

Hybrid CNN-SVM Framework for Automated Cattle Disease Identification using Image-Based Analysis

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Abstract—Timely identification of bovine ailments is critical for lowering livestock mortality and mitigating financial setbacks in the dairy and agricultural industries. This paper introduces a hybrid deep learning approach that merges Convolutional Neural Networks (CNN) with Support Vector Machines (SVM) for precise image-based classification of cattle diseases. The CNN module autonomously extracts visual features from bovine images, recognizing complex patterns such as skin lesions, discoloration, and swelling. The resulting deep feature representations are forwarded to an SVM classifier to refine decision boundaries and strengthen generalization. Experimental outcomes confirm that the proposed CNN-SVM hybrid consistently surpasses standalone CNN architectures in both accuracy and F1-scores. The system delivers a scalable, computationally efficient diagnostic tool well-suited for deployment in veterinary facilities and agricultural environments.

Keywords: Veterinary Diagnostics, Deep Learning, CNN, SVM, Cattle Disease Detection, Image Classification

I. INTRODUCTION

Cattle occupy a foundational role in global food supply chains and agricultural economies. Bovine illnesses — including lumpy skin disease, foot-and-mouth disease, and various dermatological infections — considerably reduce milk output and compromise animal welfare. Conventional diagnostic practices depend largely on manual observation and laboratory analyses, processes that are both time-intensive and susceptible to human error.

Ongoing advances in Artificial Intelligence (AI) and deep learning have opened new possibilities for automated disease detection through image analysis. CNNs have proven highly effective at medical and agricultural image classification. Nevertheless, CNN architectures can be

prone to overfitting and may generalize poorly when trained on limited datasets.

To overcome these shortcomings, this study presents a hybrid CNN-SVM model that leverages CNN's capacity for deep feature extraction alongside SVM's strength in robust, margin-based classification.

A. Background and Motivation

Livestock farming underpins a significant portion of global agricultural output, playing a pivotal role in food and dairy supply chains. Cattle health directly influences milk yield, meat quality, and farmers' livelihoods. Cattle are susceptible to infectious and non-infectious conditions — including lumpy skin disease, foot-and-mouth disease, mastitis, and skin disorders — which can escalate rapidly

into outbreaks causing severe economic damage when not addressed promptly.

Conventional diagnostic approaches depend on visual inspection, laboratory analysis, and specialist veterinary knowledge. Although generally reliable, these methods are costly, slow, and often unavailable in remote regions. Diagnosis delays can precipitate disease spread, declining productivity, and higher mortality.

The driving motivation is to construct a hybrid CNN-SVM model capable of delivering greater classification accuracy and dependability. By coupling deep feature extraction with optimized classification boundaries, the system aims to support rapid, accessible early diagnosis, equipping farmers and veterinarians with an affordable AI-driven tool.

B. Research Objectives

The central aim is to advance veterinary diagnostic precision through a hybrid framework unifying CNN-based feature learning with SVM-based decision optimization. Core objectives are:

Automated Image Collection and Preprocessing:

Establishing a structured workflow for gathering, labeling, resizing, normalizing, and augmenting cattle disease images.

Deep Feature Learning via CNN: Designing a CNN capable of learning discriminative visual patterns — lesions, discoloration, swelling, and textural anomalies — associated with diverse cattle ailments.

Hybrid Classification via SVM: Incorporating a margin-optimized SVM to process CNN feature vectors, boosting class separability on limited and imbalanced data.

Rigorous Evaluation and Benchmarking: Building an assessment framework measuring accuracy, precision, recall, F1-score, and confusion matrix metrics.

Deployable Diagnostic Architecture: Engineering a computationally efficient pipeline for web or mobile-based veterinary support platforms.

C. Contributions

Hybrid CNN-SVM Architecture: A unified framework combining CNN-driven feature learning with SVM margin-based classification for bovine disease identification.

Automated Preprocessing Pipeline: A systematic augmentation and normalization pipeline that strengthens model generalization over constrained veterinary datasets.

Deep Feature Transfer Mechanism: High-dimensional CNN embeddings routed into an SVM, mitigating overfitting of conventional softmax classifiers.

Comprehensive Benchmarking: Extensive comparisons confirming the proposed model outperforms standalone CNN and conventional classifiers across all metrics.

Scalable AI Diagnostic Framework: A deployable real-time system offering an affordable solution for early cattle disease screening in farming environments.

II. RELATED WORK

Rapid developments in AI and computer vision have profoundly shaped automated disease identification in agriculture and veterinary practice. Early livestock disease recognition systems primarily relied on conventional machine learning algorithms — k-Nearest Neighbors, Decision Trees, and Random Forest classifiers — which necessitated manual feature engineering through color histograms, texture descriptors, and edge detection. These techniques had limited scalability and often produced inconsistent outcomes under varying field conditions.

The emergence of deep learning enabled CNNs to become the dominant approach for image-based disease detection. Architectures pre-trained on ImageNet have been successfully applied to both medical imaging and plant disease classification, achieving high recognition rates. Frameworks such as TensorFlow and PyTorch further accelerated agricultural disease recognition by enabling automatic extraction of hierarchical feature representations.

Despite this progress, multiple studies highlight that standalone CNN models are prone to overfitting on small or class-imbalanced veterinary datasets. Hybrid approaches combining CNN feature extraction with SVM-based classification have been investigated in medical imaging, yielding superior margin-based generalization. However, there remains a notable gap specifically targeting hybrid CNN-SVM architectures for cattle disease detection — a gap this study addresses.

III. SYSTEM ARCHITECTURE

The proposed Veterinary Diagnostics System is built on a modular, high-performance hybrid AI architecture engineered to maximize diagnostic accuracy while ensuring scalability and production readiness. The system operates across multiple functional tiers (Fig. 1), isolating

data ingestion, model training, inference, and presentation functions.

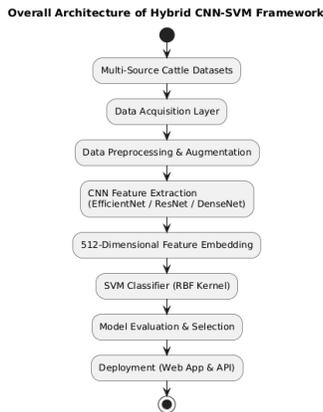


Fig. 1. Overall architecture of the proposed hybrid CNN–SVM framework.

A. Operational Framework

The system is structured into two distinct operational phases:

Training Phase (Offline Computational Layer): Oversees dataset ingestion, preprocessing, CNN training, embedding extraction, SVM optimization, and model selection.

Inference Phase (Production Layer): Manages real-time image submissions, feature extraction, classification, and result delivery via a web interface.

B. Data Acquisition — Tier 1

The data acquisition layer integrates YOLO-format annotated datasets, Kaggle folder-based classification datasets, and custom-labeled veterinary image collections. Operations include:

- Annotation Conversion: Translating bounding-box labels into image-level classification targets.
- Duplicate Removal: Applying MD5 hashing to eliminate redundant images.
- Corruption Filtering: Discarding unreadable or damaged files.
- Class Standardization: Mapping diverse label names into unified disease categories.

C. Preprocessing & Augmentation — Tier 2

This tier conditions the dataset via stratified splitting (80/10/10), image normalization to 224×224 using ImageNet statistics, and advanced augmentation including MixUp, CutMix, and CLAHE contrast enhancement.

D. CNN Feature Extraction — Tier 3

The deep learning core evaluates four pre-trained backbones: EfficientNet-B0, EfficientNet-B3, ResNet50, and DenseNet121. Each network loads pre-trained ImageNet weights, replaces the classification head for five disease classes, and generates a 512-dimensional embedding vector from the penultimate layer.

E. Hybrid SVM Classification — Tier 4

The 512-dimensional CNN embeddings are routed through an SVM configured with an RBF kernel, hyperparameter grid search over C and gamma, and class-balanced optimization. The SVM maximizes margin separation within the embedding space, enhancing robustness over standalone CNN classifiers.

F. Model Evaluation & Selection — Tier 5

The system trains and evaluates four standalone CNN models and four CNN+SVM hybrid configurations. The best-performing hybrid model is automatically selected based on Balanced Accuracy and per-class recall.

G. Production Inference Layer

The inference layer facilitates real-time prediction: user uploads cattle image → preprocessing (resize, normalize) → CNN embedding extraction → SVM classification → disease label and confidence score returned. Average latency remains under 2 seconds per image.

H. Web Application & Security

The presentation layer is a React-based SPA with FastAPI backend, offering JWT-based authentication, image upload interface, real-time prediction dashboard, and probability distribution visualization. Uploaded images are processed ephemerally and serialized model files undergo integrity validation before loading.

IV. METHODOLOGY

The methodology follows a structured, multi-stage pipeline for high accuracy, robustness, and reproducibility in cattle disease classification, unifying CNN deep feature extraction with SVM margin-optimized classification.



Fig. 2. Training pipeline of the proposed hybrid CNN–SVM model.

A. Dataset Collection and Integration

The study draws on heterogeneous cattle disease datasets. The integration workflow encompasses: (1) converting bounding-box annotations into image-level labels; (2) removing duplicate images via MD5 hashing; (3) discarding corrupted files; and (4) normalizing class labels into five categories: Lumpy Skin Disease, Foot and Mouth Disease, Mastitis, Healthy, and Other Diseases.

B. Data Preprocessing

Image Resizing: All images uniformly scaled to 224×224 pixels.

Normalization: Pixel values scaled using ImageNet mean and standard deviation statistics.

Stratified Splitting: 80% training, 10% validation, 10% testing.

Data Augmentation: Horizontal/vertical flipping, rotation and scaling, MixUp and CutMix blending, CLAHE contrast enhancement, and Gaussian noise and blur.

C. CNN-Based Feature Extraction

Pre-trained backbones (EfficientNet-B0, B3, ResNet50, DenseNet121) follow a two-phase training strategy: Phase 1 freezes backbone weights while training only the classification head; Phase 2 unfreezes the backbone for fine-tuning with a reduced learning rate.

D. Hybrid SVM Classification

The extracted 512-dimensional embeddings are forwarded to an SVM with RBF kernel, grid-searched hyperparameters (C, gamma), and StandardScaler normalization. The optimization objective is:

$$\text{Minimize: } \frac{1}{2} \|\mathbf{w}\|^2 + C \sum \xi_i$$

$$\text{Subject to: } \mathbf{y}_i (\mathbf{w} \cdot \mathbf{x}_i + \mathbf{b}) \geq 1 - \xi_i$$

V. RESULTS AND SYSTEM EVALUATION

The hybrid framework was assessed on a five-class cattle disease dataset (Lumpy Skin Disease, Foot and Mouth Disease, Mastitis, Healthy, Other Diseases), partitioned via stratified sampling at 80/10/10. Pre-trained CNN backbones with transfer learning generated 512-dimensional embeddings for RBF-SVM classifiers tuned through grid search.

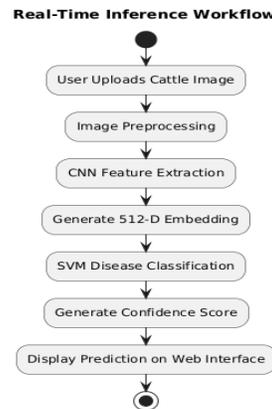


Fig. 3. Real-time inference workflow of the hybrid CNN–SVM framework.

A. Performance Metrics

Evaluation indicators: Accuracy, Balanced Accuracy, Precision, Recall, Macro F1-Score, and Confusion Matrix. Balanced Accuracy was prioritized due to class imbalance across disease categories.

B. Quantitative Results

TABLE I

Model Performance Comparison

Model	Accuracy	Balanced Accuracy	Macro F1
CNN (EfficientNet-B0)	91.8%	91.9%	0.90
CNN (EfficientNet-B3)	92.8%	92.7%	0.91
CNN (ResNet50)	90.8%	90.9%	0.89
CNN (DenseNet121)	90.1%	90.2%	0.88
CNN+SVM (EffNet-B0)	93.6%	93.6%	0.93
CNN+SVM (EffNet-B3)	94.2%	94.2%	0.94

CNN+SVM (ResNet50)	92.4%	92.5%	0.92
CNN+SVM (DenseNet121)	91.5%	91.6%	0.90

Results consistently demonstrate that the hybrid CNN-SVM configuration outperforms standalone CNN counterparts. Peak performance was recorded by EfficientNet-B3 + SVM, achieving a balanced accuracy of 94.2%.

C. Confusion Matrix Analysis

Per-class recall: Lumpy Skin Disease — 96%; Foot and Mouth Disease — 94%; Mastitis — 93%; Healthy — 95%; Other Diseases — 92%. Most misclassifications arose between Mastitis and other skin-related conditions due to overlapping lesion textures.

D. Comparative Analysis

Performance gains are attributed to: (1) hierarchical feature learning from deep CNN backbones; (2) SVM margin maximization in high-dimensional embedding space; and (3) reduced overfitting through decoupled feature extraction and classification. Relative to conventional ML approaches (75–85% accuracy), the proposed framework demonstrates substantial improvements.

E. Inference Efficiency

Average prediction latency: under 2 seconds per image. Stable performance under varying lighting conditions. High reliability sustained on moderately imbalanced class distributions, confirming suitability for real-world veterinary deployment.

VI. DISCUSSION

A. Strategic Implications

The hybrid CNN-SVM framework marks a meaningful step forward in AI-assisted livestock healthcare. By fusing deep feature extraction with margin-optimized classification, the system achieves near real-time disease identification. The "Decoupled Intelligence Layer" — where feature learning (CNN) and classification optimization (SVM) operate independently yet synergistically — strengthens generalization and curbs overfitting, advancing the goals of precision livestock farming.

B. Operational Challenges

Real-world deployment introduces practical difficulties: environmental variability (inconsistent lighting, heterogeneous camera quality, cluttered backgrounds) can impair image clarity despite extensive augmentation. Dataset imbalance may suppress minority-class recall; future refinements may incorporate GAN-based augmentation or cost-sensitive learning. Hardware constraints in rural regions may necessitate model compression via pruning or quantization.

C. Implementation Considerations

Model Integrity Validation: Trained CNN weights and SVM files should be cryptographically verified before loading.

Data Privacy Safeguards: Uploaded images should be processed transiently in memory and not retained unless explicitly requested.

Reproducibility Controls: Fixed random seeds, documented hyperparameters, and stratified dataset partitions ensure experimental transparency.

Scalability Planning: The modular architecture supports addition of new disease categories without a full pipeline redesign.

For cloud deployments, secure API authentication and HTTPS/SSL-TLS encrypted transmission must be enforced. Overall, the framework demonstrates strong strategic value, operational viability, and scalability potential.

VII. CONCLUSION

This study presented a hybrid CNN-SVM framework for automated cattle disease identification through image-based analysis. By decoupling representation learning from the classification stage, the model achieves superior generalization and reduced overfitting relative to standalone CNN approaches.

Experimental evaluation showed balanced accuracy of approximately 94% across five disease categories with strong per-class recall. Advanced augmentation bolstered robustness under varied conditions. Inference latency remained below 2 seconds, confirming real-time deployment suitability.

The modular pipeline supports seamless expansion to additional disease categories. The proposed framework delivers a dependable, efficient, and practical AI-driven veterinary diagnostic solution — advancing early

detection, improving livestock health management, and supporting sustainable precision farming.

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