Probabilistic Hierarchical Forecasting with Deep Poisson Mixtures

Kin G. Olivares † Nganba Meetei ‡ Ruijun Ma ‡ Rohan Reddy ‡ Mengfei Cao ‡ † Carnegie Mellon University ‡ Forecasting Science, Amazon, New York, NY kdgutier@cs.cmu.edu, {kigutie, meeteio, ruijunma, reddyroh, mfcao}@amazon.com

Abstract

Hierarchical forecasting problems arise when time series compose a group structure that naturally defines aggregation and disaggregation coherence constraints for the predictions. In this work, we explore a new forecast representation, the Poisson Mixture Mesh (PMM), that can produce *probabilistic*, *coherent* predictions; it is compatible with neural forecasting innovations, and defines simple aggregation and disaggregation rules capable of accommodating hierarchical structures, unknown during its optimization. We perform an empirical evaluation to compare the PMM to other hierarchical forecasting methods on Australian domestic tourism data, where we obtain a relative improvement of nearly 20 percent.

1 Introduction and Motivation

There are several cases where forecast users need simultaneous forecasts throughout a hierarchy or group structure. The coherence between the predictions is critical for the user's trust. Some examples of hierarchical forecasting problems include retail demand where sales are tracked at the product and geographic levels (Seeger et al., 2017); electricity price forecasting where predictions are needed at different regional levels (Ben Taieb & Koo, 2019; Jeon et al., 2019).

The main goal of hierarchical forecasting is to leverage available information across all the levels of the structure, to produce coherent forecasts. Prior solutions to the challenge depend on two-stage approaches where *base forecasts* are created and later reconciled. More recent work extended the generation of probabilistic hierarchical forecasts into single and simplified models. Some limitations that we find in current solutions are (i) their efficiency, as any new hierarchical structure of interest would require fitting a whole new model, (ii) their accuracy, as the models tend to use restrictive probabilistic models, to enable reconciliation. In this work we explore a *Deep Poisson Mixture Network* (DPMN) solution for hierarchical forecasting, as we envision that its following characteristics can help to tackle the mentioned challenges:

- 1. **Probabilistic Coherence**: The Poisson Mixture Mesh (PMM) defines a joint distribution across all levels of the time series structures. And by construction, it guarantees probabilistic coherence of its predictions, through its simple aggregation and disaggregation rules.
- 2. **Enhanced Accuracy**: The PMM is a flexible and powerful improvement over available hierarchical methods capable of producing coherent, probabilistic forecasts that rely on strong assumptions on the distributions, like Gaussian noise, that enable its reconciliation strategies. Additionally, the PMM representation is independent of the model that supports and can leverage the rich literature on neural forecasting models.
- 3. Efficiency: Finally, the PMM can accommodate unrevealed hierarchical structures during its optimization. Its aggregation and disaggregation rules also allow users to easily generate any new marginal distribution across the hierarchy. In contrast, other more complex approaches require the hierarchical structure in advance, and changes to the structure need to fit a whole new specialized model.

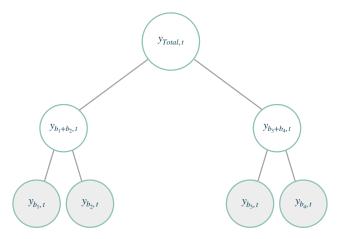


Figure 1: A simple three level time series hierarchical structure, with four bottom level variables. The disaggregated bottom variables are marked with gray background. In this description each node represents non overlapping series for a single point in time, other indices can be used like locations.

The remainder of the paper is structured as follows. Section 2 introduces notation and reviews relevant literature on hierarchical forecasting. Section 3 describes the DPMN model. In Section 4 we provide an empirical evaluation on the Australian Tourism hierarchical data, where we showcase the advantages of our method. Finally, in Sections 5 and 6 we discuss future work and conclude.

2 Literature Review

2.1 Hierarchical Forecasting Notation

A hierarchical time series, is a multivariate time series that satisfies linear aggregation constraints. The aggregation is typically depicted with a tree structure, but with group structures beyond strict hierarchies, it may not necessarily be described by a tree (Hyndman et al., 2014; Athanasopoulos et al., 2017; Spiliotis et al., 2020). A hierarchical multivariate time series can be denoted by the vector $\mathbf{y}_{[a,b],t} = [\mathbf{y}_{[a],t}^{\top} \mid \mathbf{y}_{[b],t}^{\top}]^{\top} \in \mathbb{R}^{N}$, where a,b,t stand for the single aggregate, bottom and time indices for the time series \mathbf{y} , and [a],[b],[t] listed indices respectively. With the total number of series in the hierarchy $N = N_a + N_b$, where N_a is the number of aggregated series and N_b the number of bottom series, that are at the most disaggregated level possible. The hierarchical aggregation constraints have the following convenient matrix representation:

$$\mathbf{y}_{[a,b],t} = \mathbf{S}\mathbf{y}_{[b],t} \quad \Leftrightarrow \quad \begin{bmatrix} \mathbf{y}_{[a],t} \\ \mathbf{y}_{[b],t} \end{bmatrix} = \begin{bmatrix} \mathbf{S}_{\text{sum}} \\ \mathbf{I}_{N_b} \end{bmatrix} \mathbf{y}_{[b],t}, \tag{1}$$

The matrix $\mathbf{S} \in \mathbb{R}^{(N_a+N_b)\times N_b}$ aggregates the bottom level series to the series above, it can be decomposed into a summing matrix \mathbf{S}_{sum} and an identity matrix \mathbf{I}_{N_b} . For a simple example, hierarchy described by Figure 1, where each parent node is the sum of its children. In this example the dimensions are $N_a=3$, $N_b=4$, and the hierarchical, aggregated and base series are respectively

$$y_{\text{Total},t} = y_{b_1,t} + y_{b_2,t} + y_{b_3,t} + y_{b_4,t}$$
$$\mathbf{y}_{[a],t} = [y_{\text{Total},t}, \ y_{b_1,t} + y_{b_2,t}, \ y_{b_3,t} + y_{b_4,t}]^{\mathsf{T}} \qquad \mathbf{y}_{[b],t} = [y_{b_1,t}, \ y_{b_2,t}, \ y_{b_3,t}, \ y_{b_4,t}]^{\mathsf{T}}$$

The summing matrix of the Figure 1 example can be written as:

$$\mathbf{S} = \begin{bmatrix} \mathbf{S}_{\text{sum}} \\ \mathbf{I}_{N_b} \end{bmatrix} = \begin{bmatrix} 1 & 1 & 1 & 1 \\ 1 & 1 & 0 & 0 \\ 0 & 0 & 1 & 1 \\ 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

2.2 Reconciliation Strategies

Prior solutions to the hierarchical forecasting challenge follow a two-stage process, where first a set of base forecasts $\hat{\mathbf{y}}_{[a,b],t} \in \mathbb{R}^{N_a+N_b}$ is created and then revised into coherent forecasts $\tilde{\mathbf{y}}_{[a,b],t}$ through a reconciliation method. The reconciliation can be compactly expressed by

$$\tilde{\mathbf{y}}_{[a,b],t} = \mathbf{SP}\hat{\mathbf{y}}_{[a,b],t} \tag{2}$$

where $\mathbf{S} \in \mathbb{R}^{(N_a+N_b)\times N_b}$ is the hierarchical aggregation matrix and $\mathbf{P} \in \mathbb{R}^{N_b\times (N_a+N_b)}$ is a projection matrix determined by the reconciliation strategies. The most common reconciliation methods can be classified into top-down, bottom-up and alternative reconciliation approaches.

- Bottom-up: The simple *bottom-up* strategy (NaiveBU; Orcutt et al. 1968; Dunn et al. 1976), first generates bottom level forecasts and then aggregates them to produce predictions for all the series in the hierarchical structure. Here $\mathbf{P} = [\mathbf{0}_{N_b \times N_a} \mid \mathbf{I}_{N_b}]$.
- Top-down: The *top-down* strategy (TD; Gross & Sohl 1990; Fliedner 1999), distributes the total forecast, and then disaggregates it down the hierarchy using proportions that can be historical or forecasted. In this strategy $\mathbf{P} = [\mathbf{p}_{N_b \times 1} \mid \mathbf{0}_{N_b \times (N_a + N_b 1)}]$. This method has several variants, depending on the way the proportions \mathbf{p} are created, like the *average of historical proportions*, *proportions of historical averages*, or the *forecasted proportions*.
- Alternative: The more recent middle-out strategies (MO; Hyndman & Athanasopoulos 2017), treat the second stage reconciliation as an optimization problem for the projection matrix P. These reconciliation techniques include among others the least squares reconciliation (Hyndman et al., 2011), the minimum trace reconciliation (MinT; Wickramasuriya et al. 2019) and the empirical risk minimization approach (ERM; Ben Taieb & Koo 2019), that relaxes unbiasedness assumptions in MinT.

Despite the advancements in post-processing methods, there are still fundamental limitations. First, most post-process reconciliation methods produce point rather than probabilistic forecasts, with some exceptions (Gneiting & Katzfuss, 2014; Taieb et al., 2017; Panagiotelis et al., 2020). Second, the mentioned methods learn the model parameters of the base level forecasts independently, missing the opportunity to share a common model that leverages information across all the time series in the hierarchical structure. Finally, most post-processing reconciliation strategies do not consider the model parameters obtained during the first stage but only their predictions.

2.3 Hierarchical Neural Forecasting

In the last decade, neural network-based forecasting methods have become ubiquitous in large-scale forecasting applications. Neural forecasting has transcended the boundaries of industry into academia, as it has redefined the state-of-the-art in many practical tasks, and forecasting competitions (Benidis et al., 2020; Makridakis et al., 2018, 2020). The latest contributions on neural hierarchical forecasting have incorporated the reconciliation strategies to the models. Methods like the Simultaneous Hierarchically Reconciled Quantile Regression (SHARQ; Han et al. 2021) and Hierarchically Regularized Deep Forecasting (HIRED; Paria et al. 2021) augment the training loss function with quadratic approximations to the hierarchical constraints, while the Hierarchical End-to-End learning (HierE2E; Rangapuram et al. 2021) method imposes exact hierarchical constraints to the predictions through a projection in a single end-to-end model. None of the architectures and methods mentioned are readily usable for large-scale applications. For instance, the architectures do not use the forking sequences optimization technique (Wen et al., 2017), and the HierE2E reconciliation projection scales poorly as it needs to receive all the time series of the hierarchy simultaneously.

2.4 Multi-horizon Quantile Recurrent Forecaster

Many scalability and accuracy improvements of sequence modeling have been incorporated into the MQ-Forecaster family (Wen et al., 2017). The methods of the family produce probabilistic forecasts using non parametric quantile regression (Koenker & Bassett, 1978). In spite of major advances in forecast accuracy for large-scale applications, the estimated quantiles are not additive and ensuring the hierarchical coherence of quantile regressors would potentially imply the need to fit a network for the marginal distributions associated to each hierarchical structure of interest, or coherency regularization, which encourages but does not guarantee coherence.

Our primary goal is then to create a probabilistic hierarchical forecasting model that is accurate, and efficient, by extending the proven MQ-Forecaster family.

3 Poisson Mixture Mesh Model

3.1 Poisson Mixture Mesh

The foundation of PMM model is the assumption that the joint distribution of a multivariate time series $\mathbf{y}_{[b],[t]}$, is described by a Poisson Mixture distribution. The mixture first determines a latent categorical variable K that selects a state, and conditional on the selection an independent Poisson random variable that determines the observations. Using mixing weights w_k and rate $\lambda_{b,k,t}$ parameters the joint likelihood can be parametrized:

$$P(\mathbf{y}_{[b],[t]}) = \sum_{k=1}^{N_{\lambda}} P(K = k) P(\mathbf{y}_{[b],[t]} | K = k)$$

$$= \sum_{k=1}^{N_{\lambda}} w_k \prod_{(b,t) \in \mathcal{B}} \left(\frac{(\lambda_{b,k,t})^{y_{b,t}} \exp^{-\lambda_{b,k,t}}}{(y_{b,t})!} \right)$$
(3)

We denote such Poisson Mixture distributed variable as $\mathbf{y}_{[b],[t]} \sim \mathrm{PM}(\mathbf{w}_{[k]},\,\lambda_{[b],[k],t})$. We describe properties of the Poisson Mixture distribution that make it exceptionally suited for the hierarchical forecasting task below.

3.2 Probabilistic Properties

Under conditional independence assumption and the unmixing property of the parameters $\mathbf{w}_{[k]}$ and $\lambda_{[b],[k],t}$, we show in the Appendix A that the marginal distributions follow the simple aggregation and disaggregation rules:

• Aggregation Rule: Let a multivariate time series $\mathbf{y}_{[b],[t]}$ follow the PMM then aggregated levels satisfy

$$\mathbf{y}_{[a],t} = \mathbf{S}_{\text{sum}} \mathbf{y}_{[b],t} \sim \text{PM}(\mathbf{w}_{[k]}, \mathbf{S}_{\text{sum}} \boldsymbol{\lambda}_{[b],[k],t})$$
 (4)

with $\mathbf{S}_{\text{sum}} \in \mathbb{R}^{N_a \times N_b}$ the hierarchical aggregation matrix.

• Disaggregation Rule: Let the most aggregated level time series $y_{\text{Total},t}$ follow the PMM, then l-th level series can be built with the proportions $\mathbf{p}_{[b],t}^{(l)} \in \mathbb{R}^{N_b}$ by

$$\mathbf{y}_{[b],t} = \mathbf{p}_{[b],t}^{(l)} y_{\text{Total},t} \sim \text{PM}(\mathbf{w}_{[k]}, \mathbf{p}_{[b],t}^{(l)} \odot \boldsymbol{\lambda}_{\text{Total},[k],t})$$
 (5)

with proportions $0 \le p_{b_i,t}^{(l)}, \sum_{b_i} p_{b_i,t}^{(l)} = 1$ and \odot a multiplication over the b-th dimension.

3.3 Deep Poisson Mixture Network

The PMM offers a solution to model the high dimensional regression problem

$$P(\mathbf{y}_{[b],[t:t+h]} \mid \mathbf{y}_{[b],[:t]}, \mathbf{x}_{[:t]}^{(h)}, \mathbf{x}_{[:t]}^{(f)}, \mathbf{x}^{(s)})$$
(6)

where $\mathbf{y}_{[b],[t:t+h]},\mathbf{y}_{[b],[:t]},\mathbf{x}_{[:t]}^{(h)},\mathbf{x}_{[:t]}^{(f)},\mathbf{x}^{(s)}$ represent future observations of the multivariate time series, observations of the target up until time t, the past covariates, known future information and static covariates, respectively. It is fully general and compatible with the MQ-Forecaster model family (Wen et al., 2017), for which we only need a few adjustments.

For this work, we combine the PMM with the convolutional encoder version of MQ-Forecaster (MQCNN) into the *Deep Poisson Mixture Network* (DPMN). To alleviate the computational burden of optimizing for a full joint distribution from Equation (3), simpler variants need to be devised (Varin et al., 2011). The learning objective is implied by the different versions of the PMM that we design, we describe the variants in detail in Section 3.4.

3.4 Poisson Mixture Mesh Variants

The PMM aggregation and disaggregation rules, described in Section 3.2, allow for applying any of the post-processing reconciliation strategies from Section 2.2. Yet, we can further improve the DPMN, by augmenting the model's architecture or optimization with the said strategies.

• Naive Bottom-Up:

The DPMN-NaiveBU model is estimated exclusively on the bottom-level series that are treated as conditionally independent from one another, however for a single time series its time dependencies are still estimated.

Optimization:

$$P\left(\mathbf{y}_{[b],[t]}\right) = \prod_{b \in [b]} P\left(\mathbf{y}_{b,[t]}\right) = \prod_{b \in [b]} \left(\sum_{k=1}^{N_{\lambda}} w_k \prod_{t \in [t]} \operatorname{Poisson}\left(y_{b,t} | \lambda_{b,k,t}\right)\right)$$
(7)

• Group Bottom-Up:

Given that the prediction intervals of the NaiveBU version are excessively wide, we created the GroupBU version of the model to help the PMM learn a learn a better correlation structure of the bottom-level series. The DPMN-GroupBU model learns a joint distribution for groups g of bottom-level series and uses a group sampling strategy for the network's SGD optimization. The independence treatment between the groups, maintains the tractability of the likelihood computation.

Optimization:

$$P\left(\mathbf{y}_{[b],[t]}\right) = \prod_{g \in [g]} P\left(\mathbf{y}_{[g],[t]}\right) = \prod_{g \in [g]} \left(\sum_{k=1}^{N_{\lambda}} w_k \prod_{(b,t) \in g \times [t]} Poisson\left(y_{b,t} | \lambda_{b,k,t}\right)\right)$$
(8)

During the hierarchical reconciliation stage, the DPMN's predictions of the upper levels are created using bottom-up strategy and the aggregation rule:

$$\tilde{\mathbf{y}}_{[a,b],t} \sim \text{PM}(\mathbf{w}_{[k]}, \mathbf{S}\hat{\boldsymbol{\lambda}}_{[b],[k],t})$$
 (9)

4 Empirical Evaluation

For the empirical evaluation of our proposed methods, we use a publicly available dataset from existing hierarchical forecasting literature. This allows to compare against well-performing benchmarks from the literature and test the DPMN model on real data with natural hierarchical structures.

The dataset we use is Tourism-L, a detailed Australian Tourism dataset with origin in a National Visitor Survey, managed by Tourism Research Australia¹ (Tourism Australia, Canberra, 2019). As described in Table 1, the dataset contains 555 monthly series from 1998 to 2016, it is organized by geography and purpose of travel. The geographical hierarchy comprises seven states, divided further into 27 zones and 76 regions. This dataset has been referenced by important hierarchical forecasting studies like the one of the MinT reconciliation strategy (Wickramasuriya et al., 2019).

Table 1: Australian Tourism flows (Tourism-L).

Series per Division	Series per Purpose	Total
1	4	5
7	28	35
27	108	135
76	304	380
111	444	555
	1 7 27 76	Division Purpose 1 4 7 28 27 108 76 304

¹The Tourism-L dataset can be publicly accessed in the MinT reconciliation web page.

Table 2: Empirical evaluation of probabilistic, hierarchical forecasts on the Australian Tourism Flows (Tourism-L). Mean *continuous ranked probability score* (CRPS) for predictions at each aggregation level, averaged over 8 runs. The best result is highlighted (lower is better). Only the best performing among the state-of-the-art reconciliation strategies is shown.

Level	DPMN-GroupBU	DPMN-NaiveBU	Hier-E2E	Best Reconciliation
Overall	0.1260 ± 0.0046	0.1578 ± 0.0084	0.1520 ± 0.0032	0.1609 (ARIMA-MinT-shr)
1 (geo.)	0.0411 ± 0.0110	0.1130 ± 0.0197	0.0810 ± 0.0053	0.0438 (ARIMA-MinT-shr)
2 (geo.)	0.0624 ± 0.0070	0.1189 ± 0.0161	0.1030 ± 0.0030	0.0816 (ARIMA-MinT-shr)
3 (geo.)	0.1122 ± 0.0049	0.1466 ± 0.0131	0.1361 ± 0.0024	0.1433 (ARIMA-MinT-shr)
4 (geo.)	0.1571 ± 0.0032	0.1759 ± 0.0125	0.1752 ± 0.0026	0.2036 (ARIMA-MinT-shr)
5 (prp.)	0.0747 ± 0.0056	0.1315 ± 0.0060	0.1027 ± 0.0062	0.0830 (ARIMA-MinT-shr)
6 (prp.)	0.1100 ± 0.0044	0.1416 ± 0.0058	0.1403 ± 0.0047	0.1479 (ARIMA-MinT-shr)
7 (prp.)	0.1901 ± 0.0046	0.1908 ± 0.0052	0.2050 ± 0.0028	0.2437 (ARIMA-MinT-shr)
8 (prp.)	0.2600 ± 0.0039	$\bf 0.2428 \pm 0.0045$	0.2727 ± 0.0017	$0.3406(\mathtt{ARIMA-MinT-shr})$

4.1 Evaluation Metrics

The evaluation of the model's predictions is based on the *quantile loss* (QL); consider the estimated cumulative distribution function $\hat{\mathbf{F}}_{i,t}$ for an observation $\mathbf{y}_{i,t}$, then the quantile loss is defined by:

$$QL(\hat{\mathbf{F}}_{i,t}, y_{i,t})_q = 2\left(\mathbb{1}\{y_{i,t} \le \hat{\mathbf{F}}_{i,t}^{-1}(q)\} - q\right)\left(\hat{\mathbf{F}}_{i,t}^{-1}(q) - y_{i,t}\right)$$
(10)

We further summarize the evaluation, for convenience of exposition and to ensure the comparability of our results with the existing literature, using the *continuous ranked probability score* (CRPS; Matheson & Winkler 1976). The CRPS measures the predictive power of forecast distributions and has desirable theoretical guarantees (Gneiting & Ranjan, 2011). Following notation from Laio & Tamea 2007, the CRPS² is defined as

$$CRPS(\hat{\mathbf{F}}_{i,t}, y_{i,t}) = \int_0^1 QL(\hat{\mathbf{F}}_{i,t}, y_{i,t})_q dq$$
(11)

4.2 Training Methodology and Hyperparameter Optimization

We set the training window to 202 observations (16 years) preceding the generation of the 12 (1 year) steps ahead forecast. To train the neural network, we minimize the negative log likelihood of the PMM variant, using stochastic gradient descent with *Adaptive Moments* (ADAM; Kingma & Ba 2014). The network is trained for 3,000 epochs, with 2 NVIDIA K80 GPUs and a grouped batch size of 4 regions and 4 purposes each. The DPMN is implemented using MXNet (Tianqi Chen et al., 2015) and the learning rate and random initialization hyperparameters are selected using HYPEROPT (Bergstra et al., 2011), a Bayesian optimization library that efficiently explores the hyperparameters using tree-structured Parzen estimators. The hyperparameter search process trains on data from 1998 to 2014 and evaluates the validation performance on data from 2015. Once the optimal hyperparameter values are determined, we retrain shifting the training window forward one year and predict on data from 2016 to obtain the final CRPS scores.

4.3 Forecasting Results

We compare against the next methods: (1) NaiveBU that produces univariate bottom-level time series forecasts independently and then sums them according to the hierarchical constraints, a distribution is generated using Gaussian assumptions on the errors. (2) MinT (Wickramasuriya et al., 2019) that reconciles unbiased independent forecasts and minimizes the variance of the forecast errors. (3) ERM (Ben Taieb & Koo, 2019) that improves on the unbiasedness assumption of the base forecasts in MinT and aims to minimize the bias-variance trade-off of the errors. (4) PERMBU (Taieb et al., 2017) that performs a probabilistic hierarchical aggregation and (5) HierE2E (Rangapuram et al., 2021) that combines a deep vector autoregressive approach with hierarchical constraints ³.

²In practice the evaluation of the CRPS uses numerical integration technique, that discretizes the quantiles and treats the integral with a Riemann approximation, averaging over uniformly distanced quantiles.

³The HierE2E benchmark models and experiments are available in an GluonTS library extension.

We report two variants of the DPMN model as an ablation experiment to better analyze the source of the forecast accuracy improvements. The first version, the DPMN-NaiveBU, treats the bottom level series as independent and the DPMN-GroupBU version that considers groups of time series during its likelihood estimation, both methods obtain probabilistic coherent predictions through their aggregation rule using bottom-up reconciliation.

Table 2 contains the CRPS scores for the predictions at each aggregation level through the Tourism-L hierarchy. The top row reports the overall CRPS score (averaged across all the hierarchy levels). We highlight the best result in **bolds**.

The DPMN-GroupBU improves the overall CRPS by near 17.1 percent against the second-best alternative. Its predictions show uniform performance gains across all the levels of the hierarchy. Our results confirm previous observations from the community that a shared model, capable of learning from all the time series jointly improves the predictions over those from univariate time series methods. Additionally the comparison ⁴ between the NaiveBU and the GroupBU versions of the method show that an expressive joint distribution framework capable of leveraging the hierarchical structure of the data is also beneficial for the forecast accuracy, when compared to the more restrictive Gaussian distributions based methods from the hierarchical forecasting literature.

5 Future Work

We constructed the DPMN extending the MQ-Forecaster architecture. Despite the significant improvements in computational efficiency and accuracy for large-scale applications the MQ-Forecaster family may not be suited for small datasets. Given the time constraints, we restricted our current experiments to the Tourism-L dataset. We are looking forward to perform more experiments on smaller and larger datasets to continue strengthening this paper's findings.

The current version of the GroupBU model defines the groups by simple geographical distances. However, other exciting approaches could arise when using distances over different dimensions. New ways of describing the groups could create interesting connections with nearest neighbors and clustering literature.

We envision that we can include interactions of time series beyond the predefined groups implied by the GroupBU approach to enhance the ability of the PMM to model the joint likelihood of the hierarchically structured time series while maintaining its estimation tractable. The top-down disaggregation method to learn aggregate level Poisson rates and bottom level proportions could help the model capture higher hierarchical relationships inherited from the structure's upper levels.

6 Conclusion

We have introduced a novel method for probabilistic hierarchical forecasting, the *Poisson Mixture Mesh* (PMM). Our approach is a joint probability distribution that defines simple aggregation and disaggregation rules for its predictions. By construction PMM produces coherent probabilistic forecasts. It provides a flexible and expressive representation capable of highly accurate predictions and is highly efficient since it is scalable and can accommodate hierarchical structures unrevealed during its optimization. Additionally, the method is compatible with innovations in neural forecasting as a minimal adaptation process is needed to augment existing deep forecasting architectures. We empirically showcase the advantages of PMM through its application with the DPMN model, which improves by 17.1 percent compared with previous state-of-the-art results on Australian domestic tourism data, a hierarchical dataset from the literature.

The original implementation of MinT is available in the R package hts (Hyndman et al., 2020), it automatically selects ARIMA and ETS univariate base forecast methods (Hyndman & Khandakar, 2008).

⁴Figure 2 and Figure 3 in the Appendix show a qualitative exploration of the two versions of the model.

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A Appendix

Poisson Mixture Mesh Properties

To describe the joint distribution across all levels of the hierarchical time series structure we assume that for a given time series $\mathbf{y}_{b,[t]}$, its non overlapping observations are distributed Poisson mixture, denoted $y_{b,t} \sim PM(\mathbf{w}_{[k]}, \boldsymbol{\lambda}_{b,[k],t})$, the observations are conditionally independent given the Poisson rates, the mixing weights $\mathbf{w}_{[k]}$ are shared between the observations, and their respective Poisson components do not mix⁵, under these assumptions we show in A.1, A.2 and A.4 the properties are summarized in Table A1:

A.1 Unmixing Property

To describe the joint distribution across all levels of the time series structure we assume that for a given time series $\mathbf{y}_{b,[t]}$ with non overlapping observations $y_{b,\,t} \sim PM(\mathbf{w}_{[k]}, \boldsymbol{\lambda}_{b,[k],t})$, the observations are conditionally independent once the Poisson rates are known, the mixing weights \mathbf{w} are shared between the observations, and their respective Poisson components do not mix. We prove the properties from Table A1.

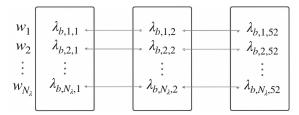


Figure 1: Representation of the un-mixing property of the Poisson components of the PMM.

A.2 Joint Distribution

Let two bottom random variables have a PM distribution with mixing weights $\mathbf{w}_{[k]}$ and rates $\lambda_{b,[k],t}, \lambda_{b,[k],t'}$, that is $y_{b,t} \sim PM(\mathbf{w}_{[k]}, \lambda_{b,[k],t})$ and $y_{b,t'} \sim PM(\mathbf{w}_{[k]}, \lambda_{b,[k],t'})$, then under the conditional independence assumption, the independence between the weights and the bottom series and the deterministic evolution property the joint distribution of the variables is:

$$P(y_{b,t},y_{b,t'}) = \sum_{k} w_k \text{Poisson}(y_{b,t}|\ \lambda_{b,k,t}) \text{Poisson}(y_{b,t'}|\ \lambda_{b,k,t'})$$

Proof

$$\begin{split} &P(y_{b,t},y_{b,t'}) \\ &= \sum_{k} \sum_{l} P(y_{b,t},\ y_{b,t'}|\ \lambda_{b,k,t},\lambda_{b,l,t'}) P(\lambda_{b,k,t}|\lambda_{b,l,t'}) P(\lambda_{b,l,t'}) \\ &= \sum_{k} P(y_{b,t}|\ \lambda_{b,k,t}) P(y_{b,t'}|\ \lambda_{b,k,t'}) P(\lambda_{b,k,t'}) \\ &= \sum_{k} w_{k} \text{Poisson}(y_{b,t}|\ \lambda_{b,k,t}) \text{Poisson}(y_{b,t'}|\ \lambda_{b,k,t'}) \end{split}$$

The first equality is the chain rule of probability, the second equality comes from the deterministic evolution property, the final part comes from the conditional independence property. By induction,

$$P(\mathbf{y}_{b,[t]}) = \sum_{k=1}^{N_{\lambda}} w_k \prod_{t=1}^{N_T} \operatorname{Poisson}(y_{b,\,t}|\ \lambda_{b,k,t})$$

⁵The un-mixing property of the Poisson components refers to the assumption that the count populations of a given Poisson component are deterministic.

A.3 Covariance Structure

From the joint distribution for $y_{b,t}$, $y_{b,t'}$ it can be shown using linearity of expectation and the Poisson distribution properties that the covariance structure of the bottom series can be described by the covariance structure of the Poisson rates:

$$Cov(y_{b,t}, y_{b,t'}) = \sum_{k=1}^{N_{\lambda}} w_k (\lambda_{b,k,t} - \bar{\lambda}_{b,[k],t}) (\lambda_{b,k,t'} - \bar{\lambda}_{b,[k],t'})$$

where $\bar{\lambda}_{b,[k],t} = \sum_k w_k \lambda_{b,k,t}$ and $\bar{\lambda}_{b,[k],t'} = \sum_k w_k \lambda_{b,k,t'}$. This shows that in spite of the strong assumptions made for the PMM the model is capable of describing a rich correlation structure from the dispersion inherited of the mixture.

A.4 Aggregation Rule

Let two bottom level random variables have a PM distribution with mixing weights $\mathbf{w}_{[k]}$ and rates $\lambda_{b,[k],t}$, $\lambda_{b',[k],t}$, that is $y_{b,t} \sim PM(\mathbf{w}_{[k]}, \lambda_{b,[k],t})$ and $y_{b',t} \sim PM(\mathbf{w}_{[k]}, \lambda_{b',[k],t})$, then under the conditional independence assumption, the independence between the weights and the bottom series and the deterministic evolution property we have the following aggregation rule:

$$\mathbf{y}_{b,t} + \mathbf{y}_{b',t} \sim PM(\mathbf{w}_{[k]}, \boldsymbol{\lambda}_{b,[k],t} + \boldsymbol{\lambda}_{b',[k],t})$$

Proof

$$\begin{split} &P(y_{b,t} + y_{b',t}) \\ &= \sum_{k} \sum_{l} P(y_{b,t} + y_{b',t} | \ \lambda_{b,k,t}, \lambda_{b',l,t}) P(\lambda_{b,k,t} | \lambda_{b',l,t}) P(\lambda_{b',l,t}) \\ &= \sum_{k} P(y_{b,t} + y_{b',t} | \ \lambda_{b,k,t}, \ \lambda_{b',k,t}) P(\lambda_{b',k,t}) \\ &= \sum_{k} w_{k} \text{Poisson}(y_{b,t} + y_{b',t} | \ \lambda_{b,k,t} + \lambda_{b',k,t}) \end{split}$$

Table A1: Summary properties of the Mixed Poisson State-Space Model

1. Marginal distribution	$y_{b,t} \sim PM(\mathbf{w}_{[k]}, \boldsymbol{\lambda}_{b,[k],t})$	
2. Assumptions		
2.1 Conditional Independence	$y_{b_1,t} \lambda_{b_1,k,t}$ $\perp \!\!\! \perp$ $y_{b_2,t} \lambda_{b_2,h,t}$ if b_1,b_2 do not overlap	
2.2 Unmixing property	$P(\lambda_{b,k,t} \lambda_{b,h,t})=\delta_{k,h}$ Poisson components do not mix where $\delta_{k,h}$ representing the Kronecker delta	
3. Joint Distribution	$P(\mathbf{y}_{b,[t]}) = \sum_{k=1}^{N_{\lambda}} w_k \prod_{t=1}^{N_T} \text{Poisson}(y_{b,t} \; \lambda_{b,k,t})$	
4. Hierarchical Coherence		
4.1 Aggregation Rule	$\mathbf{y}_{[a],t} = \mathbf{S}_{\mathrm{sum}} \mathbf{y}_{[b],t} \sim PM(\mathbf{w}_{[k]}, \mathbf{S}_{\mathrm{sum}} \boldsymbol{\lambda}_{[b],[k],t})$	
4.2 Disaggregation Rule	$\begin{array}{l} \mathbf{y}_{[b],t} = \mathbf{p}_{[b],t} y_{\mathrm{Total},t} \sim PM(\mathbf{w}_{[k]},\mathbf{p}_{[b],t} \odot \boldsymbol{\lambda}_{\mathrm{Total},[k],t}) \;) \\ \text{with shares } 0 \leq p_{b_i}, \sum_{b_i} p_{b_i,t} = \mathbb{1} \; \text{and} \; \odot \\ \text{multiplication in the b-th dimension} \end{array}$	
5. Covariance Structure	$Cov(y_{b,t}, y_{b,t'}) = \sum_{k=1}^{N_{\lambda}} w_k (\lambda_{b,k,t} - \bar{\lambda}_{b,[k],t}) (\lambda_{b,k,t'} - \bar{\lambda}_{b,[k],t'})$	

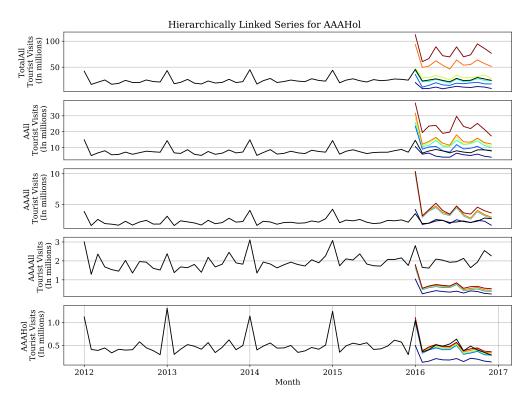


Figure 2: Example of a hierarchically linked time series from the Tourism-L dataset. The top row shows the total number of tourist visits in Australia (TotalAll), the second row shows the visits to Australia for the North South Wales state (AAll), the third row shows the holiday visits in the metropolitan area of New South Wales (AAAll), the fourth row shows the total visits to Sydney (AAAAll), the final row shows the holiday visits to Sydney (AAAHol). Quantile predictions are shown in colored lines. The DPMN-NaiveBU performs well in disaggregated series and means, but makes the predictions of the aggregated series unnecessarily wide.

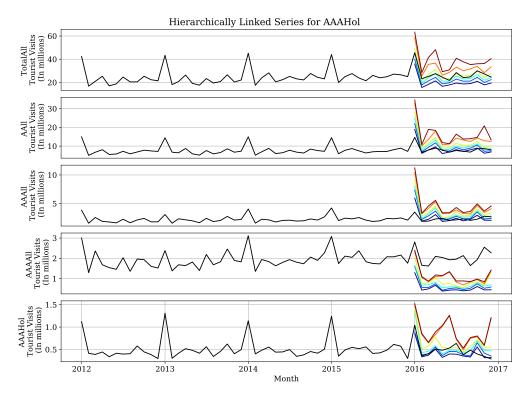


Figure 3: Example of a hierarchically linked time series from the Tourism-L dataset. The top row shows the total number of tourist visits in Australia (TotalAll), the second row shows the visits to Australia for the North South Wales state (AAll), the third row shows the holiday visits in the metropolitan area of New South Wales (AAAll), the fourth row shows the total visits to Sydney (AAAAll), the final row shows the holiday visits to Sydney (AAAHol). Quantile predictions are shown in colored lines. The DPMN-GroupBU performs well in disaggregated series and aggregate series.