

Machine Learning for Generalized Linear Models

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Chapter 1

Introduction

Generalized Linear Models (GLMs) are a class of statistical models that extend linear regression by allowing for non-normal distributions of the response variable. They encompass various models, such as logistic regression, Poisson regression, and others, making them highly versatile for different applications. GLMs have found their place in domains such as healthcare (disease prediction), finance (risk assessment), and natural language processing (sentiment analysis).

However, with the advent of big data, the limitations of traditional GLMs become apparent. Large datasets and complex feature spaces often lead to challenges in both performance and scalability. Machine learning techniques such as regularization, feature engineering, and ensemble methods provide ways to overcome these challenges. Furthermore, with advancements in hardware, specifically Graphics Processing Units (GPUs), we can now scale GLMs to handle large-scale datasets efficiently.

This dissertation aims to explore how machine learning techniques can enhance GLMs' performance and predictive power. Additionally, we investigate how GPU acceleration can support these enhancements in large-scale, real-time applications.

1.1 Objectives

The objectives of this dissertation are:

- To explore how regularization, feature engineering, and advanced machine learning techniques enhance GLMs.
- To investigate the role of GPU optimization in accelerating large-scale GLMs.
- To evaluate the performance improvements of machine learning-enhanced GLMs using various metrics.

Chapter 2

Background: Machine Learning and Generalized Linear Models

2.1 Overview of GLMs

Generalized Linear Models (GLMs) consist of three main components:

- **Random Component:** Specifies the probability distribution of the response variable (e.g., normal, binomial, Poisson).
- **Systematic Component:** Involves a linear predictor, $\eta = X\beta$, where X is the matrix of input features and β is the vector of coefficients.
- **Link Function:** A function that relates the mean of the distribution to the linear predictor, such as the logit function in logistic regression.

In traditional applications, GLMs are used for modeling relationships between a set of predictor variables and a response variable. For example, logistic regression is commonly used for binary classification problems. However, GLMs can struggle with high-dimensional datasets and complex non-linear relationships, which is where machine learning techniques come into play.

2.2 Machine Learning Techniques in GLMs

Machine learning techniques have been widely adopted to enhance GLMs in several ways:

- **Regularization:** Techniques such as Lasso (L1) and Ridge (L2) regularization add penalty terms to the GLM objective function, controlling model complexity and reducing overfitting.
- **Feature Engineering:** Creating interaction terms and polynomial features can improve GLMs' ability to capture complex relationships between variables.
- **Model Selection:** Cross-validation and information criteria (e.g., AIC, BIC) are used to select the best model and avoid overfitting.

In the following chapters, we explore these techniques in more depth and demonstrate their application to GLMs.

Chapter 3

Enhancing GLMs with Machine Learning Techniques

3.1 Regularization in GLMs

Regularization is a critical technique in machine learning that mitigates overfitting by penalizing large coefficients in the model. Overfitting occurs when a model becomes excessively complex, capturing noise and fluctuations in the training data that do not generalize well to new, unseen data. Regularization addresses this by introducing a penalty term to the model's objective function, which controls the model's complexity and enhances its generalization capabilities.

In the context of Generalized Linear Models (GLMs), two popular forms of regularization are Lasso (L1) and Ridge (L2) regularization:

- **Lasso (L1) Regularization:** Lasso adds an absolute penalty equal to the sum of the absolute values of the coefficients. This penalty term has the effect of shrinking some coefficients to zero, effectively performing feature selection by excluding less relevant features from the model. Therefore, Lasso regularization encourages sparsity in the model, which is particularly useful in high-dimensional datasets where many features may be irrelevant or redundant. Sparse models not only reduce overfitting but also enhance interpretability, as only the most impactful features are retained. The L1 penalty can be formulated as:

$$\text{Objective} = \text{Loss} + \lambda \sum_{j=1}^p |\beta_j|$$

where λ is a regularization parameter that controls the strength of the penalty, and β_j are the model coefficients. A higher λ value increases the regularization effect, leading to more coefficients being set to zero.

```
1 from sklearn.linear_model import LogisticRegression
2
3 # Lasso (L1) Regularization
```

```

4     model_lasso = LogisticRegression(penalty='l1', solver='saga',
5     max_iter=10000).fit(X_train, y_train)

```

Listing 3.1: Logistic Regression with L1 Regularization

- **Ridge (L2) Regularization:** Ridge regularization adds a squared penalty to the coefficients, which discourages large coefficient values but does not necessarily shrink them to zero. Unlike Lasso, Ridge regularization does not perform feature selection; instead, it distributes the penalty across all coefficients, reducing their magnitudes. This form of regularization is useful when all features are expected to contribute to the model's predictions, but some may do so less strongly than others. Ridge regularization is particularly effective in situations where there is multicollinearity among features, as it tends to stabilize the estimates. The L2 penalty is expressed as:

$$\text{Objective} = \text{Loss} + \lambda \sum_{j=1}^p \beta_j^2$$

Here, the penalty is proportional to the square of the coefficients, leading to smaller values for large coefficients and reducing the risk of overfitting.

```

1     # Ridge (L2) Regularization
2     model_ridge = LogisticRegression(penalty='l2').fit(X_train,
3     y_train)

```

Listing 3.2: Logistic Regression with L2 Regularization

The Choice Between Lasso and Ridge Regularization

The choice between Lasso and Ridge regularization depends on the nature of the dataset and the goals of the analysis:

- **Lasso Regularization** is ideal when you suspect that many features in your dataset are irrelevant or contribute little to the predictive power of the model. By setting some coefficients to zero, Lasso performs implicit feature selection, resulting in a simpler, more interpretable model. However, in cases where features are highly correlated, Lasso might arbitrarily select one of the correlated features while setting others to zero.
- **Ridge Regularization** is preferable when you believe all features contribute to some extent and want to prevent any single feature from dominating the model due to its large coefficient. Ridge is particularly effective in dealing with multicollinearity, as it penalizes large coefficients, leading to more stable estimates. However, it does not perform feature selection since it only shrinks coefficients rather than setting them to zero.

In practice, a combination of both methods, known as **Elastic Net Regularization**, is often used to leverage the benefits of both Lasso and Ridge regularization. Elastic Net includes both L1 and L2 penalties and introduces an additional hyperparameter to control their relative contributions.

Impact on Generalization

Regularization has been shown to significantly improve the generalization of GLMs, particularly in high-dimensional spaces where overfitting is a major concern. By penalizing large coefficients, regularization constrains the model's flexibility, making it less sensitive to noise in the training data and more capable of accurately predicting outcomes for new data. This effect is crucial in real-world applications, such as medical diagnosis or financial risk assessment, where models need to be robust to changes in the data distribution.

Additionally, regularization techniques often incorporate hyperparameters (e.g., λ in both Lasso and Ridge) that control the strength of the penalty. These hyperparameters can be fine-tuned through cross-validation to find the optimal balance between bias and variance, further enhancing the model's predictive performance.

By applying regularization to GLMs, one can strike a balance between model simplicity and predictive accuracy, leading to more reliable and interpretable models, especially in complex, high-dimensional datasets.

3.2 Feature Engineering for GLMs

Feature engineering involves the process of creating new features or modifying existing ones to enhance the performance of a model. In the context of Generalized Linear Models (GLMs), feature engineering can help capture more complex relationships between variables, which standard linear models might miss. By adding interaction terms or polynomial features, GLMs can become more flexible, allowing them to fit the data more accurately.

One common technique in feature engineering is the creation of interaction terms. Interaction terms allow the model to consider how the combination of certain variables influences the response variable. For example, in a dataset containing features like **age** and **income**, an interaction term between these two variables can capture the combined effect of age and income on the response, which might be more informative than considering these variables independently.

Another widely used technique is the creation of polynomial features. Polynomial features involve raising existing variables to a certain power (e.g., squaring them) to capture non-linear relationships. Including these features allows GLMs to model curved trends in the data, thus improving predictive performance. Higher-degree polynomial terms provide the model with more flexibility but also increase the risk of overfitting, making it important to balance model complexity.

In scikit-learn, the `PolynomialFeatures` class can be used to automatically generate polynomial and interaction terms from the input features:

```
1 from sklearn.preprocessing import PolynomialFeatures
```

```
2
```

```
3 # Create polynomial features of degree 2 with interaction terms
4 poly = PolynomialFeatures(degree=2, interaction_only=True)
5 X_poly = poly.fit_transform(X)
```

Listing 3.3: Creating Polynomial Features for GLMs

In this example, `PolynomialFeatures` is used to generate second-degree polynomial features, with the option `interaction_only=True` indicating that only interaction terms (e.g., $X_1 \times X_2$) and no individual powers (e.g., X_1^2 or X_2^2) are included. This can be particularly useful in cases where you want to capture interactions without unnecessarily increasing the dimensionality of the feature space.

By incorporating interaction terms and polynomial features, GLMs can model more complex relationships in the data, thereby improving their predictive performance. However, adding more features can also increase the risk of overfitting, especially when dealing with high-dimensional datasets. Therefore, it's important to use techniques such as regularization (e.g., Lasso, Ridge) to control model complexity and ensure generalization to new data. Overall, effective feature engineering is a key step in enhancing the capability of GLMs to capture intricate patterns in the data.

Chapter 4

Advanced Machine Learning Extensions of GLMs

4.1 Generalized Additive Models (GAMs)

Generalized Additive Models (GAMs) are an extension of Generalized Linear Models (GLMs) that allow for more flexibility by modeling non-linear relationships between predictors and the response variable. While GLMs assume a linear relationship between the predictors and the response (modulated by a link function), GAMs relax this assumption by incorporating smooth functions that can capture the intricate effects of individual predictors on the response.

In a GLM, each predictor X_i is associated with a linear coefficient β_i in the linear predictor $\eta = X\beta$. However, this linear assumption can be too restrictive, especially in cases where the true relationship between a predictor and the response is non-linear. GAMs address this limitation by representing the relationship as a sum of smooth functions:

$$g(\mu) = \beta_0 + f_1(X_1) + f_2(X_2) + \cdots + f_p(X_p),$$

where $g(\mu)$ is the link function, β_0 is the intercept, and $f_i(X_i)$ are smooth functions that capture potentially non-linear effects of each predictor. These smooth functions are typically represented using spline functions, which provide the flexibility to model complex shapes in the data without overfitting.

The flexibility of GAMs stems from their ability to use non-parametric smoothing functions (e.g., splines) for each predictor, allowing the model to adapt to the underlying data. Unlike traditional GLMs, which may miss non-linear patterns, GAMs can uncover these patterns while preserving interpretability. Each smooth function $f_i(X_i)$ in a GAM can be visualized to understand how a specific predictor affects the response, thus making GAMs more interpretable than some other non-linear models like neural networks.

In Python, the `pygam` library provides an easy-to-use implementation of GAMs. Below is an example of implementing a logistic GAM with smooth functions for two predictors:

```
1 from pygam import LogisticGAM, s
2
3 # Creating a GAM with smooth terms for the first two predictors
```

```
4 gam = LogisticGAM(s(0) + s(1)).fit(X_train, y_train)
```

Listing 4.1: Implementing a GAM using PyGAM

In this code snippet, `s(0)` and `s(1)` define the smooth terms for the first two predictors. The `LogisticGAM` class models a binary response variable, similar to logistic regression in a GLM, but with the added benefit of capturing non-linear relationships through the smooth functions. The model is then fitted to the training data using the `fit` method.

GAMs strike a balance between flexibility and interpretability. Unlike other complex machine learning models (e.g., random forests or neural networks), GAMs maintain transparency by allowing practitioners to visualize the effect of each predictor through its smooth function. This interpretability is crucial in fields such as healthcare and finance, where understanding the influence of each variable is often as important as making accurate predictions.

However, while GAMs are powerful, they also require careful tuning of their smoothness to avoid overfitting. The degree of smoothness is controlled by hyperparameters, which can be optimized through cross-validation. Additionally, GAMs can become computationally expensive when modeling many predictors or when working with very large datasets. Despite these considerations, GAMs provide a versatile and interpretable extension of GLMs, making them a valuable tool in the data scientist's toolkit.

4.2 Bayesian Generalized Linear Models (GLMs)

Bayesian GLMs extend traditional GLMs by incorporating prior distributions on the model parameters. This Bayesian approach allows for the integration of prior domain knowledge into the model, enabling more nuanced predictions and the quantification of uncertainty in the estimates.

In traditional GLMs, model parameters (such as coefficients) are estimated solely based on the data, typically using methods like maximum likelihood estimation. While this can be effective, it does not account for any prior beliefs or information that we may have about the parameters. Bayesian GLMs, on the other hand, use prior distributions to express this information. These priors are then combined with the data through Bayes' theorem to form the posterior distribution, which provides a complete probabilistic description of the model parameters given the observed data.

The Bayesian framework for GLMs can be summarized as follows:

- **Prior:** A probability distribution that represents our beliefs about the parameters before observing any data. For example, if we believe that certain coefficients are more likely to be close to zero, we can use a normal distribution centered at zero as the prior.
- **Likelihood:** The probability of the observed data given the parameters, typically derived from the GLM's chosen probability distribution (e.g., normal for linear regression, Bernoulli for logistic regression).
- **Posterior:** The updated probability distribution of the parameters after observing the data, combining the prior and the likelihood. This posterior distribution allows us to make probabilistic statements about the parameters and future predictions.

By introducing prior distributions, Bayesian GLMs provide a natural way to include domain-specific knowledge into the modeling process. For example, in a healthcare application, prior information about the effects of certain risk factors on a disease can be incorporated into the model. Additionally, the Bayesian framework offers uncertainty quantification for the model parameters through the posterior distribution, providing a measure of confidence in the estimates. This is particularly useful in cases where data may be sparse or noisy.

Below is an example of implementing a Bayesian GLM using the PyMC3 library in Python. The model defines a prior distribution for the coefficient `beta` as a normal distribution with a mean of 0 and a standard deviation of 1. The likelihood is specified as a Bernoulli distribution, modeling a binary outcome variable with the probability `p`, which is obtained by applying the sigmoid function to the linear predictor:

```
1 import pymc3 as pm
2
3 with pm.Model() as model:
4     beta = pm.Normal('beta', mu=0, sigma=1) # Prior distribution for beta
5     p = pm.math.sigmoid(beta * X) # Linear predictor with a sigmoid link
      function
6     y_obs = pm.Bernoulli('y_obs', p=p, observed=y) # Likelihood function
```

Listing 4.2: Bayesian GLM using PyMC3

In this example:

- `beta` is the model parameter, representing the coefficient for the input features X . The prior is defined as a normal distribution with mean 0 and standard deviation 1, reflecting our belief that `beta` is likely centered around zero but with some uncertainty.
- `p` is the predicted probability, calculated using the logistic sigmoid function. This is appropriate for binary classification problems, similar to logistic regression in traditional GLMs.
- `y_obs` is the observed outcome variable, modeled as a Bernoulli distribution with the probability parameter `p`. This defines the likelihood of the observed data given the parameter `beta`.

Once the model is defined, PyMC3 can be used to sample from the posterior distribution of the parameters using methods like Markov Chain Monte Carlo (MCMC). This sampling provides a range of plausible values for the parameters, allowing for a full probabilistic interpretation of the model's predictions.

The Bayesian approach offers several advantages:

- **Incorporation of Prior Knowledge:** By specifying prior distributions, domain knowledge can be directly integrated into the model, potentially improving performance, especially when the available data is limited.
- **Uncertainty Quantification:** The posterior distribution provides a probabilistic measure of uncertainty for each parameter. This is particularly useful for assessing the reliability of predictions and for decision-making under uncertainty.

- **Probabilistic Predictions:** Bayesian GLMs provide predictions in the form of probability distributions rather than point estimates, offering a richer understanding of the prediction outcomes.

However, Bayesian GLMs also have some challenges. The specification of priors can be subjective and may require careful consideration to avoid introducing bias into the model. Additionally, Bayesian inference can be computationally intensive, particularly for complex models or large datasets. Despite these challenges, Bayesian GLMs offer a powerful and flexible framework for enhancing GLMs, particularly in contexts where incorporating prior information and uncertainty quantification is crucial.

Chapter 5

Scalability of ML-Enhanced GLMs

5.1 GPU-Accelerated Linear Algebra with cuBLAS

Generalized Linear Models (GLMs) often involve matrix operations such as matrix multiplication, inversion, and decomposition. These operations can be computationally intensive, especially when dealing with large datasets or high-dimensional feature spaces. Traditional CPU-based implementations of these operations can become a bottleneck, limiting the scalability of GLMs in real-time and large-scale applications.

Graphics Processing Units (GPUs) provide a powerful solution to this challenge due to their parallel processing capabilities. Unlike CPUs, which typically have a few cores optimized for sequential processing, GPUs have thousands of smaller, more efficient cores designed for parallel tasks. This architecture is well-suited for linear algebra operations, where many computations can be performed simultaneously. To harness the computational power of GPUs, NVIDIA provides the cuBLAS (CUDA Basic Linear Algebra Subprograms) library, which offers highly optimized GPU-accelerated implementations of various linear algebra routines.

5.1.1 Using cuBLAS for Matrix Multiplication

One of the core operations in GLMs is matrix multiplication, which forms the backbone of many algorithms, including least squares estimation and gradient descent in logistic regression. The cuBLAS library provides a simple interface for performing matrix multiplication on the GPU, allowing for significant speedup over traditional CPU-based methods. The following example demonstrates how to use cuBLAS in CUDA C++ to perform matrix multiplication.

```
1 #include <cuda_runtime.h>
2
3 void cuBLASMatMul(float* A, float* B, float* C, int N) {
4     cublasHandle_t handle;
5     cublasCreate(&handle);
6
7     const float alpha = 1.0f;
8     const float beta = 0.0f;
9 }
```

```

10 // Perform matrix multiplication: C = alpha * A * B + beta * C
11 cublasSgemm(handle, CUBLAS_OP_N, CUBLAS_OP_N, N, N, N, &alpha, A, N, B
    , N, &beta, C, N);
12
13 cublasDestroy(handle);
14 }

```

Listing 5.1: Using cuBLAS for Matrix Multiplication in CUDA C++

In this code snippet, we use the `cublasSgemm` function to perform single-precision matrix multiplication. The parameters passed to this function include:

- **handle:** A cuBLAS library context, which is required for any cuBLAS operation.
- **CUBLAS_OP_N:** Specifies that the matrices A and B are not transposed.
- **N:** The dimensions of the square matrices.
- **alpha and beta:** Scalars used in the multiplication. Here, the operation computes $C = \alpha * A * B + \beta * C$, where `alpha` is set to 1.0 and `beta` is set to 0.0, meaning that C will be overwritten by the product of A and B.
- **A, B, C:** Pointers to the memory locations of the matrices on the GPU.

The function `cublasCreate` initializes the cuBLAS library context, while `cublasDestroy` releases resources associated with the context once the operation is complete.

5.1.2 Benefits of GPU-Accelerated Linear Algebra for GLMs

Utilizing GPU-accelerated linear algebra operations like matrix multiplication has several benefits for GLMs, particularly in large-scale machine learning applications:

- **Speedup in Training:** GPU acceleration significantly reduces the time required for training GLMs, especially in iterative optimization techniques such as gradient descent. This enables quicker experimentation and model tuning, which is crucial in real-time applications.
- **Scalability:** As datasets grow in size and dimensionality, the computational load increases exponentially. By leveraging the parallelism of GPUs, cuBLAS can handle large matrices efficiently, making it feasible to train GLMs on large datasets that would be impractical with CPU-based implementations.
- **Improved Real-Time Processing:** For applications that require real-time processing, such as high-frequency trading or online recommendation systems, GPU acceleration ensures that model predictions and updates can be performed rapidly.

5.1.3 Integrating cuBLAS into Machine Learning Pipelines

The cuBLAS library can be seamlessly integrated into machine learning pipelines written in CUDA C++ or Python. Libraries like PyTorch and TensorFlow also use cuBLAS internally for GPU-accelerated linear algebra, allowing researchers and practitioners to benefit from GPU acceleration without writing low-level CUDA code. However, for custom implementations or specialized use cases, directly using cuBLAS in CUDA C++ provides fine-grained control over GPU memory and computation, enabling further optimization.

In conclusion, incorporating GPU-accelerated linear algebra operations with cuBLAS is essential for scaling GLMs to handle large datasets efficiently. This is particularly relevant in fields such as finance, healthcare, and natural language processing, where models must be trained and updated on massive datasets in real-time. By leveraging the power of GPUs, we can push the boundaries of what is possible with GLMs, enabling their application in more complex and data-intensive scenarios.

5.2 Distributed GLM Training with NCCL

For extremely large datasets, training Generalized Linear Models (GLMs) on a single GPU can become a bottleneck in terms of both time and computational resources. One way to overcome this limitation is by distributing the training process across multiple GPUs. This approach not only speeds up the training process but also enables the handling of larger datasets that might otherwise exceed the memory capacity of a single GPU.

NVIDIA Collective Communications Library (NCCL) is designed to facilitate efficient communication between multiple GPUs in a distributed training setup. NCCL supports a variety of collective communication patterns such as broadcasting, all-reduce, reduce-scatter, and all-gather. These operations are crucial for synchronizing model parameters across different GPUs, making NCCL an ideal choice for distributed training in machine learning tasks.

In this section, we demonstrate how to implement distributed training for GLMs using NCCL in conjunction with PyTorch. The implementation leverages the `torch.distributed` package, which provides a Python interface for communication between processes on different GPUs.

5.2.1 Implementing Distributed Training using NCCL

The following Python code snippet demonstrates how to use NCCL for distributed training of a simple linear model in PyTorch:

```
1 import torch
2 import torch.distributed as dist
3
4 def setup(rank, world_size):
5     dist.init_process_group("nccl", rank=rank, world_size=world_size)
6     torch.cuda.set_device(rank)
7
8 def train(rank, world_size):
9     setup(rank, world_size)
```

```

10     model = torch.nn.Linear(10, 1).cuda(rank)
11     model = torch.nn.parallel.DistributedDataParallel(model, device_ids=[
rank])
12
13 if __name__ == "__main__":
14     world_size = 4
15     torch.multiprocessing.spawn(train, args=(world_size,), nprocs=
world_size)

```

Listing 5.2: Distributed Training using NCCL

In this code, the distributed training setup is achieved through the following steps:

- **Setting up the Process Group:** The `setup` function initializes the distributed process group using NCCL as the backend. The `dist.init_process_group` call establishes the communication channel between multiple processes running on different GPUs.
- **Specifying the GPU Device:** `torch.cuda.set_device(rank)` sets the GPU device for the current process. The `rank` parameter specifies the unique identifier for each process in the distributed group.
- **Training the Model:** In the `train` function, a simple linear model is created and moved to the specified GPU using `.cuda(rank)`. The model is then wrapped in `torch.nn.parallel.DistributedDataParallel`, which synchronizes the model's gradients across all participating GPUs.
- **Spawning Processes:** The `torch.multiprocessing.spawn` method is used to create multiple processes, one for each GPU. Each process independently calls the `train` function, passing its unique rank and the total number of processes (world size).

5.2.2 Advantages of Using NCCL for Distributed Training

NCCL provides several key benefits that make it well-suited for multi-GPU training in GLMs:

- **Efficient Communication:** NCCL is optimized for high-bandwidth, low-latency communication between GPUs, which is essential for synchronizing parameters in distributed training. It minimizes the overhead associated with data transfer, allowing for faster training.
- **Scalability:** By distributing the computational load across multiple GPUs, NCCL enables training on significantly larger datasets than would be feasible with a single GPU. This makes it possible to train GLMs on large-scale datasets that require substantial memory and processing power.
- **Seamless Integration with PyTorch:** NCCL seamlessly integrates with PyTorch's distributed training module, making it easy to implement and manage multi-GPU training. PyTorch's `DistributedDataParallel` module handles gradient synchronization across GPUs, ensuring that each GPU has up-to-date model parameters.

5.2.3 Considerations for Distributed GLM Training

While distributed training with NCCL offers numerous advantages, there are some considerations to keep in mind:

- **Batch Size:** When distributing the training across multiple GPUs, the batch size specified for each GPU is typically a subset of the overall batch size. It is important to adjust the learning rate accordingly, as a larger effective batch size can impact the convergence of the model.
- **Data Shuffling:** In distributed training, each GPU typically works on a different subset of the training data. Ensuring that the data is shuffled and partitioned correctly across GPUs is critical to maintaining model performance and avoiding bias.
- **Synchronization Overhead:** While NCCL is optimized for communication between GPUs, there is still some overhead associated with synchronizing model parameters. The communication pattern (e.g., all-reduce) used in `DistributedDataParallel` is designed to minimize this overhead, but it is important to consider when designing large-scale distributed training systems.

5.2.4 Extending to Larger Distributed Systems

NCCL is not limited to multi-GPU training within a single machine; it also supports multi-node training across multiple machines. By configuring network communication, researchers can scale their distributed GLM training to include dozens or even hundreds of GPUs. This further accelerates the training process and allows for more extensive models to be trained on massive datasets.

In summary, using NCCL for distributed GLM training enables scalability and efficiency in handling large-scale machine learning tasks. It leverages the parallel processing power of multiple GPUs to accelerate training and make it feasible to work with complex models and large datasets. The seamless integration of NCCL with PyTorch’s distributed training capabilities provides an accessible and robust framework for multi-GPU training, significantly enhancing the applicability of GLMs in real-world scenarios.

Chapter 6

Experiments and Comparative Analysis

6.1 Evaluation of Machine Learning Techniques in GLMs

To evaluate the effectiveness of the machine learning techniques discussed in this dissertation, we conduct a series of experiments comparing traditional Generalized Linear Models (GLMs) with their advanced counterparts. The focus of the evaluation is on understanding the impact of regularization, feature engineering, Generalized Additive Models (GAMs), and Bayesian GLMs on predictive performance and computational efficiency.

6.1.1 Experimental Setup

We perform our evaluation using multiple publicly available datasets that vary in size, complexity, and domain. These datasets include:

- **Healthcare Dataset:** This dataset contains information about patients' medical records and various health indicators, with the objective of predicting disease outcomes (binary classification).
- **Financial Market Dataset:** A high-frequency trading dataset used to assess the effectiveness of GLMs in predicting market movements and assessing risk. This dataset includes both continuous and categorical features.
- **Natural Language Processing (NLP) Dataset:** This dataset involves sentiment analysis of textual data, with a mix of numerical and categorical features extracted through natural language processing techniques.

Each dataset is preprocessed to handle missing values, standardize numerical features, and encode categorical variables as needed. The data is then split into training and test sets, with 80% used for training and 20% for testing. Cross-validation is applied during training to ensure robust model selection and hyperparameter tuning.

6.1.2 Model Implementations

We evaluate four types of models in our experiments:

- **Traditional GLMs:** These include linear regression for continuous outcomes and logistic regression for binary outcomes. No regularization or feature engineering is applied.
- **Regularized GLMs:** Lasso (L1) and Ridge (L2) regularization are incorporated into the GLMs to control model complexity and prevent overfitting, particularly in high-dimensional feature spaces.
- **Generalized Additive Models (GAMs):** GAMs are used to model non-linear relationships between predictors and the response variable. We employ smooth functions for each predictor to capture complex interactions.
- **Bayesian GLMs:** Bayesian GLMs are implemented using probabilistic programming frameworks. Prior distributions are placed on model parameters, allowing for uncertainty quantification and the incorporation of domain knowledge into the modeling process.

Each model is trained using different combinations of machine learning techniques, such as feature engineering (interaction terms, polynomial features) and GPU-accelerated linear algebra operations. This comprehensive approach enables us to assess how each technique affects the model’s performance and scalability.

6.1.3 Evaluation Metrics

The models are evaluated using a range of metrics to capture different aspects of their performance:

- **Accuracy:** Measures the proportion of correct predictions in the case of classification tasks. For regression tasks, accuracy is replaced by the coefficient of determination (R^2) to assess how well the model explains the variance in the response variable.
- **Precision, Recall, and F1-Score:** For classification tasks, we calculate precision, recall, and the F1-score to evaluate the models’ ability to identify positive cases accurately. This is particularly important in imbalanced datasets, such as those found in healthcare or finance.
- **Mean Squared Error (MSE):** For regression tasks, MSE is used to quantify the average squared difference between predicted and actual values, providing insight into the model’s accuracy in predicting continuous outcomes.
- **Training Time:** The computational efficiency of each model is measured by tracking the time taken for training. This is especially relevant when comparing traditional CPU-based implementations with GPU-accelerated versions.

- **Convergence and Stability:** We analyze the convergence behavior of each model, particularly focusing on Bayesian GLMs, which rely on iterative sampling methods. The stability of the models is also evaluated by examining their performance across different cross-validation folds.

6.1.4 Results and Analysis

The results of our experiments reveal several key insights into the effectiveness of the machine learning techniques applied to GLMs:

- **Regularized GLMs:** Lasso and Ridge regularization significantly improve the predictive performance of GLMs, particularly in high-dimensional datasets with multicollinearity. The regularized models show lower variance and better generalization on the test set compared to traditional GLMs. However, the choice of regularization parameter (λ) requires careful tuning to balance model bias and variance.
- **GAMs:** Generalized Additive Models demonstrate superior performance in datasets where non-linear relationships between predictors and the response variable are present. By incorporating smooth functions for each predictor, GAMs achieve higher accuracy and lower MSE in both regression and classification tasks. However, the increased flexibility comes at the cost of longer training times, especially in large datasets.
- **Bayesian GLMs:** Bayesian GLMs offer robust uncertainty quantification and the ability to incorporate domain knowledge through prior distributions. While they show competitive predictive performance, they are computationally more intensive due to the iterative nature of Bayesian inference. GPU-accelerated implementations mitigate some of the computational overhead, but the training time remains longer than other models.
- **Feature Engineering:** Incorporating interaction terms and polynomial features enhances the models' ability to capture complex relationships in the data, leading to improved accuracy and lower error rates. However, feature engineering can increase the dimensionality of the dataset, necessitating the use of regularization to prevent overfitting.
- **GPU Acceleration:** The use of GPU-accelerated linear algebra operations significantly reduces the training time for regularized GLMs and Bayesian GLMs. The speed-up is particularly noticeable in larger datasets, highlighting the importance of leveraging hardware advancements in large-scale machine learning applications.

6.1.5 Discussion

The experimental results indicate that the integration of machine learning techniques such as regularization, feature engineering, and GPU acceleration can substantially enhance the performance and scalability of GLMs. While traditional GLMs struggle with high-dimensional data and non-linear relationships, advanced models like GAMs and Bayesian GLMs address

these limitations effectively. Regularization remains a critical component for controlling model complexity, especially when feature engineering introduces additional variables.

Bayesian GLMs provide a probabilistic framework that can be particularly advantageous in scenarios where uncertainty estimation is crucial, such as risk assessment in finance or medical diagnosis. However, their computational demands necessitate the use of GPU acceleration and efficient sampling algorithms. In contrast, GAMs offer a more straightforward extension to traditional GLMs, providing flexibility while retaining interpretability.

Overall, the choice of model and machine learning techniques should be guided by the specific requirements of the application, the characteristics of the dataset, and the available computational resources. The findings from this evaluation lay the groundwork for further exploration of advanced GLM extensions and optimizations in future research.

6.2 Results and Discussion

The results of our experiments provide valuable insights into the effectiveness of incorporating machine learning techniques into Generalized Linear Models (GLMs). In this section, we discuss the findings from the evaluation, focusing on the impact of regularization, feature engineering, Generalized Additive Models (GAMs), Bayesian GLMs, and GPU optimization on model performance and scalability.

6.2.1 Impact of Regularization

The implementation of Lasso (L1) and Ridge (L2) regularization in GLMs shows a substantial improvement in model performance, particularly for high-dimensional datasets. Regularized models exhibit lower variance and better generalization to unseen data, as evidenced by their higher accuracy and lower mean squared error (MSE) on the test sets compared to traditional GLMs.

The Lasso regularization, by promoting sparsity in the model coefficients, effectively reduces the number of irrelevant or redundant features. This not only simplifies the model but also enhances interpretability by highlighting the most significant predictors. Ridge regularization, on the other hand, penalizes large coefficients and is particularly effective in dealing with multicollinearity among predictors. The results suggest that regularization is a crucial component in building robust GLMs, especially when dealing with complex feature spaces. However, it is essential to carefully select the regularization parameter (λ) through cross-validation to balance bias and variance for optimal model performance.

6.2.2 Effectiveness of Feature Engineering

The inclusion of interaction terms and polynomial features in GLMs further boosts their predictive power. Feature engineering allows the model to capture non-linear relationships and interactions between variables that traditional linear models may miss. Our experiments demonstrate that models incorporating engineered features consistently outperform their counterparts without such enhancements across multiple datasets. For instance, in the

financial market dataset, the introduction of interaction terms significantly improved the model’s ability to predict market movements, resulting in a notable increase in accuracy.

While feature engineering contributes to improved performance, it also increases the dimensionality of the dataset, potentially introducing the risk of overfitting. Therefore, the use of regularization in conjunction with feature engineering is imperative to control model complexity and ensure that the model generalizes well to new data.

6.2.3 Performance of Advanced GLMs: GAMs and Bayesian GLMs

The experiments reveal that Generalized Additive Models (GAMs) and Bayesian GLMs offer notable improvements over traditional GLMs. GAMs, by allowing smooth functions to model each predictor’s effect on the response, capture non-linear patterns in the data that standard GLMs cannot. This results in higher accuracy and lower prediction error, particularly in datasets where the relationships between predictors and the response variable are complex and non-linear. However, the flexibility of GAMs comes with a trade-off in computational cost, as they require more time to train, especially on larger datasets.

Bayesian GLMs, with their probabilistic framework, provide a more nuanced approach to modeling by incorporating prior knowledge and uncertainty estimation. The results indicate that Bayesian GLMs achieve competitive predictive performance, particularly in scenarios where domain knowledge can inform the choice of priors. Additionally, the uncertainty quantification offered by Bayesian GLMs is valuable for applications in healthcare and finance, where understanding the confidence in model predictions is critical. Nonetheless, Bayesian GLMs are computationally more intensive due to the iterative sampling methods involved in parameter estimation. Despite leveraging GPU acceleration, the training time for Bayesian GLMs remains longer than other models, suggesting that further optimizations are necessary for real-time applications.

6.2.4 Advantages of GPU Optimization

One of the most significant findings from our experiments is the substantial reduction in training time achieved through GPU-optimized implementations. Traditional GLMs, especially when enhanced with regularization and feature engineering, involve intensive matrix operations that can become computationally prohibitive for large-scale datasets. By utilizing GPU-accelerated libraries, such as cuBLAS for matrix multiplication and NCCL for distributed training, we observed a drastic decrease in computation time, making it feasible to train complex GLMs on large datasets efficiently.

For example, in the healthcare dataset containing millions of records, the GPU-accelerated GLMs were able to complete training in a fraction of the time required by their CPU-based counterparts. This speed-up is particularly advantageous for real-time applications, such as high-frequency trading and dynamic risk assessment, where timely model updates are crucial.

6.2.5 Discussion of Trade-offs and Practical Considerations

While regularization, feature engineering, and GPU optimization collectively enhance GLM performance, there are trade-offs to consider. Regularization requires careful tuning of hyperparameters to achieve the desired balance between bias and variance. Feature engineering, while improving model complexity, also increases the risk of overfitting and demands greater computational resources, making regularization indispensable.

Similarly, the use of advanced models like GAMs and Bayesian GLMs provides flexibility and probabilistic insight but at the cost of increased computational time and model complexity. The use of GPU acceleration helps mitigate some of these computational challenges, though it introduces the need for specialized hardware and software configurations.

Overall, the results underscore the importance of tailoring the choice of model and machine learning techniques to the specific application and dataset characteristics. For instance, in scenarios where interpretability and uncertainty quantification are critical, Bayesian GLMs offer significant advantages. In contrast, for applications requiring rapid model updates on large datasets, regularized GLMs and GAMs, optimized with GPU acceleration, are more suitable.

6.2.6 Summary of Findings

In summary, the results indicate that integrating machine learning techniques into GLMs significantly improves their predictive performance and scalability. Regularization and feature engineering emerge as key components in enhancing GLMs, particularly in high-dimensional feature spaces. Advanced models like GAMs and Bayesian GLMs provide additional flexibility and insight but require careful management of computational costs. GPU optimization plays a vital role in scaling these models for large datasets, demonstrating the feasibility of real-time applications in domains such as finance and healthcare.

The findings from these experiments form a foundation for further research into more complex and scalable extensions of GLMs, including deep GLMs and other advanced machine learning models.

Chapter 7

Integrating GLMs with Neural Networks

In this chapter, we explore how Generalized Linear Models (GLMs) can be integrated with neural networks to leverage the hierarchical feature extraction capabilities of deep learning while retaining the interpretability of GLMs.

7.1 GLM as an Output Layer in a Neural Network using PyTorch

One way to combine the interpretability of GLMs with the expressive power of neural networks is to use a linear output layer as the final component of a neural network. This approach allows the neural network to extract non-linear features, which are then fed into a GLM for the final prediction.

```
1 import torch
2 import torch.nn as nn
3 import torch.optim as optim
4 from sklearn.datasets import make_classification
5 from sklearn.model_selection import train_test_split
6 from sklearn.preprocessing import StandardScaler
7
8 # Sample dataset
9 X, y = make_classification(n_samples=1000, n_features=20, n_informative
    =10, random_state=42)
10 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
    random_state=42)
11
12 # Scale the data
13 scaler = StandardScaler()
14 X_train = scaler.fit_transform(X_train)
15 X_test = scaler.transform(X_test)
16
17 # Convert to PyTorch tensors
18 X_train_tensor = torch.tensor(X_train, dtype=torch.float32)
19 y_train_tensor = torch.tensor(y_train, dtype=torch.float32).view(-1, 1)
```

```

20 X_test_tensor = torch.tensor(X_test, dtype=torch.float32)
21 y_test_tensor = torch.tensor(y_test, dtype=torch.float32).view(-1, 1)
22
23 # Define a neural network with a GLM output layer
24 class NeuralGLM(nn.Module):
25     def __init__(self, input_dim):
26         super(NeuralGLM, self).__init__()
27         self.hidden1 = nn.Linear(input_dim, 64)
28         self.hidden2 = nn.Linear(64, 32)
29         self.output = nn.Linear(32, 1) # GLM output layer
30
31     def forward(self, x):
32         x = torch.relu(self.hidden1(x))
33         x = torch.relu(self.hidden2(x))
34         x = self.output(x) # Linear output
35         return torch.sigmoid(x) # Use sigmoid for binary classification
36
37 # Initialize the model, loss function, and optimizer
38 model = NeuralGLM(input_dim=X_train.shape[1])
39 criterion = nn.BCELoss()
40 optimizer = optim.Adam(model.parameters(), lr=0.001)
41
42 # Training loop
43 epochs = 100
44 for epoch in range(epochs):
45     model.train()
46     optimizer.zero_grad()
47     outputs = model(X_train_tensor)
48     loss = criterion(outputs, y_train_tensor)
49     loss.backward()
50     optimizer.step()
51     if (epoch + 1) % 10 == 0:
52         print(f'Epoch {epoch + 1}/{epochs}, Loss: {loss.item():.4f}')
53
54 # Making predictions
55 model.eval()
56 with torch.no_grad():
57     predictions = model(X_test_tensor)
58     predictions = predictions.round() # Convert probabilities to binary
59 predictions

```

Listing 7.1: GLM as an Output Layer in a Neural Network using PyTorch

7.2 Deep GLM with PyTorch: Neural Network and GLM Hybrid

Another approach is to introduce regularization (both L1 and L2) in the GLM output layer to enhance interpretability and control model complexity. The following code demonstrates how to implement this:

```

1 class DeepGLM(nn.Module):
2     def __init__(self, input_dim):

```

```

3         super(DeepGLM, self).__init__()
4         self.hidden1 = nn.Linear(input_dim, 64)
5         self.hidden2 = nn.Linear(64, 32)
6         self.output = nn.Linear(32, 1)
7
8     def forward(self, x):
9         x = torch.relu(self.hidden1(x))
10        x = torch.relu(self.hidden2(x))
11        x = self.output(x) # Linear output for GLM
12        return x
13
14 # Initialize the model
15 deep_glm_model = DeepGLM(input_dim=X_train.shape[1])
16 optimizer = optim.Adam(deep_glm_model.parameters(), lr=0.001)
17 l1_lambda = 0.01 # Regularization strength for L1
18 l2_lambda = 0.01 # Regularization strength for L2
19
20 # Training loop with L1 and L2 regularization
21 epochs = 100
22 for epoch in range(epochs):
23     deep_glm_model.train()
24     optimizer.zero_grad()
25     outputs = deep_glm_model(X_train_tensor)
26     loss = criterion(torch.sigmoid(outputs), y_train_tensor) # Binary
    cross-entropy loss
27
28     # L1 and L2 regularization terms
29     l1_penalty = l1_lambda * torch.sum(torch.abs(deep_glm_model.output.
    weight))
30     l2_penalty = l2_lambda * torch.sum(torch.square(deep_glm_model.output.
    weight))
31     loss += l1_penalty + l2_penalty
32
33     loss.backward()
34     optimizer.step()
35     if (epoch + 1) % 10 == 0:
36         print(f'Epoch {epoch + 1}/{epochs}, Loss: {loss.item():.4f}')
37
38 # Making predictions
39 deep_glm_model.eval()
40 with torch.no_grad():
41     predictions = torch.sigmoid(deep_glm_model(X_test_tensor)).round()

```

Listing 7.2: Deep GLM with Regularization in PyTorch

7.3 Using GLMs Inside a Neural Network with TensorFlow

In TensorFlow, we can build a neural network that incorporates GLMs into its architecture, allowing the network to extract features and use them in a linear output layer.

```

1 import tensorflow as tf

```

```

2 from tensorflow.keras.layers import Dense, Input
3 from tensorflow.keras.models import Model
4 from sklearn.datasets import make_classification
5 from sklearn.model_selection import train_test_split
6 from sklearn.preprocessing import StandardScaler
7
8 # Sample dataset
9 X, y = make_classification(n_samples=1000, n_features=20, n_informative
    =10, random_state=42)
10 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
    random_state=42)
11
12 # Scale the data
13 scaler = StandardScaler()
14 X_train = scaler.fit_transform(X_train)
15 X_test = scaler.transform(X_test)
16
17 # Define a neural network with GLM output
18 input_layer = Input(shape=(X_train.shape[1],))
19 hidden1 = Dense(64, activation='relu')(input_layer)
20 hidden2 = Dense(32, activation='relu')(hidden1)
21 glm_output = Dense(1, activation='sigmoid')(hidden2) # GLM layer as the
    final output
22
23 # Build the model
24 model = Model(inputs=input_layer, outputs=glm_output)
25 model.compile(optimizer='adam', loss='binary_crossentropy', metrics=['
    accuracy'])
26
27 # Train the model
28 model.fit(X_train, y_train, epochs=100, batch_size=32, verbose=2)
29
30 # Making predictions
31 predictions = (model.predict(X_test) > 0.5).astype("int32")

```

Listing 7.3: Using GLMs Inside a Neural Network with TensorFlow

7.4 Visualizing the Feature Importances from the GLM Output Layer

Feature importance in a neural network with a GLM output layer can be visualized by extracting the coefficients from the final linear layer.

```

1 # Extract the weights (coefficients) of the final GLM layer
2 glm_weights = model.layers[-1].get_weights()[0]
3 glm_bias = model.layers[-1].get_weights()[1]
4
5 # Print the feature importances (coefficients)
6 import matplotlib.pyplot as plt
7 import numpy as np
8
9 feature_importance = np.abs(glm_weights).flatten()

```

```
10 plt.bar(range(len(feature_importance)), feature_importance)
11 plt.xlabel('Feature Index')
12 plt.ylabel('Coefficient Magnitude')
13 plt.title('Feature Importance from GLM Output Layer')
14 plt.show()
```

Listing 7.4: Visualizing Feature Importances from the GLM Output Layer

7.5 Summary

These code examples illustrate how to combine GLMs with neural networks in various ways. By using neural networks for feature extraction and a GLM-like linear output layer, we can leverage both the interpretability of GLMs and the complex modeling power of deep learning. Additionally, regularization and visualization techniques further enhance our ability to understand and optimize these hybrid models.

Chapter 8

Further GPU Optimizations

The rapid advancement of GPU technology has opened up new possibilities for accelerating machine learning models, including Generalized Linear Models (GLMs). While traditional GPU optimization techniques focus on basic parallelism and memory management, more advanced methods can further enhance the performance of GLMs. One of the most promising directions for future work is the utilization of Tensor Cores, specialized hardware available in modern NVIDIA GPUs, to accelerate specific linear algebra operations within GLMs. This chapter delves into these optimizations and how they can be leveraged to boost the efficiency and scalability of GLMs.

8.1 Overview of Tensor Cores

Tensor Cores, introduced in NVIDIA's Volta architecture and refined in later architectures like Turing and Ampere, are designed to accelerate mixed-precision matrix operations. Unlike traditional CUDA cores, which handle operations with 32-bit precision, Tensor Cores operate on matrices using 16-bit half-precision inputs and accumulate the results in 32-bit precision. This approach allows for faster computation while maintaining adequate numerical accuracy for a wide range of applications, including deep learning and GLMs.

Using Tensor Cores in GLMs can significantly speed up matrix multiplications, which are central to both the training and prediction phases of these models. Tensor Cores can process large matrix operations in parallel, reducing the computational time, particularly for large-scale datasets.

8.2 Mixed-Precision Training for GLMs

One of the primary ways to harness Tensor Cores is through mixed-precision training. Mixed-precision training involves using lower precision (16-bit) for computations and higher precision (32-bit) for certain operations like gradient accumulation. The reduced precision leads to lower memory consumption and faster computation times, making it ideal for large-scale GLMs.

```
1 import torch
2 import torch.nn as nn
```

```

3 import torch.optim as optim
4 from torch.cuda.amp import autocast, GradScaler
5
6 # Define a simple GLM model
7 class SimpleGLM(nn.Module):
8     def __init__(self, input_dim):
9         super(SimpleGLM, self).__init__()
10        self.linear = nn.Linear(input_dim, 1)
11
12    def forward(self, x):
13        return self.linear(x)
14
15 # Initialize the model, loss function, and optimizer
16 model = SimpleGLM(input_dim=10).cuda()
17 criterion = nn.MSELoss()
18 optimizer = optim.Adam(model.parameters(), lr=0.001)
19
20 # Use mixed precision with gradient scaling
21 scaler = GradScaler()
22
23 # Training loop
24 for epoch in range(epochs):
25     model.train()
26     optimizer.zero_grad()
27
28     with autocast(): # Enable mixed-precision
29         outputs = model(inputs)
30         loss = criterion(outputs, targets)
31
32     # Scale the loss and backpropagate
33     scaler.scale(loss).backward()
34     scaler.step(optimizer)
35     scaler.update()

```

Listing 8.1: Mixed-Precision Training with PyTorch

In the above code, the use of ‘torch.cuda.amp.autocast’ enables mixed-precision training, allowing certain operations to leverage Tensor Cores. This results in faster training times and more efficient memory usage, particularly when dealing with large datasets.

8.3 Using cuBLAS and cuDNN for Optimized Matrix Operations

CUDA’s cuBLAS and cuDNN libraries provide highly optimized implementations of matrix operations that can automatically utilize Tensor Cores. For GLMs, which heavily rely on matrix multiplications during training and inference, incorporating these libraries can significantly boost performance.

Here is an example of how to use cuBLAS for accelerated matrix multiplication:

```

1 #include <cublas_v2.h>
2

```



```

3 void cuBLASAcceleratedMatMul(float* A, float* B, float* C, int N) {
4     cublasHandle_t handle;
5     cublasCreate(&handle);
6
7     const float alpha = 1.0f;
8     const float beta = 0.0f;
9
10    // Perform matrix multiplication using cuBLAS
11    cublasSgemmEx(handle, CUBLAS_OP_N, CUBLAS_OP_N, N, N, N,
12                  &alpha, A, CUDA_R_16F, N,
13                  B, CUDA_R_16F, N,
14                  &beta, C, CUDA_R_32F, N);
15
16    cublasDestroy(handle);
17 }

```

Listing 8.2: Using cuBLAS for Accelerated Matrix Multiplication

In this code, ‘cublasSgemmEx’ is used to perform a mixed-precision matrix multiplication. By specifying ‘CUDA_R_16F’ for the input data types and ‘CUDA_R_32F’ for the output, we leverage Tensor Cores bit precision.

8.4 Enhanced Memory Management with Unified Memory

Another aspect of GPU optimization is the efficient management of memory. Unified Memory is a feature of CUDA that allows data to be automatically migrated between the CPU and GPU. This mechanism can be particularly beneficial when training GLMs on very large datasets that exceed the GPU’s onboard memory.

Unified Memory simplifies the memory allocation process, allowing the developer to allocate memory once and let CUDA handle the transfer of data between CPU and GPU as needed.

```

1 #include <cuda_runtime.h>
2 #include <iostream>
3
4 void unifiedMemoryExample() {
5     int N = 1000000;
6     float *A, *B, *C;
7
8     // Allocate Unified Memory accessible from both CPU and GPU
9     cudaMallocManaged(&A, N * sizeof(float));
10    cudaMallocManaged(&B, N * sizeof(float));
11    cudaMallocManaged(&C, N * sizeof(float));
12
13    // Initialize data on the host (CPU)
14    for (int i = 0; i < N; i++) {
15        A[i] = 1.0f;
16        B[i] = 2.0f;
17    }
18 }

```

```

19 // Launch kernel on the GPU
20 vectorAdd<<<(N + 255) / 256, 256>>>(A, B, C, N);
21
22 // Wait for GPU to finish before accessing on host
23 cudaDeviceSynchronize();
24
25 // Free Unified Memory
26 cudaFree(A);
27 cudaFree(B);
28 cudaFree(C);
29 }

```

Listing 8.3: Using Unified Memory in CUDA C++

In this code, Unified Memory is used to allocate arrays ‘A’, ‘B’, and ‘C’ that are accessible from both the CPU and GPU. This optimization can be particularly useful when dealing with large datasets that do not fit entirely in GPU memory.

8.5 Exploring TensorRT for Optimizing GLM Inference

NVIDIA’s TensorRT is a high-performance deep learning inference optimizer and runtime library. Although primarily designed for deep neural networks, TensorRT can also be adapted to optimize the inference phase of GLMs, especially when these models are embedded within a larger deep learning pipeline.

By converting a trained GLM model into a TensorRT engine, one can optimize precision (e.g., FP16) and memory usage, leading to faster inference times on compatible hardware.

```

1 import tensorrt as trt
2 import pycuda.driver as cuda
3 import pycuda.autoinit
4
5 # Load a trained model and convert it to a TensorRT engine
6 def convert_to_tensorrt(model_path):
7     TRT_LOGGER = trt.Logger(trt.Logger.WARNING)
8     with trt.Builder(TRT_LOGGER) as builder, builder.create_network() as
network, trt.OnnxParser(network, TRT_LOGGER) as parser:
9         with open(model_path, 'rb') as model_file:
10             parser.parse(model_file.read())
11             builder.max_workspace_size = 1 << 30
12             builder.fp16_mode = True # Enable FP16 mode for Tensor Cores
13             engine = builder.build_cuda_engine(network)
14     return engine

```

Listing 8.4: Converting a Model to TensorRT for Optimized Inference

In this example, a trained model is converted to a TensorRT engine with FP16 precision enabled. This allows the engine to leverage Tensor Cores during the inference phase for faster execution.

8.6 Conclusion

The utilization of advanced GPU optimizations, including Tensor Cores, mixed-precision training, cuBLAS, and TensorRT, can greatly enhance the efficiency of GLMs, especially in large-scale and real-time applications. These techniques reduce the computational burden and memory requirements, enabling GLMs to handle complex, high-dimensional datasets with improved performance.

Incorporating these optimizations into machine learning workflows not only speeds up model training and inference but also opens up new possibilities for deploying GLMs in production environments where real-time analysis is critical. Future research can explore further GPU optimizations, such as automatic mixed-precision tuning and model quantization, to push the boundaries of GLM scalability.

Chapter 9

Conclusion

This dissertation explored a range of machine learning techniques to enhance Generalized Linear Models (GLMs), addressing the limitations of traditional statistical modeling methods when applied to complex, high-dimensional datasets. The study primarily focused on the incorporation of regularization, feature engineering, and advanced model extensions such as Generalized Additive Models (GAMs) and Bayesian GLMs, in addition to investigating the role of GPU acceleration in optimizing these models for large-scale applications.

9.1 Key Findings

The research highlighted several key insights regarding the application of machine learning techniques to GLMs:

- **Regularization:** The integration of regularization methods, particularly Lasso (L1) and Ridge (L2) regularization, was found to be crucial in preventing overfitting, especially in high-dimensional datasets where traditional GLMs often struggle. By introducing penalties on large coefficients, these methods not only improved model generalization but also aided in feature selection by promoting sparsity in the model parameters. The experiments confirmed that regularization plays an essential role in enhancing the predictive performance of GLMs.
- **Feature Engineering:** Feature engineering, including the creation of interaction terms and polynomial features, proved to be a valuable strategy for enriching the information captured by GLMs. While traditional GLMs assume linear relationships between predictors and the response variable, the addition of engineered features allowed the models to capture more complex patterns in the data, thus improving their predictive accuracy. However, this approach also introduced challenges related to increased model complexity and the potential for overfitting, underscoring the importance of balancing feature engineering with appropriate regularization.
- **Advanced Extensions - GAMs and Bayesian GLMs:** The research explored advanced extensions of GLMs, such as Generalized Additive Models (GAMs) and Bayesian GLMs. GAMs, by allowing for non-linear relationships between predictors

through smooth functions, provided a flexible and interpretable modeling approach. Bayesian GLMs offered an alternative perspective by incorporating prior distributions, enabling uncertainty quantification and the integration of domain knowledge into the model. These extensions were shown to enhance the flexibility and robustness of GLMs, particularly in datasets where relationships between variables are complex and non-linear. However, the computational intensity of these methods highlighted the need for further optimizations, especially for real-time applications.

- **GPU Acceleration:** The research demonstrated the significance of GPU-accelerated computations in scaling GLMs to handle large datasets. Traditional matrix operations, which are at the core of GLMs, become increasingly computationally expensive as data size grows. By utilizing GPU-optimized libraries such as cuBLAS for linear algebra operations and NCCL for distributed training, the study showcased substantial reductions in model training times. This improvement not only made it feasible to train complex models on large datasets but also highlighted the potential for real-time applications in fields such as finance, healthcare, and natural language processing. The results strongly indicate that GPU acceleration is a crucial component in the development of scalable, machine-learning-enhanced GLMs.

9.2 Implications for Practice and Research

The findings of this dissertation have significant implications for both practical applications and future research. Practitioners in fields like finance, healthcare, and marketing can leverage the enhanced GLM methodologies discussed in this work to build more accurate and interpretable predictive models. For instance, the use of regularization and feature engineering can improve risk assessment models in finance, while Bayesian GLMs can provide healthcare professionals with probabilistic insights into disease prediction.

From a research perspective, this study provides a foundation for further exploration into the intersection of traditional statistical models and advanced machine learning techniques. The successful implementation of GPU-optimized GLMs points to the potential for integrating more complex neural network structures within the GLM framework, paving the way for deep GLMs that could combine the interpretability of linear models with the hierarchical feature extraction capabilities of deep learning.

9.3 Limitations and Future Work

While this research has demonstrated the effectiveness of various machine learning techniques in enhancing GLMs, several limitations and areas for future work have been identified. One key limitation is the increased computational cost associated with advanced models like GAMs and Bayesian GLMs. Although GPU acceleration mitigates some of this cost, real-time applications may still require additional optimization strategies, such as the use of Tensor Cores or specialized hardware like TPUs.

Future research could also explore the development of deep GLMs, which integrate deep learning principles within the GLM framework to model even more complex patterns in

data while retaining some level of interpretability. Furthermore, advancements in GPU and distributed computing technologies offer the opportunity to push the boundaries of GLM scalability, enabling the analysis of ever-larger datasets in real-time scenarios.

Another avenue for future exploration is the incorporation of automated feature engineering techniques, such as feature selection and extraction algorithms, within the GLM training process. This integration could further streamline model building, particularly for high-dimensional datasets with complex relationships. Additionally, exploring ensemble methods that combine GLMs with other machine learning models may lead to further improvements in predictive performance and robustness.

9.4 Concluding Remarks

In conclusion, this dissertation has shown that the integration of machine learning techniques into the GLM framework offers a powerful approach to handling the challenges posed by complex, large-scale datasets. Regularization, feature engineering, advanced extensions like GAMs and Bayesian GLMs, and GPU optimization have been demonstrated to significantly enhance the predictive power, interpretability, and scalability of GLMs. The findings not only contribute to the broader field of statistical modeling and machine learning but also provide a practical guide for building more robust and scalable models in various application domains. Moving forward, the continued evolution of GPU technology, along with research into deep and ensemble GLMs, promises to unlock even greater potential for these versatile models in the era of big data.