

The Problem: Drug Toxicity Is the #1 R&D Barrier

- 90% overall clinical failure rate; toxicity is a leading cause of attrition (preclinical to Phase I)
 - \$2.6B average cost to bring a drug to market; 10-12 years timeline
 - Late-stage toxicity discoveries cause costly trial terminations and portfolio delays
 - Complex, non-linear biological interactions are hard to capture with classical models alone
 - Imbalanced and noisy datasets reduce signal detection in early screening
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The Solution: Quantum Machine Learning f

- End-to-end QML pipeline leveraging **QVC** (Variational Classifier) and **QSVM** (Quantum SVM)
 - Rich quantum feature embeddings + fidelity-based kernels for complex interaction capture
 - Balanced training via **BorderlineSMOTE** to address minority toxic class
 - NISQ-ready** design: dimensionality reduction to **6 qubits** while preserving variance
 - Transparent model selection and reporting for regulatory-friendly evidence
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Technology Stack

- Quantum: Qiskit 1.2.4, AerSimulator, Sampler; **ZZFeatureMap** (reps=3), **TwoLocal** ansatz
 - ML & Data: scikit-learn (PCA, feature selection, metrics), imbalanced-learn (BorderlineSMOTE)
 - Optimization: COBYLA (stable for noisy landscapes), Optuna-ready for HPO
 - Engineering: Python/Colab, reproducible artifacts (models, preprocessing, results .pkl/.csv)
 - Governance: Deterministic splits, stratified sampling, robust scaling, exportable reports
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Key Features

- Advanced preprocessing:** Variance thresholding, correlation filtering, mutual information, and feature importance
 - Class imbalance handling:** BorderlineSMOTE for improved minority class detection and balance
 - Quantum-ready scaling:** RobustScaler + MinMax to $[0, 2\pi]$ range for optimal quantum circuit performance
 - Dimensionality reduction:** PCA to 6 components (qubits) with variance tracking for NISQ compatibility
 - Comprehensive evaluation:** Accuracy, Precision, Recall, F1, MCC metrics + confusion matrix analysis
 - Model persistence:** Saved pipelines and artifacts for seamless deployment and reproducibility
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Model Architecture: Quantum Circuits

Data-to-Circuit Flow

- Selected features (**20 features** after importance ranking)
- Robust scaling + MinMax normalization (**0 to 2π** range)
- Class balancing with BorderlineSMOTE (**minority class augmentation**)
- Dimensionality reduction with PCA (**6 components = 6 qubits**)
- Quantum circuits (encoding data into quantum states)

Quantum Algorithms

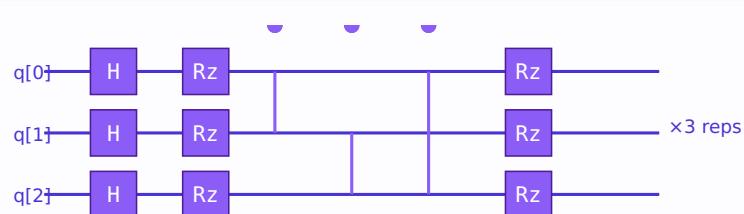
QVC

ZZFeatureMap → TwoLocal ansatz → COBYLA optimizer

QSVM

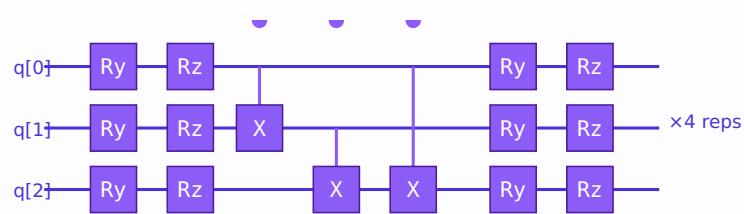
FidelityQuantumKernel with ZZFeatureMap → margin maximization

Feature Map (ZZFeatureMap, reps=3)



ZZFeatureMap with full entanglement

TwoLocal Ansatz (ry/rz + cx, reps=4)



TwoLocal ansatz with rotation + entangling layers

Fidelity Kernel (QSVM)

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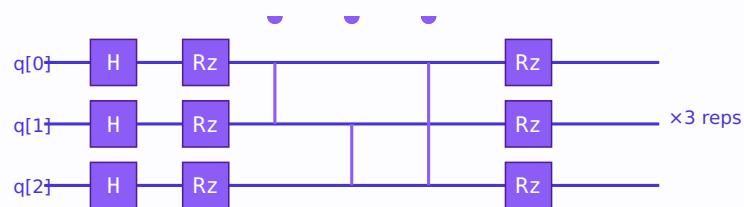
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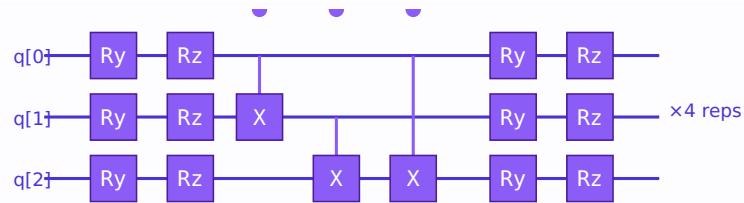
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Results & Performance

Model Comparison Metrics

Key Performance Highlights:

Best Model: QSVM (MCC: 0.734)

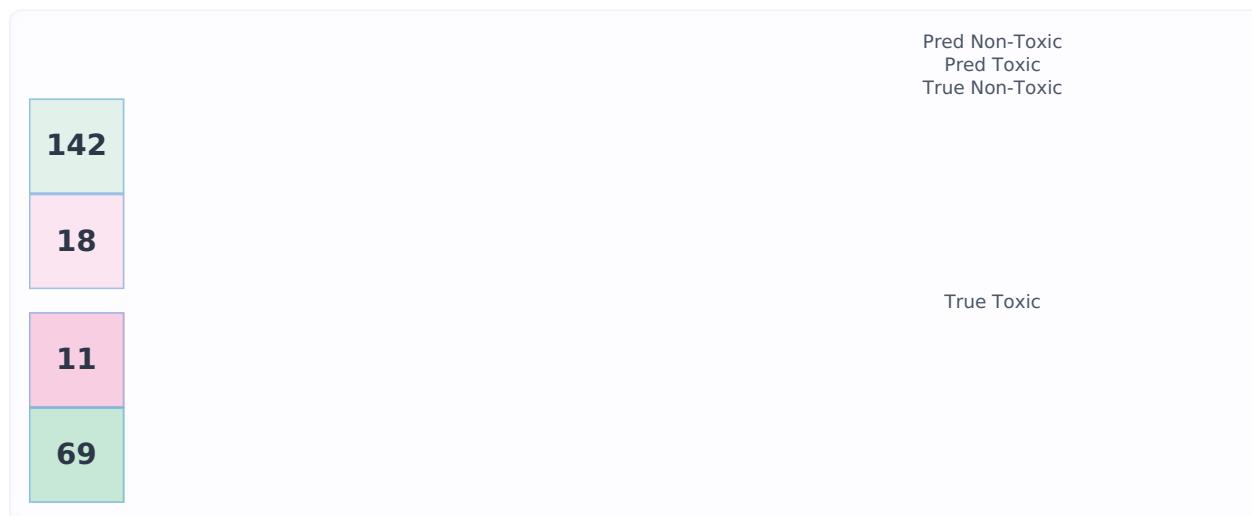
Highest Precision: QSVM (0.841)

Best F1-Score: QSVM (0.852)

MCC is prioritized for class-imbalanced toxicity datasets, as it balances true positive/negative rates.

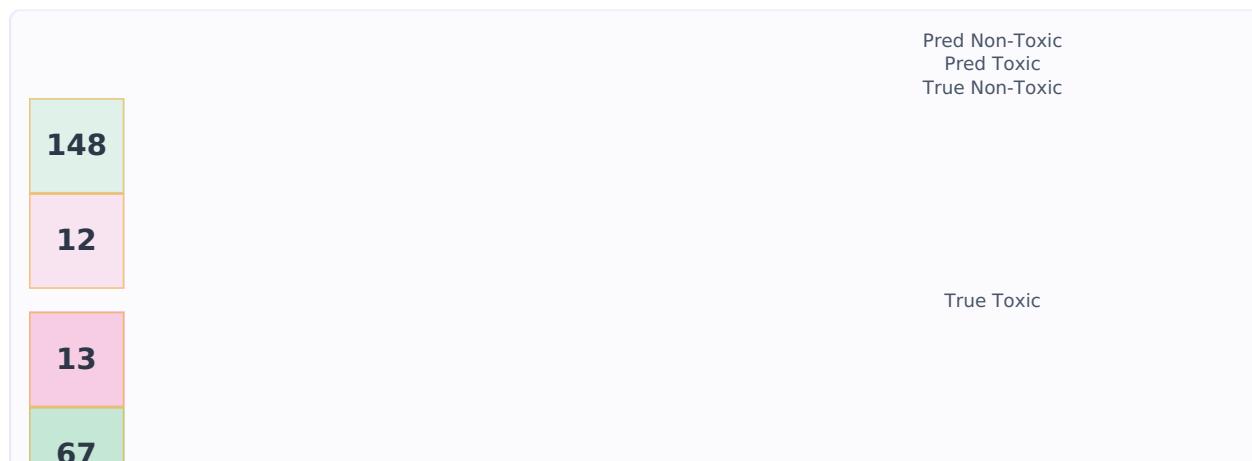
Confusion Matrices (Test Set)

QVC Confusion Matrix



Accuracy: 0.879

QSVM Confusion Matrix



Competitive Advantage

- Quantum feature maps capture **high-order interactions** beyond many classical kernels
 - Expressive yet **parameter-efficient** circuits reduce overfitting on small datasets
 - Fidelity-based kernels** provide powerful similarity measures for complex manifolds
 - Hybrid training (QVC) balances expressivity and **optimization stability**
 - NISQ-aligned** design now; hardware acceleration path as quantum devices mature
 - Complements classical baselines; **ensemble-ready** for best-of-both worlds
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Applications in Drug Discovery

- **Early toxicity screening** for large compound libraries - rapidly identify and filter out toxic compounds early in the discovery pipeline
 - **Lead optimization:** flag off-target risks and cardiotoxicity concerns (e.g., hERG inhibition) during lead optimization
 - **ADMET profiling augmentation** to improve success probability - complement traditional approaches with AI-powered predictions
 - **De-risking IND packages** with transparent metrics and comprehensive reports for regulators
 - **Drug repurposing:** reassess legacy compounds with improved toxicity prediction to identify new applications
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Impact & ROI

- **Time:** compress toxicology signal-finding from months to weeks via prioritized screening
 - **Cost:** reduce late-stage attrition; shift failure left—millions saved per asset
 - **Portfolio:** improve hit triage, elevate likelihood of technical success (LoTS)
 - **Operations:** scalable simulations today; future hardware speed-ups
 - **KPI examples:** ↓ time-to-insight, ↓ cost-per-screen, ↑ precision in toxic class
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Team & Technical Expertise

- **Quantum ML Scientist:** Qiskit circuits, VQC/QSVM design, quantum kernel methods
 - **Computational Chemist/ADMET Lead:** domain features, assay integration, interpretability
 - **Data Engineer/MLOps:** pipelines, model versioning, secure data workflows
 - **Pharma Advisor:** translational strategy, regulatory alignment
 - **Partners & Ecosystem:** quantum hardware providers; academic collaborators
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Next Steps & Call to Action

Engagement Options:

- **4-6 week pilot** on your historical assays (confidential, secure)
- **POC integration** with your cheminformatics stack (RDKit/ELN/LIMS)
- **Joint research** or co-development for indication-specific toxicity

Asks:

- Data access for benchmarking, feedback from tox and DMPK teams
- Investment/partnership to scale models and evaluate on proprietary datasets

Contact & Timeline:

Week 1:

Data onboarding & validation

Weeks

2-3: Model adaptation & benchmarking

Weeks

4-6: Results review, decision, and scale plan
