

# Harris MRSA Model - Comprehensive Description

## Overview

This is an Agent-Based Model (ABM) simulating MRSA transmission dynamics in a hospital setting. The model tracks patient admissions, discharges, transfers between ICU and ward, healthcare worker (HCW) visits to patients.

## Next Steps

- finish visit process including transmission to patient, from patient.
- finish disease model especially transitions from C to I, I to R
- address death vs. discharge

## Model Entry Point and Initialization

### Builder.build() - The Main Entry Point

The simulation begins when Repast Symphony calls `Builder.build(Context<Object> context)`. This method orchestrates the entire initialization process.

#### Step-by-step initialization:

##### 1. Get the Simulation Schedule

- Retrieves the Repast Symphony schedule that manages all time-based events
- The schedule represents time as a double precision numeric. Each day is **1** unit, and events may be scheduled at any point between one “tick” and the next. **10.0** is the beginning of the 10th day. **10.5** is the middle of the 10th day.

##### 2. Create the Hospital

- Creates the main `Hospital` object with configured capacity
- Hospital capacity: **120** beds total (configurable via `hospitalCapacity`)
- ICU capacity: **20** beds (configurable via `icuCapacity`)
- Ward capacity: **100** beds (`hospitalCapacity - icuCapacity`)
- The Hospital is added to the simulation context

##### 3. Initialize Hospital Internal State

During Hospital construction, the following are initialized:

- **Patient containers:**
  - `patients` - all patients currently in the hospital

- `inIcu` - patients currently in ICU
- `notInIcu` - patients currently in ward
- `patientsNeedingOt/Pt/Rt` - patients requiring therapy visits
- **Discharge processes:**
  - `icuDischarger` - handles ICU patient discharges using `LogNormal(scale=0.820, shape=0.916)`
  - `nonIcuDischarger` - handles ward patient discharges using `LogNormal(scale=0.768, shape=1.253)`
- **Transfer process:**
  - `transferer` - handles ICU-to-Ward transfers using `LogNormal(scale=1.0, shape=0.5)`
- **Data collection buffers:**
  - `visitData` - `StringBuffer` accumulating all HCW visit records
  - `admissionData` - `StringBuffer` accumulating all admission records
  - `dischargedPatients` - `ArrayList` storing `DischargedPatient` records
- **Network for HCW-Patient assignments:**
  - `hospitalnet` - network tracking which HCWs are assigned to which patients

#### 4. Start the Admission Process

- Creates an `Admission` process with mean inter-arrival time of **0.05** days
- The admission process uses an Exponential distribution
- `admissionProcess.start()` schedules the first admission event

#### 5. Schedule the Builder's Recurring Methods

- Schedules `Builder.daily()` to run every **1.0** tick starting at tick **1.0**
- Schedules `Builder.perShiftOperations()` to run every **0.5** ticks starting at tick **0.5**
- Schedules `Builder.endOfRun()` to run once at tick **365**

#### 6. Build All Healthcare Workers

- Calls `buildHealthCareWorkers()` to create the HCW workforce

#### 7. Create Networks

- Creates two network structures for visualization/analysis (ICU and wards)

### Healthcare Worker Creation

The `buildHealthCareWorkers()` method creates all HCWs and assigns each a `PatientVisit` process:

## Ward Doctors

- **Count:**  $(\text{hospitalCapacity} - \text{icuCapacity}) \times \text{physiciansPerPatient} = 100 \times 0.2 = 20$  doctors
- **Type:** DOCTOR
- **Visit pattern:** Each doctor gets a PatientVisit process that:
  - Uses Gamma(shape=0.52, scale=90.7) distribution for inter-visit intervals (in minutes)
  - Adds 6.6 minutes for room visit duration
  - Starts immediately with `pv.start()`
- **Infection control attributes:**
  - Hand hygiene compliance (pre-visit): 0.5
  - Hand hygiene compliance (post-visit): 0.5
  - PPE/glove compliance: 0.5
- **Assignment:** Added to `wardContext` with `icu=false`

## ICU Doctors

- **Count:**  $\text{icuCapacity} \times \text{icuPhysiciansPerPatient} = 20 \times 0.3 = 6$  doctors
- **Type:** DOCTOR
- **Visit pattern:**
  - Uses Gamma(shape=0.52, scale=35.3) distribution (shorter intervals than ward)
  - Starts immediately
- **Assignment:** Added to `icuContext` with `icu=true`

## Ward Nurses

- **Count:**  $(\text{hospitalCapacity} - \text{icuCapacity}) \times \text{nursesPerPatient} = 100 \times 0.2 = 20$  nurses
- **Type:** NURSE
- **Visit pattern:**
  - Uses Gamma(shape=0.54, scale=55.1) distribution
  - Starts immediately
- **Assignment:** Added to `wardContext` with `icu=false`

## ICU Nurses

- **Count:**  $\text{icuCapacity} \times \text{icuNursesPerPatient} = 20 \times 0.5 = 10$  nurses
- **Type:** NURSE
- **Visit pattern:**
  - Uses  $\text{Gamma}(\text{shape}=0.54, \text{scale}=20)$  distribution (much more frequent visits)
  - Visit check interval: 1/3 day (8 hours)
- **Assignment:** Added to `icuContext` with `icu=true`

## ICU Respiratory Therapists

- **Count:**  $\text{icuCapacity} \times \text{icuRtsPerPatient} = 20 \times 0.1 = 2$  ICU RTs
- **Type:** ICURT
- **Visit pattern:**
  - Uses  $\text{Gamma}(\text{shape}=0.54, \text{scale}=20)$  distribution
  - Visit check interval: 1/3 day
- **Assignment:** Added to `icuContext` with `icu=true`

## Ward Therapists

All three types (RT, PT, OT) currently have staffing ratios set to 0.1 (default), so: - **Respiratory Therapists (RT):**  $\text{hospitalCapacity} \times 0.1 = 12$  RTs - **Physical Therapists (PT):**  $\text{hospitalCapacity} \times 0.1 = 12$  PTs - **Occupational Therapists (OT):**  $\text{hospitalCapacity} \times 0.1 = 12$  OTs

**Key difference for therapists:** - They visit patients from specific “needs” lists: `hospital.patientsNeedingRt/Pt/Ot` - Visit pattern:  $\text{Gamma}(\text{shape}=0.62, \text{scale}=61.7)$  distribution

## Simulation Execution - The Event Loop

Once initialization completes, the Repast Symphony scheduler begins executing events. The simulation runs for 365 ticks (days).

## Recurring Scheduled Events

### 1. **Builder.perShiftOperations()** - Every 0.5 Ticks

**When:** Runs at ticks 0.5, 1.0, 1.5, 2.0, etc. (twice per day, representing shift changes)

**What it does:** - Calls `hospital.setPatientNurseAssignments()` to reassign nurses to patients

**Nurse Assignment Algorithm:**

1. **Clear all existing nurse-patient assignments**
  - Removes all edges in the hospital network between nurses and patients
  - Doctor-patient assignments remain intact
2. **For each patient in the hospital:**
  - a. **Get appropriate nurse pool**
    - If patient is in Ward: get all ward nurses from `wardContext`
    - If patient is in ICU: get all ICU nurses from `icuContext`
  - b. **Sort nurses by current workload**
    - Nurses are sorted by their network degree (number of assigned patients)
    - This implements load balancing
  - c. **Find nurses with minimum workload**
    - Find the minimum degree among available nurses
    - Create list of all nurses tied for minimum degree
  - d. **Assign two nurses to the patient**
    - If 2+ nurses tied for minimum: randomly select 2
    - If only 1 nurse at minimum: assign that nurse, then find next-lowest workload nurse
    - Creates edges in `hospitalnet` connecting nurses to patient
    - Ensures each patient has 2 nurse assignments (when possible)

### 2. **Builder.daily()** - Every 1.0 Tick

**When:** Runs at ticks 1.0, 2.0, 3.0, etc. (once per day)

**What it does:** - Calls `hospital.resetTherapyNeeds()` to refresh therapy assignment lists

**Therapy Needs Reset:**

1. **Clear existing therapy lists**
  - `patientsNeedingOt.clear()`

- `patientsNeedingPt.clear()`
- `patientsNeedingRt.clear()`

## 2. Rebuild therapy lists from current patient population

- Loop through all current patients
- If `patient.needsOt` is true, add to `patientsNeedingOt`
- If `patient.needsPt` is true, add to `patientsNeedingPt`
- If `patient.needsRt` is true, add to `patientsNeedingRt`

**Why daily reset?** Ensures therapy assignment lists stay synchronized with current patient population, removing discharged patients and adding newly transferred patients.

## Stochastic Event Processes

In addition to the regular scheduled methods, several stochastic processes fire at random intervals:

### 3. Admission Process

**Frequency:** Exponential(0.05) distribution - mean of 0.05 days (~72 minutes) between admissions

**Process flow:**

#### 1. Fire event

- Increments total admissions counter
- Calls `hospital.createAndAdmitPatient()`
- Reschedules itself for next admission

#### 2. Create and Admit Patient

##### Capacity check

- Only admit if `patients.size() < bedCount (120)`

##### Patient creation

- Create new `Patient()` object
- Patient gets unique `agentId` from Agent base class
- Patient gets new `AgentDisease` object tracking disease state

##### ICU vs Ward admission

- Random draw: probability `icuAdmitProbability (0.15)`

- Additional check for ICU: ICU must have available beds (`inIcu.size() < icuBedCount`)
- **Default: 85%** of admissions go to Ward

#### **If admitted to Ward:**

##### **a. Disease importation check**

- Probability `admitImportationInfectionProbability` (0.01)
- If true: patient starts with `INFECTED` disease state
- Mark as imported, record infection date, start disease process

##### **b. Update patient attributes**

- Add to tracking lists: `patients`, `notInIcu`
- Set location: `admitLocation="Ward"`, `currentLocation="Ward"`
- Set admission time to current tick
- Mark `icuAdmit=false` attribute

##### **c. Schedule discharge**

- `nonIcuDischarger.scheduleDischarge(p)` creates one-time event
- Time drawn from `LogNormal(scale=0.768, shape=1.253)`
- Schedules call to `hospital.dischargePatient(p)`

##### **d. Assign therapy needs**

- Randomly assign based on probabilities:
  - `needsOt`: probability `needsOt` (0.1)
  - `needsRt`: probability `needsRt` (0.1)
  - `needsPt`: probability `needsPt` (0.1)

##### **e. Add to ward context**

- Add patient to `wardContext` for ward-specific operations

##### **f. Record admission**

- Append to `admissionData`: `patientId`, `admitTime`, `icuAdmit=false`, importation status

#### **If admitted to ICU:**

Process differs from Ward admission in these ways:

##### **a. Disease importation**

- Uses `admitImportationInfectionProbabilityICU` (0.01) - same probability as ward

##### **b. Tracking lists and location**

- Added to `inIcu` list instead of `notInIcu`
- Location set to "ICU" instead of "Ward"
- Mark `icuAdmit=true` attribute

c. **Discharge schedule**

- Uses `icuDischarger` with different distribution: `LogNormal(scale=0.820, shape=0.916)`

d. **Additional: Schedule potential transfer**

- **Key difference:** ICU patients get a transfer event scheduled
- `transferer.scheduleTransfer(p)` creates one-time event
- Time drawn from `LogNormal(scale=1.0, shape=0.5)`
- Schedules call to `hospital.transferPatient(p)`
- Note: Both transfer and discharge are scheduled; whichever fires first will execute

e. **Therapy needs**

- Different probabilities than ward:
  - `needsOt`: probability `needsOtIcu (0.1)` - same as ward
  - `needsRt`: probability `needsRtIcu (1.0)` - **ALL ICU patients need RT**
  - `needsPt`: probability `needsPtIcu (0.1)` - same as ward

f. **Context assignment**

- Added to `icuContext` instead of `wardContext`

**Add to therapy lists if needed**

- If patient needs OT: add to `patientsNeedingOt`
- If patient needs PT: add to `patientsNeedingPt`
- If patient needs RT: add to `patientsNeedingRt`

**Assign doctor**

- Calls `setPatientDoctorAssignments(p)`
- Uses similar load-balancing algorithm as nurse assignment
- Finds doctor (from appropriate context) with fewest current patients
- Adds edge in `hospitalnet` connecting doctor to patient
- Doctor assignment persists until patient discharge (unlike nurses who reassign every shift)

#### **4. PatientVisit Process**

**One process instance per HCW** - Each doctor, nurse, and therapist has their own visit process

**Frequency:** Varies by HCW type and location, using Gamma distributions

**Process flow:**

1. **Start**



- Calculate next event time from distribution
- Schedule one-time event to call `fire()`

## 2. Fire

- Calls `hgw.makeAVisit()`
- Reschedules itself with `start()` - creating continuous loop

## 3. Next Event Time Calculation

- Sample from HCW-specific Gamma distribution (in minutes)
- Add fixed room visit duration: **6.6** minutes
- Convert to days: multiply by `TimeUtils.MINUTE` constant
- Return: `currentTime + elapsedTime`

## 4. Make a Visit - Implementation varies by HCW type

The base class `HealthCareWorker.makeAVisit()` is overridden in each subclass:

### Doctors:

- Get the `hospitalnet` network from `hospital`
- Check if doctor has any assigned patients (`getDegree(this) > 0`)
- If yes: select random patient from those connected to this doctor (`getRandomAdjacent(this)`)
- **Key point:** Doctors only visit their ASSIGNED patients (assigned at patient admission)
- Check for transmission
- Record visit to `hospital.visitData`

### Nurses:

- Identical logic to doctors
- Get `hospitalnet` network
- Check if nurse has assigned patients
- Select random patient from those connected to this nurse
- **Key point:** Nurses only visit their ASSIGNED patients (assigned every 0.5 ticks at shift change)
- Check for transmission
- Record visit

### Therapists (RT/PT/OT):

- Check if `needsArray` has any patients (size > 0)
- `needsArray` is set to `hospital.patientsNeedingOt/Pt/Rt` for each therapist type
- Select random patient from the needs list
- **REMOVE patient from needs list** after selection

- **Key point:** Each patient is visited at most once per day by each therapist type
- The needs list is repopulated daily by `Builder.daily()` → `hospital.resetTherapyNeeds()`
- Check for transmission
- Record visit

#### ICU Respiratory Therapists (`IcuRt`):

- Get list of all current ICU patients (`hospital.inIcu`)
- If list is empty, return (no visit)
- Select random patient from ALL ICU patients (not using assignment network)
- **Key point:** ICU RTs can visit any ICU patient, not limited to assigned patients
- Check for transmission
- Record visit

#### Transmission check:

- Logic for all HCW types:
  - If HCW not contaminated AND patient not colonized/infected: no transmission possible
  - If HCW contaminated AND patient clean: potential HCW→patient transmission
  - If HCW clean AND patient colonized/infected: potential patient→HCW transmission
  - If both contaminated/infected: no additional transmission
- Note: Actual transmission logic is not fully implemented

#### Visit recording - All HCW types append to `hospital.visitData`:

- Format: `hcwId`, `hcwType`, `hcwDiseaseState`, `patientId`, `patientDiseaseState`, `patientLocation`, `visitTime`
- Creates comprehensive audit trail of all visits

## 5. Discharge Process

### Scheduled individually for each patient upon admission

**Timing:** - ICU patients: `LogNormal(scale=0.820, shape=0.916)` - Ward patients: `LogNormal(scale=0.768, shape=1.253)`

#### Process flow:

##### 1. Schedule discharge

- Called when patient is admitted
- Sample discharge time from distribution

- Schedule one-time event: `hospital.dischargePatient(p)`

## 2. Discharge patient

### Collect timing data

- Record admit time and current time for length-of-stay calculation

### Remove from hospital

- Remove from main context
- Remove from `patients` list
- Remove from `inIcu` or `notInIcu` list

### Create discharge record

- Create `DischargedPatient` object with:
  - `agentId`
  - `admitTime`
  - `dischargeTime` (current tick)
  - `died` flag (always false in current implementation)
  - `icuAdmit` flag (whether admitted to ICU)
  - `transferTime` (if transferred from ICU)
  - `admitLocation` (“ICU” or “Ward”)
  - `dischargeLocation` (current location at discharge)

### Store record

- Add to `dischargedPatients` list
- Add to context (for potential Repast probes/data collection)

## 6. Transfer Process

### Scheduled only for ICU-admitted patients

**Timing:** `LogNormal(scale=1.0, shape=0.5)`

### Process flow:

#### 1. Schedule transfer

- Called when patient is admitted to ICU
- Sample transfer time from distribution
- Schedule one-time event: `hospital.transferPatient(p)`

#### 2. Transfer patient

### Update location lists

- Remove from `inIcu` list
- Add to `notInIcu` list

#### Update patient state

- Set `currentLocation` = "Ward"
- Record `transferTime` (current tick)

**Note:** Patient's original discharge event (scheduled upon ICU admission) is NOT cancelled. If the ICU discharge event fires after transfer, it will still discharge the patient from their current location (Ward).

## Model Termination and Output

### Builder.endOfRun() - At Tick 365

**When:** Single execution at tick 365

**What it does:**

#### 1. Write output files

- Calls `writeSingleRunFiles()`

#### 2. End simulation

- Calls `RunEnvironment.getInstance().endRun()`
- Stops the scheduler

## Output File Generation

The model creates three output files:

### 1. discharged\_patients.txt

**Format:** CSV with header

```
agentId,admitTime,dischargeTime,icuAdmit,transferTime,admitLocation,dischargeLocation
```

**Content:** - One row per discharged patient - `transferTime` is 0.0 if patient was never transferred - `icuAdmit` indicates if patient was initially admitted to ICU - Enables length-of-stay analysis, transfer analysis, ICU vs ward utilization

**Generation:** - Loops through `hospital.getDischargedPatients()` list - Calls `DischargedPatient.toString()` for each record

## 2. visit\_data.txt

**Format:** CSV with header

hcwId,hcwType,hcwDiseaseState,patientId,patientDiseaseState,patientLocation,visitTime

**Content:** - One row per HCW visit to a patient - Records disease states of both HCW and patient at time of visit - Includes visit location (ICU or Ward) and exact time - Enables visit pattern analysis, HCW workload analysis, transmission chain reconstruction

**Generation:** - Writes the accumulated `hospital.visitData` StringBuffer - This buffer is appended to throughout the simulation

## 3. admission\_data.txt

**Format:** CSV with header

patientId,admitTime,icuAdmit,importation

**Content:** - One row per patient admission - `icuAdmit`: true if admitted to ICU, false if admitted to Ward - `importation`: true if patient arrived with imported MRSA infection - Enables admission pattern analysis and importation tracking

**Generation:** - Writes the accumulated `hospital.admissionData` StringBuffer - This buffer is appended to during admission

## Key Model Parameters

All parameters are defined in `Builder.java` and accessible via Repast GUI:

### Hospital Structure

- `hospitalCapacity` = 120 beds
- `icuCapacity` = 20 beds
- Ward capacity = 100 beds (derived)

## Admission-Discharge-Transfer (ADT)

- `admissionsRate` = 0.05 (mean inter-arrival time in days)
- Ward discharge: `LogNormal(scale=0.768, shape=1.253)`
- ICU discharge: `LogNormal(scale=0.820, shape=0.916)`
- `icuAdmitProbability` = 0.15 (15% of admissions go to ICU)
- `icuTransferProbability` = 0.1 (affects transfer timing, not probability)
- ICU→Ward transfer: `LogNormal(scale=1.0, shape=0.5)`

## Therapy Needs

- `needsRt` = 0.1 (ward), `needsRtIcu` = 1.0 (ICU) - respiratory therapy
- `needsPt` = 0.1 (both ICU and ward) - physical therapy
- `needsOt` = 0.1 (both ICU and ward) - occupational therapy

## Staffing Ratios (HCWs per bed)

- Ward nurses: 0.2
- ICU nurses: 0.5
- Ward doctors: 0.2
- ICU doctors: 0.3
- ICU respiratory therapists: 0.1
- Ward therapists (RT/PT/OT): 0.1 each

## HCW Visit Patterns (Gamma distributions, shape and scale parameters)

All times in minutes, converted to days via `TimeUtils.MINUTE`

**Ward:** - Nurse visits: `Gamma(0.54, 55.1)` + 6.6 min room time - Doctor visits: `Gamma(0.52, 90.7)` + 6.6 min room time - Therapist visits: `Gamma(0.62, 61.7)` + 6.6 min room time

**ICU:** - Nurse visits: `Gamma(0.54, 20)` + 6.6 min room time - Doctor visits: `Gamma(0.52, 35.3)` + 6.6 min room time

## Disease Parameters

- `admitImportationInfectionProbability` = 0.01 (ward)
- `admitImportationInfectionProbabilityICU` = 0.01 (ICU)
- `hhAdherenceBase` = 0.5 (hand hygiene compliance for all HCW types)
- `ppeAdherenceIfCp` = 0.5 (PPE/glove compliance)

## Disease Transmission Model (Partially Implemented)

The model includes disease tracking infrastructure but transmission logic is not fully implemented:

### Disease States (`DiseaseStates.java`)

- SUSCEPTIBLE - not colonized or infected
- COLONIZED - carrying MRSA but asymptomatic
- INFECTED - active MRSA infection

### Current Implementation

- Patients can be imported with INFECTED state on admission
- HCWs can be marked as contaminated
- Visit checking logic exists in `HealthCareWorker` class
- **However:** Actual transmission during visits is not fully implemented
  - `checkTransmissionToPatient()` returns true if HCW performs hand hygiene
  - `checkTransmissionToHcw()` returns false
  - `checkTransmission()` is empty

### Data Collection for Transmission Analysis

Even though transmission is not implemented, the model records: - HCW contamination status at each visit - Patient disease state at each visit - Visit timing and location This allows post-hoc transmission analysis or model extension.

### Summary of Model Dynamics

The model simulates a continuous flow hospital over **365** days:

1. **Patients arrive** stochastically (~20 admissions per day on average)
2. **Admitted to ICU (15%) or Ward (85%)** based on probability and bed availability
3. **Assigned to doctors** (once, at admission) using load-balancing
4. **Assigned to nurses** (twice daily, at shift changes) using load-balancing
5. **Visited by HCWs** throughout their stay according to stochastic schedules
6. **ICU patients may transfer** to ward after some time
7. **All patients eventually discharge** after length-of-stay drawn from location-specific distributions
8. **Visit data** and **discharge data** accumulate throughout simulation

## 9. **Output files** written at day **365** for analysis

The model generates realistic admission-discharge-transfer patterns and HCW workload distributions that can be analyzed using the R/Quarto scripts in the R/ directory.