A COUPLED ETAS-I\textsuperscript{2}GMM POINT PROCESS WITH APPLICATIONS TO SEISMIC FAULT DETECTION

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Abstract Epidemic-type aftershock sequence (ETAS) point processes are a common model for the occurrence of earthquake events. ETAS models consist of a stationary background Poisson process modeling spontaneous earthquakes and a triggering kernel representing the space-time-magnitude distribution of aftershocks. Two popular non-parametric methods for estimation of the background intensity include histograms and kernel density estimators. While these methods are able to capture local spatial heterogeneity in the intensity of spontaneous events, they do not capture patterns resulting from fault-line structure over larger spatial scales. Here we propose a two-layer infinite Gaussian mixture model for clustering of earthquake events into fault-like groups over intermediate spatial scales. We introduce a Monte-Carlo expectation-maximization (EM) algorithm for joint inference of the ETAS-I\textsuperscript{2}GMM model and then apply the model to the Southern California earthquake Catalog. We illustrate the advantages of the ETAS-I\textsuperscript{2}GMM model in terms of both goodness of fit of the intensity and recovery of fault line clusters in the Community Fault Model 3.0 from earthquake occurrence data.

1. Introduction.

1.1. Background on point-process models of seismicity. The Epidemic-Type Aftershock Sequence (ETAS) model of earthquake occurrence [Oga88a, Oga98] is a self-exciting point-process model where the conditional intensity \( \lambda(t, x, y|H_t) \) of events is determined by a stationary Poisson intensity generating spontaneous earthquake events along with a dynamic term representing a branching process of aftershocks:

\[
\lambda(t, x, y|H_t) = \mu(x, y) + \sum_{i:t_i < t} g(t - t_i, x - x_i, y - y_i, m_i).
\]

Here \((x, y)\) is the epicenter of an earthquake event described by longitude and latitude in decimal degrees of the WGS 84 coordinate system, \(m\) is its magnitude on the Richter scale computed using a body-wave magnitude
formula \([SSC89]\), \(H_t = \{(t_i, x_i, y_i, m_i) : t_i < t\}\) is the history of all earthquake events up to time \(t\) in a catalog, and \(\mu(x, y)\) is the background intensity reflecting spatial heterogeneity of spontaneous earthquakes and the fact that earthquake catalogs with aftershocks removed are approximately Poisson in time \([GK74]\).

The space-time-magnitude distribution of parent-offspring events in the branching process given by the function \(g(t, x, y, m)\) is called the triggering kernel, typically following Omori’s law \([Uts61]\),

\[
g(t - t_i, x - x_i, y - y_i, m_i) = \frac{K_0 e^{a(m_i - m_0)}}{(t - t_i + c)(1 + \omega)(1 + w)(1 + \rho)(1 + \omega)(1 + w)(1 + \rho)}
\]

where \(m_0\) is the cutoff magnitude of the considering dataset following \([Oga88b]\), \(K_0, a, c, \omega, d, \rho > 0\) are the parameters under estimation. Estimation of Equation 1 typically consists of constructing a non-parametric estimate for \(\mu(x, y)\) along with finding maximum likelihood estimators for the parameters of the triggering kernel in Equation 2. Methods for maximizing the likelihood include quasi-Newton \([Oga88a]\) and expectation-maximization (EM) \([VS08]\), and the most common estimators for \(\mu(x, y)\) are spatial histograms \([ML08, VS08]\) or isotropic kernel density estimators \([ZOJ02, AC15]\).

1.2. A New Model: Coupled ETAS-\(I^2\)GMM. Earthquakes cluster at multiple scales, as earthquakes cluster locally through aftershock activity but also over larger scales along fault lines (see Figure 1). While there is research on the reconstruction of aftershock clusters from event data \([ZOJ02, ZGKBW08]\), existing point-process models of earthquake activity fail to capture clustering patterns at the larger scale of fault lines. In particular, histograms and kernel density estimators are able to capture spatial heterogeneity in the risk of spontaneous earthquakes, but the methods capture variation over only one scale. To our knowledge, our work here is the first to attempt to reconstruct the community fault model \([PSB07]\) with a statistical model based on earthquake event data.

In this paper we introduce a new type of ETAS model that can capture multiscale clustering in earthquake patterns. In particular, we propose using an infinite mixture of infinite Gaussian mixtures (\(I^2\)GMM) \([YRD14]\) to estimate the background rate of earthquakes \(\mu(x, y)\). The \(I^2\)GMM uses a different Dirichlet process mixture of Gaussians (DPMG) for each cluster that simultaneously predicts the number of clusters along with performing model inference. While \(I^2\)GMM has been introduced for high-dimensional clustering and ETAS is well known in seismology, what is brand new in this paper is the use of \(I^2\)GMM for modeling the intensity of a point process and
the coupling of these two techniques for multiscale modeling of space-time event patterns. Through the use of an expectation-maximization algorithm, the benefit of our approach is that earthquakes are assigned membership to aftershock clusters in addition to a larger-scale fault-line cluster.

Another advantage of our approach is that multi-modal and skewed clusters are more accurately captured. In the case of spatial earthquake patterns, each fault may be considered as a separate cluster with multi-modality and skewness that the $I^2$GMM can handle better than histograms and KDE estimators. One additional advantage of the $I^2$GMM model is that earthquakes are assigned membership to clusters inferred by the model. In this research, we explore the relevance of cluster membership to automatic detection of fault lines within the ETAS-$I^2$GMM framework.

1.3. Outline of the paper. In Section 2 we describe our methodology, including an overview of the $I^2$GMM model and details on a Monte-Carlo EM algorithm for joint inference of the ETAS-$I^2$GMM model. In Section 3 we present results for several experiments where the ETAS-$I^2$GMM is applied to a Southern California earthquake catalog [sce]. We compared the goodness of fit of the estimated intensity of the model to a baseline approach. In Section 3, we also use the Community Fault Model 3.0 [PSB+07] to explore the ETAS-$I^2$GMM model’s ability to detect fault locations and event-fault linkage from space-time event data.

2.1. **Infinite mixture of infinite Gaussian mixtures.** The finite Gaussian mixture model (GMM) uses a single Gaussian for each cluster and requires the number of clusters to be specified. In the infinite version of GMM (IGMM) [Fer73], the number of components is estimated along with the component mean vectors and covariances. Both GMM and IGMM are used for clustering problems, albeit with limited success, as these techniques often overestimate the number of mixture components so as to more accurately estimate the density of the underlying dataset. However, a more accurate estimation of the density does not necessarily translate into a more accurate estimation of cluster distributions, as density estimation does not readily solve the problem of many-to-one mappings between components and clusters. Two different approaches are considered in the literature to overcome this limitation.

The first approach replaces Gaussian mixture components by student-t [PM00, SB05, AV07, AM12] or skewed-t [LM14] distributions and their Pearson-type extensions [FW14, SKG10] in an effort to better model cluster distributions with heavy tails. Although closed-form solutions for maximum-likelihood estimations of parameters do not in general exist under these settings, extensions of the EM algorithm can still be derived for this family of mixture models by placing certain restrictions on the original model. This line of models has proved quite effective in clustering datasets with skewed distributions but are less ideal for clusters with multi-mode distributions.

The second approach generates a large number of Gaussian components and merges them according to various metrics in an effort to recover true cluster distributions. The study in [FJ02] initializes the model with a large number of components and uses the concept of minimum message length to merge components. Another technique uses the Bayesian information criterion (BIC) to choose the initial number of components and merges components to minimize entropy [BRC+10]. Other options for assigning components to clusters include clustering modes of components [GS12] and ridgeline analysis and Bhattacharyya dissimilarity [Hen10]. Compared to mixtures of student-t or skewed-t distributions, this line of models is more flexible in terms of the type of distribution they can model. However, the main limitation of these techniques is the independence assumption made during component estimation that makes EM derivations possible. These techniques assume that all components are generated independently, which is not a very realistic assumption in a setting where some clusters are known to be multi-mode, and components originating from the same cluster are more likely to share certain latent parameters than two random components. Another limitation of these techniques is the computational complexity that increases with the square of the number of Gaussian components, as the
decision to merge two components requires evaluating the metric for every possible pair of components. This puts a constraint on the maximum number of components that can be used to model datasets.

When clustering datasets with skewed/multi-mode cluster distributions, two dependent subproblems, namely density estimation and component clustering, need to be addressed jointly. The two-layer non-parametric GMM (I²GMM) model, which can grow arbitrarily large in the number of components and clusters generated, was introduced earlier to more accurately cluster datasets with multi-mode and skewed cluster distributions [YRD14].

In I²GMM the lower layer estimates the density of the overall dataset by clustering individual data points to components, while the upper layer associates components with clusters to allow for cluster recovery. More specifically, the generative model is a two-layer hierarchical Dirichlet process mixture (DPM) model where the lower layer uses one DPM for each cluster and the upper layer uses a global DPM for modeling cluster shapes and sizes. The dependency between the two layers is achieved by centering the base distributions of DPMs in the lower layer on a unique parameter distributed according to the global DPM. Inference, which involves sampling component indicator variables for individual data points and sampling cluster indicator variables for components, is performed by a collapsed Gibbs sampler, enabling optimization of two subproblems simultaneously.

We believe that I²GMM has three unique features that would make it very suitable for the estimation of background intensity $\mu(x, y)$ in the ETAS model.

- As a two-layer non-parametric model, I²GMM allows the number of clusters and the number of mixture components in each cluster to grow arbitrarily large, offering great flexibility in modeling clusters with multi-mode/skewed distributions. This is the main feature that distinguishes I²GMM from other model-based clustering techniques that use one component for each cluster.
- As a Bayesian model, I²GMM has hyper-parameters that can be tuned to recover clusters with varying shapes and different levels of rarity without facing singularities during model estimation. This distinguishes I²GMM from purely data-driven techniques such as finite mixture of Gaussians and t-distributions that rely on EM and its extensions during model learning.
- As a hierarchical model, I²GMM can share parameters not only across different clusters but also across different components of the same cluster. In other words, I²GMM assumes that components are generated independently only when conditioned on the unique parameter defining
their clusters of origin. This differentiates the proposed work from other techniques that estimate a large number of Gaussian components and merges them sequentially to recover clusters, thus violating component dependence.

In Figure 2 we provide an illustration of the generative model for I²GMM: where \( t_{kl} \) indicates the \( l^{th} \) component in the \( k^{th} \) cluster \( C_k \); \( x_{kl} \) indicates the \( i^{th} \) data point in the \( l^{th} \) component in the \( k^{th} \) cluster. In the generative process, \( t_{kl} \) is a Gaussian distribution and \( C_k \) is a Gaussian mixture defined by its components. We will use the top-level label (i.e., \( C_k \)) to identify different clusters of spontaneous earthquakes.
The generative model for I²GMM is given by

$$H = NIW(\mu, \Sigma | \mu_0, \Psi_0, \kappa_0, m) = N(\mu | \mu_0, \Sigma \kappa_0^{-1})W^{-1}(\Sigma | \Psi_0, m)$$

$$G \sim DP(\gamma H)$$

$$(\mu_k, \Sigma_k) = \theta_k \sim G$$

1. $$H_k = N(\mu_k, \kappa_1^{-1}\Sigma_k)$$
2. $$G_k = DP(\alpha H_k)$$
3. $$\mu_{kl} \sim G_k$$
4. $$x_{kl} \sim N(\mu_{kl}, \Sigma_k)$$

In this generative model $DP(\gamma H)$ is a global Dirichlet process with normal-inverse-Wishart base distribution $H$ and a concentration parameter $\gamma$. $G$ is a discrete mixing measure sampled from the global DP. Center $\mu_k$ and covariance $\Sigma_k$ of the cluster $k$ are drawn from $G$. For each cluster generated by the global DPM, a local DPM is defined with base distribution $H_k$ and concentration parameter $\alpha$. All $H_k$ are Gaussian distributions with centers $\mu_k$ and covariances $\kappa_1^{-1}\Sigma_k$. $G_k$ is the cluster-specific discrete mixing measure drawn from the local DP. The components in cluster $k$ are generated with mean vectors $\mu_{kl}$ drawn from $G_k$. Data points $x_{kl}$ are generated from the Gaussian components with mean vectors $\mu_{kl}$ and covariance matrices $\Sigma_k$.

User-specified hyper-parameters ($\mu_0, \Psi_0, \kappa_0, \kappa_1, \alpha, \gamma, m$) are listed in Table 1 along with their descriptions.

To perform inference with the I²GMM model using spatial event data, we first initialize the cluster and component indicators for each event to some arbitrary values (for example put all data in the same component of a cluster) and then use a collapsed Gibbs sampler to infer values for indicator variables one at a time, given all other indicator variables [YRD14]. Conditioned on the indicator variables, the location vectors and scale matrices are determined by maximizing the complete data log-likelihood and have closed form solutions. One sweep of the Gibbs sampler will go over all events in the dataset; convergence typically requires several hundred to thousand Gibbs sweeps.

2.2. EM inference for ETAS. The ETAS model given in Equation 1 can be viewed as a branching process where first-generation events occur according to a Poisson process with intensity $\mu(x, y)$. Events (from all generations) each give birth to direct offspring events determined by the triggering kernel $g(t - t_i, x - x_i, y - y_i, m_i)$.

Given an initial guess for the parameters of the triggering kernel in Equation 2 and the background rate $\mu(x, y)$, the branching structure along with
Expected mean vector for each cluster. This is usually set to the mean of the overall dataset.

$\Psi_0$ is the expected covariance matrix for clusters. This is usually set to identity.

$\kappa_0$ A positive scaling constant that adjusts the separation among clusters. The smaller the $\kappa_0$, the more separated the clusters will be from each other.

$\kappa_1$ A positive scaling constant that adjusts the separation among components of a given cluster. The smaller the $\kappa_1$, the more separated the components will be from each other, and clusters will tend to emerge with multi-modal distributions.

$\alpha$ The concentration parameter for the local DPMs that controls the expected number of components and their sizes within a cluster.

$\gamma$ The concentration parameters for the global DPM that controls the expected number of clusters and their sizes.

Degree of freedom for the inverse Wishart that controls the degree of deviation of actual component covariances from the expected covariance. The higher the $m$, the less the deviation and the more similar component shapes will be.

<table>
<thead>
<tr>
<th>Hyper-parameters for $\hat{P}_{\text{GMM}}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\mu_0$ Expected mean vector for each cluster. This is usually set to the mean of the overall dataset.</td>
</tr>
<tr>
<td>$\Psi_0$ is the expected covariance matrix for clusters. This is usually set to identity.</td>
</tr>
<tr>
<td>$\kappa_0$ A positive scaling constant that adjusts the separation among clusters. The smaller the $\kappa_0$, the more separated the clusters will be from each other.</td>
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<td>$\alpha$ The concentration parameter for the local DPMs that controls the expected number of components and their sizes within a cluster.</td>
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<td>$\gamma$ The concentration parameters for the global DPM that controls the expected number of clusters and their sizes.</td>
</tr>
<tr>
<td>$m$ Degree of freedom for the inverse Wishart that controls the degree of deviation of actual component covariances from the expected covariance. The higher the $m$, the less the deviation and the more similar component shapes will be.</td>
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</table>

The model parameters of the triggering kernel can be estimated using an EM algorithm [VS08, MSB+11]. In the E-step of the EM algorithm the probability $p_{ij}$ that event $i$ is a direct offspring of event $j$ is estimated, along with the probability $p_i^b$ that the event was generated by the Poisson process $\mu$.

$$p_{ij} = \frac{g(t_i - t_j, x_i - x_j, y_i - y_j)}{\lambda(t_i, x_i, y_i)},$$

(4)

$$p_i^b = \frac{\mu(x_i, y_i)}{\lambda(t_i, x_i, y_i)},$$

(5)

Given the probabilistic estimate of the branching structure, the complete data log-likelihood is then maximized in the M-step (using standard methods for estimating a Pareto distribution) [VS08], providing an estimate of the model parameters.

2.3. **Joint inference of the ETAS-$\hat{P}_{\text{GMM}}$**. We propose three variants for inferring the joint ETAS-$\hat{P}_{\text{GMM}}$ model.

**ETAS-$\hat{P}_{\text{GMM}}$ 1.** In the first variant we start by clustering all events spatially using $\hat{P}_{\text{GMM}}$. We then evaluate the convex hull of each cluster and enforce $\mu(x, y)$ to be constant for each cluster in the convex hull. With
the support of $\mu(x, y)$ fixed, we then proceed to estimate ETAS using the EM algorithm in Section 2.2.

**ETAS-I$^2$GMM 2.** In the next variant we perform joint inference using a Monte-Carlo EM algorithm. In particular, at each EM iteration we perform the following steps:

i. (I$^2$GMM-step) Sample background events from probabilistic branching structure $p_{ij}$. Estimate $\mu(x, y)$ based on the clusters generated by $I^2GMM$ on the sampled background events only.

ii. (E-step) Estimate probabilistic branching structure and model parameters of triggering kernel as in Section 2.2.

**ETAS-I$^2$GMM 3.** In the last variant we use a weighted $I^2GMM$ algorithm in place of the i step in variant two above. Instead of estimating $\mu(x, y)$ based on sampled background events, we estimate $\mu(x, y)$ using weighted $I^2GMM$ on all events whose weights are estimated by (5). In the first EM iteration, since $p^b_i$ does not exist we initialized the weight to 1.

2.4. Baseline ETAS model with histogram estimator. We use the histogram estimator proposed in [VS08] as a baseline model for comparison. In particular, we let the background rate $\mu(x, y)$ be a constant

$$
\mu(x, y) = \mu_k \text{ if (x,y) is in cell } k, k \in 1, \ldots, K
$$

over each rectangular cell of a regular grid. There are then $K + 6$ parameters $\theta = (\mu_1, \ldots, \mu_K, a, c, d, w, \rho, K_0)$ that we need to estimate (assuming there are K cells in the grid) and for that purpose we use the EM algorithm in [VS08].

3. Experiments and Results.

3.1. Experiment 1: goodness of fit of ETAS-$I^2$GMM applied to CA earthquakes 3.5 and greater since 2000. We apply our models to the California earthquake event data filtered by year (greater than 2000) and magnitude (greater than 3.5). The geographic bounds range from 46.116 > latitude > 29.615 and -113.581 > longitude > -130.427. The dataset is divided into training and testing using time point 2010-01-01 00:00:00 for cutoff. All events before this timestamp are placed in the training dataset while all events after it are placed in the testing dataset. We performed experiments
with the following six models to analyze how the performance varies by adopting different modeling strategies:

1. ETAS-I²GMM 1.
2. ETAS-I²GMM 2.
3. ETAS-I²GMM 3.
4. 4×4 grid baseline model.
5. 3×4 grid baseline model.
6. 3×3 grid baseline model.

Note that experiments 1 to 3 are repeated 10 times and the means of the likelihood are recorded. For I²GMM we run 400 Gibbs sweeps; the hyperparameters are set as follows: \( \mu_0 = [36.4603 - 119.3265] \) the mean of the data set; \( \Sigma_0 = [21.4972 \quad 0 \quad 0 \quad 23.1351] \) the diagonal of the covariance of the data set; \( m = 22 \); \( \kappa_0 = 0.1 \); \( \kappa_1 = 0.5 \). The values of \( m, \kappa_0, \kappa_1 \) are tuned to let I²GMM generate equivalent number of clusters as the 4×4 grid baseline model.

We use the log-likelihood function

\[
\log L = \sum_{i=1}^{N} \log(\lambda(t_i, x_i, y_i)) - \int_{0}^{T} \int_{S} \lambda(t, x, y) dxdydt
\]

to evaluate the competing models for the background intensity. The results are shown in Table 2.

<table>
<thead>
<tr>
<th>Model</th>
<th>logL</th>
<th>( \sum_i \log(\lambda_i) )</th>
<th>( \int \lambda )</th>
</tr>
</thead>
<tbody>
<tr>
<td>ETAS-I²GMM 1</td>
<td>-4619</td>
<td>-1206</td>
<td>3413</td>
</tr>
<tr>
<td>ETAS-I²GMM 2</td>
<td>-4686</td>
<td>-1283</td>
<td>3403</td>
</tr>
<tr>
<td>ETAS-I²GMM 3</td>
<td>-4716</td>
<td>-1319</td>
<td>3397</td>
</tr>
<tr>
<td>4×4 Grid</td>
<td>-4980</td>
<td>-1590</td>
<td>3390</td>
</tr>
<tr>
<td>3×4 Grid</td>
<td>-4937</td>
<td>-1590</td>
<td>3330</td>
</tr>
<tr>
<td>3×3 Grid</td>
<td>-5023</td>
<td>-1582</td>
<td>3440</td>
</tr>
</tbody>
</table>

Table 2: Log-likelihood model comparison.

In Table 2 we find that all three ETAS-I²GMM models outperform ETAS with a histogram estimator. Between the three ETAS-I²GMM variants, the best-performing model is variant 1, where I²GMM is first estimated and the EM algorithm is run separately to estimate the triggering kernel parameters. It is worthwhile to note that finer clusters do not necessarily yield better results. Even though the 4×4 grid model generates more clusters it produces lower likelihood than the 3×4 grid model.
3.2. Experiment 2: ETAS-I²GMM for event-fault linkage from space-time event data. Next, we investigate the extent to which the ETAS-I²GMM model can learn fault structure from space-time event data. For this purpose we use the Community Fault Model 3.0 [PSB⁺07], which is a three dimensional representation (latitude, longitude, and elevation) of faults in Southern California. The CFM is a collaborative project undertaken by scientists of the Southern California Earthquake Center (SCEC) for studying active faults and earthquake phenomena and to improve regional earthquake hazard assessments. Our goal here is to assess how well ETAS-I²GMM recovers a 2D projection of the CFM 3.0 using only space-time-magnitude earthquake incident data as input. In particular, we generate a fault label for each event in the dataset by assigning fault membership as the nearest fault in CFM 3.0 (see Figure 3).

The ETAS-I²GMM-predicted label is taken from the first layer of the I²GMM model clusters, where offspring events are assigned to the cluster of their nearest neighbor among background events. To allow for comparison to the CFM 3.0, we restrict the geographic bounds of the CA earthquake event data to $36.958 > \text{latitude} > 31.518$ and $-113.719 > \text{longitude} > -121.176$, but we expand the magnitude threshold down to 2.5.

Since there are 145 actual fault lines in CFM 3.0 but all the six models we used in previous experiments generate at most 26 clusters, we added two additional models in the experiments for fault recovery:

- A I²GMM with parameters tuned to generate approximately 145 clus-
ters on average, and name this version as ETAS-I$^2$GMM 145 in our experiments.

- A $16 \times 15$ grid model that contains 143 non-empty clusters.

<table>
<thead>
<tr>
<th>Model</th>
<th>Accuracy</th>
<th>$\overline{\text{Acc}_{10}}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>ETAS-I$^2$GMM 145</td>
<td>0.46</td>
<td>0.52</td>
</tr>
<tr>
<td>ETAS-I$^2$GMM 1</td>
<td><strong>0.50</strong></td>
<td><strong>0.67</strong></td>
</tr>
<tr>
<td>ETAS-I$^2$GMM 2</td>
<td>0.45</td>
<td>0.46</td>
</tr>
<tr>
<td>ETAS-I$^2$GMM 3</td>
<td>0.41</td>
<td>0.33</td>
</tr>
<tr>
<td>Grid 16x15</td>
<td>0.45</td>
<td>0.47</td>
</tr>
<tr>
<td>Grid 4x4</td>
<td>0.37</td>
<td>0.37</td>
</tr>
<tr>
<td>Grid 3x4</td>
<td>0.36</td>
<td>0.28</td>
</tr>
<tr>
<td>Grid 3x3</td>
<td>0.35</td>
<td>0.32</td>
</tr>
</tbody>
</table>

**Table 3**

*Accuracy comparison of fault classification.*

Given that the number of clusters estimated by I$^2$GMM may be different from the number of faults in the CFM 3.0, we evaluate the success of fault-cluster recovery by considering the percentage of correctly classified data points. In addition to the overall accuracy, we evaluate the mean accuracy for the 10 largest faults, which contains 67% of data points across 145 faults. In Table 3 we present the accuracy for the six models listed in 3.1 as well as the two additional models. To calculate the accuracy, we first align the generated clusters with the ground-truth classes using the Hungarian algorithm [Kuh55, Ste00], and then calculate the percentage of the data points that fall into their classes of origin. We adopt accuracy for its simplicity and its invariance to potential switching between ground truth and predicted class labels. Mean accuracy is calculated as below:

\[
\overline{\text{Acc}} = \frac{1}{|C^*|} \sum_{C_k^* \in C^*} \frac{|C_k^* \cap C_k|}{|C_k^*|}
\]

where $C^*$ contains all fault clusters under consideration; $C_k$ is the predicted cluster corresponding to $C_k^*$ after alignment; $C_k^* \cap C_k$ indicates data points in both $C_k^*$ and $C_k$; $|S|$ denote the cardinality of the set $S$. To compute the mean accuracy for the ten largest faults $\overline{\text{Acc}_{10}}$, we set $C^*$ to contain the ten largest true fault clusters in the above equation.

Here again we see that ETAS-I$^2$GMM 1 performs best both in terms of accuracy and $\overline{\text{Acc}_{10}}$. In Figure 4 we plot the clusters recovered corresponding to ETAS-I$^2$GMM 145, ETAS-I$^2$GMM 1, $16 \times 15$ grid, $4 \times 4$ grid, and the true clusters for a better understanding of this outcome. Despite the fact that
Figure 4. True and predicted CFM fault groupings. Events with same labels are shown by the same color.
I²GMM generated only 26 unique clusters on average compared to 145 actual fault lines in CFM 3.0, a meaningful accuracy of 0.5 was achieved. Results suggest that a majority of events in fault clusters that tend to have elongated, skewed, and in some cases multi-mode shapes are clustered correctly by I²GMM. In contrast, the clusters in the two histogram models have abrupt boundaries formed from the grid irrespective of the shape of the underlying faults, as shown in Figure 4. Moreover, when we adjust the parameters of the I²GMM to get approximately the same number of clusters as the true number of fault clusters, we observe that the accuracy does not improve owing to erroneous splitting of events belonging to larger fault lines into multiple clusters. This is also true for the $16 \times 15$ grid model that generates 143 non-empty clusters. Although the accuracy improves with this model compared to the grid model with a smaller number of clusters, overall accuracy achieved by this model is still less than that achieved by I²GMM with 26 clusters (0.45 vs 0.50). The difference in accuracy between the two models increases in favor of I²GMM when we take into account only the largest ten fault clusters (0.47 vs 0.67). This is a natural result of the grid model arbitrarily splitting fault clusters compared to the more effective handling of elongated fault cluster shapes by I²GMM.

We illustrate this over-splitting problem in Figure 5 by plotting the clustering results of the ten largest faults. From the $\text{Acc}_{10}$ results in Table 3 and Figure 5 we can see that ETAS-I²GMM 1 did the best by achieving a mean accuracy of 0.67 across ten faults while recovering several of them by an accuracy of over 0.9. On the other hand, for the grid models the $\text{Acc}_{10}$ values are consistent with their corresponding overall accuracies.

4. Discussion. We introduced a coupled ETAS-I²GMM model for jointly estimating multi-scale clustering in earthquake data with parameters governing earthquake productivity and self-excitation. We also introduced what we believe is a novel machine-learning task for statistical seismology, namely estimating CFM fault clusters using unlabeled space-time-magnitude event data. Improving upon algorithms aimed at solving this task could aid in the development of future versions of CFM, as well as fault models in other regions of the world.

We also believe that the I²GMM model may have applications to point processes beyond those arising in seismology. Space-time self-exciting point processes arise in the study of crime [MSB+11, Moh14, MSM+15], conflict [LM11], and terrorism [PW+12, M+13, WPM13, WP14], as well as in social-network event dynamics, for example in social media [LMY+14, SJ12, ZEH+15]. In the case of crime, clusters arise naturally from the su-
Figure 5. Ten largest CFM faults and recovered clusters. Events with same labels are shown by the same color.
perposition of events committed by different offenders with different modus operandi. Similar clusters may arise from the operations of different terrorist groups within a geographic region. P²GMM is a flexible model for capturing this type of clustering in the intensity of events of a point process.

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