

The introduction of the Simplification Logic, named \mathbf{SL}_{FD} , [5] opened the door to the development of automated reasoning methods directly based on its novel axiomatic system. \mathbf{SL}_{FD} considers reflexivity as axiom scheme

$$\text{[Ref]} \quad \overline{A \rightarrow A}$$

together with the following inference rules called Fragmentation, Composition and Simplification respectively.

$$\text{[Frag]} \quad \frac{A \rightarrow BC}{A \rightarrow B} \quad \text{[Comp]} \quad \frac{A \rightarrow B, C \rightarrow D}{AC \rightarrow BD} \quad \text{[Simp]} \quad \frac{A \rightarrow B, C \rightarrow D}{A(C \setminus B) \rightarrow D}$$

Later, in [15] we presented an attribute closure method closely tied to the Simplification logic axiomatic system. Apart from having a strong base, the main advantage of our method is that its output is twofold: besides the maximal set constituting the closure of the input, it also renders a reduced set of implications which enclose the semantics that is outside the set A^+ . We would like to remark that this two inputs are computed in linear time, overtaking the hard cost of a data mining process if we were interested in extracting the new set of implication for the reduced dataset after each search step.

This characteristics provides a key information to further inferences in an iterative search process. This is the core of our proposal to design a conversational recommendation based on our attribute closure operator. The recommendation process will go along the following points:

0. We depart from the premise that we have a dataset containing items and attributes, and the set of implications that holds on it. This is considered point zero and, as we have mentioned, it does not requieres any information from the user to be started.
1. Once we count on this information, the user interacts with the system by selecting an attribute we wish an item to fit.
2. Then, the process flows into the closure algorithm calculating both the set closure for this attribute and above all, the set of implications that remains outside the closure and complete them.
3. Once the closure algorithm has finished, a new reduced dataset is shown. At this point, we can stop the interaction whether we are already satisfied with the result or we can go ahead trying to get a more suitable recommendation. The improvement here goes as follows. For further queries, we have reduced the number of available attributes

deleting those included in the closure set. Even that this could be accomplished by classic closure algorithms, the major point of our method is that, *at the same time*, we also reduce the number of implications, and so, in every refining-attempt we do not need to start the process from the beginning but continuing from here, where both attributes and implications have been decreased. Consequently, the process maintains its linear complexity and the interaction becomes truly faster.

4. In this way, we select a new attribute and resume the search.
5. We carry on selecting attributes until we get a satisfying recommendation or we run out of attributes.

4 Conclusion and future works

In this paper we propose to approach the cold-start problem. We mining the dataset containing the product information to get a set of attribute implications. This set is managed by using the inference system of simplification logic to guide the search of new users.

As a future work, we propose to study the impact of simplification closure in the performance of our approach. Our method allows to get, in an iterative way, intermediate closure set of attributes and the corresponding reduced set of implications. This characteristics allows to proceed step by step and, at the same time, accelerate the search.

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Apéndice C

Keys for the fusion of heterogeneous information

Keys for the fusion of heterogeneous information

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Abstract

The management of heterogeneous information is a current topic which demands the use of intelligent techniques to deal with data semantics. In this work we approach this problem by using Simplification Logic. It has a sound and complete inference system conceived to treat implications and functional dependencies. The automatic processing of functional dependencies allows to develop methods and tools to tackle most classical problems in database and information processing. In this work, we use Simplification Logic to design a method to enumerate all minimal keys of a data repository inferring them from a set of functional dependencies. We also illustrates how this method provides a successful way to solve some outstanding problems in data processing in linked data.

Key words: Heterogenous information, integration, keys, logic.

1 Introduction

The notion of a key is fundamental to any data model, including Codd's relational model of data [Codd, 1970]. A key of a relation schema is composed by a subset of attributes playing the role of a *domain* in a given function whose *image* is the whole set of attributes. This way, the table is viewed as its extensional definition. These functions are described by means of a *Functional Dependency (FD)* which specifies a constraint between two subset of attributes, denoted $A \rightarrow B$, ensuring us that for any two tuples in a table, if they agree on A , they also agree on B .

Besides primary keys, unique constraints (candidate keys) provide a more complete understanding of the model. Keys are not only fundamental to data design, they are also considered powerful tools to solve several problems related to a lot of fields of information management as mentioned before. Indeed, in [Sismanis et al., 2006] the authors affirm

that “*identification of keys is a crucially important task in many areas of modern data management, including data modeling, query optimization (provide a query optimizer with new access paths that can lead to substantial speedups in query processing), indexing (allow the database administrator to improve the efficiency of data access via physical design techniques such as data partitioning or the creation of indexes and materialized views), anomaly detection, and data integration*”.

Identifying properly the keys of a relation schema is a crucial task for a lot of modern areas of information management (data modeling [Simson and Witt, 2005], query optimization [Kemper and Moerkotte, 1991], indexing [Manolopoulos et al., 1999], etc).

A very outstanding characteristic of keys is their minimality. We denote a key as minimal when every attribute contained in its attribute set is necessary to keep the property of key, i.e. keys with no superfluous attributes. Thus, in the literature, the key finding problem is focused on minimal keys. Giving an attribute set A, the cardinality of the set 2^A forces us to consider in applying special techniques that lead the search of the sets being candidates to become minimal keys.

Keys constraints specify the sets of attributes of a relation such that their projection univocally identifies each tuple of the relation. The key finding problem consists in finding all the attribute subsets which make up a minimal key as of a set of FDs occurring within a schema of a relational model table.

In Table 1, we illustrate its semantics by the following basic example.

Table 1: Movie table

Title	Year	Country	Director	Nationality	Star
Pulp Fiction	1994	USA	Quentin Tarantino	USA	John Travolta
Pulp Fiction	1994	USA	Quentin Tarantino	USA	Uma Thurman
Pulp Fiction	1994	USA	Quentin Tarantino	USA	Samuel L. Jackson
King Kong	2005	New Zealand	Peter Jackson	New Zealand	Naomi Watts
King Kong	2005	New Zealand	Peter Jackson	New Zealand	Jack Black
King Kong	1976	USA	De Laurentiis	IT	Jessica Lange
King Kong	1976	USA	De Laurentiis	IT	Jeff Bridges
Django Unchained	2012	USA	Quentin Tarantino	USA	Jamie Foxx
Django Unchained	2012	USA	Quentin Tarantino	USA	Samuel L. Jackson

From the information in Table 1, we may ensure that the following FDs are satisfied: $Title, Year \rightarrow Country$; $Title, Year \rightarrow Director$; $Director \rightarrow Nationality$. Moreover, the table has only one minimal key: $\{Title, Year, Star\}$

It is imperative to state that though we are dealing with FDs, it is not the responsibility of this work to extract them from a relation schema. There are several techniques that provide us this work [Huhtala et al., 1999], [Yao et al., 2002]. Therefore, we will begin with given sets of FDs and then go ahead finding minimal keys within them.

In the same way, it is necessary to clarify that what we are researching is not a matter of data mining techniques [Fayyad et al., 1996]. Giving an overall view, data mining could be considered as a computational process of discovering patterns in large data sets and its

goal goes to extract information from a data set and transform it into an understandable structure for further use [Witten et al., 2011]. Nevertheless, what we are studying and developing are mechanisms and algorithms for deriving all minimal keys so they can help us performing a more intelligent and efficient way to manage the information stored in relational schemes.

In this work we will concentrate our efforts on those algorithms guided by logic, and most specifically, those using Tableaux paradigm [Morgan, 1992] [Risch and Schwind, 1992] for deriving keys of a relation schema using inference systems.

The problem of the key finding methods arises when we try to deal with a huge amount of information since the tableaux's building mechanism produces such an explosion of the search space that we go beyond the machine capabilities, even with small problems.

2 Background

Before going further, we need to introduce the necessary terms from relational database theory. Due to space limitation, we refer those readers non familiar with the basic notions of FDs and relational databases to previously visit [Elmasri and Navathe, 2010]. The notion of FDs are well-known in other areas as Formal Concept Analysis with the concept of implications.

Definition 1 (Attribute) *An attribute a is an identifier for an element of some domain D . We use letters a, b, c, d, \dots for attributes. Let U be a set of attributes. An attribute set X over U is a subset of U . We use capital letters X, Y, Z, V, \dots for attribute sets.*

Definition 2 (Functional dependency) *Let U be a set of attributes. A functional dependency (FD) over U is an expression of the form $X \rightarrow Y$, where X, Y are attribute sets. It is satisfied in a table R if for every two tuples of R , if they agree on X , then they agree on Y .*

Definition 3 (Relation schema) *A relation schema $R = \langle U, F \rangle$ is an ordered pair consisting of an attribute set U and a set F of FDs over U .*

A key of a relational table is a subset of attributes that allows us to uniquely characterize each row. It may be defined by means of FDs as follows:

Definition 4 (Key) *Given a table R over the set of attributes U , we say that K is a key in R if the functional dependency $K \rightarrow U$ holds in R .*

Definition 5 (Minimal Key) *Given the table R , the attribute set $K \subset U$ is said to be a minimal key if it is a key of R and for all attribute $k \in K$ the subset $K - \{k\}$ is not a key of R .*

3 Simplification Logic for finding-key problem

3.1 SL_{FD}

Now, we summarize the axiomatic system of SL_{FD} [Cordero et al., 2013]. The inference system for SL_{FD} is equivalent to the well-known Armstrong's axioms as the first complete inference system for functional dependencies. It avoids the use of transitivity and is guided by the idea of simplifying the set of FDs by removing redundant attributes efficiently.

SL_{FD} is defined as the pair (L_{FD}, S_{FD}) where S_{FD} has the following axiom scheme and inference rules. The third rule is named Simplification rule and it is the core of SL_{FD} :

$$[Axiom] \quad \frac{Y \subseteq X}{X \rightarrow Y}$$

$$[Frag] \quad \frac{X \rightarrow Y}{X \rightarrow Y'}, \quad [Comp] \quad \frac{X \rightarrow Y \quad U \rightarrow V}{XU \rightarrow YV}$$

$$[Simp] \quad \frac{X \rightarrow Y \quad U \rightarrow V}{(U - Y) \rightarrow (V - Y)'} \quad X \subseteq U, X \cap Y \neq \emptyset$$

3.2 SST Method

The method we will use to find all minimal keys in schemes provided from two heterogeneous sources is SST method (see [Cordero et al., 2014] for more details). The input of the tableaux method is a set of attributes Ω and a set of formulas F . We build a tree as follows:

1. The root of the tree will be $(\Omega \rightarrow \Omega, F)$.
2. For each node $(U \rightarrow \Omega, F)$ with $U \neq \emptyset$ and each minimal formula $A \rightarrow B \in F$, a new children node $(U' \rightarrow \Omega, F')$ is added where:
 - $U' \rightarrow \Omega$ is obtained applying [lSimp] to $A \rightarrow B$ and $U \rightarrow \Omega$. That is, $U' = A \cup (U \setminus B)$.
 - F' is computed by applying [sSimp] to $A \rightarrow B$ and every formula in F . Moreover, in the new set of formulas, Union equivalence is applied and degenerated formulas are removed.
3. The method renders $Minimal\{U | (U \rightarrow \Omega, \emptyset)\}$ is a leaf of the tree}.

4 Applications of keys in Linked Data

There are a lot of fields of knowledge in computer science where counting on efficient techniques for data management is crucial. Generally, it is an engineering work to establish and choose the keys as a part of the normalization process of the schema. The challenge is to figure out those attributes of the schema that allow identifying univocally each tuple of the relation. Lets show an easy example.

Example 1 Assume that we have stored in a table the main data of an enterprise crew such as: Name, Age, ID Number, Phone Number. *A priori*, we notice several alternatives in order to identify each employee.

As a key we could consider: ID Number, the pair (Name, Phone Number) or even (ID Number, Name). However, from all of that possibilities, ID Number arises as the best one since the other ones contain information that is not absolutely necessary to identify each person. Two people with different names could share the same phone number (members of a same home will share the same phone number). Pairs (ID Number, Name) will also identify properly each person, but the Name attribute is not indispensable since ID Number will be definitely enough as identification.

So, even in this trivial example we can easily notice that deriving keys is a task with an indisputable importance. Lets go deeper in details with a bit more realistic problems.

Linked Data is about connecting pieces of related data and information coming from different sources. In computing, linked data describes a method of publishing structured data so that it can be interlinked and become more useful. It builds upon standard Web technologies such as HTTP, RDF¹ and URIs, but rather than using them to serve web pages for human readers, it extends them to share information in a way that can be read automatically by computers. This enables data from different sources to be connected and queried [Bizer et al., 2009].

A typical case of a large Linked Dataset is DBpedia, which, essentially, makes the content of Wikipedia available in RDF. The importance of DBpedia is not only that it includes Wikipedia data, but also that it incorporates links to other datasets on the Web, e.g., to Geonames. By providing those extra links (in terms of RDF triples) applications may exploit the extra (and possibly more precise) knowledge from other datasets when developing an application; by virtue of integrating facts from several datasets, the application may provide a much better user experience.

For instance, the DBpedia resource for Brussels (<http://dbpedia.org/resource/Brussels>) can be linked to the one maintained by the Statistics Belgium

¹RDF, the Resource Description Framework, is one of the key ingredients of Linked Data, and provides a generic graph-based data model for describing things, including their relationships with other things. RDF data can be written down in a number of different ways, known as serialisations. Examples of RDF serialisations include RDF/XML, Notation-3 (N3), Turtle, N-Triples, RDFA, and RDF/JSON.

(<http://location.testproject.eu/so/au/AdministrativeUnit/STATBEL/24000>).

Linking these two data resources allows us to get richer information about Brussels. Besides that, the attributes of the data entity for the city of Brussels is country. This attribute reveals that a city is positioned in/belongs to a country. In our case the value for country is Belgium. There are different options for encoding this information.

One way would be to include the value for country as text, e.g. a literal or a string. This option however cannot take us too far and can suffer from different writings, different languages and even spelling errors. The Linked Data approach in this case opts for replacing the text value with a URI pointing to the specific country, i.e. to Belgium (the URI of DBpedias resource for Belgium is <http://dbpedia.org/resource/Belgium>). The Linked Data option allows us to unambiguously refer to Belgium and also navigate through the links in order to collect more information about Brussels.

Due to these considerations, we need to find out the minimal set of properties necessary to make up the appropriate connections between the data in both schemes. Therefore, we need the minimal keys from each one of the sets so we can decide later which element will be our joining one.

In the following, we outline how the heterogeneous information is linked in two dataset with information about films.

Example 2 *The repositories <http://www.imdb.com> and <https://www.filmaffinity.com> contain information about films with different structure. The goal is to obtain the keys in order to connect the knowledge stored in both datasets.*

The method to make the fusion of information is summarized in the next items:

- *To select a group of films.*
- *To take the topics from IMDB repository.*
- *To extract the implications using the tool for Formal Concept Analysis named ConceptExplorer.*
- *To calculate the minimal keys for IMBD schema.*
- *To take the topics from Filmaffinity repository.*
- *To extract the implications using the tool for Formal Concept Analysis named ConceptExplorer.*
- *To calculate the minimal keys for Filmaffinity schema.*
- *To repeat the same with the joining of both implication sets.*

The result of applying SST Method to this two heterogeneous dataset is the following:

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IMDB

Topics: Action Comedy Crime Drama Fantasy Romance Thriller

Little City (1998) Comedy Romance
Driver, The (1978) Action Crime Thriller
Father of the Bride (1950) Comedy Romance
Bio-Dome (1996) Comedy
Fast Runner, The (2001) Drama Fantasy
Overboard (1987) Comedy Romance
Get Rich or Die Tryin? (2005) Crime Drama

Implications:

Romance -> Comedy
Thriller -> Action Crime
Action -> Crime Thriller

Keys:

Romance Thriller
Romance Action

Filmaffinity

Topics: Action Comedy Crime Drama Family Film-noir Nature Romance Survival

Little City (1998) Comedy Drama
Driver, The (1978) Action Crime Film-noir
Father of the Bride (1950) Comedy Romance Family
Bio-Dome (1996) Comedy
Fast Runner, The (2001) Drama Nature Survival
Overboard (1987) Comedy
Get Rich or Die Tryin? (2005) Drama

Implications:

Action -> Crime Filmnoir
Crime -> Action Filmnoir
Family -> Comedy Romance
Filmnoir -> Action Crime
Romance -> Comedy Family
Survival -> Drama Nature
Nature -> Drama Survival

KEYS FOR FUSIONING HETEROGENEOUS INFORMATION

Keys:

Action Family Survival
Action Family Nature
Action Romance Survival
Action Romance Nature
Family Filmnoir Survival
Family Filmnoir Nature
Filmnoir Romance Survival
Filmnoir Romance Nature
Crime Family Survival
Crime Family Nature
Crime Romance Survival
Crime Romance Nature

The joining of both topics:

Topics: Action Comedy Crime Drama Family Fantasy Film-noir Nature Romance Survival Thrill

Little City (1998) Comedy Drama Romnace
Driver, The (1978) Action Crime Film-noir Thriller
Father of the Bride (1950) Comedy Romance Family
Bio-Dome (1996) Comedy
Fast Runner, The (2001) Drama Fantasy Nature Survival
Overboard (1987) Comedy Romance
Get Rich or Die Tryin? (2005) Crime Drama

Implications:

Comedy Drama -> Romance
Family -> Comedy Romance
Fantasy -> Drama Nature Survival
Nature -> Drama Fantasy Survival
Romance -> Comedy
Survival -> Drama Fantasy Nature
Thriller -> Action Crime Filmnoir
Action -> Crime Filmnoir Thriller
Filmnoir -> Action Crime Thriller

Keys:

Family Filmnoir Fantasy
Family Action Fantasy

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Family Thriller Fantasy
Family Filmnoir Nature
Family Action Nature
Family Thriller Nature
Family Filmnoir Survival
Family Action Fantasy
Family Thriller Survival

5 Conclusions

The fusion of heterogeneous information is an emergent problem in which the use of Logic is adequate in order to incorporate automated reasoning mechanism. We have proposed the use of Simplification Logic to manipulate functional dependencies (implications).

SST Method is based of Simplification Logic and allow us to enumerate all minimal keys of a data repository inferring them from a set of functional dependencies. We illustrate how the method can be used to solve problems related with the integration/fusion of heterogeneous information in linked data.

For a extension of this work, we are envolved in the introduction of parallelism techniques in the implementation of the algorithms that could help us dealing with problems containing a substantial amount of information at the input. We think that the tableaux paradigm used in logic matches perfectly the design of the parallel versions of the algorithms. This strategy could be used in order to resolve complex input problems. We truly need to go further searching for strategies in order to find out good BOVs for the partial implementation process.

Besides, it is necessary to go ahead in the design of a *benchmark* that takes into account the different aspects and nature of these problem and algorithms in order to direct us searching the best strategies.

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Apéndice D

Increasing the Efficiency of Minimal Key Enumeration Methods by Means of Parallelism

Increasing the Efficiency of Minimal Key Enumeration Methods by Means of Parallelism

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Abstract: Finding all minimal keys in a table is a hard problem but also provides a lot of benefits in database design and optimization. Some of the methods proposed in the literature are based on logic and, more specifically on tableaux paradigm. The size of the problems such methods deal with is strongly limited, which implies that they cannot be applied to big database schemas. We have carried out an experimental analysis to compare the results obtained by these methods in order to estimate their limits. Although tableaux paradigm may be viewed as a search space guiding the key finding task, none of the previous algorithms have incorporated parallelism. In this work, we have developed two different versions of the algorithms, a sequential and a parallel one, stating clearly how parallelism could naturally be integrated and the benefits we get over efficiency. This work has also guided future work guidelines to improve future designs of these methods.

1 INTRODUCTION

Identifying properly all the keys of a table in a relational schema is a crucial task for several areas in information management: data modeling (Simsion and Witt, 2005), query optimization (Kemper and Mörkotte, 1991), indexing (Manolopoulos et al., 1999), etc. Key constraints specify sets of attributes in a relation such that their projection univocally identifies each tuple of the relation. Therefore, each key is composed by a subset of attributes playing the role of a *domain* in a given function whose *image* is the whole set of attributes. This way, the table is viewed as its extensional definition. These functions are described by means of a *Functional Dependency* (*FD*) which specifies a constraint between two subset of attributes, denoted $A \rightarrow B$, ensuring us that for any two tuples in a table, if they agree on A , they also agree on B .

All functional dependencies satisfied in a given table may be deduced from its dataset using data mining techniques (Appice et al., 2011; Huhtala et al., 1999), or may be provided by database designers. It is out the scope of this work to extract FDs from relational tables, since it becomes a data mining problem (Fayyad et al., 1996). Minimal key problem consists in finding all the attribute subsets which make up a minimal key given a set of FDs occurring within a schema of a relational table.

Nowadays, several algorithms are capable to solve this problem using different classical techniques (Lucchesi and Osborn, 1978; Yu and Johnson, 1976; Saiedian and Spencer, 1996; Zhang, 2009; Armstrong, 1974) (see Section 2 for further details). Recently, some alternative methods have been introduced using logic. In this work we will concentrate on algorithms guided by logic, and most specifically, those using tableaux paradigm (Morgan, 1992) for deriving keys of a relation schema using inference systems. As we shall see later, tableaux might be considered a flexible and powerful framework to design automated deduction methods to solve complex problems in a flexible and effective way.

In previous works, several tableaux-like methods have been introduced (Wastl, 1998a; Cordero et al., 2013). Efficient versions of both of them have been implemented. Nevertheless, tableaux-like methods generate wide search spaces and, in many cases, the same solution (same minimal key) appears at the end of several branches of the tree representing the search space. These intrinsic characteristics of tableaux-like methods establish a strong limitation in the size of the problems to be treated by them and, usually, discourage their use.

Indeed, sequential implementation of these methods produces such an explosion of search space that we go beyond machine capabilities, even with small

problems (20+ FDs for K's method (Wastl, 1998a) as it has been demonstrated in previous studies of this work (Cordero et al., 2013)). However, a very interesting property within search spaces induced by tableaux methods, is the fully independence of their branches, so we can directly consider them as separated sub-problems leading to several solutions of the original problem. It is in this spirit that tableaux paradigm supplies us an optimal guide to build parallel algorithms finding all minimal keys in a table, since the building of the tableaux tree directs the parallel and independent processing.

In this work, we have developed parallel implementations of tableaux-like methods to solve minimal keys finding problems. As shown in the following, they have significantly increased the size of the problem these methods are able to handle. Intentionally, we have executed them under different hardware configurations trying to discover tendencies in which the efficiency within method could be influenced. As already implied above, a processing cluster will be available for us to engage problems with a substantial size at the input so we can deal with them in terms of storage capacity and execution time.

The paper is organized as follows: In Section 2 we introduce several background. A brief study of the state-of-the-art is exposed in Section 3. Section 4 introduces us into sequential tableaux-like methods showing some experimental results. Parallelism implementations enter the scene in Section 5 presenting their way to proceed, and showing the high benefits obtained. Several conclusions are then given in Section 6.

2 BACKGROUND

We begin this section with three brief definitions of basic concepts.

Definition 1 (Functional dependency). *Let U be a set of attributes. A functional dependency (FD) over U is an expression of the form $X \rightarrow Y$, where X, Y are attribute sets. It is satisfied in a table R if for every two tuples of R , if they agree on X , then they agree on Y .*

A key of a relational table is a subset of attributes that allows us to uniquely characterize each row. It may be defined by means of functional dependencies as follows:

Definition 2 (Key). *Given a table R over the set of attributes U , we say that K is a key in R if the functional dependency $K \rightarrow U$ holds in R .*

Definition 3 (Minimal Key). *Given the table R , the attribute set $K \subset U$ is said to be a minimal key if it is a key of R and for all attribute $k \in K$ the subset $K - \{k\}$ is not a key of R .*

Due to space limitation, we refer those readers non familiar with the formal notions of FDs, keys and relational tables to (Elmasri and Navathe, 2010). In Table 1, we just illustrate its semantics by a basic example.

From the information in Table 1, we may ensure that the following FDs are satisfied: $Title, Year \rightarrow Country$, $Title, Year \rightarrow Director$ and $Director \rightarrow Nationality$. Moreover, the table has only one minimal key: $\{Title, Year, Star\}$

Inferring minimal keys from a set of FDs has been well studied. The algorithm of Lucchesi and Osborn (Lucchesi and Osborn, 1978) to find all the keys in a relational schema is considered the first deep study around this problem. Yu and Johnson (Yu and Johnson, 1976) established that the number of keys is limited by the factorial of the number of dependencies, so, there does not exist a polynomial algorithm for this problem.

3 TABLEAUX-LIKE METHODS

Some authors have used several techniques to solve this problem. Saiedian and Spencer (Saiedian and Spencer, 1996) propose an algorithm for computing the candidate keys using attribute graphs. Zhang in (Zhang, 2009) uses Karnaugh maps to calculate all the keys. Nevertheless, we are interested in the use of artificial intelligence techniques and, more specifically, in the use of logic. Armstrong's axiomatic system (Armstrong, 1974) is the former system introduced to manage FDs in a logic style. In (Wastl, 1998b), for the first time a Hilbert-styled inference system for deriving all keys of a relation schema was introduced. Alternatively, in (Cordero et al., 2013) the authors tackle the key finding problem with another inference rule inspired by the Simplification Logic for Functional Dependencies. These two papers constitute the target algorithms to be compared in this work. We refer the reader to the original papers for further details.

Both works are strongly based on tableaux paradigm. Tableaux-like methods represent the search space as a tree, where its leaves contain the solutions (keys). Tree building process begins with an initial root and from there on, inference rules generate new branches labeled with nodes representing simpler instances of the parent node. The very best advantage of this process goes to its versatility, since developing new inference systems –which is not a trivial task indeed– allows us to design a new method. Com-

Table 1: Movie table.

Title	Year	Country	Director	Nationality	Star
Pulp Fiction	1994	USA	Quentin Tarantino	USA	John Travolta
Pulp Fiction	1994	USA	Quentin Tarantino	USA	Uma Thurman
Pulp Fiction	1994	USA	Quentin Tarantino	USA	Samuel L. Jackson
King Kong	2005	New Zealand	Peter Jackson	New Zealand	Naomi Watts
King Kong	2005	New Zealand	Peter Jackson	New Zealand	Jack Black
King Kong	1976	USA	De Laurentiis	IT	Jessica Lange
King Kong	1976	USA	De Laurentiis	IT	Jeff Bridges
Django Unchained	2012	USA	Quentin Tarantino	USA	Jamie Foxx
Django Unchained	2012	USA	Quentin Tarantino	USA	Samuel L. Jackson

parisons between tableaux-like methods can be easily drawn as its efficiency goes hand-in-hand with the size of the generated tree.

Although \mathbb{K} and SL_{FD} are the two inference systems which are the basis of two tableaux-like methods, they are very different. \mathbb{K} is the former basis for a Hilbert-styled method and it deals with unitary functional dependencies, which produce a significant growth of the input set. SL_{FD} avoids the use of fragmentation by using generalized formulas. It is guided by the idea of simplifying the set of FDs by removing redundant attributes efficiently. Moreover, SL_{FD} incorporates a pre-processing task at a first step which prunes the input up to an algebraic characterization of the problem by providing significant reduction of the problem's size to be treated by the tableaux-like method, which is the hardest task.

4 LIMITS OF SEQUENTIAL TABLEAUX-LIKE METHODS

As we mentioned in the introduction, in this section we show the strong limitation that sequential implementation of tableaux-like methods will face to solve minimal keys.

We have developed two efficient implementations of both methods and they have been executed over a high performance architecture sited in the Supercomputing and Bioinnovation Center¹. In these experiments we take into account the following parameters: execution time in seconds (named [Ti]), number of nodes in the tableaux search space (named [No]) and number of redundant keys (named [RK]). This last parameter shows the impact of extra branches, i.e., the number of duplicated keys computed by the algorithm.

Execution times may be considered a parameter linked to the architecture we are using for running the experiments. Number of nodes and redundant keys

represent the size of the search space and the repeated operations respectively. They are independent of the implementation, providing a fair comparison between methods in the future.

4.1 First Experiment: Benchmarking Problems

Although there not exists a benchmarking for functional dependency problems, our first intention was to execute the methods over a set of different and characteristic problems. Thus, we began building a battery of problems gathered from several related papers around (Saiedian and Spencer, 1996; Wastl, 1998a) conforming an initial suitable set of problems in a benchmarking style. Results obtained for this battery of problems are shown in Table 2.

As shown in Table 2 only one of the entry problems needs more than one second to finalize and it is in the case of \mathbb{K} method. This is due to the small sets handled by these academic problems. Results for SL_{FD} are better than \mathbb{K} except for saiedian3 problem. Indeed, there are cases where we need just one node to finish the algorithm, corresponding to those problems where the algebraic characterization used by SL_{FD} reduces the problem to its canonical version.

4.2 Second Experiment: Random Problems

In this subsection we deal with larger randomly generated problems. We have built several examples varying two parameters: number of attributes and number of FDs. We have not established a correlation between both parameters in the generation because different ratios between them produce problems with significant differences. Moreover, observe that number of minimal keys is not directly influenced by these two parameters.

The size of the problems in Table 3 is greater than those presented in previous section. They may be considered *medium-size* problems, having param-

¹<http://www.scbi.uma.es>.

Table 2: Sequential executions over benchmarking problems.

Problem	Attrib	FDs	Keys	\mathbb{K}			SL_{FD}		
				[Ti]	[No]	[RK]	[Ti]	[No]	[RK]
saiedian1	6	5	1	0	12	5	0	1	0
saiedian2	6	5	3	0	7	0	0	10	3
saiedian3	7	7	4	0	139	64	0	674	284
derivation5	9	4	5	0	17	4	0	41	20
a3rojo	7	5	2	0	81	29	0	4	0
elmasri1	6	3	1	0	1	0	0	1	0
wastl2	7	3	1	0	2	0	0	1	0
wastl10	3	3	1	0	5	2	0	1	0
wastl13	4	4	3	0	10	2	0	13	5
example001	10	7	8	20	55.768	24.174	0	1.090	448

Table 3: Sequential executions over random problems.

Problem	Attrib	FDs	Keys	\mathbb{K}			SL_{FD}		
				[Ti]	[No]	[RK]	[Ti]	[No]	[RK]
med1	10	10	3	155	1.463.228	723.372	0	902	404
med2	5	17	4	0	1.906	1.029	0	1.149	772
med3	15	7	2	40	432.104	220.230	0	143	71
med4	30	5	5	12	802.770	300.485	12	23	7
med5	20	10	3	180	2.038.746	1.012.651	3	1.102	666
med6	5	50	5	20	218.892	179.444	1	129.508	117.461
med7	40	10	2	204	1.130.539	467.512	0	122	53
med8	15	15	7	38	6.715.949	3.023.693	5	77.922	41.070
med9	7	50	5	325	32.219.336	21.357.930	18	2.760.961	2.227.596
med10	10	20	4	835	496.380.119	218.275.528	8	20.442	9.966

eters which properly match with real tables in software engineering and execution times are quite reasonable, particularly for SL_{FD} method (less than one minute). Nevertheless, we notice that as far as problem's size grows (even just a little), results go considerably beyond. Therefore, we are absolutely limited when dealing with real problems, where the input size would be worthy of consideration. So the challenge is to figure out whether parallelism will help us solving these kind of problems, and even greater ones.

5 PARALLELIZATION OF TABLEAUX-LIKE METHODS

As a conclusion of the experiments presented in previous section, parallel strategy and big hardware resources will be totally indispensable if we want to compare tableaux-like methods from one to another. Our intention is to take advantage of tableaux design to split the original problem into *atomic* instances that may be solved within a reasonable time and resources. Then, we combine the solutions of all these sub-problems to enumerate all the minimal keys.

Thus, our parallel implementations of the algorithms will work in two steps:

1. Splitting step: It executes the tableaux-like meth-

ods but it will stop in a determinate level that we will introduce at the process call, generating a set of sub-problems. The level is induced by the size of the root (the number of attributes in this level).

2. Parallel step: In this second stage we execute parallel task solving those sub-problems and, at the end, we combine all the solutions to get all minimal keys.

In order to test parallel versions we run another battery of problems whose results are retrieved in Table 4. This time we include several new columns gathering parallel implementation's parameters: Break-off value [L] and number of generated sub-problems [Sp].

It is imperative to state some critical considerations concerning the limit size of the atomic problems. In one hand, we have observed that the greater this limit is, the better will be the improvement by parallelism, since it *would* generate a higher number of sub-problems. However, as far as we try to make this improvement to be better by a wide limit value, the longer the partial version will take to split the entry problem.

In addition, this is not an independent parameter among the algorithms, we need to choose a different break-off value depending on the method we are using as each one of them will need a particular amount of resources.

Table 4: Parallel executions over random problems.

Prob	Attr	FDs	Keys	\mathbb{K}					SL_{FD}				
				[L]	[Sp]	[Ti]	[No]	[RK]	[L]	[Sp]	[Ti]	[No]	[RK]
cp01	7	50	5	6	87	0	32,219,336	21,357,930	45	50	3	2,760,961	2,227,596
cp02	10	20	4	9	44	0	496,380,119	218,275,528	10	268	11	20,442	9,966
cp03	15	20	3	11	32	1	17,917,662	9,340,225	15	78	4	1,405,153	814,026
cp04	20	20	4	10	27,284	31	3,145,751,761	1,424,991,475	15	47	3	587,765	313,513
cp05	20	30	45	12	1,696	3	1,563,813,853	677,457,455	24	98	136	271,402,277	162,828,760
cp06	10	35	5	8	2,358	3	121,396,806	65,571,971	30	158	8	3,215,995	1,686,149
cp07	25	15	15	14	25,836	28	3,975,400,144	1,980,982,101	8	802	39	220,047	914,80
cp08	15	40	1	9	39,708	46	837,341,359	433,068,418	0	0	1	1	0
cp09	25	20	25	14	33,146	38	123,283,772,804	59,975,556,886	12	18,999	1,077	47,014,652	23,418,562
cp10	35	10	37	22	1,370	5	101,429,265,443	68,138,197,993	7	219	10	27,985	13,436

As an example of this last point, we would like to check execution times for cp09 and cp10 in the case of \mathbb{K} , where the huge difference is due to a wiser splitting process.

As a general conclusion, execution times are pretty reasonable considering the dimension of these entry problems (several minutes were enough to resolve most of cases).

On a separate issue, we can notice that results are pretty huge for this kind of problems using \mathbb{K} 's method. The hugest number of nodes of the tableaux overtakes up to 123 billions of nodes. So, efficiency of \mathbb{K} 's method is so far to be accepted. SL_{FD} reaches better times and dimensions tableaux in the very most of cases. Thus, several noteworthy outcomes come up.

First, problem cp04 needs a tableaux of over half-million nodes with SL_{FD} , while \mathbb{K} goes up to 3 billions! This is due to the high number of FDs after the hard fragmentation rule inherent to \mathbb{K} . A similar situation involves cp09 and cp10 problems.

Finally, if we care about useless computing time, we notice the number of redundant keys is terrible; the difference here between both methods is as impressive as in the rest of parameters. For instance, the resulting set of minimal keys for cpx04 problem contains just 4 minimal keys. However, \mathbb{K} generates 1,424,991,475 redundant keys and SL_{FD} 313,513. This is indeed, a huge waste of space and time.

6 CONCLUSIONS

The first point we want to state clear is that the concept of parallelism we are dealing with refers to a *hardware parallelism*. We mean that the benefits we are obtained from parallelism are due to a cluster of cores where we can deliver each of our jobs continuously.

In order to resolve real problems where the size of the input is substantial, it is imperative to count on the

participation of a great amount of resources. Moreover, it is difficult, at a first look, to estimate whether a given problem will result in a difficult or an easy one. In some sense we may say that it is a chaotic and unpredictable problem.

A glance is enough to easily realize that \mathbb{K} algorithm needs more time, more nodes and more redundant keys to reach the solution than SL_{FD} in the very most of cases. Indeed, the differences are not trivial so far. Concerning the size of the tableaux, \mathbb{K} builds up billion of nodes whereas SL_{FD} generates a reasonable amount of nodes. A similar conclusion may be established for the number of redundant keys.

Finally, establishing an appropriate limit to split up the entry problem in the parallel versions of the algorithms is not an easy issue so far. We have to run several experiments to reach a good value which will not spend so much time splitting the entry but it should spend time enough to take advantage of the parallelism.

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*-¿Qué te parece desto, Sancho? -Dijo Don Quijote-
¿Hay encantos que valgan contra la verdadera valentía?
Bien podrán los encantadores quitarme la ventura,
pero el esfuerzo y el ánimo, será imposible.*

El Ingenioso Hidalgo Don Quijote de la Mancha

Miguel de Cervantes

