Traffic state prediction using Markov chain models

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Abstract—Motorway traffic management and control relies on models that estimate and predict traffic conditions. In this paper, a methodology for the identification and short-term prediction of the traffic state is presented. The methodology combines model-based clustering, variable-length Markov chains and nearest neighbor classification. An application of the methodology for short-term speed prediction in a freeway network in Irvine, CA, shows encouraging results.

I. INTRODUCTION

Simulation of traffic dynamics is a fundamental component of dynamic traffic management and control. Mesoscopic and macroscopic simulation models usually represent traffic dynamics using speed-density relationships, fluid representation of traffic flow, elements of queuing theory, etc. Emerging DTA systems, used in real-time applications for traffic estimation and prediction, are simulation-based and use mesoscopic simulation models to capture traffic dynamics (DynaMIT [1], DYNASMART [2], and RENAISSANCE [3]).

Traffic is volatile and can change states quickly, thus making modeling a challenging task. State identification would allow state-specific modeling of traffic, which could in turn result in more accurate simulation models, as statespecific models could then be employed, presumably with superior performance. For example, [4] present an application of multimodal regression to speed-flow data, while [5] use cluster analysis to segment speed-density data and determine the regime boundaries for classic (tworegime and three-regime) speed-density models. [6] combined clustering and locally weighted regression to develop multi-state alternative models to the typical speed-

Manuscript received December 5, 2006. The research was partially supported by the EU Marie Curie project MIRG-CT-2005-036590 and NSF projects CMS-0339005 and CMS-0339108 and. The authors would like to thank Prof. Wilfred Recker from University of California, Irvine, for providing the data used in this research.

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density relationship, while [7] compared machine learning approaches for more flexible modeling of traffic dynamics models.

Key methodological elements used in this research are presented in the next section, in particular model-based clustering, variable-length Markov chains, and nearest neighbor classification. An application of this framework using data from a freeway network in Irvine, CA, follows. The performance and potential benefits of the presented framework in short-term speed prediction is demonstrated. The paper concludes with directions for further research.

II. METHODOLOGY

A. Model-based clustering

Clustering and classification are tasks that are rather well researched as they have extensive applications in many practical and research fields. As a result, a range of approaches and algorithms are available, often based on heuristics. One rigorous approach to cluster analysis is based on probability modes (see [8-9] for a survey). Some of the most popular heuristics used for clustering are approximate estimation methods for appropriately defined probability models [10]. For example, standard k-means clustering [11] is equivalent to known procedures for approximately maximizing the multivariate normal classification likelihood when the covariance matrix is the same for each component and proportional to the identity matrix [10].

Finite mixture models have been proposed and studied in the context of clustering [12-17], often as a statistical approach to shed some light into practical questions that arise from the application of clustering methods [18-21]. Each component probability distribution in finite mixture models corresponds to a cluster. The problems of determining the number of clusters and of choosing an appropriate clustering method can be recast as statistical model choice problems, and models that differ in numbers of components and/or in component distributions can be compared. Outliers are handled by adding one or more components representing a different distribution for outlying data.

1) Mixture models

Given data y with independent multivariate observations

 y_1, \ldots, y_n the likelihood for a mixture model with G components is:

$$L_{MIX}(\theta_1, \cdots, \theta_G; \tau_1, \cdots, \tau_G \mid \mathbf{y}) = \prod_{i=1}^n \sum_{k=1}^G \tau_k f_k(\mathbf{y}_i \mid \theta_k)$$
(1)

where f_k and θ_k are the density and parameters of the *k*th component in the mixture and τ_k is the probability that an observation belongs to the *k*th component.

Data generated by mixtures of multivariate normal densities are characterized by groups or clusters centered at the means, with ellipsoidal surfaces of constant density. The geometric features (shape, volume, orientation) of the clusters are determined by the covariances Σ_k , which may be further parameterized to impose cross-cluster constraints. In the simplest case of spherical clusters of the same size $\Sigma_k = \lambda I$, while in the case of clusters with the same geometry (but not necessarily spherical) $\Sigma_k = \Sigma$ [22]. Only one parameter is needed to capture the covariance structure of the mixture when $\Sigma_k = \lambda I$, while for d-dimensional data d(d+1)/2 and G(d(d+1)/2)parameters are needed for constant Σ_k and unrestricted Σ_k [15]. [19,23] proposed more flexible and general frameworks for geometric cross-cluster constraints in multivariate normal mixtures by parameterizing covariance matrices through eigenvalue decomposition.

2) Cluster analysis

The purpose of cluster analysis is to classify data of previously unknown structure into meaningful groupings. A strategy for cluster analysis based on mixture models is outlined next [10]. The strategy comprises three core elements: (i) initialization via model-based hierarchical agglomerative clustering, (ii) maximum likelihood estimation via the expectation-maximization (EM) algorithm, and (iii) selection of the model and the number of clusters using approximate Bayes factors with the BIC (Bayesian Information Criterion) [24] approximation.

Model-based hierarchical agglomerative clustering is an approach to computing an approximate maximum for the classification likelihood:

$$L_{CL}(\boldsymbol{\theta}_{1},\cdots,\boldsymbol{\theta}_{G};\boldsymbol{\ell}_{1},\cdots,\boldsymbol{\ell}_{n}\mid\mathbf{y}) = \prod_{i=1}^{n} f_{\boldsymbol{\ell}_{i}}(\mathbf{y}_{i}\mid\boldsymbol{\theta}_{\boldsymbol{\ell}_{i}})$$
(2)

where ℓ_i are labels indicating a unique classification of each observation, taking the value *k* if \mathbf{y}_i belongs to the kth component. In the mixture likelihood (eq. 1), each component is weighted by the probability that an observation belongs to that component. The presence of the class labels in the classification likelihood (eq. 2)

introduces a combinatorial aspect that makes exact maximization impractical [10].

[23] successfully applied model-based agglomerative hierarchical clustering to problems in character recognition using a multivariate normal model, with volume and shape held constant across clusters. This approach was generalized by [19] to other models and applications.

In hierarchical agglomeration, each stage of merging corresponds to a unique number of clusters and a unique partition of the data. A given partition can be transformed into indicator variables, which can then be used as conditional probabilities in an M step of EM for parameter estimation, initializing an EM algorithm. This, combined with Bayes factors as approximated by BIC for model selection, yields a comprehensive clustering strategy:

- Determine a maximum number of clusters, M, and a set of mixture models to consider.
- Perform hierarchical agglomeration to approximately maximize the classification likelihood for each model and obtain the corresponding classifications for up to M groups.
- Apply the EM algorithm for each model and each number of clusters 2, ..., M, starting with the classification from hierarchical agglomeration.
- Compute BIC for the one-cluster case for each model and for the mixture model with the optimal parameters from EM for 2, ..., M clusters.

B. Variable-length Markov chains

One of the most general models for a stationary categorical process taking values in a finite categorical space X, is a full Markov chain (of possibly high, but finite order). [25]. The only implicit assumption aside from stationarity is the finite memory of the process. A stationary full Markov chain of order p exists whenever the transition mechanism has no specific structure; that is the state space is the entire X_p. While such general models may be theoretically attractive, they also have practical limitations. For example, a full Markov chain is rather inflexible in terms of the number of parameters that it can represent. For a model with 4 states, chains with 0 to 5 parameters have dimensions of 3, 12, 48, 192 and 768, respectively. Markov chains can only be fitted in these "intervals", thus reducing the model flexibility (e.g. if 48 parameters are not enough, one needs to estimate 192 parameters; intermediate values are not possible.) This issue introduces another problem with the full Markov chain model, the "dimensionality curse", as the dimension of the model increases exponentially with the order p.

Markov processes have found many applications in a diverse number of fields. For example, [26] propose an analytical methodology for prediction of the platoon arrival profiles and queue length along signalized arterials using Markov decision processes (an extension of Markov chains). Other applications of Markov processes in transport-related literature include indicatively pavement management [27] and bridge maintenance management [28-29]. [30] model mobile terminals communication with their base station using hidden Markov models in combination with clustering algorithms.

Variable length Markov chains (vlmc) address both issues introduced above and provide a natural and elegant way to avoid (some of) the difficulties mentioned. The idea is to allow the memory of the Markov chain to have a variable length, depending on the observed past values [31]. Fitting a vlmc from data involves estimation of the structure of the variable length memory, which can be reformulated as a problem of estimating a tree, using the so-called context algorithm [32], which can be implemented very efficiently.

A variable length Markov chain is a potentially high order Markov chain, taking values in a finite categorical space, with a natural parsimonious structure for the transition probabilities. Let $(X_t)_{t\in Z}$ be a stationary process with values $X_t \in X$. A function called *preliminary context function* encapsulates the information that reflects which vectors from the infinite past of the process are relevant. If a number p exists, such that the cardinality of the preliminary context function is not infinite for the entire domain, then $(X_t)_{t\in Z}$ is called a stationary variable length Markov chain of order p. The context function $c(\cdot)$ can be reconstructed from the context tree τ_c which is nothing else than the minimal state space of the underlying variablelength Markov chain.

Fitting a variable-length Markov chain can be done with a version of the tree structured context algorithm [32]. The variable length memory is usually represented with an estimated context tree. Thus, tree-structured models were fitted, where every terminal node (as well as some internal nodes) represents a state in the Markov chain and is equipped with corresponding transition probabilities. The context algorithm grows a large tree and prunes it back subsequently. The pruning part requires specification of a tuning parameter, the so-called cutoff. The cutoff K is a threshold value when comparing a tree with its subtree by pruning away one terminal node; the comparison is made with respect to the difference of deviance from the two trees. A large cutoff has a stronger tendency for pruning and yields smaller estimated context trees, i.e. a smaller dimension of the model.

C. Nearest neighbors classification

Clustering methods are usually accompanied by a classification algorithm so that they can be applied.

Nearest neighbor classification is one of the standard classification methods. During the application phase, standard methods, such as the k-nearest neighbor approach, can be used to classify new traffic measurements (characterized by e.g. flow and density) to the most appropriate cluster. k-nearest neighborhood learning is the most basic instance-based method, and assumes that all instances (or observations) correspond to points in the n-dimensional space [11]. The nearest neighbors of an instance are defined in terms of the standard Euclidean distance.

Let an observation x be described by the feature tuple $\langle a_1(x), a_2(x),..., a_n(x) \rangle$ where $a_r(x)$ denotes the values of the r^{th} attribute of x. In the context of traffic dynamics, the attributes of x could be density and flow, but also other parameters, such as time of day, prevailing weather conditions, and traffic mix. The distance between two instances x_i and x_j is then defined to be:

$$d(x_{i}, x_{j}) = \sqrt{\sum_{r=1}^{n} \left(a_{r}(x_{i}) - a_{r}(x_{j})\right)^{2}}$$
(3)

In nearest-neighbor learning the target function may be either discrete-valued or real-valued. In the discrete case, such as this one, where the goal is to assign a cluster to each new instance x_q , the algorithm selected the k instances from the training set that are nearest to x_q (as defined by the distance above), and returns

$$\hat{f}(x_q) \leftarrow \operatorname*{argmax}_{u \in V} \sum_{i=1}^{k} \delta(u, f(x_i))$$
(4)

where δ is the Kronecker operator

$$\delta(a,b) = \begin{cases} 1 & \text{if } a = b \\ 0 & \text{otherwise} \end{cases}$$
(5)

III. APPLICATION

A. Data

The Irvine data set includes five days of sensor data from freeway I-405. Four days of data were used for the model based clustering, Markov chain estimation, and k-nearest neighbor training, while the data from the fifth day was used for the validation of the approach. Morning period (04:00am to 10:00am) data have been used. Speed, occupancy and flow data over 2-minute intervals were available for calibration and validation. (Local) density was obtained from the occupancy measurements.

One of the few explicit assumptions of Markov chain modeling is the stationarity of data, which is not in general true for traffic data. Fig. 1 demonstrates the nonstationarity of the data used in this research by showing the autocorrelation functions for up to a lag of 300 intervals. The long series of slowly decaying autocorrelation fractions is a typical evidence of non-stationarity.



One way to transform non-stationary data to stationary data is to difference them (once or more). Fig. 2 shows the autocorrelation functions for the (one-step) differenced data. Note that this time autocorrelation fractions for up to 50 lags is shown. Simply differencing the data once results in stationary data, that can be modeled using Markov processes.



Fig. 2. Autocorrelation functions of differenced data

B. Goodness-of-fit statistics

The following appropriate statistics have been used [33]:

- Normalized root mean square error (RMSN) [34-35]
- Root mean square percent error (RMSPE) [36]
- Mean percent error (MPE) [36]
- Theil's U coefficient, as well as its bias, variance and covariance components [37]

The purpose of using multiple statistics is that they can capture different aspects of the obtained results. The normalized root mean square error (RMSN) and root mean square percent error (RMSPE) quantify the overall error of the simulator. These measures penalize large errors at a higher rate than small errors. The formula for calculating the RMSN value is:

$$RMSN = \frac{\sqrt{N\sum_{n=1}^{N} \left(Y_{n}^{s} - Y_{n}^{o}\right)^{2}}}{\sum_{n=1}^{N} Y_{n}^{o}}$$
(6)

where N is the number of observations, Y_n^{o} is an observation and Y_n^{s} is an estimated/predicted value at time *n*.

RMSPE is calculated based on the following formula:

$$RMSPE = \sqrt{\frac{1}{N} \sum_{n=1}^{N} \left[\frac{Y_n^s - Y_n^o}{Y_n^o} \right]^2}$$
(7)

The mean percent error (MPE) statistic indicates the existence of systematic under- or over-prediction in the estimated measurements and is calculated by:

$$MPE = \frac{1}{N} \sum_{n=1}^{N} \left[\frac{Y_n^s - Y_n^o}{Y_n^o} \right]$$
(8)

Percent error measures are often preferred to their absolute error counterparts because they provide information on the magnitude of the errors relative to the average measurement. Another measure that provides information on the relative error is Theil's inequality coefficient, given by:

$$U = \frac{\sqrt{\frac{1}{N}\sum_{n=1}^{N} (Y_n^s - Y_n^o)^2}}{\sqrt{\frac{1}{N}\sum_{n=1}^{N} (Y_n^s)^2} + \sqrt{\frac{1}{N}\sum_{n=1}^{N} (Y_n^o)^2}}$$
(9)

U is bounded and takes values between zero and one $(0 \le U \le 1$, where U=0 implies perfect fit between observed and simulated measurements). Theil's inequality coefficient may be decomposed into three proportions of inequality: the bias (U^M) , the variance (U^S) and the

covariance (U^C) proportions given, respectively by:

$$U^{M} = \frac{\left(\overline{Y}^{s} - \overline{Y}^{o}\right)^{2}}{\frac{1}{N} \sum_{n=1}^{N} \left(Y_{n}^{s} - Y_{n}^{o}\right)^{2}}$$
(10)

$$U^{S} = \frac{(s^{o} - s^{o})}{\frac{1}{N} \sum_{n=1}^{N} (Y_{n}^{s} - Y_{n}^{o})^{2}}$$

$$U^{C} = \frac{2(1 - \rho)s^{s}s^{o}}{\frac{1}{N} \sum_{n=1}^{N} (Y_{n}^{s} - Y_{n}^{o})^{2}}$$
(11)
(12)

where ρ is the correlation between the two sets of measurements, s^s and s^o are the standard deviations of the average simulated and observed measurements, respectively. By definition, the three proportions sum to 1 $(U^{M}+U^{S}+U^{C}=1)$. The bias proportion reflects the systematic error. The variance proportion indicates how well the simulation model is able to replicate the variability in the observed data. These two proportions should be kept as close to zero as possible. The covariance proportion measures the remaining error and therefore should be close to 1. Note that since the various measurements are taken from non-stationary processes, the proportions can only be viewed as rough indicators to the sources of error.

The application of this research was performed within the R Software for Statistical Computing v.2.4.0 [38] using the Mclust package [10] for model-based clustering and the vlmc package [31] for estimation of variable-length Markov chains.

C. Clustering and classification

The best functional form of the mixtures to be considered for clustering, and the optimal number of clusters were sought using the model-based clustering algorithm [10] on the stationary/differenced data. The following different mixture models were considered (in increasing order of flexibility and complexity):

- "EII": spherical, equal volume
- "VII": spherical, unequal volume
- "EEI": diagonal, equal volume and shape
- "VEI": diagonal, varying volume, equal shape
- "EVI": diagonal, equal volume, varying shape
- "VVI": diagonal, varying volume and shape
- "EEE": ellipsoidal, equal volume, shape, and orientation
- "EEV": ellipsoidal, equal volume and equal shape
- "VEV": ellipsoidal, equal shape

• "VVV": ellipsoidal, varying volume, shape, and orientation

Fig. 3 shows the obtained curves for the different functional forms. The best one -indicated by the "top" line, i.e. the largest BIC, in the figure- is obtained by the most general model (VVV) with 5 clusters.



However, the marginal benefit of 5 clusters over 3 clusters is minimal, and it is considered to use the more parsimonious model with only 3 clusters. The classification obtained with the 5 clusters is compared with the classification obtained by 3 clusters in Table 1. 98% of the observations are clustered "correctly" from the five-mixture model to the three-mixture model. The interpretation of this mapping is that clusters 1 and 5 of the 5-cluster model are mapped onto cluster A of the parsimonious 3-cluster model, clusters 2 and 3 are mapped onto cluster B of the 3-cluster model, and cluster 4 is mapped onto cluster C. The parsimonious 3-cluster model is retained for the remainder of this application.

			TABLE	E 1					
CORRESPONDENCE OF CLUSTERS IN 3- AND 5-MIXTURE MODELS									
			3 clusters						
			Α	В	С				
	ers	1	1641	62	0				
		2	3	1217	0				
	nst	3	0	440	0				
	2 cl	4	7	0	172				
	47	5	57	0	0				

D. Markov chain training

Variable length Markov Chain computes a huge tree and then prunes it. The cutoff for this pruning is an input parameter, which –in this case- was obtained using a linesearch for the value that provided the highest value for the Akaike Information Criterion (AIC) [39]. It appears from Fig. 4 that the optimal value for the cutoff parameter is 3.6, which is actually picked as the cutoff for the remainder of this application. The minimum AIC obtained with this cutoff value is equal to 4699.09.

E. Speed prediction

2

3

4

5

Fig. 4. AIC vs cutoff for VLMC

6

Cutoff

7

9

10

8

5600

4800

AIC 5200

Traffic state identification can have many applications in the field of traffic management and control. In this section, it is indicatively applied to speed prediction. Locally weighted regression (loess) [11] is trained on the training data.

Given the estimated variable-length Markov process, for each observation, the Markov chain model is used to predict the state of the traffic for the next interval. After each "new" instance is classified to a cluster, a speed value is estimated using the appropriate locally weighted regression (which was trained with data only from the relevant cluster.)

Table 2 presents the results of the speed prediction using the two approaches, using several measures of goodnessof-fit. Percent improvement due to the more elaborate technique is also presented. A clear improvement is indicated by all statistics. The percent improvement ranges between 5 and 10 percent for RMSN, RMSPE, and U, while it exceeds 60% for MPE and U^M.

		TABLE 2				
COMPARISON OF GOODNESS-OF-FIT STATISTICS						
	Estimation	Reference prediction	Markov- based prediction	Prediction improve- ment		
RMSN	0.0449	0.0883	0.0801	9.3%		
RMSPE	0.0504	0.1049	0.0989	5.7%		
MPE	0.0044	0.0085	0.0033	60.7%		
U	0.0221	0.0434	0.0396	8.8%		
$\mathbf{U}^{\mathbf{M}}$	0.0177	0.0046	0.0017	63.6%		

In recent research, [6-7] found that loess outperforms other machine learning approaches and the typical speeddensity relationship for speed estimation. Therefore, this performance should be considered as an improvement over an already powerful reference case.

F. Model diagnostics

Fig. 5 shows the residuals of the fitted variable-length Markov Chain model against the contexts, i.e., produces a boxplot of residuals for all contexts used in the model fit. Intuitively, a context is a "case" that is not pruned from the context tree. The number of observations per context state is also illustrated, along the x-axis of the figure. Furthermore, the width of each boxplot is proportional to the square root of the number of observations that it represents. For a few contexts (such as 2 and 2222) the residuals are zero (indicated by lack of a boxplot and a small square along the x-axis). This implies that this past state seems fairly well predictable, based on this dataset.

IV. CONCLUSION

A methodology for the identification and short-term prediction of traffic state has been presented. The methodology comprises state-of-the-art components, such as model-based clustering, variable length Markov chains and nearest neighbor classification. An application of the methodology to short-term speed prediction in a freeway network in Irvine, CA, provides encouraging results.

Traffic state identification and prediction has many possible applications in the field of motorway traffic surveillance and control, including automated incident detection and capacity estimation. Incident detection can be achieved when the observed traffic state differs from the state that was expected, based on short-term predictions. A deviation from this expectation may suggest that some special event has disrupted the normal flow of traffic. In terms of capacity, there are no direct ways to measure or observe it. Instead, modelers and operators rely on indirect evidence on capacity. Information on the expected vs. observed traffic state may provide some further evidence.

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Residuals vs. Context



Fig. 5. Residuals vs. Context plot for vlmc

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