

Model-based Exception Mining for Object-Relational Data

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Abstract This paper develops model-based exception mining and outlier detection for the case of object-relational data. Object-relational data represent a complex heterogeneous network, which comprises objects of different types, links among these objects, also of different types, and attributes of these links. We follow the well-established Exceptional Model Mining (EMM) framework, which has been previously applied for subgroup discovery in propositional data; our novel contribution is to develop EMM for relational data. EMM leverages machine learning models for exception mining: A object is exceptional to the extent that a model learned for the object data differs from a model learned for the general population. In relational data, EMM can therefore be used for detecting single outlier or exceptional objects. We apply EMM with state-of-the-art statistical-relational model discovery methods that construct a graphical model (Bayesian network), that compactly represents probabilistic associations in the data. We investigate several outlierness metrics, based on the learned object-relational model, that quantify the extent to which the association pattern of a potential outlier object deviates from that of the whole population. Our method is validated on synthetic datasets and on real-world data sets about soccer and hockey matches, IMDb movies and mutagenic compounds. Compared to baseline methods, the EMM approach achieved the best detection accuracy when combined with a novel outlierness metric. An empirical evaluation on soccer and movie data shows a strong correlation between our novel outlierness metric and success metrics: Individuals that our metric marks out as unusual tend to have unusual success.

Keywords Outlier Detection · Exception Mining · Statistical-Relational Learning · Bayesian network · Likelihood Ratio · Network Data

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1 Introduction: Exception Mining for Relational Data

Exception mining and outlier detection are important data analysis tasks in many domains. For relational data, exception mining can be applied to support outlier detection, where statistical deviations are viewed as due to a node or entity being genuinely exceptional, rather than due to statistical noise in the data. Statistical approaches to unsupervised outlier detection leverage a generative model of the data [2]. The generative model represents normal behavior. An individual object is deemed an outlier if the model assigns sufficiently low likelihood to generating it. We describe a new method, based on the Exceptional Model Mining (EMM) framework [15], for extending statistical outlier detection with generative models to the case of object-relational data.

Data Type. The object-relational data model is one of the main data models for structured data [31]. The main characteristics of objects that we utilize in this paper are the following.

- Object Identity. Each object has a unique identifier that is the same across contexts. For example, a player has a name that identifies him in different matches.
- Class Membership. An object is an instance of a class, which is a collection of similar objects. Objects in the same class share a set of attributes. For example, van Persie is a player object that belongs to the class striker, which is a subclass of the class player.
- Object Relationships. Objects are linked to other objects. Both objects and their links have attributes.

A common type of object relationship is a component relationship between a complex object and its parts. For example, a match links two teams, and each team comprises a set of players for that match. A difference between relational and vectorial data is therefore that an individual object is characterized not only by a list of attributes, but also by its links and by attributes of the objects linked to it. We refer to the substructure comprising this information as the *object data*. Equivalent terms are “egonet” from network analysis [4] and “interpretation” [34]. The *relational outlier detection problem is to identify objects whose object data differ from the general population/class*. Our approach to this problem combines EMM with statistical-relational model discovery [21].

Approach: Exceptional Model Mining + Statistical-Relational Learning. In the EMM framework, a subgroup is exceptional to the extent that a model learned from data for the subgroup deviates from a model learned for the general population. Figure 1 illustrates an EMM blueprint. EMM therefore leverages the extensive work on model learning for exception mining. Relational EMM leverages the extensive work on statistical-relational model learning for relational exception mining. In the propositional setting with i.i.d. data (independent and identically distributed), each object is represented by a single data row, and it is meaningless to learn a model for a single object. Instead, EMM is applied to

identify exceptional subgroups of objects. With relational data, each object is represented by its own dataset, and it is meaningful to apply EMM to identify single exceptional objects—a subgroup of size 1. Therefore *relational EMM can be utilized for relational outlier detection*; this paper evaluates relational EMM for this task.

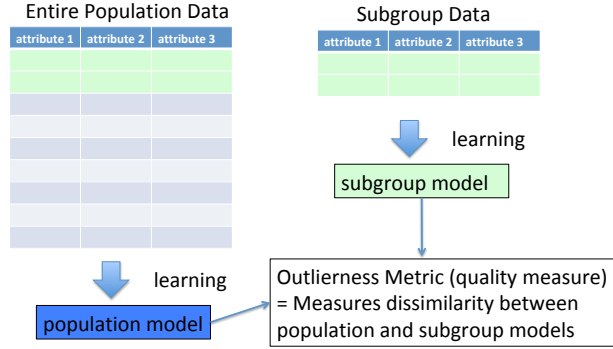


Fig. 1 A general schema for Exceptional Model Mining for propositional i.i.d. data

Model Type. EMM is an inclusive framework in that any model type of interest can be utilized. In this paper we focus on first-order Bayesian networks (BNs) [58, 42, 28], but our definitions apply to other log-linear models such as Markov logic networks [14]. A class-model Bayesian network (BN) structure is learned with data for the entire population. The nodes in the BN represent attributes for links, of multiple types, and attributes of objects, also of multiple types. To learn the BN model, we apply techniques from statistical-relational learning, a recent field that applies and extends AI and machine learning to relational data [21, 50, 13]. For a learned BN structure, we compare two parameter vectors (see Figure 6 below): The *class model parameters* are estimated from the entire population data, and the *object model parameters* are estimated from the object data.

Quality Measure = Outlierness Metric. For a given model type, quantifying the extent to which an object model deviates from the population class model is the main research question in EMM [15]. A computational method for measuring this extent is called a *quality measure*; when we apply EMM to relational outlier detection, we also refer to it as an *outlierness metric*. We propose utilizing the relational log-linear likelihood function [28, 48] that measures how well a parametrized statistical-relational model fits an input database. The *likelihood ratio* is the difference in log-likelihood between the class model and the object model, when applied to the object data. Assuming maximum likelihood estimates, it is equivalent to the Kullback-Leibler divergence (KLD) between class and object models. While our evaluation indicates that KLD is a good outlierness metric, we propose improving it further by applying two transformations: (1) a mutual information decomposition, and

(2) replacing log-likelihood differences by log-likelihood distances. We refer to the resulting novel score as the *expected log-likelihood distance*. A Taylor series analysis shows that this score is approximated by the L1-norm (total variation distance) between the object and class data distributions.

Evaluation. Our code and datasets are available on-line at [44]. We evaluate relational EMM outlierlieness metrics in three ways.

Detection Accuracy. Our performance evaluation follows the design of previous outlier detection studies [19, 2], where the methods are scored against a test set of known outliers. We use three synthetic and four real-world datasets, from the UK Premier Soccer League, the Internet Movie Database (IMDb), the National Hockey League, and Mutagenesis. On the synthetic data we have known ground truth. For the real-world datasets, we use a one-class design, where one object class is designated as normal and objects from outside the class are the outliers. For example, we compare goalies as outliers against the class of strikers as normal objects. Compared to six baseline methods, the EMM-based scores (likelihood ratio and expected log-likelihood distance) achieve the top detection accuracy on all three synthetic datasets, and on 4 out of 5 real-world datasets.

Case Studies. We also offer case studies where we assess whether individuals that our score ranks as highly unusual in their class are indeed unusual. The case studies illustrate that the EMM scores are *easy to interpret*, because the Bayesian network provides a local decomposition of log-likelihood differences. Interpretability is very important for users of an outlier detection method as there is often no ground truth to evaluate outliers.

Correlation with Success. We compare the likelihood-based outlierlieness metrics to independent success metrics for a given domain. Success rankings are one of the most interesting features to users. Our reasoning is that high success is an independent metric that indicates an unusual individual. So a correlation between an outlierlieness metric and success is an independent validation of the metric, and also shows that it points to meaningful and interesting outliers.

Contributions Our main contributions may be summarized as follows.

1. We develop the first application of the EMM framework to outlier detection for structured relational data based on a probabilistic model.
2. We define a new model-based outlierlieness score, derived from a novel model likelihood comparison, the expected log-likelihood distance.

Paper Organization We review related work on outlier detection for structured data, then background about Bayesian networks for relational data. Then we introduce likelihood-ratio based outlierlieness scores for relational EMM, including our novel log-likelihood distance. Empirical evaluation compares EMM and aggregation-based approaches to relational outlier detection, with respect to three synthetic and four real-world datasets.

2 Related Work

Outlier detection is a densely researched field, for a survey please see [2, 4]. Figure 2 provides a tree picture of where our method is situated with respect to other outlier detection methods and other data models. Our method falls in the category of *unsupervised* statistical model-based approaches. To our knowledge, ours is the first model-based method tailored for object-relational data. Like EMM and other model-based approaches, it detects *global outliers*. Aggarwal [2] defines a global outlier to be a data point that notably deviates from the rest of the population. We review relevant approaches from two different data models, the most common atomic object model—where data is represented by vectors—and structured data models. A preliminary version of this paper appeared in the Proceedings of the IEEE SSCI series [45].

a) Attribute Vector Data Model: By far most work on outlier detection considers atomic objects with flat feature vectors. This leads to an impedance mismatch: The required input format for these outlier detection methods is a single data matrix, not a structured dataset. For example, one cannot provide a relational database as input. This mismatch is not simply a question of choosing a file format, but instead reflects a different underlying data model: complex objects with both attributes and component objects vs. atomic objects with attributes only. It is possible to “flatten” structured data by converting it to unstructured feature vectors, for instance by using aggregate functions. We evaluated the aggregation approach in this paper by applying three standard methods for outlier detection.

Work on atomic contextual outliers [55] is like ours in that it considers the distinctness of a target individual from a reference class. A reference class is not specified for each object, but is constructed as part of outlier detection. Our work could be combined with a class discovery approach by providing a score of how informative the inferred classes are.

b) Structured Data Models: We discuss related techniques in three types of structured data models: SQL (relational), XML (hierarchical), and OLAP (multi-dimensional).

For relational data, many outlier detection approaches aim to discover rules that represent the presence of anomalous associations for an individual or the absence of normal associations [34, 19]. In these approaches what we call the object data is usually referred to as an interpretation, so the problem is to score outlier interpretations [34]. The survey by [38] unifies within a general rule search framework three important related tasks: i) exception mining, which looks for associations that characterize unusual cases, ii) subgroup mining, which looks for associations characterizing important subgroups, and iii) contrast space mining, which looks for differences between classes. Another rule-based approach uses Inductive Logic Programming techniques [6]. While local rules are informative, they are not based on a global statistical model and do not provide a single outlier score for each individual. As we show in our case studies (Section 8.3), the conditional probability parame-

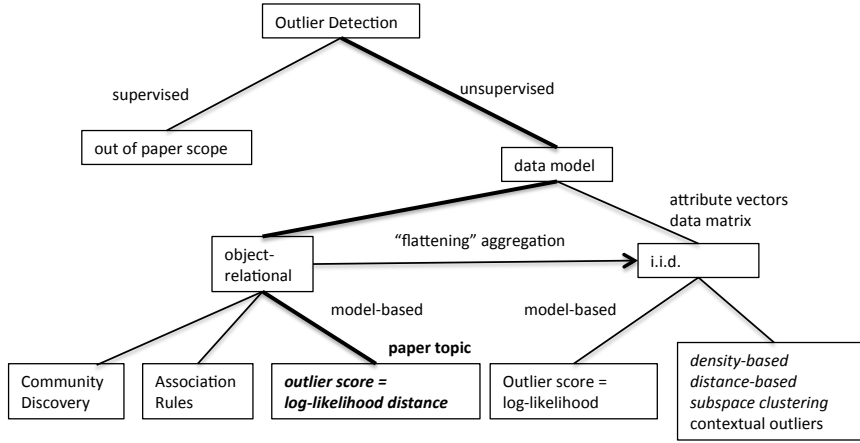


Fig. 2 A tree structure for related work on outlier detection for structured data. A path specifies an outlier detection problem, the leaves list major approaches to the problem. Approaches in *italics* appear in experiments.

ters in a Bayesian network can be used to extract rules from the BN model (of the form *parent_values* \rightarrow *child_value*). The distance-based algorithms designed for outlier detection can be applied to the relational case. Horvath *et al.* introduced a similarity measure for the first-order instance-based learner RIBL that uses the edit distance to compute the distance between lists and terms [25]. This distance has been used for the clustering task in [29]. To the best of our knowledge, distance-based relational learning has not been used for outlier detection. One-class classification can be viewed as a special case of outlier detection. Khot *et al.* introduced a non-parametric relational one-class classification based on first-order trees. They proposed a tree-based distance metric to discover new relational features and to differentiate relational examples [27]. Khot *et al.* emphasize that their method is not appropriate for outlier detection because outlier detection makes different assumptions about the data than they do (e.g. outliers are far from other points and isolated).

Propositionalization summarizes the multi-relational data into a single data table and can be used for outlier detection and classification tasks [32, 41, 33, 16, 5].

A latent variable approach in information networks ranks potential outliers in reference to the latent communities inferred by network analysis [19]. Our model also aggregates information from entities and links of different types, but does not assume that different communities have been identified.

Koh *et al.* [30] propose a method for hierarchical structures represented in XML document trees. Their aim is to identify feature outliers, not class outliers as in our work. Also, they use aggregate functions to convert the object hierarchy into feature vectors. Their outlier score is based on local correlations, and they do not construct a model.

The multi-dimensional data model defines numeric measures for a set of dimensions. The differences in the two data models mean that multi-dimensional outlier detection models [47] do not carry over to object-relational outlier detection. (1) The object data model allows but does not require any numeric measures. In our datasets, all features are discrete. Nor do we assume that it is possible to aggregate numeric measures to summarize lower-level data at higher levels. (2) In scoring a potential outlier object, our method considers other objects *both* below and above the target object in the component hierarchy. OLAP exploration methods consider only cells below or at the same level as the target cell. For example, in scoring a player, our method would consider features of the player’s team. Also, the log-likelihood distance outlier score of an object is not determined by the outlier scores of its components, in contrast to approaches derived from Sarawagi *et al.* [47]. They use values such as the most unusual cell that is below a target cell. (3) Our approach models a joint distribution over features, exploiting correlations among features. Most of the OLAP-based methods consider only a single numeric measure at a time, not a joint model.

3 Background: Bayesian Networks for Relational Data

We adopt the Parametrized Bayesian network (PBN) formalism [42] that combines Bayesian networks with logical syntax for expressing relational concepts.

3.1 Relational Data

We adopt a term-based notation for combining logical and statistical concepts [42, 28]. Table 1 summarizes our notation. A functor is a function or predicate symbol. Each functor has a set of values (constants) called the **domain** of the functor. The range of a **predicate** is $\{T, F\}$. Predicates are usually written with uppercase Roman letters, other terms with lowercase letters. A predicate of arity at least two is a **relationship** functor. Relationship functors specify which objects are linked. Other functors represent **features** or **attributes** of an object or a tuple of objects (i.e., of a relationship). A **population** is a set of objects. A **term** is of the form $f(\sigma_1, \dots, \sigma_k)$ where f is a functor and each σ_i is a first-order variable or a constant denoting an object. A term is **ground** if it contains no first-order variables; otherwise it is a first-order term. In the context of a statistical model, we refer to first-order terms as **Parametrized Random Variables** (PRVs) [28]. A **grounding** replaces each first-order variable in a term by a constant; the result is a ground term. A grounding may be applied simultaneously to a set of terms. A relational database \mathcal{D} specifies the values of all ground terms.

Consider a joint assignment $P(\mathbf{V} = \mathbf{v})$ of values to a set of PRVs \mathbf{V} . The *grounding space* of the PRVs is the set of all possible grounding substitutions, each applied to all PRVs in \mathbf{V} . The *count* of groundings that satisfy the assignment with respect to a database \mathcal{D} is denoted by $\#_{\mathcal{D}}(\mathbf{V} = \mathbf{v})$. The **database**

Symbol	Definition
a, b, a_1, b_1, \dots	Constant
A, B, T, M, \dots	First-order variable
f, g, \dots	Functor, function symbol
$f(A_1, \dots, A_k)$	First-order term
$f(A = a_1, \dots, A_k = a_k)$	Ground term
\mathcal{D}	Relational database
\mathcal{D}_C	Database for the entire class of objects
\mathcal{D}_o	Restriction of the input database to the target object
$V(A, \dots, B)$	A parametrized random variable
\mathbf{V}	A set of first-order random variable
$\mathbf{V} = \mathbf{v}$	Joint assignment of values to a set of PRVs
$P(\mathbf{V} = \mathbf{v}) \equiv P(\mathbf{v})$	Joint probability that each variable V_i takes on value \mathbf{v}_i
$\#_{\mathcal{D}}(\mathbf{V} = \mathbf{v})$	Count of groundings that satisfy the assignment
$A \setminus v$	Ground a first-order variable
B	A Bayesian network structure
B_C	A Bayesian network structure learned with \mathcal{D}_C as the input database
θ_C	Parameters learned for B_C using \mathcal{D}_C as the input database
θ_o	Parameters learned for B_C using \mathcal{D}_o as the input database
pa_i	Parent of node i

Table 1 Notation and Definition

frequency $P_{\mathcal{D}}(\mathbf{V} = \mathbf{v})$ is the grounding count divided by the number of all possible groundings.

Example. The Opta dataset represents information about Premier League data (Sec. 7.2). The basic populations are teams, players, matches, with corresponding first-order variables T, P, M . As shown in Table 2, the groundings count can be visualized in terms of a groundings table [50], also called a universal schema [46]. The first three column headers show first-order variables ranging over different populations. The remaining columns represent terms. Each row represents a single grounding and the values of the ground terms defined by the grounding. In terms of the grounding table, the grounding count of a joint assignment is the number of rows that satisfy the conditions in the joint assignment. The database frequency is the grounding count, divided by the total number of rows in the groundings table. Table 4 shows an example of computing frequencies. Counts are based on the 2011-2012 Premier League Season. We count only groundings ($team, match$) such that $team$ plays in $match$. Each team, including Wigan Athletic, appears in 38 matches. The total number of team-match pairs is $38 \times 20 = 760$.

A novel aspect of our paper is that we learn model parameters for specific objects as well as for the entire population. The appropriate **object data table** is formed from the population data table by restricting the relevant first-order variable to the target object. For example, the object database for target Team *WiganAthletic*, forms a subtable of the data table of Table 2 that contains only rows where $\text{TeamID} = \text{WA}$; see Table 3. In database terminology, an object database is like a view centered on the object.

MatchId M	TeamId T	PlayerId P	TimePlayed(P, M)	ShotEff(T, M)	result(T, M)
117	WA	McCarthy	90	0.53	win
148	WA	McCarthy	85	0.57	loss
15	MC	Silva	90	0.59	win
...

Table 2 Sample Population Data Table (Soccer).

MatchId M	TeamId $T = WA$	PlayerId P	TimePlayed(P, M)	ShotEff(WA, M)	result(WA, M)
117	WA	McCarthy	90	0.53	win
148	WA	McCarthy	85	0.57	loss
...	WA

Table 3 Sample Object Data Table, for team $T = WA$; all rows have WA in the T column.

Database	Count or $\#_D(\mathbf{V} = \mathbf{v})$	Frequency or $P_D(\mathbf{V} = \mathbf{v})$
Population	76	$76/760 = 0.10$
Wigan Athletic	7	$7/38 = 0.18$

Table 4 Example of Grounding Count and Frequency in Premier League Data, for the conjunction $passEff(T, M) = hi, shotEff(T, M) = hi, Result(T, M) = win$.

3.2 Bayesian Networks

A Bayesian network (BN) structure B is a Directed Acyclic Graph (DAG) whose nodes comprise a set of random variables [39]. Depending on context, we interchangeably refer to the nodes and variables of a BN, collectively denoted by $\mathbf{V} = \{V_1, \dots, V_n\}$. The possible values of V_i are enumerated as $\{v_{i1}, \dots, v_{ir_i}\}$. The notation $P(V_i = v) \equiv P(v)$ denotes the probability of variable V_i taking on value v . We also use the vector notation $P(\mathbf{V} = \mathbf{v}) \equiv P(\mathbf{v})$ to denote the joint probability that each variable V_i takes on value \mathbf{v}_i .

The conditional probability parameters of a Bayesian network specify the distribution of a child node given an assignment of values to its parent nodes. For an assignment of values to its nodes, a BN defines the joint probability as the product of the conditional probability of the child node value given its parent values, for each node in the network. This means that the log-joint probability can be *decomposed* as the node-wise sum

$$\ln P(\mathbf{V} = \mathbf{v}; B, \boldsymbol{\theta}) = \sum_{i=1}^n \ln \theta(\mathbf{v}_i | \mathbf{v}_{\mathbf{pa}_i}) \quad (1)$$

where \mathbf{v}_i is the assignment of values to node V_i , and $\mathbf{v}_{\mathbf{pa}_i}$ is the assignment of values to the parents of V_i , as determined by the assignment \mathbf{v} . To avoid difficulties with $\ln(0)$, here and below we assume that joint distributions are positive everywhere. Since the parameter values $\boldsymbol{\theta}$ for a Bayesian network define a joint distribution over its nodes, they therefore entail a marginal, or unconditional, probability for a single node. We denote the **marginal probability** that node V has value v as $P(V = v; B, \boldsymbol{\theta}) \equiv \theta(v)$. In the following we use the term Bayesian network model to refer to a network structure with

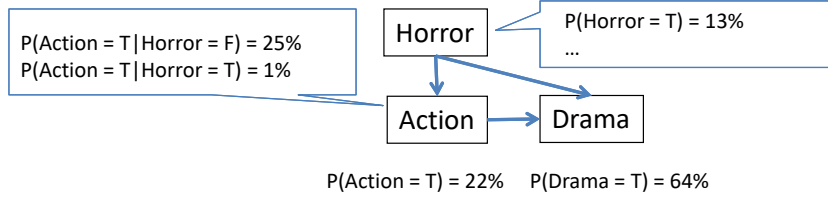


Fig. 3 Example of joint and marginal probabilities computed from a simple Bayesian network structure that was learned from the *Movies* table in the IMDb dataset described in Section 7.2. The parameters were estimated from the IMDb dataset. The conditional probability parameters for the *Drama* node are not shown. The Bayesian network shows that movie genres are largely but not completely exclusive. For instance, among horror movies, only 1% are also classified as action movies. The marginal probabilities are the base rate frequencies of horror, action, and drama movies, which are respectively 13%, 22%, 64%.

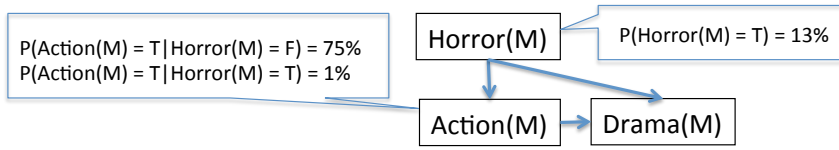


Fig. 4 The Bayesian network from Figure 3, where the node names are expanded using the syntax of parametrized random variables.

parameters (i.e., a pair (B, θ)); for brevity, we also use the terms “Bayesian network” or “model”.

Example. Figure 3 shows an example of a Bayesian network model and associated conditional and marginal probabilities.

3.3 Bayesian Networks for Relational Data

A **Parametrized Bayesian Network Structure** (PBN) is a Bayesian network structure whose nodes are PRVs [42]. The relationships and features in an object database define a set of nodes for a Parametrized Bayesian network. I.i.d. data represented in a single table can be viewed as a special limiting case of multi-relational data with no relationships [36]. Syntactically, this means that columns in an i.i.d. data table represent unary functors, where the relevant population is assumed to be clear from the context rather than explicitly specified as a first-order variable. Figure 4 illustrates how the usual syntax for i.i.d. Bayesian networks is a special case of the PBN syntax.

Figure 5 shows a Parametrized Bayesian network for the Premier League domain that is truly relational in that the functors depend on more than one population variable. For example, shot efficiency does not depend on a match only, but also depends on specifying a team. The BN product formula (1) can be applied to any PBN to compute (estimated) frequencies. In the case of a

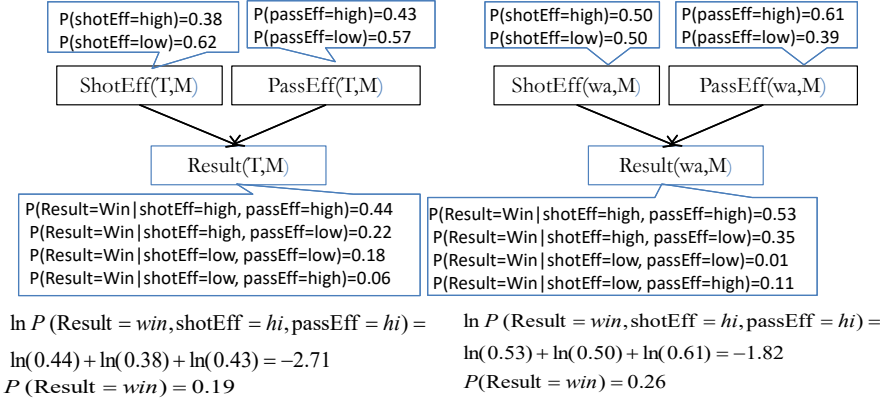


Fig. 5 Example of joint and marginal probabilities computed from a toy Bayesian network structure. The parameters were estimated from the Premier League dataset. (left): A class model Bayesian network B_c for all teams with class parameters θ_c . The first-order variable T ranges over teams, and the first-order variable M over matches. For compactness, conditional probabilities omit the first-order variables and constants. (right): The same Bayesian network structure with object parameters θ_o learned for Wigan Athletic ($T = WA$).

truly relational PBN, as in Figure 5, the PBN can be viewed as representing database frequencies (rather than data table frequencies as in Figure 4). Using Getoor’s terminology, the PBN can be viewed as a Statistical-Relational Model (SRM) [20, 51, 49].

3.4 Likelihood Score for Parametrized Bayesian Networks.

A standard method for applying a generative model assumes that the generative model represents normal behavior since it was learned from the entire population. An object is deemed an outlier if the model assigns sufficiently low likelihood to generating its features [10]. This likelihood method is an important baseline for our investigation. The other outlier scores we consider can be viewed as improved variants of the likelihood approach. Defining a likelihood for relational data is more complicated than for i.i.d. data, because an object is characterized not only by a feature vector, but by an object database. We employ the previously defined normalized relational log-likelihood score [48, 59], which can be computed as follows for a given Bayesian network and database.

$$LOG(\mathcal{D}, B, \theta) = \sum_{i=1}^n \sum_{j=1}^{r_i} \sum_{\mathbf{pa}_i} P_{\mathcal{D}}(v_{ij}, \mathbf{pa}_i) \ln \theta(v_{ij} | \mathbf{pa}_i) \quad (2)$$

Equation (2) has the same form of the standard BN log-likelihood function for vectorial i.i.d. data [11], except that parent-child instantiation counts

are standardized to be proportions [48]. For the notations please refer to Table 1. The condition under which the score of Equation (2) sums to 1 over all relational datasets are discussed in [48].¹ The equation can be read as follows.

1. For each parent-child configuration, use the conditional probability of the child given the parent.
2. Multiply the logarithm of the conditional probability by the database frequency of the parent-child configuration.
3. Sum this product over all parent-child configurations and all nodes.

The maximum of the relational log-likelihood (2) is given by the empirical database frequencies [48, Prop.3.1.]. In all our experiments we use these maximum likelihood parameter estimates.

Example. Consider the class-level BN of Figure 5(left). The family configuration

$$passEff(T, M) = high, shotEff(T, M) = high, Result(T, M) = win$$

contributes one term to the log-likelihood score For the population database \mathcal{D} , this term is

$$\begin{aligned} P_{\mathcal{D}}(passEff(T, M) = high, shotEff(T, M) = high, Result(T, M) = win) \times \\ \ln P(Result(T, M) = win | passEff(T, M) = high, shotEff(T, M) = high) = \\ 0.1 \times \ln(0.44) = -0.08 \end{aligned}$$

where the database frequency 0.1 is computed as in Table 4 and the conditional probability 0.44 is the parameter specified by the Bayesian network of Figure 5.

For the Wigan Athletic data \mathcal{D}_{WA} , the corresponding term is

$$\begin{aligned} P_{\mathcal{D}_{WA}}(passEff(WA, M) = high, shotEff(WA, M) = high, Result(WA, M) = win) \times \\ \ln P(Result(WA, M) = win | passEff(WA, M) = high, shotEff(WA, M) = high) = \\ 0.18 \times \ln(0.44) = -0.14 \end{aligned}$$

where the database frequency 0.18 is computed as in Table 4 and the conditional probability 0.44 is the parameter specified by the Bayesian network of Figure 5(left).

4 Exceptional Model Mining for Relational Data

This section describes our approach to applying the EMM framework to relational data, using the following notation.

- \mathcal{D}_C is the database for the entire class of objects; cf. Table 2. This database defines the **class distribution** $P_C \equiv P_{\mathcal{D}_C}$.

¹ Briefly, if the Parametrized Bayesian Network is viewed as a template for an unrolled ground network, then (1) the ground network cannot contain cycles, and (2) each ground network cannot have multiple parent instantiations; see also [23].

- \mathcal{D}_o is the restriction of the input database to the target object; cf. Table 3. This database defines the **object distribution** $P_o \equiv P_{\mathcal{D}_o}$.
- B_C is a Bayesian network structure learned with \mathcal{D}_C as the input database. Note that we use the entire population data to learn the Bayesian network structure. Therefore, the Bayesian network structure is the same across different individuals.
- θ_C resp. θ_o are parameters learned for B_C using \mathcal{D}_C resp. \mathcal{D}_o as the input database.

Figure 6 illustrates these concepts and the system flow for computing an outlier score. First, we learn a Bayesian network structure B_C for the entire population using a previous learning algorithm (see Section 7.4). We then compute an *outlierness metric* that evaluates how much the object model deviates from the class model. For a given model type, quantifying the extent to which an object model deviates from the population class model is the main research question in EMM [15]. In this paper we examine several outlierness metrics that are based on the likelihood ratio between the object model and the class model. A novel metric that we propose is the **expected log-likelihood distance** (*ELD*). It is defined as follows for each feature i ; the total score is the sum of feature-wise scores. Section 5.2 provides example computations.

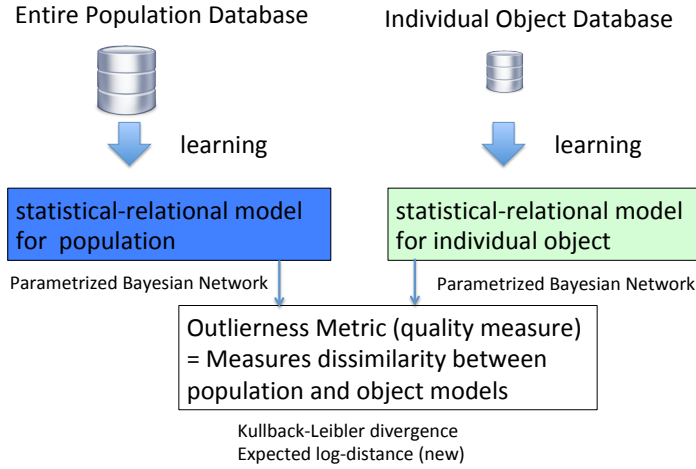


Fig. 6 Exceptional Model Mining for statistical-relational models (cf. Figure 1). The model class we utilize in this paper are Parametrized Bayesian networks, with a log-linear likelihood function. As outlierness metrics we consider the standard Kullback-Leibler divergence, and the novel divergence *ELD* described in the text.

$$ELD_i = \sum_{j=1}^{r_i} P_o(v_{ij}) \left| \ln \frac{\theta_o(v_{ij})}{\theta_C(v_{ij})} \right| + \quad (3)$$

$$\sum_{j=1}^{r_i} \sum_{\mathbf{pa}_i} P_o(v_{ij}, \mathbf{pa}_i) \left| \ln \frac{\theta_o(v_{ij}|\mathbf{pa}_i)}{\theta_o(v_{ij})} - \ln \frac{\theta_C(v_{ij}|\mathbf{pa}_i)}{\theta_C(v_{ij})} \right|. \quad (4)$$

4.1 Interpretation

The first sum (3) is the **single-feature** component, where each feature is considered independently of all others. It computes the expected log-distance with respect to the single feature value probabilities between the object and the class models. The second *ELD* sum (4) is the **mutual information component**, based on the mutual information among all features. It computes the expected log-distance between the object and the class models with respect to the mutual information of feature value assignments. Intuitively, the first sum measures how the models differ if we treat each feature in isolation. The second sum measures how the models differ in terms of how strongly parent and child features are associated with each other.

4.2 Motivation

The motivation for the mutual information decomposition is two-fold. (1) *Interpretability*. This is very important for outlier detection. The single-feature components are easy to interpret since they involve no feature interactions. Each parent-child local factor is based on the average relevance of parent values for predicting the value of the child node, where relevance is measured by

$$\ln \frac{\theta(v_{ij}|\mathbf{pa}_i)}{\theta(v_{ij})}.$$

This relevance term is basically the same as the widely used lift measure [57], hence is an intuitively meaningful quantity. The *ELD* score compares how relevant a given parent condition is in the object data with how relevant it is in the general class.

(2) *Avoiding cancellations*. For different child-parent configurations, the different components of the *ELD* sum may have different signs. This leads to a partial cancelling of differences between the class and object distributions. Since our goal is to assess the distinctness of an object, *we do not want differences to cancel out*. Taking distances as in Equations (3) and (4) avoids the undesirable loss of information. The general point is that averaging differences is appropriate when considering costs, or utilities, but not appropriate for assessing the distinctness of an object. In contrast, the absolute values add differences regardless of their sign. The next section provides comparison scores

and example computations that illustrate the cancelling phenomenon that occurs without adding absolute values. The subsequent section uses Taylor series analysis for a theoretical understanding of the cancelling phenomenon: We show that without the absolute values (as with LR), the first-order term in the Taylor series approximation vanishes, whereas with the absolute values (as with ELD), the first-order term is equivalent to the total variation distance (L1 norm).

5 Comparison Scores

We introduce several alternative likelihood-ratio based outlier scores, following a lesion design where different scores omit different components of our main ELD proposal. To illustrate their essential difference with ELD , we give toy examples before we evaluate them on full datasets.

5.1 Definition of Comparison Outlier Scores

Log-likelihood Ratio Score. Our first comparison score omits the absolute values from the ELD score:

$$LR_i = \sum_{j=1}^{r_i} P_o(v_{ij}) \ln \frac{\theta_o(v_{ij})}{\theta_C(v_{ij})} + \sum_{j=1}^{r_i} \sum_{\mathbf{pa}_i} P_o(v_{ij}, \mathbf{pa}_i) \left(\ln \frac{\theta_o(v_{ij}|\mathbf{pa}_i)}{\theta_o(v_{ij})} - \ln \frac{\theta_C(v_{ij}|\mathbf{pa}_i)}{\theta_C(v_{ij})} \right).$$

By using the **mutual information decomposition**:

$$\ln \frac{\theta_o(v_{ij}|\mathbf{pa}_i)}{\theta_C(v_{ij}|\mathbf{pa}_i)} = \ln \frac{\theta_o(v_{ij}|\mathbf{pa}_i)}{\theta_o(v_{ij})} - \ln \frac{\theta_C(v_{ij}|\mathbf{pa}_i)}{\theta_C(v_{ij})} + \ln \frac{\theta_o(v_{ij})}{\theta_C(v_{ij})}, \quad (5)$$

it can be shown that the ELD score without the absolute values is equivalent to the likelihood ratio, or **log-likelihood difference**:

$$LR(\mathcal{D}_o, B_C, \theta_o) \equiv LOG(\mathcal{D}_o, B_C, \theta_o) - LOG(\mathcal{D}_o, B_C, \theta_C). \quad (6)$$

Assuming maximum likelihood parameter estimation, LR is equivalent to the Kullback-Leibler divergence between the class-level and object-level parameters [11]. The log-likelihood difference compares how well the class-level parameters fit the object data with respect to a particular object, vs. how well the object parameters fit the object data. In terms of the conditional probability parameters, it measures how much the log-conditional probabilities in the class distribution differ from those in the object distribution.

The Feature Divergence Score. The feature divergence outlier score FD uses only part (3) of the ELD score. That is, it considers each feature independent of all others. FD computes the expected log-distance with respect to the single feature value probabilities between the object and the class models. This can be computed using the following formula:

$$FD_i = \sum_{j=1}^n \sum_{k=1}^{r_i} P_o(v_{ij}) \left| \ln \frac{\theta_o(v_{ij})}{\theta_C(v_{ij})} \right| \quad (7)$$

Log-Likelihood Score. In previous work on applying Bayesian networks to outlier detection for i.i.d. (independent and identically-distributed) non-relational data, the outlier metric used was the log-likelihood of a data point [10]. This means that an object was deemed an outlier if the model assigns sufficiently low likelihood to generating its features. For relational data, we use the relational log-likelihood (2) of the target *database*:

$$LOG(\mathcal{D}_o, B_C, \theta_C) = \sum_{i=1}^n \sum_{j=1}^{r_i} \sum_{\mathbf{pa}_i} P_{\mathcal{D}}(v_{ij}, \mathbf{pa}_i) \ln \theta(v_{ij} | \mathbf{pa}_i). \quad (8)$$

5.2 Score Computation Examples

We provide three simple examples with only two variables/features that illustrate the computation of the outlier scores. They are designed so that outliers and normal objects are easy to distinguish, and so that it is easy to trace the behavior of an outlier score. The examples therefore serve as thought experiments that bring out some key strengths and weaknesses of the model-based outlier scores we evaluate. Figure 7 describes the BN representation of the examples. For intuition, we can think of a soccer setting, where each match assigns a value to each attribute F_1 and F_2 for each player.

Table 5 illustrates the computation of the scores. Scores for the F_2 feature are computed conditional on $F_1 = 1$. Expectation terms are computed first for $F_2 = 1$, then $F_2 = 0$. The table shows the cancelling effects in LR . In the high correlation scenario 7(a), the outlier object has a lower probability than the normal class distribution of $Match_Result = 0$ given that $Shot_Efficiency = 1$. Specifically, 0.5 vs. 0.9. The outlier object exhibits a higher probability $Match_Result = 1$ than the normal class distribution, conditional on $Shot_Efficiency = 1$; specifically, 0.5 vs. 0.1. In line 1, column 2 of Table 5 the log-ratios $\ln(0.5/0.9)$ and $\ln(0.5/0.1)$ therefore have different signs. In the low correlation scenario 7(b), the cancelling occurs in the same way, but with the normal and outlier probabilities reversed. The cancelling effect is even stronger for attributes with more than two possible values.

While log-likelihood LOG is a good baseline score for detecting outliers, it fails to detect some clear outliers, as shown in Figure 8. In that example the strength of the correlation among the attributes is the same in both normal and outlier examples and the only difference is in their feature distributions.

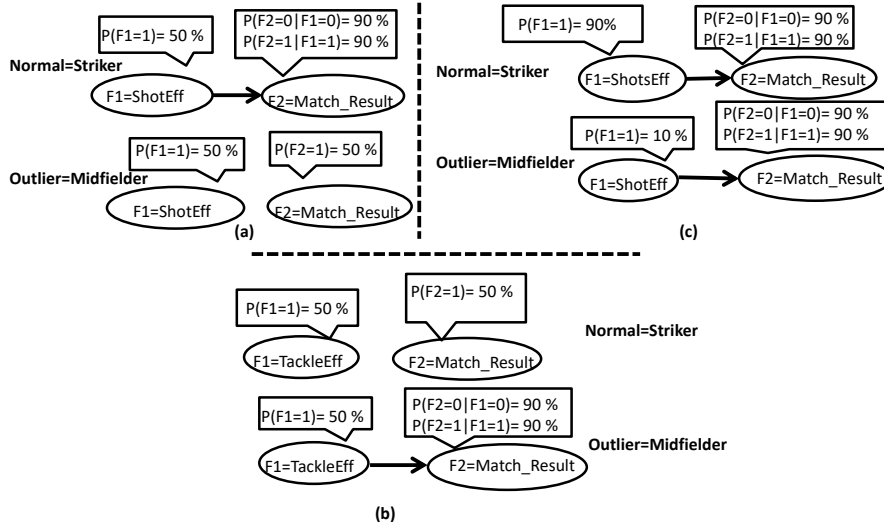


Fig. 7 Illustrative Bayesian networks with two nodes. The networks are not learned from data, but hand-constructed to be plausible for the soccer domain. (a) *High Correlation*: Normal individuals exhibit a strong association between their features, outliers no association. Both normals and outliers have a close to uniform distribution over single features. (b) *Low Correlation*: Normal individuals exhibit no association between their features, outliers have a strong association. Both normals and outliers have a close to uniform distribution over single features. (c) *Single Attributes*: Both normal and outlier individuals exhibit a strong association between their features. In normals, 90% of the time, feature 1 has value 1. For outliers, feature 1 has value 1 only 10% of the time.

Table 5 Example computation of different outlier scores for outliers given the distributions of Figure 7(a),(b). Negative terms are highlighted by boxes; part of their impact cancels out with positive terms. Our new *ELD* metric contains no negative terms due to the use of absolute values.

Score	$F1 = 1$ Computation	$F2 F1 = 1$ Computation	Result
<i>LR</i>	$1/2 \ln(0.5/0.5) = 0$	$1/4 \ln(0.5/0.9) + 1/4 \ln(0.5/0.1)$	0.36
<i>FD</i>	$ \ln(0.5/0.5) = 0$	$1/2 \ln(0.5/0.5) + 1/2 \ln(0.5/0.5) $	0
<i>ELD</i>	0 (no parents)	$1/2 \ln(0.5/0.5) + 1/2 \ln(0.5/0.5) + 1/4 \ln(0.5/0.5) - \ln(0.9/0.5) + 1/4 \ln(0.5/0.5) - \ln(0.1/0.5) $	0.79

Table 5(a): High Correlation Case. Figure 7(a).

Score	$F1 = 1$ Computation	$F2 F1 = 1$ Computation	Result
<i>LR</i>	$1/2 \ln(0.5/0.5) = 0$	$0.5 \cdot 0.9 \ln(0.9/0.5) + 0.5 \cdot 0.1 \ln(0.1/0.5)$	0.26
<i>FD</i>	$ \ln(0.5/0.5) = 0$	$1/2 \ln(0.5/0.5) + 1/2 \ln(0.5/0.5) $	0
<i>ELD</i>	0 (no parents)	$1/2 \ln(0.5/0.5) + 1/2 \ln(0.5/0.5) + 0.5 \cdot 0.9 \ln(0.9/0.5) - \ln(0.5/0.5) + 0.5 \cdot 0.1 \ln(0.1/0.5) - \ln(0.5/0.5) $	0.50

Table 5(b): Low Correlation Case. Figure 7(b).

	Feature1		Feature2		
Normal	0	0.5	0	0	0.9
	1	0.5	0	1	0.1
			1	0	0.1
			1	1	0.9

$$\begin{aligned} \text{Log}(D_{\text{normal}}, B, \theta) &= -(0.5 \times \log 0.5 + 0.5 \log 0.5 + \\ &0.5 \times 0.9 \log 0.9 + 0.5 \times 0.1 \log 0.1 + 0.5 \times 0.1 \log 0.1 + 0.5 \times 0.9 \log 0.9) = 0.44 \\ FD_{\text{Normal}} &= 0.5 \left| \log \frac{0.5}{0.5} \right| + 0.5 \left| \log \frac{0.5}{0.5} \right| + 0.5 \left| \log \frac{0.5}{0.5} \right| + 0.5 \left| \log \frac{0.5}{0.5} \right| = 0 \end{aligned}$$

Outlier	0	0.9	0	0	0.9
	1	0.1	0	1	0.1
			1	0	0.1
			1	1	0.9

$$\begin{aligned} \text{Log}(D_{\text{outlier}}, B, \theta) &= -(0.9 \log 0.5 + 0.1 \log 0.5 + \\ &0.9 \times 0.9 \log 0.9 + 0.9 \times 0.1 \log 0.1 + 0.1 \times 0.1 \log 0.1 + 0.1 \times 0.9 \log 0.9) = 0.44 \\ FD_{\text{outlier}} &= 0.5 \left| \log \frac{0.9}{0.5} \right| + 0.5 \left| \log \frac{0.1}{0.5} \right| + 0.5 \left| \log \frac{0.5}{0.5} \right| + 0.5 \left| \log \frac{0.5}{0.5} \right| = 0.46 \end{aligned}$$

Fig. 8 An example of normal and outlier individuals and their conditional probability tables created using Bayesian network shown in Figure 7(c). Log-likelihood assigns the same score to the normal and individuals in this example, while FD is able to differentiate between these two individuals.

	Feature1		Feature2		
Normal	0	0.5	0	0	0.9
	1	0.5	0	1	0.1
			1	0	0.1
			1	1	0.9

$$\begin{aligned} FD_{\text{Normal}} &= 0.5 \left| \log \frac{0.5}{0.5} \right| + 0.5 \left| \log \frac{0.5}{0.5} \right| + 0.5 \left| \log \frac{0.5}{0.5} \right| + 0.5 \left| \log \frac{0.5}{0.5} \right| = 0 \\ LR(D_{\text{normal}}, B, \theta) &= 0.5 \log \frac{0.5}{0.5} + 0.5 \log \frac{0.5}{0.5} + \\ &0.5 \times 0.9 \log \frac{0.9}{0.9} + 0.5 \times 0.1 \log \frac{0.1}{0.1} + 0.5 \times 0.1 \log \frac{0.1}{0.1} + 0.5 \times 0.9 \log \frac{0.9}{0.9} = 0 \end{aligned}$$

Outlier	0	0.5	0	0	0.5
	1	0.5	0	1	0.5
			1	0	0.5
			1	1	0.5

$$\begin{aligned} FD_{\text{outlier}} &= 0.5 \log \left| \frac{0.5}{0.5} \right| + 0.5 \log \left| \frac{0.5}{0.5} \right| + 0.5 \log \left| \frac{0.5}{0.5} \right| + 0.5 \log \left| \frac{0.5}{0.5} \right| = 0 \\ LR(D_{\text{outlier}}, B, \theta) &= 0.5 \times \log \frac{0.5}{0.5} + 0.5 \log \frac{0.5}{0.5} + \\ &0.5 \times 0.9 \log \frac{0.5}{0.9} + 0.5 \times 0.1 \log \frac{0.5}{0.1} + 0.5 \times 0.1 \log \frac{0.5}{0.1} + 0.5 \times 0.9 \log \frac{0.5}{0.9} = -0.17 \end{aligned}$$

Fig. 9 An example of normal and outlier individuals and their conditional probability tables created using Bayesian network shown in Figure 7(a). FD assigns the same score to the normal and individuals in this example, while LR is able to differentiate between these two individuals.

Conversely, if there is a correlation among the attributes of individuals, the feature divergence score FD fails to take it into account and therefore fails to differentiate between normal and outlier individuals. Figure 9 shows an example where the normal individual has a correlation among its attributes while the outlier object does not have a correlation. The FD metric cannot separate those two individuals.

6 Theoretical Analysis and Comparison

The mathematical analysis of this section relates the KLD and ELD divergences defined in the above sections to other well-known scores. Readers who wish to proceed to the empirical evaluation can omit this section without loss of generality. A well-known Taylor series argument shows that KLD can be approximated by Pearson's χ^2 statistic [37]. We use a similar Taylor series approximation to show that our ELD divergence can be approximated by the total variation distance (TVD), also known as the L1-norm, whose statistical properties are well understood [7, 8]. Our analysis uses the f -divergence framework, which unifies all the standard divergence definitions for probability distributions. The f -divergence analysis shows the generality of the cancellation phenomenon: the first-order Taylor series term vanishes for all f -divergences whose generator f is differentiable at $\lambda = 1$, which includes all standard f -divergences (including KLD), except for those that use absolute values.

The χ^2 approximation for f -divergences. To focus the mathematics on the essential insights, we discuss the case of divergences defined for a single discrete random variable V with possible values v_1, \dots, v_m . The results carry over to the joint BN distributions over discrete variables considered in this paper, by applying the approximation to each parent state. Given two probability distributions P_1 and P_2 over the values of V , an f -divergence is of the form

$$I_f(P_1||P_2) \equiv \sum_{i=1}^m P_1(v_i) f\left(\frac{P_2(v_i)}{P_1(v_i)}\right) \quad (9)$$

where the **generator** $f : R \rightarrow R$ is a convex function such that $f(1) = 0$. The generator transforms the ratio of the two compared probabilities for each possible value; the f -divergence is the weighted average of the transformed ratios. Common divergences can be represented by choosing a suitable generator. For example, KLD is generated by $f(u) = -\ln(u)$, the χ^2 statistic by $f(u) = (1 - u)^2/u$, and the TVD by $f(u) = |u - 1|$ [37].

An f -divergence I_f can be approximated by replacing f with its Taylor series expansion around the point $u = 1$. Assuming that the derivatives $f^{(i)}(1)$ of any order exist, the Taylor series for f takes the form $f(u) = \sum_{i=0}^{\infty} f^{(i)}(1)/i! [u - 1]^i$, where the $i = 0$ term is 0 because $f(1) = 0$. If the generator f is twice differentiable, the first-order term with $i = 1$ vanishes, and the remaining second-order term with $i = 2$ is a scaled χ^2 statistic [37]:

Proposition 1 (Nielsen and Nock 2014) *Assume that the generator f for I_f is twice differentiable.*

1. *The $i = 1$ Taylor series term for I_f around $u = 1$ is 0.*
2. *The $i = 2$ Taylor series term for I_f around $u = 1$ is a scaled χ^2 divergence.*

Therefore the second-order Taylor series approximation to I_f is scaled χ^2 -divergence:

$$I_f(P_1||P_2) \approx \frac{f''(1)}{2} \chi^2(P_1||P_2) = \frac{f''(1)}{2} \sum_{i=1}^m \frac{(P_2(v_i) - P_1(v_i))^2}{P_1(v_i)}.$$

Proof Outline. The first-order term reduces to $f'(1)[\sum_{i=1}^m P_2(v_i) - \sum_{i=1}^m P_1(v_i)]$, which vanishes because both probability measures sum to 1. The second-order term reduces to $\frac{f''(1)}{2} \chi^2(P_1||P_2)$. The details are given in the Appendix 12.

The TVD approximation for ELD. Our ELD metric is an example of transforming an f -divergence with absolute values by replacing the generator f with $|f|$; in our case, replacing $-\ln(u)$ by $|\ln(u)|$ (cf. Equation (6)). Because $|x|$ is not differentiable at 0, the absolute value generator $|f|$ is generally not differentiable at $u = 1$, so Proposition 1 does not apply. To apply the Taylor series analysis, we can rewrite the absolute value divergence as a positive sum where $f(u) > 0$ and a negative sum where $f(u) < 0$.

$$I_{|f|}(P_1||P_2) \equiv \sum_{i: f(\frac{P_2(v_i)}{P_1(v_i)}) > 0} P_1(v_i) f\left(\frac{P_2(v_i)}{P_1(v_i)}\right) - \sum_{i: f(\frac{P_2(v_i)}{P_1(v_i)}) < 0} P_1(v_i) f\left(\frac{P_2(v_i)}{P_1(v_i)}\right)$$

In the Taylor series approximation for ELD (using $f = -\ln(u)$ in Equation (6)), cancellation is avoided, so the first-order term does *not* vanish, and is in fact equivalent to the Total Variation Distance:

Proposition 2 *The $i = 1$ term in the Taylor series approximation for ELD around $u = 1$ is total variation distance. Therefore the first-order Taylor series approximation for ELD is total variation distance:*

$$ELD(P_1||P_2) \approx TVD = \sum_i |P_2(v_i) - P_1(v_i)|$$

7 Experimental Evaluation and Comparison

All the experiments were performed on a 64-bit Centos machine with 4GB RAM and an Intel Core i5-480 M processor. The likelihood-based outlier scores were computed with SQL queries using JDBC, JRE 1.7.0. and MySQL Server version 5.5.34. We utilized the synthetic datasets and real-world datasets from the soccer, hockey, movie and biology domains.

Premier League Statistics		IMDB Statistics		NHL Statistics	
Number of Teams	20	Number of Movies	3060	Number of Teams	30
Number of Players	484	Number of Directors	220	Number of Players	921
Number of Matches	380	Number of Actors	98690	Number of Matches	720
avg player per match	26.01	avg actor per movie	36.42	avg player per match	29

Table 6 Summary Statistics for the IMDB and PL and NHL data sets

7.1 Synthetic Datasets

We generated three synthetic datasets with normal and outlier players using the distributions represented in the three Bayesian networks of Figure 7. The main goal of designing synthetic experiments is to test the methods on easy to detect outliers. We used the *mlbench* package in *R* to generate synthetic features in matches, following these distributions for 240 normal players and 40 outliers. (There were 280 players in our Premier League dataset.) Each player participates in 38 matches, similar to the Premier League’s real-world data. Each match assigns a value to each feature F_1 and F_2 for each player. This design yields the following three synthetic datasets.

High Correlation Normal individuals exhibit a strong association between their features, outliers no association. Both normals and outliers have a close to uniform distribution over single features. See Figure 7(a).

Low Correlation Normal individuals exhibit no association between their features, outliers have a strong association. Both normals and outliers have a close to uniform distribution over single features. See Figure 7(b).

Single Features Both normal and outlier individuals exhibit a strong association between their features. In normals, 90% of the time, feature 1 has value 1. For outliers, feature 1 has value 1 only 10% of the time. See Figure 7(c).

7.2 Real-world Datasets

Our datasets and code are available on-line [44]. The following is the list of the real-world datasets we used in this work:

Premier League Data The Opta data were released by Manchester City. It lists box scores, that is, counts of all the ball actions within each game by each player, for the 2011-2012 season. For each player in a match, our data set contains eleven player features. For each team in a match, there are five features computed as player feature aggregates, as well as the team formation and the result (win, tie, loss). There are two relationships, *Appears_Player*(P, M), *Appears_Team*(T, M).

Table 7 lists the populations and features. Table 6 shows summary statistics for three of real-world datasets.

Individuals	Features
PL-Player per Match	<i>TimePlayed, Goals, SavesMade, ShotEff, PassEff, WinningGoal, FirstGoal, PositionID, TackleEff, DribbleEff, ShotsOnTarget</i>
PL-Team per Match	<i>Result, TeamFormation, $\sum Goals, \mu ShotEff, \mu PassEff, \mu TackleEff, \mu DribbleEff$.</i>
IMDB-Actor	<i>Quality, Gender</i>
IMDB-Director	<i>Quality, avgRevenue</i>
IMDB-Movie	<i>year, isEnglish, Genre, Country, RunningTime, Rating by User</i>
IMDB-User	<i>Gender, Occupation.</i>
NHL-Player per Match	<i>Goals, Assists, Points, PowerPlayTime, PlusMinus, Penalties, Shots, Hits, BlockedShots, Giveaways, ShotHandedTime, TimeOnIce</i>
NHL-Team per Match	<i>Goals, GoalDifference</i>

Table 7 Attribute Features.

IMDb Data The Internet Movie Database (IMDb) is an on-line database of information related to films, television programs, and video games. The IMDb website offers a dataset containing information on cast, crew, titles, technical details and biographies into a set of compressed text files. We preprocessed the data like [40] to obtain a database with seven tables: one for each population and one for the three relationships *Rated*(*User*, *Movie*), *Directs*(*Director*, *Movie*), and *ActsIn*(*Actor*, *Movie*).

National Hockey League Data The NHL provides information about the sequences of play-by-play events. We used the data crawled by [52] that consists of the player game statistics for the seasons 2009-2013. The box scores summarize player statistics for each match, a total of 13 features.

Mutagenesis Data This dataset contains mutagenic activity of 188 compounds. 125 of these compounds have positive levels of log mutagenicity that are labelled “active”. The remaining compounds are labelled “inactive” and constitute the source of negative examples. In this dataset we considered examples of active compounds as the normal population and the inactive ones as the outlier.

Normal class	#Normal	Outlier	#Outlier class
Striker	153	Goalie	22
Midfielder	155	Striker	74
Drama	197	Comedy	47
Defender	186	Forward	38
Positive Compound	125	Negative Compound	63

Table 8 Outlier/normal Objects in Real-World Datasets.

7.3 Evaluation Techniques for Outlier Detection

Measuring the effectiveness of outlier detection methods is not often an easy task. Most of the time ground truth information, that shows which data points are outliers, is unavailable.

Several techniques have been employed in the literature to evaluate the performance of outlier detection methods:

1. Intuitive evaluation: case studies have been extensively used in the literature to evaluate outliers [2]. In Section 8.3 we use this method of evaluation for the top n ranked detected outliers and we try to explain and make sense of the detected outliers.
2. Synthetic data generation: another approach to evaluate anomaly detection methods is generating synthetic data and injecting synthetic outliers into the data [2]. We have designed and developed three synthetic datasets as discussed in section 7.1.
3. Anomaly injection: anomalies are injected into real-world datasets. Outlier detection methods are expected to detect the injected data points as outliers [4]. We employ this approach in our real-world datasets. In real-world data, there is no ground truth about which objects are outliers. To address this issue, we employ a one-class design: we learn a model for the class distribution, with data from that class only. Then we rank all individuals from the normal class together with all objects from a contrast class treated as outliers, to test whether an outlier score recognizes objects from the contrast class as outliers. Table 8 shows the normal and contrast classes for three different datasets. In-class outliers are possible, e.g. unusual strikers are still members of the striker class. Our case studies describe a few in-class outliers. In the soccer data, we considered only individuals who played more than 5 matches out of a maximum 38. For the three synthetic datasets, the scores are used to rank all 280 synthetic players, 240 of which are normal and 40 are outliers. The disadvantage of this evaluation metric is that the real-world data may also contain anomalies, known as in-class outliers. However, this metric treats only the injected data points as true positives and will score anything other than those as false positives.

7.4 Methods Compared

We compare three types of approaches, based on relational model likelihood, aggregation, and distance.

Likelihood-based Outlier Scores. The first approach evaluates the likelihood-based outlier scores described in Section 5. For relational Bayesian network structure learning we utilize the previous learn-and-join algorithm (LAJ), which is a state-of-the-art BN structure learning method for relational data [50]. The LAJ algorithm employs an iterative deepening strategy, which can be described as a search through a lattice of table joins. For each table join, different BNs are learned and the learned edges are propagated from smaller to larger table joins. For a full description, complexity analysis, and learning time measurements, please see [50]. We used the implementation of the LAJ algorithm due to its creators [26]. Of the likelihood-based scores, the likelihood ratio LR and our ELD score fit in the EMM framework as they compare an object model with a population model.

Aggregation-based Methods. The second approach first “flattens” the structured data into a matrix of feature vectors, then applies standard matrix-based outlier detection methods. We refer to such methods as **aggregation-based** (cf. Figures 2). For example, this was the approach taken by Breunig *et al.* for identifying anomalous players in sports data [9]. Following their paper, for each continuous feature in the object data, we use the average over its values, and for each discrete feature, we use the occurrence count of each feature value in the object data. Aggregation tends to lose information about correlations. Our experiments address the empirical question of whether this loss of information affects outlier detection. We evaluated three standard matrix-based outlier detection methods: Density-based LOF [9], distance-based $KNNOutlier$ [43] and subspace analysis $OutRank$ [35]. These represent common, fundamental approaches for vectorial data. Like ELD , subspace analysis is sensitive to correlations among features. We used the available implementation of all three data matrix methods from the state of the art data mining software $ELKI$ [1]. We used $PRO-CLUS$ as the clustering function for $OutRank$ as recommended by [35].

Relational Distance-based Method. The third approach is based on a first-order distance measure that was developed and first used for the instance-based learning system RIBL2 [25]. This measure has proven successful in several applications [29, 24]. We employ this metric and compute the distance between any two individuals in our population domain. Then, we rank the individuals based on their mean distance to the normal population.

8 Empirical Results

We present results regarding computational feasibility, predictive performance, and case studies.

8.1 Computational Cost of the *ELD* Score.

Table 9 shows that the computation of the *ELD* value for a given target object is feasible. On average, it takes a quarter of a minute for each soccer player, and one minute for each movie. This includes the time for parameter learning from the object database. Learning the class model BN takes longer, but needs to be done only once for the entire object class. *The BN model provides a crucial low-dimensional representation of the distribution information in the data.*

Dataset	Class Model	Average per Object
PL: Strikers vs. Goalies	4.14	0.25
PL: Midfielder vs. Goalies	4.02	0.25
IMDb: Drama vs. Comedy	8.30	1.00
PL: Forward vs. Defender	5.30	0.35
Mutagenesis: Positive Compound vs. Negative Compound	1.40	0.1

Table 9 Time (min) for computing the *ELD* score.

8.2 Detection Accuracy

Several accuracy scores have been developed to measure the performance of outlier ranking methods [2]. Arguably the most widely used score is the *area under curve (AUC)* of the receiver operating characteristic *ROC* curve [17, 10, 35]. The relationship between false positive rate (1- Specificity) and true positive rate (Sensitivity) is captured by the *ROC* curve. Ideally, the best performance is achieved when we have the highest sensitivity and the highest specificity. The maximum values for *AUC* is, 1.0 indicating a perfect ranking with 100% sensitivity and 100% specificity. In order to compute the *AUC* value, we used the *R* package *ROCR* [53]. Given a set of outlier scores, one for each object, this package returns an *AUC* value.

AUC Accuracy in Synthetic Datasets. Table 10 shows the *AUC* values for probabilistic methods and the outlier detection methods. On the synthetic data, it ought to be easy to distinguish the outliers and most methods detect most outliers well. However, the EMM methods *LR* and *ELD* are the only scores that achieves perfect or near perfect detection across all three synthetic datasets. *RIBL* is the only method that fails in detecting outliers. *RIBL* computes the distance between individuals by computing the distance of each

Dataset	<u>ELD</u>	<i>LOG</i>	<u>LR</u>	<i>FD</i>	<i>RIBL</i>	<i>LOF</i>	<i>OutRank</i>	<i>KNN Outlier</i>
High Correlation	1.00	0.99	0.97	0.89	0.50	0.68	0.99	0.97
Low Correlation	1.00	0.97	0.99	0.42	0.50	0.58	0.83	0.97
Single Feature	1.00	0.79	1.00	1.00	0.50	0.63	0.88	0.86

Table 10 AUC of the *ELD* Outlier detection methods in Synthetic datasets. The EMM methods are underlined.

Dataset	<u>ELD</u>	<i>LOG</i>	<u>LR</u>	<i>FD</i>	<i>RIBL</i>	<i>LOF</i>	<i>OutRank</i>	<i>KNN Outlier</i>
PL: Strikers vs. Goalies	0.89	0.61	0.65	0.61	0.71	0.61	0.60	0.61
PL: Midfielders vs. Strikers	0.66	0.45	0.55	0.59	0.78	0.76	0.71	0.58
IMDb: Drama vs. Comedy	0.70	0.66	0.66	0.64	0.70	0.51	0.68	0.68
NHL: Defender vs. Forward	0.78	0.58	0.87	0.79	0.71	0.73	0.73	0.66
Mutagenesis: Positive vs Negative	0.86	0.64	0.81	0.70	0.70	0.51	0.57	0.53

Table 11 AUC outlier detection methods in real-world datasets.

instance of an individual to its nearest instance from other instances of the other individual. When comparing two individuals with only two binary features, *RIBL* results in the same distance from the center of normal community for each individual.

AUC Accuracy in Real-world Datasets. Table 11 shows the *AUC* values for aggregation-based methods compared to *ELD* and *RIBL* in the real-world datasets. *RIBL*’s recursion depth bound is set to 2. The top performance of *RIBL* and the model-based methods is substantially better than aggregation-based methods on all datasets, confirming that it is important to develop outlier detection methods based on relational statistics for the relational data. The EMM score *ELD* works better than *RIBL* or equally good in all the datasets except Midfielder vs. Strikers. In that dataset *ELD* finds many in-class exceptional midfielders. However, *RIBL* seems to perform better in distinguishing these two classes from each other. *RIBL*’s behavior in synthetic and real-world datasets suggests that when we have a small number of discrete features in our dataset, *RIBL* may not be a good candidate to compute distance and hence outlier detection. But for numeric features it seems to be a good alternative.

Comparing the two EMM methods *ELD* and *LR*, using distance rather than expectation allows *ELD* to achieve a substantially higher AUC than *LR* on all datasets except for the NHL. On the NHL dataset, the *LR* metric outperforms both *RIBL* and *ELD*, which is an example of how taking absolute values does not always result in better detection performance.

On the synthetic data, we compare the two EMM methods further with precision@r% another metric used in a previous study on relational outlier detection [19], which is computed as follows: 1) Sort the outlier scores obtained by an outlier ranking method in descending order. 2) Return the fraction of the top *r* percent that are in fact outliers. Precision@r% focuses on the ability to detect clear outliers. As Table 12 shows, the *ELD* score is the best for this metric.

Conclusion. The outlier detection methods with the best over all performance are the two based on the Exceptional Model Mining Framework, using

Table 12 Precision@r% of Outlier detection methods in Synthetic datasets.

Dataset	Percentage	<i>ELD</i>	<i>LOG</i>	<i>LR</i>	<i>FD</i>	<i>LOF</i>	<i>OutRank</i>	<i>KNN Outlier</i>
High-Correlation	1%	1.00	0.91	0.73	0.47	0.11	0.53	0.48
	5%	1.00	0.95	0.85	0.65	0.22	0.50	0.65
Low-Correlation	1%	1.00	0.93	0.87	0.14	0.10	0.00	0.06
	5%	1.00	0.95	0.90	0.25	0.25	0.10	0.57
Single-Feature	1%	1.00	0.81	1.00	1.00	0.46	1.00	0.51
	5%	1.00	0.89	1.00	1.00	0.55	1.00	0.54

the traditional log-likelihood ratio (or Kulback-Leibler divergence), and our novel expected log-distance metric *ELD*. While the *ELD* score does not uniformly outperform *LR*, the improvement is frequent and substantial enough to recommend it as an alternative for practical applications.

8.3 Case Studies

For a case study, we examine the three top outliers as ranked by *ELD*, shown in Table 13. The aim of the case study is to provide a qualitative sense of the outliers indicated by the scores. Also, we illustrate how the BN representation leads to an interpretable ranking. Specifically, we employ a *feature-wise decomposition* of the score combined with a *drill down* analysis:

1. Find the node V_i that has the highest ELD_i divergence score for the outlier object.
2. Find the parent-child combination that contributes the most to the ELD_i score for that node.
3. Decompose the *ELD* score for the parent-child combination into feature and mutual information component.

We present strong associations—indicated by the *ELD*’s mutual information component—in the intuitive format of association rules. This analysis can also be applied to *LR* using the form (5); we focus on *ELD* for compactness.

Strikers vs. Goalies In real-world data, a rare object may be a *within-class outlier*, i.e., highly anomalous even within its class. In an unsupervised setting without class labels, we do not expect an outlier score to distinguish such an in-class outlier from outliers outside the class. An example is the striker Edin Dzeko. He is a highly anomalous striker who obtains the top *ELD* divergence score among both strikers and goalies. His *ELD* score is highest for the Dribble Efficiency feature. The highest *ELD* score for that feature occurs when Dribble Efficiency is low, and its parents have the following values: Shot Efficiency high, Tackle Efficiency medium. Looking at the single feature divergence, we see that Edin Dzeko is indeed an outlier in the Dribble Efficiency subspace: His dribble efficiency is low in 16% of his matches, whereas a randomly selected striker has low dribble efficiency in 50% of their matches. Thus, Edin Dzeko is an unusually good dribbler. Looking at the mutual information component of

ELD, i.e., the parent-child correlations, for Edin Dzeko the confidence of the rule

$$\text{ShotEff} = \text{high}, \text{TackleEff} = \text{medium} \rightarrow \text{DribbleEff} = \text{low}$$

is 50%, whereas in the general striker class it is 38%.

Midfielders vs. Strikers For the single feature score, Robin van Persie is recognized as a clear striker because of the *ShotsOnTarget* feature. It makes sense that strikers shoot on target more often than midfielders. Robin van Persie achieves a high number of shots on targets in 34% of his matches, compared to 3% for a random midfielder. The mutual information component shows that he also exhibits unusual correlations. For example, the confidence of the rule

$$\text{ShotEff} = \text{high}, \text{TimePlayed} = \text{high} \rightarrow \text{ShotsOnTarget} = \text{high}$$

is 70% for van Persie, whereas for strikers overall it is 52%.

The most anomalous midfielder is Scott Sinclair. His most unusual feature is *DribbleEfficiency*: For feature divergence, he achieves a high dribble efficiency 50% of the time, compared to a random midfielder with 30%.

Drama vs. Comedy The top outlier rank is assigned to the within-class outlier *BraveHeart*. Its most unusual feature is *ActorQuality*: In a random drama movie, 42% of actors have the highest quality level 4, whereas for *BraveHeart* 93% of actors achieve the highest quality level.

The *ELD* score identifies the comedies *BluesBrothers* and *AustinPowers* as the top out-of-class outliers. In a random drama movie, 49% of actors have casting position 3, whereas for *AustinPowers* 78% of actors have this casting position, and for *BluesBrothers* 88% of actors do.

Defender vs. Forward The first two players in the ranking belong to forward group and are identified as outliers for their unusual high value for points. Eric Staal has *points* = 2 in 30% of his matches, while an average player has that value for points only on 6% of his matches. Dustin Byfuglien is discovered as a within-class outlier. His most unusual feature is *PowerPlayTime*. While an average player in the population has *PowerPlayTime* = 2 for only 33% of the times, he has that value for 79% of the times.

9 Limitation of Model-based outlier detection

The main limitations of the work presented in this paper are the following:

1. Limitation of Approach:

- (a) Our proposed method ranks potential outliers, but does not set a threshold for a binary identification of outlier vs. non-outlier.
- (b) Our current Bayesian network learning method can only be applied to discrete data. Prior to learning the model, we take an extra step in data preprocessing and convert continuous data into discrete, which naturally causes some information loss.

Strikers (Normal) vs. Goalies (Outlier)					
PlayerName	Position	ELD Rank	ELD Max Node	ELD Node Score	FD Max feature Value
Edin Dzeko	Striker	1	DribbleEfficiency	83.84	DE=low
Paul Robinson	Goalie	2	SavesMade	49.4	SM=Medium4
Michel Vorm	Goalie	3	SavesMade	85.9	SM=Medium
Midfielders (Normal) vs. Strikers (Outlier)					
PlayerName	Position	ELD Rank	ELD Max Node	ELD Node Score	FD Max feature Value
Robin Van Persie	Striker	1	ShotsOnTarget	153.18	ST=high
Wayne Rooney	Striker	2	ShotsOnTarget	113.14	ST=high
Scott Sinclair	Midfielder	6	DribbleEfficiency	71.9	DE=high
Drama (Normal) vs. Comedy (Outlier)					
MovieTitle	Genre	ELD Rank	ELD Max Node	ELD Node Score	FD Max feature Value
Brave Heart	Drama	1	ActorQuality	89995.4	a_quality=4
Austin Powers	Comedy	2	Cast_Position	61021.28	Cast_Num=3
Blue Brothers	Comedy	3	Cast_Position	24432.21	Cast_num=3
Defender (Normal) vs. Forward (Outlier)					
PlayerName	Position	ELD Rank	ELD Max Node	ELD Node Score	FD Max feature Value
Eric Staal	Forward	1	Points	49.57	Points=2
Phil Kessel	Forward	2	Points	43.34	Points=2
Dustin Byfuglien	Defender	3	Power_PlayTime	25.65	PP_time=2

Table 13 Case study for the top outliers returned by the log-likelihood distance score *ELD*

- (c) Our generative model-based methods learn a generic Bayesian network structure for the entire population, so the detected outliers are global outliers. However, there are more complex outliers that locally deviate from their subgroups and can be detected only by subgroup comparison. One direction for future work is to first detect subgroups in the population and then apply outliers within subgroups.²

2. Limitation of Data Analysis:

In this work, to simplify the outlier detection task, we used only part of the full information available in our rich datasets. The model-based outlier detection can be extended in future work to take advantage of the full information, in the following manner.

- In the Premier League dataset, players are naturally related to one another and modeling the interaction between players can be another way to detect anomalous players. The graph-based features, such as detecting near-clique nodes and star nodes, proved to be efficient in discovering patterns for anomaly detection task as shown in ODDBALL [3].
- In this paper we did not use the temporal information available in the data. In the learning process we do not give a higher weight (importance) to the more recent action (performance) of an individual. This point is especially important when applying the methods to dynamic data or the data that are collected over long periods of time.
- In the datasets that we used for the experiments, we did not have the missing value problem. Therefore, we did not incorporate ways to estimate missing values in our modeling. However, real-world datasets may involve arbitrary patterns of missing data. Maximum likelihood density estimation is a way to estimate such values. We leave this feature for future work.

² We are indebted to Daniel Lowd for this point.

10 Correlation with Success

The aim of this section is to compare the *ELD* metric with other meaningful metrics for comparing individuals. Our reference metrics are success rankings of individuals selected for a specific domain, shown in Table 14. We use the same data as in our other experiments, described in Section 7 except Mutagenesis as there was no meaningful success metrics we could find in that dataset.

Success rankings are one of the most interesting features to users. Strong correlations between the *ELD* metric and meaningful success metrics provide evidence that the *ELD* metric is meaningful as well. We measure correlation strength by the standard Pearson correlation coefficient ρ . The coefficient ranges from -1 to 1, where 0 means no correlation and 1 or -1 indicates maximum strength [18].

The observed correlations are remarkable in at least two respects. 1) the strength of the correlation between the *ELD* metric and salary is high as shown in Table 16 : coefficients range from 0.45 to 0.82. 2) We observe this phenomenon across different domains, different types of individuals and different success metrics.

Dataset	Success Metric	Min	Max	Standard Dev.	Mean
IMDb	Sum of Rating	1.0	14795	1600.22	1057.58
PL-Player	TimePlayed	5.0	3420	1015.69	1484.00
PL-Player	Normalized Salary	0.007	0.28	0.62	0.10
PL-Player	Sum of Shot Efficiency	0	82	9.87	6.53
PL-Team	Standing	1.0	20	5.91	10.50
NHL-Player	Power Play Time	0	669	106.78	84.38
NHL-Player	Time on Ice	4	2099	278.03	1187.31
NHL-Player	Assists	0	4	0.49	0.20

Table 14 Success metrics and their distributions.

For a population with a diverse set of skills and resources, being different from the generic class can be interpreted as both exceptionally better or worse than normal population. In the domains we study in this data, we found that higher *ELD* scores indicate exceptionally good individuals but not exceptionally bad individuals. Our interpretation of the correlation between *ELD* and success, rather than failure, is that our domains featured skilled individuals, such that the average is quite successful already. For example, in Premier League we expect most players to be in the range of good players. Therefore, deviating from the rest of the population is a signal for detecting exceptionally good players. Our *ELD*-success scatterplots below provide empirical evidence for this interpretation: we typically see a large cluster of individuals around the origin, meaning that their success level is normal and their *ELD* score is low, see Figure 11.

Team	Standing
Top Teams	-0.71
Bottom Teams	-0.33
All Teams	-0.13

Table 15 Correlation between *ELD* metric and standing of Teams. The best standing is place 1.

Class	Time Played	Salary	Saves Made	Shots On target	Pass Efficiency
Strikers	0.86	0.82	NA	0.79	NA
Midfielders	0.80	0.45	NA	NA	0.77
Goalies	0.77	NA	0.74	NA	NA
All players	0.18	0.56	NA	NA	NA

Table 16 Correlation between *ELD* metric and success metrics of Soccer Players.

Genre	Sum of Rating	Average of Rating	Number of Rating
Action	0.67	0.30	0.72
Drama	0.76	0.29	0.81
Comedy	0.85	0.41	0.84
All Movies	0.56	0.17	0.60

Table 17 Correlation between *ELD* metric and success metric of Movies.

10.1 Methodology

We report the correlations between the *ELD* metric and metrics of success for a specific domain. We also focus on some unusually successful individuals as case studies. In considering the correlation between *ELD* and success, it is useful to investigate subgroups of individuals to ensure an apples-to-apples comparison [54]. For instance, the attributes that lead to success are different for strikers and goalies. Accordingly, we report correlations for subgroups as well as entire classes of individuals. We found that *ELD* shows the strongest correlation with success for all metrics and subgroups, except for the log-likelihood metric; the details are given below in Section 10.3.

10.2 Correlations between the *ELD* outlier metric and success

The next three tables summarize the observed correlations between success and *ELD* metrics: Teams in Table 15, Players in Table 16, Movies in Table 17.

10.2.1 Soccer Teams

Team Standing The most successful team has Standing=1 and the least successful team has Standing=20 in the 2011-2012 Season. For the top teams, there is a very strong negative correlation emerges between *ELD* and standing: teams with higher *ELD* achieve a better (lower) standing.

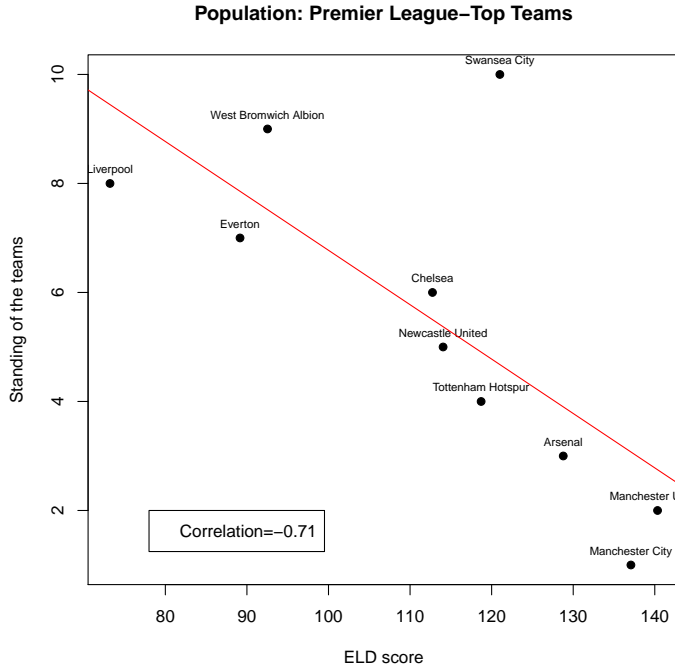


Fig. 10 Teams: Team Standing vs. *ELD* for the top teams in Premier League.

Figure 10 shows the correlation of *ELD* with team success metrics in a scatter plot. The top two teams Manchester City and Manchester United stand out very strongly in terms of the *ELD* metric (bottom right corner).

10.2.2 Soccer Players

Players Time Played is the total time that a player played over all matches in the season. This metric was shown to correlate strongly with other success metrics, such as salary, in soccer data [56]. For each subgroup, there is a strong positive correlation with *ELD*, meaning that atypical players with higher *ELD* tend to play more minutes.

Salary is probably the most obvious, and at the same time often the most misleading way to measure success of the players. Previous studies suggest that salary of the players does not always follow their performance in many sports such as Baseball and Soccer [22,12]. They show that pay cannot be explained only by past performance and there are other factors that are hard to quantify and have great effects on the salaries.

We manually collected salaries of 120 players that we could find on-line. Table 16 and Figures 11 and 12 show the correlation between *ELD* and this

success metric. The correlation is high, especially for Strikers. We found salary data for only 5 goalies. We discuss the relatively weaker salary correlation for midfielders in more detail below.

Shots on Target applies to strikers only. This is defined as any shot attempt that would or does enter the goal if left unblocked. We record the total number of these shots over all matches of the strikers only. This metric was shown to correlate strongly with *ELD* (see Table 16, Figure 11(b)).

Figure 11 plots *ELD* against striker success metrics. We observe a large cluster around the origin, which points to a large base of normal strikers with both salaries and low *ELD* scores.

Saves Made applies to Goalies only. It is defined as the total number of saves that goalies had made over all the matches. This metric shows a strong correlation with *ELD* as well (see Table 16, Figure 12(b)).

Figure 12 shows the correlation of *ELD* with Goalie success metrics in a scatter plot. Goalies do not vary much in terms of the time they play. Wayne Hennessey has the highest number of Saves Made and also an unusually high *ELD* score, although not the highest.

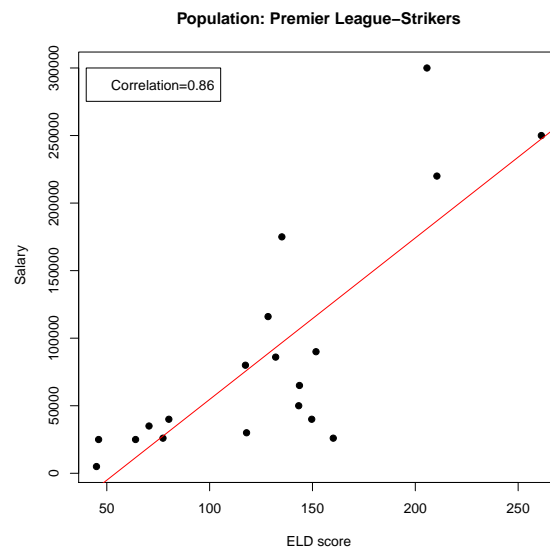
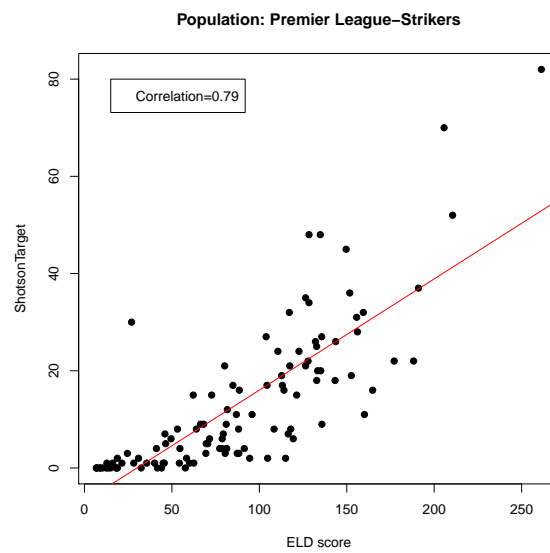
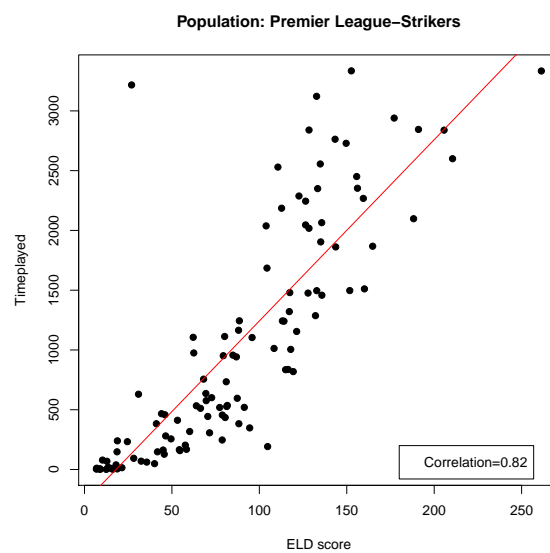
Midfielder Salary We omit a scatterplot for midfielder salary vs. *ELD* because it is less informative due to the weaker correlation (0.45). To investigate the reason for the weaker correlation, we picked two midfielders: 1) Stephane Sessegnon who has been ranked second in the *ELD* ranking but does not draw a large salary. 2) Steven Gerrard who is a very well known player and ranked second in the Salary ranking but according to the *ELD* score, he has been ranked 21. Based on domain knowledge, we picked some of the features that are relevant to midfield performance from the raw data and compared the feature statistics for these two players. Table 18 shows the details of their appearances in different matches. Sessegnon scored higher than Gerrard in three out of the four categories (passes and Time Played). However, his salary was much lower than Gerrard's. Indeed his next contract with West Bromwich Albion netted him a club record fee. This is an example of how our *ELD* metric can identify players who are underpaid relative to their potential.

Name	Team	age	Salary Ranking	<i>ELD</i> Ranking	Time Played	Unsuccessful Passes	Successful Long Passes	Successful corners
Steven Gerrard	Liverpool	31	2	21	1212 min	244	52	25
Stephane Sessegnon	Sunderland	26	22	2	3133 min	231	82	15

Table 18 Comparison of two midfielders.

10.2.3 Hockey Players

For the hockey players the features that we have selected as success metrics are as follows: *TimeonIce*, that can be defined as the total amount of time a

(a) Strikers: Salary vs *ELD*.(b) Strikers: Shots On Target vs *ELD*(c) Strikers: Time played vs *ELD*.**Fig. 11** Correlations in Strikers population

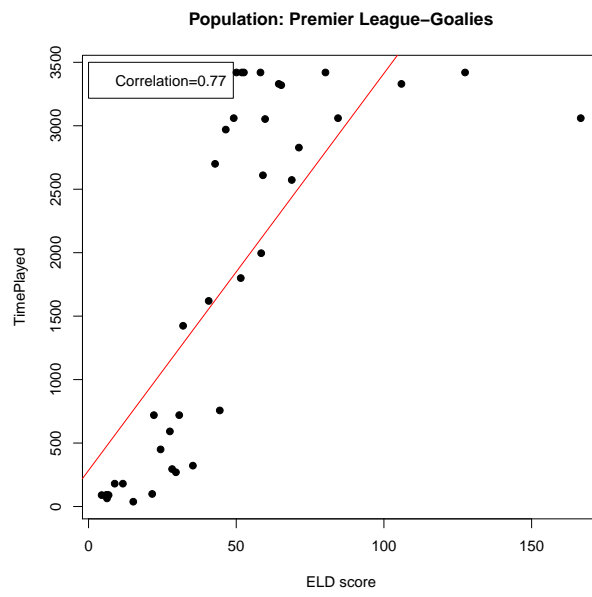
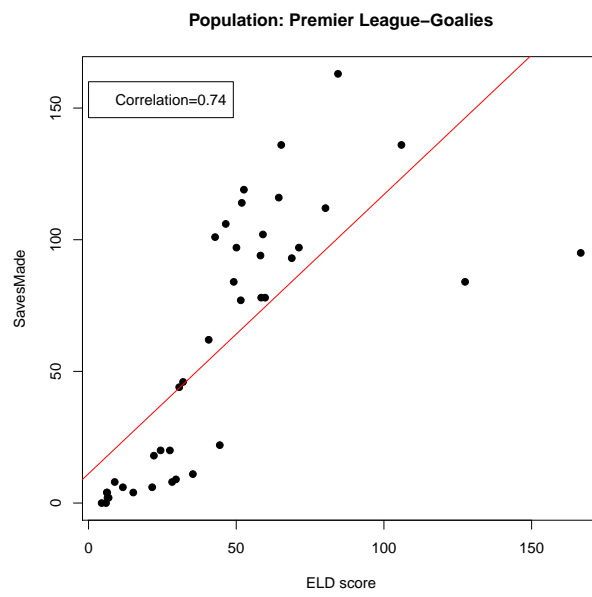
(a) Goalies: sum of time played vs *ELD*.(b) Goalies: sum of saves made vs *ELD*

Fig. 12

Class	Time On Ice	Power Play Time	Assists	Goals
Forwarder	0.79	0.78	0.81	0.78
Defender	0.66	0.57	0.60	0.40

Table 19 Correlation between *ELD* metric and success metrics of NHL Players.

player has played over the course of a season. *PowerPlayTime* which is the total amount of power player time for a given player in a season. *Assists* is the total number of any actions that led to a goal. And finally total number of *Goals* that a given player has scored. The correlation between these features and *ELD* is shown in Table 19. *ELD* has high correlations in features of both categories, however, it is stronger in Forwarder subgroup.

Movie Sum of Ratings

10.2.4 Movies

Movie Sum of Ratings is the number of user ratings of a movie. Table 17 shows a high correlation with the *ELD* metric. The highest correlation obtains for the genre Comedy (0.84). The correlation between a movie and the sum of its ratings is equally strong, but the correlation with its average rating is much weaker. Thus the *ELD* score is related mainly with how many users have rated the movie rather than with how they have rated it. The number of ratings is a meaningful success metric as it indicates the number of people who have gone to see a movie.

10.3 Correlations between other outlier metrics and success

In Section 3.4 and 5 we introduced other metrics that could be used in order to detect outliers. In this section we discuss the correlation between those metrics and success. The correlation between *ELD* and success is always stronger than *FD* and *LR* in all the datasets and subgroups, therefore we omit those results. The Log-likelihood is the only metric that results in a stronger correlation in some datasets and subgroups. Table 20 shows the subgroups that Log-likelihood has a stronger correlation than the *ELD* metric. In a way a high correlation with success can be interpreted as detecting in-class outliers and this results shows that Log-likelihood could be an alternative score to detect those type of outliers.

11 Conclusion and Future Work

We presented a new approach for applying Bayesian networks to object-relational outlier detection, a challenging and practically important topic for machine learning. The key idea is to apply the Exceptional Model Mining framework

Subgroup	Success metric	<i>ELD</i> correlation	Log-likelihood correlation
Comedy	Sum of Rating	0.85	0.87
Drama	Sum of Rating	0.78	0.82
PL-Midfielder	Pass Efficiency	0.77	0.89
PL-Midfielder	Time Played	0.80	0.86
PL-Goalies	Time Played	0.77	0.87
PL-Goalies	Saves Made	0.74	0.85
NHL-Forward	Time on Ice	0.79	0.95
NHL-Forward	Goals	0.78	0.83

Table 20 Subgroups and success metrics for which the log-likelihood metric’s correlates with success stronger than *ELD*.

as follows. First, use statistical-relational learning to construct from relational data a graphical model. Then learn one set of parameter values that represent class-level associations, another set to represent object-level associations, and compare how well each parametrization fits the relational data that characterize the target object. The classic metric for comparing two parametrized models is their likelihood ratio. As an novel alternative, we define a new relational log-likelihood distance metric via two transformations: (1) a mutual information decomposition, and (2) replacing log-likelihood differences by log-likelihood distances. This metric combines a single feature component, where features are treated as independent, with a correlation component that measures the deviation in the features’ mutual information.

In experiments on three synthetic and four real-world outlier sets, the EMM methods based on likelihood ratio and the log-likelihood distance achieved the best detection accuracy. On all but one real-world dataset, log-likelihood distance outperformed the likelihood ratio. As an alternative to model-based EMM, converting the structured data to a flat data matrix via aggregation had a negative impact on outlier detection. Case studies showed that the EMM scores lead to easily interpreted rankings. We found that the log-distance score correlated with success metrics to a surprising degree, across different domains and classes of individuals. The correlation with metrics of independent interest corroborates that the log-distance score produces meaningful and interesting results.

There are several avenues for future work. (i) A limitation of our current approach is that it ranks potential outliers, but does not set a threshold for a binary identification of outlier vs. non-outlier. (ii) Our divergence uses expected L1-distance for interpretability, but other distance scores like L2 could be investigated as well. (iii) Extending the expected L1-distance for continuous features is a useful addition. (iv) Compare our metric with the interestingness measures that have been developed for relational exception mining. (v) In the movie and soccer domains, our metric identified exceptionally successful individual objects, but not exceptionally unsuccessful ones. Our hypothesis was that in these domains, individuals have gone through a rigorous selection process, so the normal baseline performance is high. While we provided evidence for this hypothesis, it can be further investigated, by applying our outlier

detection to datasets that feature a range of skills, rather than professional performance. (vi) The theoretical properties of our *ELD* metric are important to understand. For example, when is the Taylor series approximation of *ELD* by Total Variation Distance sufficient close that theoretical guarantees for TVD hold also for *ELD*

In sum, outlier metrics based on model likelihoods are a new type of structured outlier score for object-relational data. Our evaluation indicates that model-based scores provide informative, interpretable, and accurate rankings of objects as potential outliers.

Acknowledgements

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12 Appendix: Proofs

Assume that the generator f is twice differentiable for f -divergence I_f . Then $I_f(P_1||P_2) \approx \frac{f''(1)}{2} \chi^2(P_1||P_2) = \frac{f''(1)}{2} \sum_{i=1}^m \frac{(P_2(v_i) - P_1(v_i))^2}{P_1(v_i)}$.

Proof Truncating the Taylor series for f at $i = 2$, the f -divergence expression 9 becomes

$$\begin{aligned} & \sum_{i=1}^m P_1(v_i) \left[f'(1) \left(\frac{P_2(v_i)}{P_1(v_i)} - 1 \right) + \frac{f''(1)}{2} \left(\frac{P_2(v_i)}{P_1(v_i)} - 1 \right)^2 \right] \\ &= \sum_{i=1}^m P_1(v_i) \left[f'(1) \left(\frac{P_2(v_i) - P_1(v_i)}{P_1(v_i)} \right) + \frac{f''(1)}{2} \left(\frac{P_2(v_i) - P_1(v_i)}{P_1(v_i)} \right)^2 \right] \\ &= f'(1) \left[\sum_{i=1}^m P_2(v_i) - \sum_{i=1}^m P_1(v_i) \right] + \sum_{i=1}^m \frac{f''(1)}{2} \frac{(P_2(v_i) - P_1(v_i))^2}{P_1(v_i)} \\ &= 0 + \frac{f''(1)}{2} \chi^2(P_1||P_2) \end{aligned}$$

The first-order Taylor series approximation for *ELD* is total variation distance:

$$ELD(P_I||P_{\mathcal{D}}) \approx TVD.$$

Proof Truncating the Taylor series for $f = -\ln(u)$ at $i = 1$, the $|f|$ -divergence expression 6 becomes

$$\begin{aligned}
& \sum_{i: f(\frac{P_2(v_i)}{P_1(v_i)}) > 0} P_1(v_i) f'(1) \left(\frac{P_2(v_i)}{P_1(v_i)} - 1 \right) - \sum_{i: f(\frac{P_2(v_i)}{P_1(v_i)}) < 0} P_1(v_i) f'(1) \left(\frac{P_2(v_i)}{P_1(v_i)} - 1 \right) \\
&= \sum_{i: f(\frac{P_2(v_i)}{P_1(v_i)}) > 0} P_1(v_i) f'(1) \left(\frac{P_2(v_i) - P_1(v_i)}{P_1(v_i)} \right) - \sum_{i: f(\frac{P_2(v_i)}{P_1(v_i)}) < 0} P_1(v_i) f'(1) \left(\frac{P_2(v_i) - P_1(v_i)}{P_1(v_i)} \right) \\
&= - \sum_{i: P_2(v_i) < P_1(v_i)} P_2(v_i) - P_1(v_i) + \sum_{i: P_2(v_i) > P_1(v_i)} P_2(v_i) - P_1(v_i) \quad (10) \\
&= \sum_i |P_2(v_i) - P_1(v_i)|
\end{aligned}$$

Equation (10) holds for $f = -\ln$ because then $f(\frac{P_2(v_i)}{P_1(v_i)}) > 0$ if and only if $P_2(v_i) < P_1(v_i)$ and also $f'(1) = -1$.

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