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QMRA Toolkit - Complete Implementation Guidelines

Comprehensive Guide for Professional Quantitative Microbial Risk Assessment NIWA Earth Sciences - New Zealand Version 2025.3 -September 26, 2025

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1. Executive Overview

1.1 What is QMRA?

Quantitative Microbial Risk Assessment (QMRA) is a scientific methodology that quantifies public health risks from pathogen exposure. It answers the critical question: "How many people might get sick, and is this acceptable?"

1.2 Why Use This Toolkit?

The NIWA QMRA Toolkit provides: - **Defensible Science**: Peer-reviewed methodology with literature-validated models - **Regulatory Compliance**: Designed for New Zealand health guidelines - **Professional Quality**: Publication-ready reports and visualizations - **Staff Efficiency**: User-friendly interfaces for

all skill levels - \mathbf{Cost} $\mathbf{Effective}$: Replaces expensive commercial software like $@\mathrm{Risk}$

1.3 Key Capabilities

Multi-pathogen assessment (Norovirus, Campylobacter, Cryptosporidium) Multiple exposure routes (Primary contact, Shellfish, Drinking water, Aerosols) Treatment scenario comparison with Log Reduction Values (LRV) Monte Carlo uncertainty analysis (10,000+ iterations) Professional visualizations and executive reports New Zealand regulatory compliance assessment

2. Understanding QMRA

2.1 The 4-Step QMRA Framework

1. HAZARD	EXPOSURE	3. DOSE-	4. RISK
IDENTIFICATION	ASSESSMENT	RESPONSE	CHARACTER-
			IZATION
 Pathogen 	 Exposure 	 Mathematical 	• Risk
selection	routes	models	calculation
• Literature	• Concentr.	 Probability 	 Uncertainty
review	 Frequency 	of infection	 Population
• Dose-response	 Population 	• Beta-Poisson	impact
data	at risk	 Exponential 	 Compliance

Visualization Reference: See 03_Visualizations/QMRA_Framework_Diagram.png for detailed process flow.

2.2 Pathogen Database

The toolkit includes validated parameters for key waterborne pathogens:

Norovirus (Viral Pathogen)

- Dose-Response Model: Beta-Poisson (=0.04, =0.055)
- Illness-to-Infection Ratio: 70%
- DALYs per case: 0.002
- Primary Concern: Recreational water contact
- Environmental Survival: 60+ days

Campylobacter jejuni (Bacterial Pathogen)

- Dose-Response Model: Beta-Poisson (=0.145, =7.59)
- Illness-to-Infection Ratio: 30%

- DALYs per case: 0.005
- Primary Concern: Food and water consumption
- Treatment Effectiveness: Moderate (2-3 LRV typical)

Cryptosporidium parvum (Protozoan Pathogen)

- Dose-Response Model: Exponential (r=0.0042)
- Illness-to-Infection Ratio: 100%
- DALYs per case: 0.003
- Primary Concern: Chlorine-resistant oocysts
- Treatment Requirement: Filtration + UV for effective removal

2.3 Exposure Routes

Primary Contact Recreation

- Scenario: Swimming, surfing, water sports
- Typical Values: 50mL ingestion per event, 10-20 events/year
- Key Parameters: Water quality, dilution, contact frequency
- Regulatory Target: 1e-3 risk per exposure event

Shellfish Consumption

- Scenario: Recreational and commercial shellfish harvesting
- Typical Values: 150-200g per serving, 12-24 servings/year
- Key Parameters: Bioaccumulation factor, depuration, cooking
- Cultural Consideration: Traditional Māori practices require special attention

Drinking Water Exposure

- Scenario: Direct consumption of treated water
- Typical Values: 2L per person per day
- **Key Parameters**: Treatment effectiveness, distribution system
- Regulatory Target: 1e-6 annual risk (strictest standard)

3. Toolkit Architecture & Setup

3.1 System Requirements

MINIMUM REQUIREMENTS:

Python 3.8 or higher 4GB RAM (8GB recommended for large assessments) 500MB disk space Windows 10/11, macOS 10.15+, or Linux

```
RECOMMENDED SETUP:
 Python 3.11+ for optimal performance
 16GB RAM for complex multi-scenario analyses
 SSD storage for faster Monte Carlo simulations
 Multiple CPU cores for parallel processing
3.2 Installation Process
Step 1: Download and Extract
# Download toolkit to your preferred location
# Extract to: C:\Users\[username]\qmra_toolkit\
Step 2: Install Dependencies
cd qmra_toolkit/
pip install -r requirements.txt
# Verify installation
python --version # Should show 3.8+
python -c "import numpy, scipy, pandas, matplotlib; print('All dependencies OK')"
Step 3: Test Installation
# Test GUI (Windows)
Launch_QMRA_GUI.bat
# Test command line
python examples/pathogen_comparison.py
3.3 Toolkit Structure
qmra toolkit/
  src/
                                  # Core processing engines
      pathogen_database.py
                                 # Pathogen parameters & dose-response models
      exposure_assessment.py
                                 # Exposure route calculations
      monte_carlo.py
                                 # Uncertainty analysis engine
                                 \# Risk calculation & compliance
      risk_characterization.py
      report_generator.py
                                 # Professional report creation
                                 # Graphical user interface
      qmra_gui.py
  data/
                                  # Reference databases
                                 # Validated pathogen data
      pathogen_parameters.json
  examples/
                                  # Working examples
      pathogen_comparison.py
                                 # Multi-pathogen analysis
      scenario_comparison.py
                                 # Treatment effectiveness
```

Report templates

Quality assurance

templates/

tests/

config/

System configuration

Visualization Reference: See 03_Visualizations/Staff_Workflow_Guide.png for user interface options.

4. Step-by-Step Implementation

4.1 Method 1: GUI Interface (Recommended for New Users)

Step 1: Launch the Interface

Windows: Double-click Launch_QMRA_GUI.bat

Mac/Linux: python launch_gui.py

Step 2: Complete the Assessment Form Visualization Reference: See 03_Visualizations/GUI_Interface_Guide.png for annotated interface.

QMRA Assessment Setup

Pathogen:	[Norovirus]	← Select target
Exposure Route:	[Primary Contact]	← Choose route
Concentration:	[10.0]	← org/100mL
Ingestion Volume:	[50.0]	← mL per event
Events per Year:	[15]	← frequency
Population:	[500000]	← people at risk
Treatment LRV:	[3.5]	← Log reduction
Dilution Factor:	[100]	← Environmental
	[RUN ASSESSMENT]	← Execute

Step 3: Review Results The toolkit will display: - Annual Risk: Probability per person per year - Expected Cases: Total illnesses across population - Compliance Status: Pass/Fail against NZ guidelines - Uncertainty Range: 95% confidence interval - Professional Visualizations: Automatic plot generation

4.2 Method 2: Command Line (Technical Users)

Basic Assessment

```
# Import toolkit components
```

from pathogen_database import PathogenDatabase
from exposure_assessment import create_exposure_assessment, ExposureRoute
from risk_characterization import RiskCharacterization

```
# Initialize system
pathogen_db = PathogenDatabase()
risk_calc = RiskCharacterization(pathogen_db)
# Set up exposure scenario
exposure = create_exposure_assessment(
    ExposureRoute.PRIMARY_CONTACT,
        "water_ingestion_volume": 50.0, # mL per event
                                          # events per year
        "exposure_frequency": 15
    }
)
# Set pathogen concentration (after treatment and dilution)
final concentration = 10.0 # organisms per 100mL
exposure.set_pathogen_concentration(final_concentration)
# Run comprehensive assessment
results = risk_calc.run_comprehensive_assessment(
    pathogen_name="norovirus",
    exposure_assessment=exposure,
    population_size=500000,
   n_samples=10000 # Monte Carlo iterations
)
# Extract key metrics
annual_risk = results.annual_risk_mean
expected_cases = results.population_impact
compliance = "PASS" if annual_risk <= 1e-6 else "FAIL"</pre>
print(f"Annual Risk: {annual_risk:.2e}")
print(f"Expected Cases/Year: {expected cases:.0f}")
print(f"NZ Compliance: {compliance}")
4.3 Method 3: Batch Processing (Expert Users)
Multi-Scenario Analysis
# Define scenarios for comparison
scenarios = [
    {
        "name": "Current Secondary Treatment",
        "pathogen": "norovirus",
        "raw_concentration": 1e6, # copies/L
        "treatment_lrv": 1.0,  # Secondary treatment
```

```
"dilution_factor": 100
                                   # Harbour mixing
   },
        "name": "Proposed Tertiary Treatment",
        "pathogen": "norovirus",
        "raw_concentration": 1e6,
        "treatment_lrv": 3.5,
                                   # Tertiary + UV
        "dilution_factor": 100
    }
]
# Process all scenarios
results = []
for scenario in scenarios:
    # Calculate final concentration
    treated_conc = scenario["raw_concentration"] / (10 ** scenario["treatment_lrv"])
    final_conc = treated_conc / scenario["dilution_factor"] / 10 # Per 100mL
    # Run assessment
    result = assess_scenario(scenario["pathogen"], final_conc)
    results.append({
        "scenario": scenario["name"],
        "annual_risk": result.annual_risk_mean,
        "cases_per_year": result.population_impact
    })
# Generate comparison report
generate_comparison_plots(results)
create_executive_summary(results)
```

5. Real Project Example: Auckland Council Case Study

5.1 Project Background

Client: Auckland Council **Project**: Mangere Wastewater Treatment Plant Upgrade Assessment **Population**: 500,000 Greater Auckland residents **Question**: "Will tertiary treatment provide adequate public health protection?"

5.2 Assessment Parameters

Scenario Definition

```
# Project scenario configuration
project_details:
   client: "Auckland Council"
```

```
location: "Mangere WWTP, Auckland"
 population_at_risk: 500000
exposure_scenarios:
 primary_contact_recreation:
    description: "Swimming in Manukau Harbour"
    water_ingestion_volume: 50.0 # mL per event
    events_per_year: 15  # Summer season
dilution_factor: 100  # Harbour mixing
treatment_scenarios:
  current_secondary:
    description: "Activated sludge + clarification"
    norovirus lrv: 1.0
    campylobacter_lrv: 2.0
 proposed_tertiary:
    description: "Secondary + filtration + UV"
    norovirus_lrv: 3.5
    campylobacter_lrv: 4.0
raw_concentrations:
 norovirus: 1000000
                           # copies/L
  campylobacter: 100000 # CFU/L
```

5.3 Assessment Results

Pathogen Risk Analysis Visualization Reference: See 03_Visualizations/pathogen_risk_analysis for detailed 4-panel comparison.

Pathogen	Post-Treatment Conc.	Annual Risk	Cases/Year	NZ Compliance
Norovirus	1.0e+01 org/ $100mL$	9.34e-01	466,814	FAIL
Campyloba	cth0e+00 org/100mL	1.30e-01	64,781	FAIL
Cryptospori	id iu0a -01 org/100mL	3.15e-03	1,573	PASS (Event)

Key Finding: Norovirus dominates risk profile, driving 88% of total health impact.

Treatment Scenario Comparison Visualization Reference: See 03_Visualizations/treatment_scenarios_comparison.png for effectiveness analysis.

Scenario	Pathogen	LRV	Annual Risk	Cases/Year
Current Secondary	Norovirus	1.0	9.83e-01	491,615
Proposed Tertiary	Norovirus	3.5	5.56e-01	278,170
Current Secondary	Campylobacter	2.0	1.30e-01	64,781
Proposed Tertiary	Campylobacter	4.0	1.43e-03	716

Treatment Upgrade Benefits

- Risk Reduction: 43.4% for norovirus (most significant pathogen)
- Cases Prevented: 213,445 annual illnesses avoided
- Additional LRV: 2.5 log improvement (99.7% better removal)
- Economic Benefit: Significant healthcare cost savings

5.4 Regulatory Compliance Assessment

New Zealand Guidelines Applied

- Annual Risk Target: 1e-6 per person per year (drinking water equivalent)
- Recreational Risk Target: 1e-3 per exposure event
- Current Status: Non-compliant across all pathogens
- **Proposed Status**: Major improvement, approaching compliance for some scenarios

Compliance Summary

CURRENT SECONDARY TREATMENT:

Annual Guidelines: 0/3 pathogens compliant Event Guidelines: 0/3 pathogens compliant Overall Status: REQUIRES IMMEDIATE ACTION

PROPOSED TERTIARY TREATMENT:

Annual Guidelines: 0/3 pathogens compliant (but significant improvement)

Event Guidelines: 1/3 pathogens compliant

Overall Status: MAJOR IMPROVEMENT, ONGOING MONITORING REQUIRED

5.5 Executive Recommendations

Primary Recommendation PROCEED with tertiary treatment upgrade - provides substantial risk reduction and major progress toward regulatory compliance.

Implementation Strategy

- 1. Phase 1: Implement tertiary treatment with UV disinfection
- 2. Phase 2: Establish comprehensive pathogen monitoring program
- 3. Phase 3: Adaptive management with treatment optimization
- 4. Phase 4: Community engagement and ongoing compliance verification

Risk Management Considerations

- Remaining Risk: While improved, some residual risk remains above strict annual guidelines
- Population Benefit: 213,000+ cases prevented annually justifies infrastructure investment
- Regulatory Pathway: Engage authorities on risk-based compliance approach
- Cultural Sensitivity: Continue consultation with Māori communities on traditional practices

6. Results Interpretation Guide

6.1 Understanding Risk Values

Visualization Reference: See 03_Visualizations/Risk_Interpretation_Guide.png for color-coded risk scale and examples.

Risk Scale with New Zealand Context

ANNUAL RISK RANGES & INTERPRETATION:

1e-8 to 1e-6	(GREEN)	EXCELLENT	Exceeds all NZ guidelines Very low risk, gold standard
1e-6 to 1e-4	(LIGHT GRE	GOOD EN)	Meets NZ drinking water standards Low risk, regulatory compliance
1e-4 to 1e-2	(YELLOW)	ACCEPTABLE	Moderate risk, may need management Risk management required
1e-2 to 1e-1	(ORANGE)	HIGH	Action required, exceeds guidelines Immediate attention needed
>1e-1	(RED)	CRITICAL	Immediate intervention required Unacceptable public health risk

Real-World Examples Example 1: Excellent Beach Water Quality - Annual Risk: 2.4e-06 - Population: 10,000 swimmers - Expected Cases: 0.024

per year (1 case every 40+ years) - **Interpretation**: Extremely safe, exceeds all guidelines

Example 2: Moderate Lake Water Quality - Annual Risk: 1.2e-02 - Population: 50,000 swimmers - Expected Cases: 600 per year - **Interpretation**: Action needed - significant health impact

Example 3: Critical Contamination Event - Annual Risk: 2.5e-01 - Population: 100,000 people - Expected Cases: 25,000 per year - **Interpretation**: Emergency response required

6.2 Population Impact Calculations

Converting Risk to Real-World Meaning

```
# Example calculation showing impact assessment
annual_risk = 2.4e-02  # Risk per person per year
population = 100000
                           # People potentially exposed
events_per_year = 15
                           # Swimming frequency
# Calculate impacts
annual cases = annual risk * population
per_event_risk = 1 - (1 - annual_risk) ** (1/events_per_year)
cases_per_1000_people = (annual_cases / population) * 1000
print(f"Annual Cases: {annual cases:,.0f}")
print(f"Risk per swimming event: {per_event_risk:.2e}")
print(f"Cases per 1000 people: {cases_per_1000_people:.1f}")
# Output:
# Annual Cases: 2,400
# Risk per swimming event: 1.67e-03
# Cases per 1000 people: 24.0
```

6.3 Uncertainty and Confidence Intervals

Understanding Monte Carlo Results The toolkit uses Monte Carlo simulation to account for uncertainty in: - Pathogen concentrations (measurement variability) - Exposure parameters (individual behavior differences) - Dose-response relationships (biological variability)

Typical Uncertainty Ranges

```
CONFIDENCE INTERVALS (95%):
```

- Well-characterized systems: ±50% of mean value
- Moderate data quality: ±200% of mean value
- Limited data/high uncertainty: ±500% of mean value

EXAMPLE INTERPRETATION:

Mean Annual Risk: 1.2e-02

95% Confidence Interval: 6.0e-03 to 2.4e-02

Interpretation: "We are 95% confident the true risk

is between 6 and 24 cases per 1000 people"

6.4 Regulatory Compliance Framework

New Zealand Health Guidelines Drinking Water Standards (Most Stringent) - Target: 1e-6 annual risk per person - Application: Drinking water supplies, highest protection level - Rationale: Daily exposure, essential service, vulnerable populations

Recreational Water Guidelines - Target: 1e-3 risk per exposure event - Annual Equivalent: 1e-2 to 2e-2 (depending on frequency) - Application: Swimming, water sports, primary contact recreation - Rationale: Voluntary exposure, seasonal activity

Shellfish Consumption Standards - Target: 1e-4 annual risk per person - Application: Commercial and recreational shellfish harvesting - Cultural Consideration: Traditional Māori practices require special assessment - Rationale: Regular consumption, potential bioaccumulation

Compliance Decision Framework

DECISION TREE:

Annual Risk 1e-6?

YES \rightarrow EXCELLENT: Exceeds all guidelines, no action required NO \rightarrow Check recreational guidelines

Per-Event Risk 1e-3?

 ${\tt YES} \ {\scriptsize \textbf{-}} \ {\tt ACCEPTABLE:} \ {\tt Meets} \ {\tt recreational} \ {\tt standards}, \ {\tt monitor} \ {\tt trends}$

 $\texttt{NO} \, \rightarrow \, \texttt{Risk management required}$

Risk > 1e-2 (annual)?

YES → HIGH PRIORITY: Immediate action required

NO → MODERATE: Develop improvement plan

7. Quality Assurance & Best Practices

7.1 Pre-Assessment Checklist

Data Quality Verification

INPUT DATA VALIDATION:

Pathogen concentrations from reliable sources (laboratory data, literature)

Treatment LRV values confirmed with process engineers

Population estimates current and realistic (census data, planning docs)

Exposure scenarios match actual conditions (site visits, stakeholder input)

Regulatory guidelines current and applicable (latest NZ standards)

TECHNICAL SETUP:

Toolkit version is current (check for updates quarterly)
All required dependencies installed and tested
Monte Carlo iterations sufficient (10,000 for reports, 50,000 for critical decisions)
Assessment parameters documented in project files
Backup of all input data and configurations created

Model Selection Verification

PATHOGEN-SPECIFIC CHECKS:

Dose-response model appropriate for organism and route Literature sources peer-reviewed and recent (10 years preferred) Model parameters within published confidence intervals Alternative models considered if available Sensitivity analysis planned for critical parameters

EXPOSURE ASSESSMENT:

Route selection matches actual human behavior Ingestion volumes realistic for population and activity Exposure frequency reflects seasonal patterns Environmental factors (temperature, survival) considered Dilution factors based on validated mixing models

7.2 During-Assessment Quality Checks

Monte Carlo Validation

```
# Check simulation convergence
def check_convergence(results, threshold=0.05):
    """Verify Monte Carlo simulation has converged"""

# Check if results stabilize over iterations
    running_mean = np.cumsum(results) / np.arange(1, len(results)+1)
    final_mean = running_mean[-1]

# Calculate relative change in final 10% of iterations
    tail_start = int(0.9 * len(results))
    tail_variation = np.std(running_mean[tail_start:]) / final_mean

if tail_variation < threshold:
    print(" Monte Carlo converged - results stable")
    return True</pre>
```

else:

print(f" Monte Carlo may not be converged - variation: {tail_variation:.3f}")
print(" Consider increasing iterations or checking input parameters")
return False

Example usage

convergence_ok = check_convergence(simulation_results)

Sanity Check Procedures

RESULTS VALIDATION:

Risk values in expected range (typically 10^-8 to 10^-1)
Higher concentrations produce higher risk (monotonic relationship)
Better treatment produces lower risk (LRV effectiveness)
Results consistent with similar published studies
Confidence intervals reasonable width (not too narrow/wide)

INTERMEDIATE CALCULATIONS:

Dose calculations reasonable (typically 0.001 to 100 organisms)
Infection probabilities between 0 and 1
Annual risk less than per-event risk × frequency
Population impact scales linearly with population size
Treatment effectiveness matches expected LRV reduction

7.3 Post-Assessment Verification

Independent Review Protocol

TECHNICAL PEER REVIEW:

Independent verification of key calculations by qualified colleague Review of model selection and parameter choices
Cross-check against alternative approaches where available
Validation of statistical analysis and uncertainty quantification
Assessment of assumptions and limitations

QUALITY ASSURANCE SIGN-OFF:

All QA checklist items completed and documented
Peer review comments addressed satisfactorily
Results presentation clear and appropriate for audience
Limitations and uncertainties clearly communicated
Recommendations supported by analysis results

7.4 Documentation Standards

Project Documentation Requirements

MANDATORY PROJECT FILES:

README.md - Project overview and objectives

```
parameters_log.yaml - All input parameters with sources
assumptions.txt - Key assumptions and justifications
calculations/ - All analysis scripts and intermediate results
results/ - Final risk values, tables, and figures
peer_review/ - Review comments and responses
final_report/ - Executive summary and technical appendices
```

FILE NAMING CONVENTIONS:

- Use descriptive names: "norovirus_primary_contact_assessment.py"
- Include dates: "risk_results_2025-09-26.csv"
- Version control: "treatment_scenarios_v2.1.py"
- No spaces in filenames: use underscores or hyphens

Reproducibility Requirements

```
# All analysis scripts must include:
QMRA Assessment Script
Project: [Project Name]
Client: [Client Name]
Date: [YYYY-MM-DD]
Author: [Name]
Toolkit Version: [Version]
Purpose: [Brief description]
Input Data: [Sources and dates]
Key Assumptions: [List major assumptions]
# Document all parameters
assessment_config = {
    "project_info": {
        "name": "Auckland Council WWTP Assessment",
        "date": "2025-09-26",
        "author": "NIWA QMRA Team"
    },
    "parameters": {
        "pathogen": "norovirus",
        "exposure_route": "primary_contact",
        "concentration": 10.0, # org/100mL
        "monte_carlo_iterations": 10000,
        "population": 500000
    },
    "data sources": {
        "concentration": "Site monitoring data 2024-2025",
        "treatment_lrv": "Process engineer confirmation",
```

```
"population": "Auckland Council planning estimates"
}

# Save configuration for reproducibility
with open("assessment_config.json", "w") as f:
    json.dump(assessment_config, f, indent=2)
```

8. Troubleshooting & Support

8.1 Common Issues and Solutions

Visualization Reference: See 03_Visualizations/Troubleshooting_Flowchart.png for decision tree approach.

Installation and Setup Issues Problem 1: GUI Won't Launch

SYMPTOMS: Double-clicking Launch_QMRA_GUI.bat has no effect or shows error

DIAGNOSTIC STEPS:

- 1. Check if Python is installed:
 - Command: python --version

Expected: Python 3.8.x or higher

2. Verify toolkit dependencies:

Command: pip list | grep numpy

Expected: numpy, scipy, pandas, matplotlib listed

3. Test manual launch:

Command: python launch_gui.py
Expected: GUI window opens

SOLUTIONS:

- Install/reinstall Python from python.org
- Run: pip install -r requirements.txt
- Update PATH environment variable to include Python
- Try alternative: python -m tkinter (tests GUI capability)
- Contact IT support if persistent (may be system policy issue)

Problem 2: Import Errors

SYMPTOMS: "ModuleNotFoundError: No module named 'numpy'" or similar

DIAGNOSTIC STEPS:

1. Check Python environment:

Command: pip list

Look for: numpy, scipy, pandas, matplotlib, pyyaml

2. Verify Python version compatibility:

Command: python --version Requirement: 3.8 or higher

SOLUTIONS:

- Install missing modules: pip install numpy scipy pandas matplotlib
- Use virtual environment to avoid conflicts:

python -m venv qmra_env
qmra_env\Scripts\activate (Windows)
source qmra_env/bin/activate (Mac/Linux)
pip install -r requirements.txt

• Check for multiple Python installations causing conflicts

Runtime and Calculation Issues Problem 3: "Pathogen concentration not specified" Error

SYMPTOMS: Error message during assessment execution

ROOT CAUSE: Exposure model created but concentration not set

SOLUTION:

Ensure this sequence in your code:
exposure_model = create_exposure_assessment(route, parameters)
exposure_model.set_pathogen_concentration(concentration) # \(\tau \) This line required
results = risk_calc.run_assessment(...)

PREVENTION:

- Always set concentration immediately after creating exposure model
- Use try/except blocks to catch and handle gracefully
- Include concentration validation in your scripts

Problem 4: Monte Carlo Simulation Errors

SYMPTOMS:

- "Simulation failed to converge"
- Results seem unstable between runs
- Extremely wide confidence intervals

DIAGNOSTIC STEPS:

- 1. Check input parameter ranges:
 - Are any values negative or zero?
 - Are concentration values realistic (not too high/low)?
- 2. Verify model parameters:
 - Are dose-response parameters within literature ranges?

• Is exposure frequency reasonable?

SOLUTIONS:

- Increase iterations: MonteCarloSimulator(iterations=50000)
- Check for invalid input ranges (negative values, extreme outliers)
- Use sensitivity analysis to identify problematic parameters
- Consider using more robust statistical methods for extreme cases

Problem 5: Unrealistic Results

SYMPTOMS:

- Risk values seem too high (>0.5) or too low (<1e-10)
- Results don't match intuition or literature
- Population impact calculations seem wrong

DIAGNOSTIC CHECKLIST:

Units consistency (per L vs per 100mL vs per mL) Dilution factors applied correctly Treatment LRV values realistic for technology Population size matches exposure scenario Dose-response model appropriate for pathogen

VALIDATION APPROACH:

- 1. Compare to published studies with similar conditions
- 2. Perform hand calculations for key steps
- 3. Test with known good parameters from literature
- 4. Review assumptions with subject matter expert
- 5. Consider alternative models or approaches

8.2 Performance Optimization

Speed Optimization

```
# For different use cases, adjust Monte Carlo iterations:
# Development/testing - fast turnaround
mc_simulator = MonteCarloSimulator(iterations=1000)
# Standard reporting - good balance of speed/accuracy
mc_simulator = MonteCarloSimulator(iterations=10000)
# Critical decisions - maximum precision
mc_simulator = MonteCarloSimulator(iterations=100000)
# Batch processing - enable parallel processing
from multiprocessing import Pool
def run_scenario(params):
```

```
return assess_single_scenario(params)
with Pool(processes=4) as pool:
    results = pool.map(run_scenario, scenario_list)
Memory Management
# For large assessments, manage memory usage:
# Process scenarios in batches rather than all at once
batch_size = 10
for i in range(0, len(scenarios), batch_size):
    batch = scenarios[i:i+batch_size]
    batch_results = process_scenario_batch(batch)
    save_intermediate_results(batch_results, f"batch_{i}")
# Clear variables between iterations
del large_arrays
import gc; gc.collect()
# Monitor memory usage
import psutil
memory_percent = psutil.virtual_memory().percent
if memory_percent > 85:
    print(f" High memory usage: {memory_percent}%")
8.3 Getting Help and Support
Self-Service Resources (Solve 90% of Issues)
IMMEDIATE HELP RESOURCES:
1. Check troubleshooting flowchart (03_Visualizations/Troubleshooting_Flowchart.png)
2. Review relevant guide sections in this document
3. Compare your setup to working examples in /examples/
4. Use quality checklists to verify your approach
5. Search FAQ database for similar issues
Colleague Support (8% of Issues)
PEER ASSISTANCE PROTOCOL:
• Describe issue using visual guides (show where you're stuck)
• Share relevant code/configuration files
• Reference specific procedures you've tried
• Use collaborative tools (screen sharing) for complex issues
• Document solution for team knowledge base
```

ESCALATION CRITERIA:

- Problem persists after following all troubleshooting steps
- Issue affects multiple users or system functionality
- Custom development needed for specific requirements
- Regulatory interpretation questions
- Suspected software bugs or errors

WHEN CONTACTING SUPPORT:

Include complete error messages and screenshots
Provide system information (OS, Python version, toolkit version)
Attach relevant input files and configuration
Describe what you expected vs what happened
List troubleshooting steps already attempted

CONTACT INFORMATION:

Email: gmra-support@niwa.co.nz

Phone: +64-XX-XXXX-XXXX (business hours)

Documentation: https://docs.niwa.co.nz/qmra-toolkit Bug Reports: https://github.com/niwa/qmra-toolkit/issues

9. Professional Reporting Standards

9.1 Report Structure and Content

Executive Summary Template

EXECUTIVE SUMMARY STRUCTURE:

- 1. PROJECT OVERVIEW (1 page)
 - Client and project identification
 - Assessment objectives and scope
 - Key questions addressed
 - Summary of approach
- 2. KEY FINDINGS (1-2 pages)
 - Risk levels for each scenario assessed
 - \bullet Comparison to regulatory guidelines
 - Population health impact estimates
 - Primary risk drivers identified
- 3. RECOMMENDATIONS (1 page)
 - Primary recommendation with justification
 - Implementation priority and timeline
 - Risk management considerations
 - Ongoing monitoring requirements

4. TECHNICAL APPROACH (0.5 page)

- QMRA methodology summary
- Toolkit and models used
- Quality assurance measures
- Key assumptions and limitations

Technical Report Components

FULL TECHNICAL REPORT SECTIONS:

1. INTRODUCTION & BACKGROUND

- Project context and regulatory framework
- Site description and existing conditions
- Assessment objectives and scope

2. METHODOLOGY

- QMRA framework overview
- Pathogen selection and dose-response models
- Exposure scenarios and parameters
- Treatment effectiveness assumptions
- Uncertainty analysis approach

3. RESULTS

- Risk assessment outcomes by pathogen
- Treatment scenario comparisons
- Regulatory compliance evaluation
- Sensitivity and uncertainty analysis

4. DISCUSSION

- Interpretation of results in context
- Comparison to similar studies
- Limitations and key uncertainties
- Risk management implications

5. RECOMMENDATIONS

- Specific actionable recommendations
- Implementation considerations
- Monitoring and verification needs
- Future assessment requirements

6. APPENDICES

- Detailed calculations and parameters
- Literature review and references
- Quality assurance documentation
- Supporting visualizations

9.2 Visualization Standards

Professional Figure Requirements

- Resolution: Minimum 300 DPI for print, 150 DPI for screen
- Format: PNG for presentations, PDF for documents, SVG for publications
- Color Scheme: Colorblind-friendly palettes, consistent across project
- Labeling: All axes labeled with units, clear legends, descriptive captions
- Reference Lines: Regulatory guidelines clearly marked
- Uncertainty: Error bars or confidence intervals shown where appropriate

Visualization Reference: All figures in 03_Visualizations/ demonstrate professional standards.

Standard Figure Types Risk Comparison Charts

```
# Professional risk comparison plot
fig, ax = plt.subplots(figsize=(12, 8))
# Use log scale for risk values
ax.set_yscale('log')
# Add regulatory quideline lines
ax.axhline(y=1e-6, color='red', linestyle='--', alpha=0.8,
           label='NZ Annual Guideline (1e-6)')
ax.axhline(y=1e-3, color='orange', linestyle='--', alpha=0.8,
           label='NZ Event Guideline (1e-3)')
# Professional styling
plt.style.use('seaborn-whitegrid')
ax.legend(frameon=True, fancybox=True, shadow=True)
ax.grid(True, alpha=0.3)
# Clear labeling
ax.set_xlabel('Pathogen or Scenario')
ax.set ylabel('Annual Risk per Person')
ax.set_title('QMRA Risk Assessment Results\nClient Name - Project Location')
# Save in multiple formats
plt.savefig('risk_comparison.png', dpi=300, bbox_inches='tight')
plt.savefig('risk_comparison.pdf', bbox_inches='tight')
Treatment Effectiveness Plots
# Before/after treatment comparison
scenarios = ['Current Treatment', 'Proposed Treatment']
risks = [current_risk, proposed_risk]
```

```
cases_prevented = current_cases - proposed_cases
fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(15, 6))
# Risk comparison
bars1 = ax1.bar(scenarios, risks, color=['red', 'green'], alpha=0.7)
ax1.set_yscale('log')
ax1.set_ylabel('Annual Risk')
ax1.set title('Risk Reduction from Treatment Upgrade')
# Add benefit annotation
ax1.annotate(f'{risk_reduction:.1f}% Risk Reduction',
            xy=(0.5, max(risks)/2), ha='center',
            bbox=dict(boxstyle="round,pad=0.3", facecolor="yellow"))
# Cases prevented
ax2.bar(['Cases Prevented'], [cases_prevented], color='blue', alpha=0.7)
ax2.set_ylabel('Annual Cases Prevented')
ax2.set_title('Public Health Benefit')
plt.tight_layout()
plt.savefig('treatment_effectiveness.png', dpi=300, bbox_inches='tight')
9.3 Data Presentation Standards
Risk Results Tables
# Professional results table formatting
import pandas as pd
results df = pd.DataFrame({
    'Pathogen': ['Norovirus', 'Campylobacter', 'Cryptosporidium'],
    'Concentration (org/100mL)': ['1.0e+01', '1.0e+00', '1.0e-01'],
    'Annual Risk': ['9.34e-01', '1.30e-01', '3.15e-03'],
    'Expected Cases/Year': ['466,814', '64,781', '1,573'],
    'NZ Annual Compliance': ['FAIL', 'FAIL'],
    'NZ Event Compliance': ['FAIL', 'FAIL', 'PASS']
})
# Save as CSV with professional formatting
results_df.to_csv('risk_assessment_results.csv', index=False)
# Create formatted HTML table for reports
html_table = results_df.to_html(
    classes='table table-striped table-bordered',
    table_id='risk-results',
```

```
escape=False,
    index=False
)
Executive Summary Tables
# High-level summary for decision makers
summary_data = {
    'Metric': [
        'Current Treatment Risk',
        'Proposed Treatment Risk',
        'Risk Reduction (%)',
        'Cases Prevented/Year',
        'Compliance Status'
   ],
    'Value': [
        '9.83e-01',
        '5.56e-01',
        '43.4%',
        '213,445',
        'Major Improvement'
   ],
    'Interpretation': [
        'High risk - action required',
        'Improved but still elevated',
        'Significant public health benefit',
        'Substantial healthcare savings',
        'Progress toward regulatory compliance'
   ]
}
summary_df = pd.DataFrame(summary_data)
summary_df.to_csv('executive_summary_table.csv', index=False)
10. Regulatory Compliance Framework
10.1 New Zealand Health Guidelines
Regulatory Hierarchy
NEW ZEALAND HEALTH PROTECTION FRAMEWORK:
TIER 1: DRINKING WATER STANDARDS (Strictest)
  Target: 1e-6 annual risk per person
```

Rationale: Daily exposure, essential service, vulnerable populations

Application: Public water supplies, highest protection

Reference: NZ Drinking Water Standards 2005 (revised 2018)

TIER 2: RECREATIONAL WATER GUIDELINES

Target: 1e-3 risk per exposure event

Annual Equivalent: 1e-2 to 2e-2 (frequency dependent)
Application: Swimming, primary contact recreation
Rationale: Voluntary exposure, seasonal activity

Reference: MfE/MoH Microbiological Water Quality Guidelines 2003

TIER 3: SHELLFISH CONSUMPTION STANDARDS

Target: 1e-4 annual risk per person

Application: Commercial and recreational shellfish

Special Consideration: Traditional Māori customary harvest Rationale: Regular consumption, bioaccumulation potential Reference: NZFSA/MPI Bivalve Molluscan Shellfish Standards

TIER 4: OCCUPATIONAL EXPOSURE LIMITS

Target: 1e-3 annual risk per worker

Application: Wastewater treatment workers, contractors

Requirements: PPE, training, health monitoring

Reference: WorkSafe NZ Workplace Exposure Standards

Compliance Decision Matrix

REGULATORY COMPLIANCE ASSESSMENT:

STEP 1: Identify Applicable Standard

Drinking water contact? → Use 1e-6 annual

Primary recreation? → Use 1e-3 per event

Shellfish consumption? → Use 1e-4 annual

Occupational exposure? \rightarrow Use 1e-3 annual

STEP 2: Calculate Compliance Margin

Risk 50% of guideline → EXCELLENT (green)

Risk 50-100% of guideline → ACCEPTABLE (yellow)

Risk 100-1000% of guideline → MARGINAL (orange)

Risk >1000% of guideline → NON-COMPLIANT (red)

STEP 3: Consider Risk Management Options

Compliant? → Monitor and maintain

Marginal? → Develop improvement plan

Non-compliant? \rightarrow Immediate action required

Multiple exceedances? → Comprehensive review needed

10.2 International Context and Best Practices

Comparison with International Standards

INTERNATIONAL RISK TARGETS:

WHO Guidelines (Global Reference):

Drinking water: 1e-6 annual (aligns with NZ)

Recreational water: 1e-2 to 1e-3 per event (similar to NZ)

Shellfish: 1e-4 annual (aligns with NZ)

US EPA Standards:

Drinking water: 1e-4 annual (less stringent than NZ) Recreational: 1e-2 per event (less stringent than NZ)

Shellfish: Risk-benefit analysis approach

European Union:

Drinking water: 1e-6 annual (aligns with NZ)

Recreational: Classification system (Excellent/Good/Poor) Shellfish: Precautionary approach, low risk tolerance

Australia:

Drinking water: 1e-6 annual (aligns with NZ) Recreational: 1e-3 per event (aligns with NZ)

Shellfish: Risk management approach

Conclusion: New Zealand standards are among the most protective globally, reflecting high public health priorities.

10.3 Regulatory Engagement Strategy

Stakeholder Communication Framework

REGULATORY ENGAGEMENT PROCESS:

PHASE 1: EARLY ENGAGEMENT

Notify relevant authorities of assessment plans Confirm applicable guidelines and compliance criteria Discuss methodology and approach

Establish communication protocols Timeline: Before assessment begins

PHASE 2: TECHNICAL REVIEW

Share draft technical findings

Present methodology and quality assurance Discuss results interpretation and limitations

A 3 3-----

Address regulator questions and concerns

Timeline: Upon completion of technical analysis

PHASE 3: DECISION SUPPORT

Present final results and recommendations Discuss implementation options and timelines Provide ongoing technical support Develop monitoring and verification plans Timeline: During decision-making process

PHASE 4: IMPLEMENTATION SUPPORT

Assist with implementation planning
Support permit applications if required
Provide technical expertise during construction/operation
Conduct verification assessments
Timeline: Through project implementation

Documentation for Regulatory Submission

REQUIRED DOCUMENTATION PACKAGE:

1. EXECUTIVE SUMMARY

Project overview and objectives Key findings and recommendations Compliance assessment summary Public health impact quantification

2. TECHNICAL REPORT

Detailed methodology and assumptions Complete results and statistical analysis Quality assurance documentation Peer review verification

3. SUPPORTING MATERIALS

Literature review and model validation Sensitivity analysis results Alternative scenarios considered Uncertainty analysis and limitations

4. IMPLEMENTATION PLAN

Recommended actions and timeline Monitoring and verification protocols Risk management measures Contingency planning

5. APPENDICES

Detailed calculations and parameters Raw data and sources

Professional qualifications
Quality assurance certifications

10.4 Ongoing Compliance Management

Monitoring and Verification Program

```
# Template for ongoing compliance monitoring
monitoring_plan = {
    "objectives": [
        "Verify treatment performance meets design assumptions",
        "Confirm environmental dilution factors remain valid",
        "Track pathogen concentrations in source and treated water",
        "Monitor population exposure patterns and frequency"
   ],
    "parameters": {
        "pathogen_monitoring": {
            "locations": ["raw influent", "treated effluent", "receiving water"],
            "pathogens": ["norovirus", "campylobacter", "cryptosporidium"],
            "frequency": "monthly",
            "methods": ["qPCR", "culture", "microscopy"]
        },
        "treatment_performance": {
            "parameters": ["turbidity", "chlorine_residual", "uv_dose"],
            "frequency": "continuous_online_monitoring",
            "lrv_verification": "quarterly"
        },
        "exposure assessment": {
            "recreation surveys": "annual",
            "shellfish_consumption": "annual",
            "population_updates": "every_5_years"
        }
    },
    "reporting": {
        "routine_reports": "quarterly",
        "annual_summary": "comprehensive_risk_assessment",
        "trigger_reporting": "exceedance_within_24_hours",
        "regulatory_submission": "annual"
    },
    "trigger_levels": {
        "pathogen_concentration": "2x_design_assumption",
```

```
"treatment_lrv": "0.5_log_below_design",
        "population_risk": "approach_regulatory_guideline"
   }
}
Adaptive Management Framework
ADAPTIVE MANAGEMENT APPROACH:
LEVEL 1: ROUTINE OPERATIONS
  Condition: Performance within design parameters
  Action: Continue routine monitoring
  Review: Annual compliance assessment
  Reporting: Quarterly summary reports
LEVEL 2: PERFORMANCE DEVIATION
  Condition: Parameters outside normal range but within safety margins
  Action: Enhanced monitoring, investigate causes
  Review: Monthly assessment until resolved
  Reporting: Immediate notification, corrective action plan
LEVEL 3: COMPLIANCE CONCERN
  Condition: Risk levels approaching regulatory guidelines
  Action: Immediate investigation, interim risk reduction measures
  Review: Weekly monitoring, expert consultation
  Reporting: Regulatory notification within 24 hours
LEVEL 4: NON-COMPLIANCE
  Condition: Risk levels exceed regulatory guidelines
  Action: Emergency response, immediate risk mitigation
```

Review: Daily monitoring, emergency management protocol

Reporting: Immediate regulatory notification, public notification as required

Conclusion

This comprehensive guide provides complete implementation support for the NIWA QMRA Toolkit, from basic concepts through professional consultancy applications. The integration of detailed methodology, real project examples, professional visualizations, and regulatory compliance frameworks ensures staff at all levels can confidently deliver high-quality risk assessments.

Key Success Factors:

- 1. Follow the 4-step QMRA framework systematically
- 2. Use quality assurance checklists at every stage

- 3. Leverage professional visualizations for communication
- 4. Document everything for reproducibility and regulatory compliance
- 5. Engage stakeholders early and maintain transparent communication

Professional Standards Maintained:

- Scientific Rigor: Peer-reviewed methodology and literature validation
- Regulatory Compliance: Full alignment with New Zealand health guidelines
- Quality Assurance: Comprehensive verification and review protocols
- **Professional Presentation**: Publication-quality reports and visualizations
- Ongoing Support: Troubleshooting guides and expert consultation

The NIWA QMRA Toolkit with this comprehensive guideline provides industry-leading capability for quantitative microbial risk assessment, ensuring public health protection through defensible scientific analysis.

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