

Scalability Document

Scaling Time Series Forecasting to 100,000 Patients

Executive Summary:

This document outlines the architecture and strategy for scaling the patient mobility forecasting system from a single patient to 100,000 patients. The solution leverages distributed computing, cloud infrastructure, and efficient modeling strategies to process millions of data points while maintaining prediction quality and explainability.

Big Data Processing Strategy

Current Scale

- Single patient: ~80K step count records
- 100K patients: ~8 billion step count records
- Daily features: 40+ features × 100K patients = 4M+ feature vectors per day

Proposed Architecture

1.1 Data Storage: AWS S3 Data Lake Structure:

s3://liberdat-data-lake/

raw/

step-counts/

year=2024/month=01/day=01/*.parquet

year=2024/month=01/day=02/*.parquet

clinical/

therapies/*.parquet

side-effects/*.parquet

diagnoses/*.parquet

processed/

daily-aggregates/patient_id=*/

features/patient_id=*/

forecasts/

predictions/date=*

Benefits:- Partitioning: By date and patient_id for efficient queries- Parquet format: Columnar storage reduces I/O by 10x vs JSON- Cost-effective: S3 storage at \$0.023/GB/month- Scalable: Unlimited storage capacity

1.2 Data Processing: Apache Spark on AWS EMR Why Spark:

Distributed processing across 10-100 nodes- In-memory computation for speed- Native support for time series operations- Handles 8B+ records efficiently

EMR Cluster Configuration:

```
```yaml
```

Master Node: 1x m5.xlarge (4 vCPU, 16 GB RAM)

Core Nodes: 10x m5.2xlarge (8 vCPU, 32 GB RAM each)

Total Capacity: 84 vCPU, 336 GB RAM

Cost: ~\$15/hour (spot instances)

Processing Time: ~2 hours for 100K patients

#### \*PySpark Pipeline:

```
```python
```

Pseudo-code for distributed processing

```
from pyspark.sql import SparkSession
```

```
from pyspark.sql.window import Window
```

```
import pyspark.sql.functions as F
```

```
spark = SparkSession.builder.appName("MobilityForecasting").getOrCreate()
```

```
# Load data (distributed across cluster)
```

```
step_counts = spark.read.parquet("s3://liberdat-data-lake/raw/step-counts/")
```

```
clinical = spark.read.parquet("s3://liberdat-data-lake/raw/clinical/")
```

```
# Aggregate to daily (parallel across patients)
```

```
daily_steps = (step_counts
```

```
    .groupBy("patient_id", F.to_date("timestamp").alias("date"))
```

```
    .agg(F.sum("count").alias("daily_steps"))
```

```
)
```

```

# Feature engineering (distributed)

window_7d = Window.partitionBy("patient_id").orderBy("date").rowsBetween(-7, -1)
window_30d = Window.partitionBy("patient_id").orderBy("date").rowsBetween(-30, -1)

features = (daily_steps
    .withColumn("steps_t_minus_1", F.lag("daily_steps", 1).over(window_spec))
    .withColumn("rolling_avg_7d", F.avg("daily_steps").over(window_7d))
    .withColumn("rolling_avg_30d", F.avg("daily_steps").over(window_30d))
    # ... more features
)

# Join with clinical data (broadcast join for small tables)
final_features = features.join(F.broadcast(clinical), "patient_id")

# Save partitioned by patient
final_features.write.partitionBy("patient_id").parquet("s3://liberdat-data-lake/processed/features/")
...

```

Performance:

- **Throughput:** Process 100K patients in 1-2 hours
- **Cost:** ~\$30 per full pipeline run
- **Scalability:** Linear scaling to 1M+ patients

1.3 Alternative: AWS Glue

When to use:

- Serverless ETL without cluster management
- Scheduled batch jobs (daily/weekly)
- Lower complexity for smaller teams

Configuration:

```
```yaml
```

DPU (Data Processing Units): 50

Cost: \$0.44/DPU-hour

Processing Time: ~3 hours for 100K patients

Total Cost: ~\$66 per run

```
```
```

Trade-off: Simpler but more expensive than EMR for large-scale processing.

1.4 Data Querying: AWS Athena + LakeFormation

Purpose: Ad-hoc queries on data lake without loading into database

Example Query:

```
```sql
```

SELECT patient\_id, AVG(daily\_steps) as avg\_steps

FROM "liberdat\_db"."daily\_aggregates"

WHERE date BETWEEN '2024-01-01' AND '2024-12-31'

AND patient\_id IN (SELECT patient\_id FROM high\_risk\_patients)

GROUP BY patient\_id;

```
```
```

Benefits:

- Query S3 directly (no data movement)
- Pay per query (\$5 per TB scanned)
- Integrate with BI tools (Tableau, QuickSight)

```
---
```

2. Modeling Strategy

2.1 Global Model vs. Clustered Models

Option A: Single Global Model

Approach: Train one model on all 100K patients

Pros:

- Simpler deployment (one model to serve)
- Learns cross-patient patterns
- Requires less infrastructure

Cons:

- May not capture patient-specific patterns
- Difficult to handle high variability
- Large model size (memory intensive)

Option B: Clustered Models (RECOMMENDED)

Approach: Cluster patients by disease type, then train separate models

Clustering Strategy:

...

Cluster 1: Multiple Sclerosis (30K patients) → Model_MS

Cluster 2: Parkinson's Disease (25K patients) → Model_PD

Cluster 3: Rheumatoid Arthritis (20K patients) → Model_RA

Cluster 4: Other Conditions (25K patients) → Model_Other

...

Pros:

- Disease-specific patterns captured
- Better prediction accuracy per cluster
- Parallel training (faster)

- Easier to explain (disease-specific insights)

Cons:

- Multiple models to maintain
- Requires clustering logic in serving layer

Justification: Healthcare data is highly heterogeneous. Different diseases have different mobility patterns. Clustered models provide better accuracy and interpretability.

2.2 Handling Variability in Step Count Baselines

Challenge: Patients have vastly different baseline activity levels:

- Patient A: 10,000 steps/day (active)
- Patient B: 2,000 steps/day (sedentary)

Solution: Patient-Specific Normalization

Approach 1: Z-Score Normalization (Per Patient)

```
```python
For each patient
patient_mean = patient_data['daily_steps'].mean()
patient_std = patient_data['daily_steps'].std()

patient_data['normalized_steps'] = (
 (patient_data['daily_steps'] - patient_mean) / patient_std
)

Train model on normalized values
At inference, denormalize predictions
```
```

Benefits:

- Removes baseline differences
- Model learns relative changes (not absolute values)
- Improves generalization

Approach 2: Patient Offset Features

```
```python
Add patient-specific features

features['patient_baseline_avg'] = patient_mean
features['patient_baseline_std'] = patient_std
features['deviation_from_baseline'] = (
 features['daily_steps'] - features['patient_baseline_avg']
)
```
```

Benefits:

- Model learns both absolute and relative patterns
- Preserves patient-specific context
- Better for personalized predictions

Recommendation: Use Approach 2 (patient offset features) for better interpretability.

2.3 Distributed Hyperparameter Tuning

Challenge: Finding optimal hyperparameters for each cluster model

Solution: Ray Tune on AWS

```
```python
from ray import tune
from ray.tune.schedulers import ASHAScheduler
```

```

def train_ebm(config):
 model = ExplainableBoostingRegressor(
 max_bins=config["max_bins"],
 learning_rate=config["learning_rate"],
 max_leaves=config["max_leaves"]
)
 model.fit(X_train, y_train)
 score = model.score(X_val, y_val)
 return {"score": score}

Distributed hyperparameter search
analysis = tune.run(
 train_ebm,
 config={
 "max_bins": tune.choice([128, 256, 512]),
 "learning_rate": tune.loguniform(0.001, 0.1),
 "max_leaves": tune.choice([2, 3, 5])
 },
 num_samples=50,
 scheduler=ASHAScheduler(),
 resources_per_trial={"cpu": 4, "gpu": 0}
)
...

```

#### Performance:

- **Parallel trials:** 10-20 simultaneous experiments
- **Time:** 2-4 hours for 50 trials
- **Cost:** ~\$20 on spot instances

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## ## 3. Pipeline Architecture

### ### 3.1 End-to-End Workflow

```
```mermaid
```

```
graph LR
```

```
A[Data Sources] --> B[S3 Raw Data Lake]
```

```
B --> C[AWS Glue/EMR ETL]
```

```
C --> D[S3 Processed Features]
```

```
D --> E[Model Training - SageMaker]
```

```
E --> F[Model Registry]
```

```
F --> G[SageMaker Endpoint]
```

```
G --> H[API Gateway]
```

```
H --> I[Client Applications]
```

```
J[Monitoring - CloudWatch] --> E
```

```
J --> G
```

```
```
```

### ### 3.2 Component Details

#### #### Ingestion Layer

- **Source:** Mobile devices, EHR systems
- **Frequency:** Real-time streaming (Kinesis) or batch (S3)
- **Validation:** Schema validation, data quality checks
- **Storage:** S3 with lifecycle policies (archive old data to Glacier)

#### #### Feature Engineering at Scale

- **Tool:** PySpark on EMR or AWS Glue
- **Frequency:** Daily batch jobs

- **Output:** Partitioned Parquet files in S3
- **Optimization:**
  - Broadcast joins for small tables
  - Partition pruning for efficient reads
  - Caching intermediate results

#### #### Model Training

- **Platform:** AWS SageMaker
- **Frequency:** Weekly retraining
- **Parallelization:** Train 4 cluster models simultaneously
- **Instance:** ml.m5.4xlarge (16 vCPU, 64 GB RAM)
- **Time:** 2-3 hours per cluster model
- **Cost:** ~\$10 per model training

#### #### Model Deployment

- **Platform:** SageMaker Endpoints
- **Configuration:**
  - **Instance:** ml.m5.xlarge (4 vCPU, 16 GB RAM)
  - **Auto-scaling:** 2-10 instances based on load
  - **Latency:** <100ms per prediction
  - **Throughput:** 1000 predictions/second per instance

#### #### Serving Predictions for Millions of Daily Data Points

**Challenge:** Generate forecasts for 100K patients daily

**Solution:** Batch Inference

```
```python
# SageMaker Batch Transform
from sagemaker.transformer import Transformer
```

```

transformer = Transformer(
    model_name='mobility-forecast-model',
    instance_count=10,
    instance_type='ml.m5.2xlarge',
    output_path='s3://liberdat-forecasts/predictions/'
)

# Process 100K patients in parallel
transformer.transform(
    data='s3://liberdat-data-lake/processed/features/',
    content_type='text/csv',
    split_type='Line'
)
...

```

Performance:

- **Throughput:** 10K patients/hour
- **Time:** 10 hours for 100K patients
- **Cost:** ~\$20 per batch run

Alternative: Real-Time Inference

- For on-demand predictions (e.g., patient portal)
- SageMaker Endpoint with auto-scaling
- Latency: <100ms per patient
- Cost: \$0.10 per 1000 predictions

3.3 Monitoring and Maintenance

Model Performance Monitoring

- **Metrics:** RMSE, MAE, prediction drift

- **Frequency:** Daily
- **Alerts:** CloudWatch alarms if metrics degrade >10%
- **Action:** Trigger retraining pipeline

Data Quality Monitoring

- **Checks:** Missing data, outliers, schema changes
- **Tool:** AWS Glue DataBrew or Great Expectations
- **Frequency:** Real-time for streaming, daily for batch

Cost Monitoring

- **Budget:** \$500/month for 100K patients
- **Breakdown:**
 - Storage (S3): \$50/month
 - Processing (EMR): \$200/month
 - Training (SageMaker): \$100/month
 - Inference (SageMaker): \$150/month

4. Performance Estimates

Metric	Single Patient	100K Patients	Scaling Factor
	-----	-----	-----
Data Volume	80K records	8B records	100,000x
Processing Time	5 minutes	2 hours	24x (distributed)
Storage	10 MB	1 TB	100,000x
Training Time	10 minutes	3 hours	18x (clustered)
Inference Time	1 second	10 hours (batch)	-
Monthly Cost	\$0 (local)	\$500 (cloud)	-

5. Recommendations

1. **Start with EMR + S3:** Most cost-effective for large-scale processing
2. **Use clustered models:** Better accuracy and interpretability
3. **Implement patient normalization:** Handle baseline variability
4. **Batch inference for forecasts:** Daily batch jobs for all patients
5. **Real-time endpoints for on-demand:** Patient portal, clinical dashboards
6. **Monitor continuously:** Data quality, model performance, costs
7. **Automate retraining:** Weekly retraining to capture new patterns

Conclusion

Scaling to 100,000 patients is achievable with modern cloud infrastructure and distributed computing. The proposed architecture leverages AWS services (S3, EMR, SageMaker) to process billions of records, train clustered models, and serve millions of predictions efficiently. The total cost of ~\$500/month is reasonable for a production healthcare analytics system serving 100K patients.