

On Contemporary Models for Actuarial Use II: Principles

Angus S Macdonald^a and Stephen J Richards^b

Longevity 19, Amsterdam

17 September 2024

^aHeriot-Watt University and the Maxwell Institute for Mathematical Sciences

^bLongevitas Ltd.

Contents

1. Discrete time \Rightarrow complicated events!
2. Breaking down events — the Bernoulli ‘atom’
3. Building up events — the product integral
4. Data — the stochastic switch $Y(t)$
 - Survival models
 - Pseudo-Poisson models
 - True Poisson models

Contents

1. Discrete time \Rightarrow complicated events!
2. Breaking down events — the Bernoulli ‘atom’
3. Building up events — the product integral
4. Data — the stochastic switch $Y(t)$
 - Survival models
 - Pseudo-Poisson models
 - True Poisson models

Models: q -type and μ -type

Forfar, D.O., McCutcheon, J.J. & Wilkie, A.D. (1988). *On Graduation by Mathematical Formula*. Journal of the Institute of Actuaries, **115**, 1–149.

FMW graduated models using estimators of three parameters:

- q_x the one-year probability of death;
- μ_x the hazard rate*; or
- m_x the central rate of mortality.

* ‘force of mortality’ if you prefer

Models: q -type and μ -type

Two different roads to estimation:

1. q -type models:

- Inspired by life table *probability* q_x .
- Obvious statistical model Binomial . . .
- . . . or is it?

2. μ -type models:

- Inspired by *hazard rate* μ_x .
- Obvious statistical model Poisson . . .
- . . . or is it?

Both models have flaws, but those of the Binomial are more serious.

Models: q -type and μ -type

Flaws with simple models.

1. q -type Model

- Assumes E_x persons exposed for a whole year BUT ...
- ... some will leave before the year-end ...
- ... while others will join part-way through the year ...
- ... so we can't have a Binomial distribution.

2. μ -type Model

- Knowing we have M individuals in the study (as we usually do) ...
- ... the probability of more than M deaths is zero ...
- ... so we can't have a Poisson distribution.

Models: q -type and μ -type

Flaws with simple models.

1. q -type Model

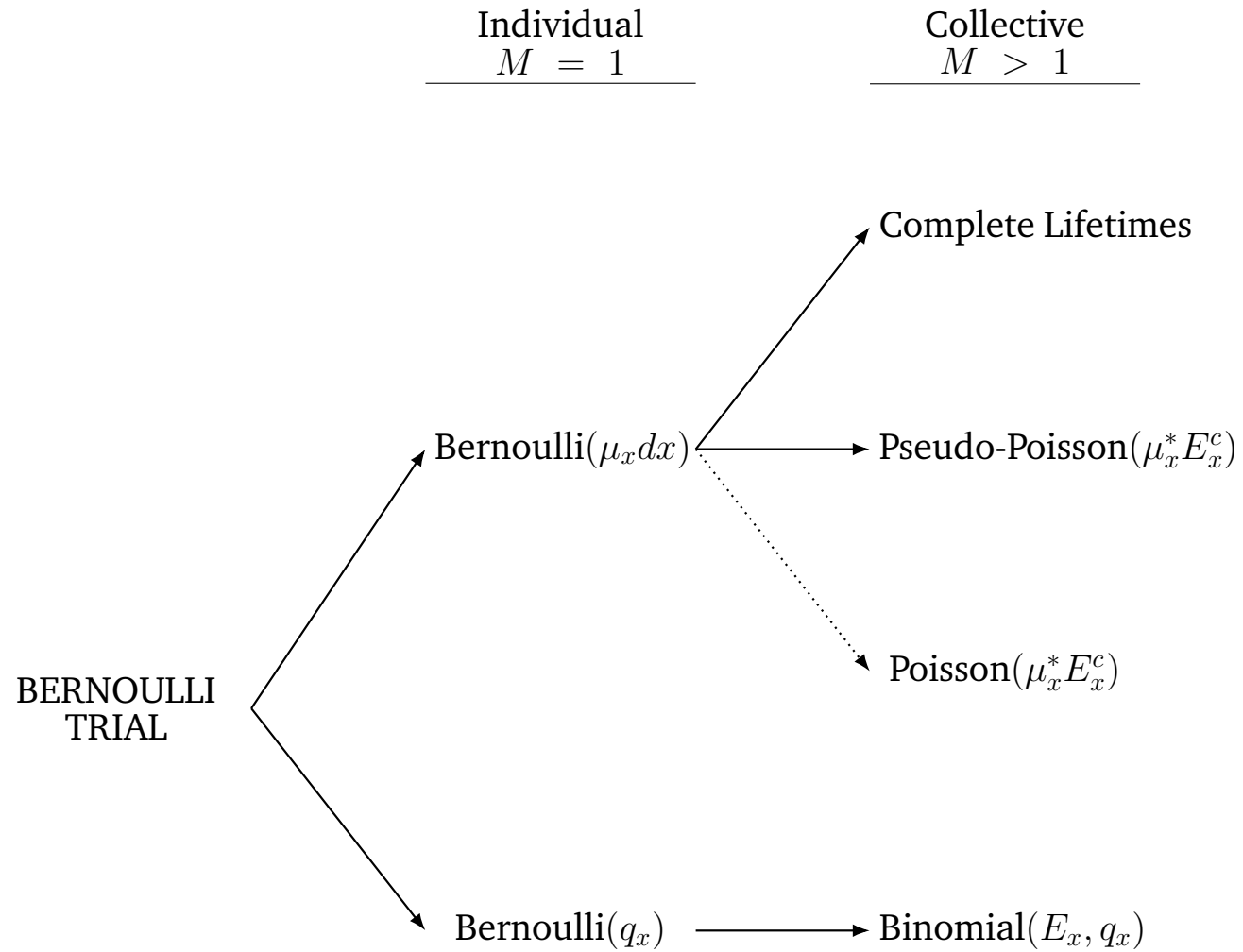
- Assumes E_x persons exposed for a whole year BUT ...
- ... some will leave before the year-end ...
- ... while others will join part-way through the year ...
- ... so we can't have a Binomial distribution.

2. μ -type Model

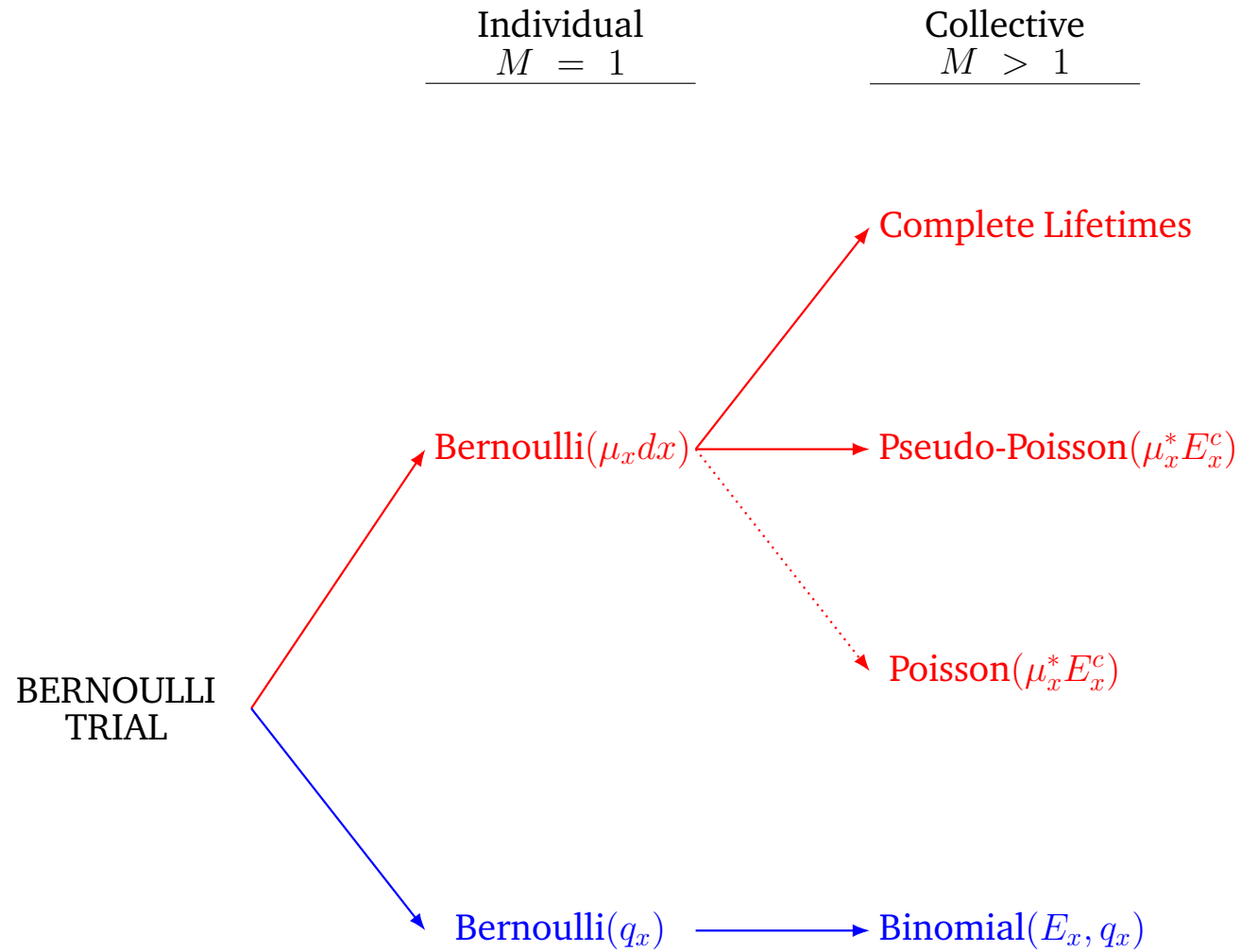
- Knowing we have M individuals in the study (as we usually do) ...
- ... the probability of more than M deaths is zero ...
- ... so we can't have a Poisson distribution.

'Fixing' the Binomial model leads us further into the weeds. Fixing the Poisson model leads to enlightenment!

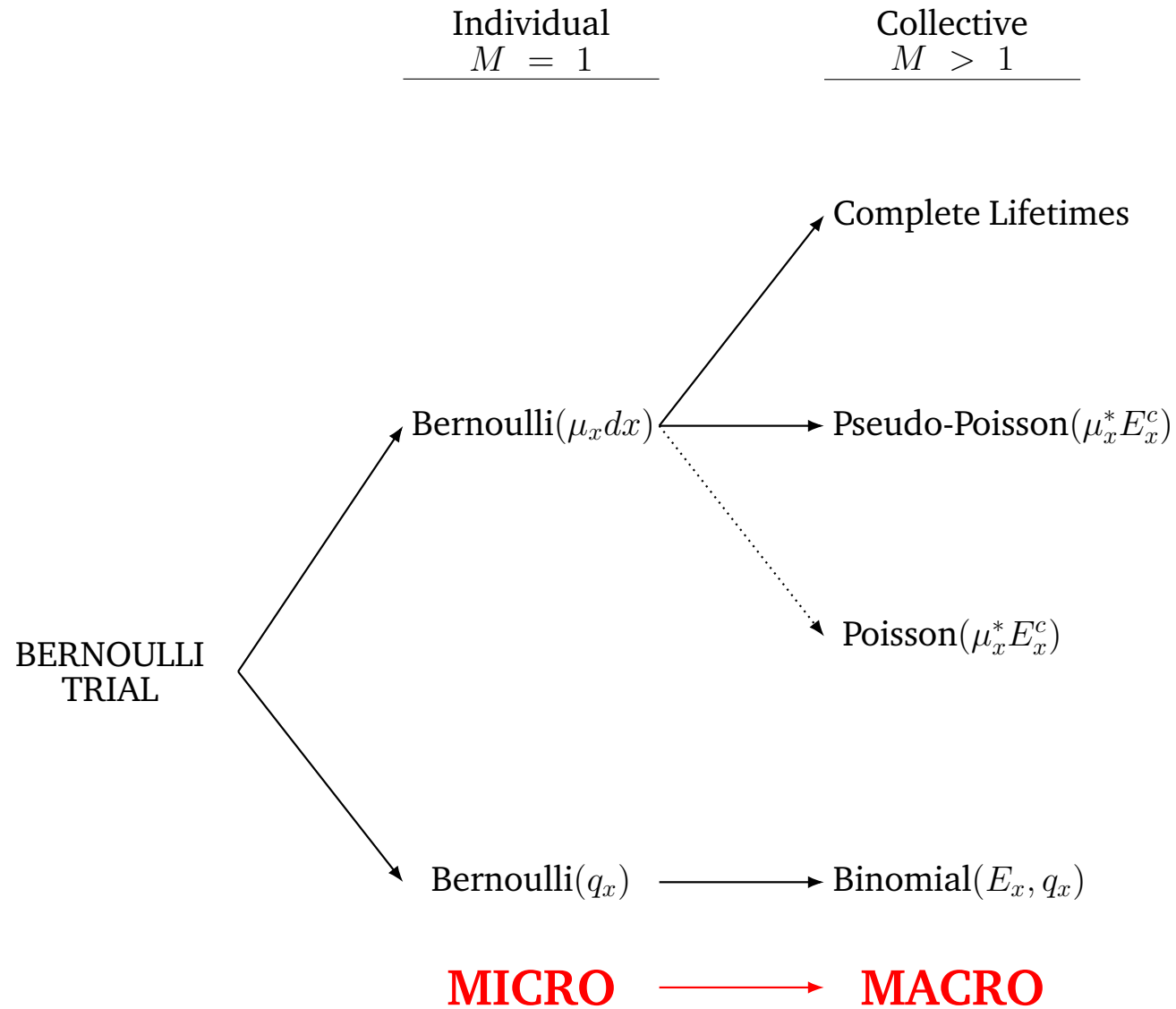
Bernoulli Family Tree



Bernoulli Family Tree



Bernoulli Family Tree



The Lower Branch: The Binomial/**Bernoulli** Model

Observe M lives $i = 1, 2, \dots, M$ for one year, define ‘indicator’ of death d_i :

$$d_i = \begin{cases} 1 & \text{if life } i \text{ dies} \\ 0 & \text{if life } i \text{ survives} \end{cases}$$

Binomial likelihood is:

$$\begin{aligned} L_i &\propto (1 - q_x)^{M - \sum d_i} (q_x)^{\sum d_i} \\ &= \prod_{i=1}^M (1 - q_x)^{1 - d_i} (q_x)^{d_i}. \end{aligned}$$

... a product of **Bernoulli** likelihoods for each life.

... But THIS Bernoulli Model is Still a Complicated Thing!

Define T_x = random lifetime of (x) and consider $p_x = P[T_x > 1]$:

Event $\{T_x > 1\}$ is highly **composite**:

$$\begin{aligned} p_x &= 1p_x \\ &= 0.5p_x \times 0.5p_{x+0.5} \\ &= 0.25p_x \times 0.25p_{x+0.25} \times 0.5p_{x+0.5} \\ &= 0.125p_x \times 0.125p_{x+0.125} \times 0.25p_{x+0.25} \times 0.5p_{x+0.5} \dots \\ &= \dots \text{and so on, } ad \text{ infinitum.} \end{aligned}$$

(Apologies to Zeno!)

In fact, event $\{T_x > 1\}$ is **infinitely composite**. Survival happens from moment to moment. And $q_x = P[T_x \leq 1]$ is **worse**.

Contents

1. Discrete time \Rightarrow complicated events!
2. Breaking down events — the Bernoulli ‘atom’
3. Building up events — the product integral
4. Data — the stochastic switch $Y(t)$
 - Survival models
 - Pseudo-Poisson models
 - True Poisson models

The Basic 'Atom' — An Infinitesimal Bernoulli Trial

The idea of the hazard rate μ_t is the infinitesimal:

$$P[t < T \leq t + dt \mid T > t] = \mu_t dt + o(dt) \approx \mu_t dt.$$

For convenience (re)define the indicator:

$$d_i = \Delta N_i(t) = \begin{cases} 1 & \text{if } t < T_i < t + dt \\ 0 & \text{otherwise} \end{cases}$$

$$P[\text{Obs. in } dt] = (1 - \mu_t dt)^{(1 - \Delta N_i(t))} (\mu_t dt)^{\Delta N_i(t)} = \text{Bernoulli trial.}$$

We have the **infinitesimal Bernoulli trial**. Not quite right yet, but let's pursue it . . .

Contents

1. Discrete time \Rightarrow complicated events!
2. Breaking down events — the Bernoulli ‘atom’
3. Building up events — the product integral
4. Data — the stochastic switch $Y(t)$
 - Survival models
 - Pseudo-Poisson models
 - True Poisson models

The Product Integral

Revision: $\lim_{n \rightarrow \infty} \left(1 + \frac{1}{n}\right)^n = e$ or $\lim_{n \rightarrow \infty} \left(1 - \frac{1}{n}\right)^n = e^{-1}$.

Let $(a_i, b_i]$ be the time interval under observation by life i . Then:

$$\begin{aligned}
 P[\text{Observation}_i] &= \underbrace{\prod_{(a_i, b_i]} (1 - \mu_t dt)^{(1 - \Delta N_i(t))} (\mu_t dt)^{\Delta N_i(t)}}_{\text{Product Integral}} \\
 &= \left(\prod_{(a_i, b_i]} (1 - \mu_t dt)^{(1 - \Delta N_i(t))} \right) \times (\mu_{b_i} dt)^{\Delta N_i(b_i)} \\
 &= \exp \left(- \int_{a_i}^{b_i} \mu_t dt \right) (\mu_{b_i} dt)^{\Delta N_i(b_i)}.
 \end{aligned}$$

Contents

1. Discrete time \Rightarrow complicated events!
2. Breaking down events — the Bernoulli ‘atom’
3. Building up events — the product integral
4. Data — the stochastic switch $Y(t)$
 - Survival models
 - Pseudo-Poisson models
 - True Poisson models

Data: The Stochastic Switch $Y(t)$

Define the process $Y^i(t)$:

$$Y^i(t) = \begin{cases} 1 & \text{if alive and under observation at time } t^- \\ 0 & \text{otherwise} \end{cases}$$

$Y^i(t)$ acts as a stochastic ‘switch’ depending on the status of (x) .

For example, $Y^i(t) \mu_t$ is a stochastic hazard rate.

$$Y^i(t) \mu_t = \begin{cases} \mu_t & \text{if alive and under observation at time } t^- \\ 0 & \text{otherwise} \end{cases}$$

Data: The Product Integral Likelihood

$$Y^i(t) = I_{\{\text{Life } i \text{ alive and under observation}\}} \cdot$$

$$(1 - Y^i(t) \mu_t dt)^{1 - \Delta N_i(t)} (Y^i(t) \mu_t dt)^{\Delta N_i(t)}$$

MICRO: The ‘atom’ of all Poisson-type likelihoods:

$$\begin{aligned} L_i = P[\text{Observation}_i] &= \prod_{[0, \infty)} (1 - Y^i(t) \mu_t dt)^{(1 - \Delta N_i(t))} (Y^i(t) \mu_t dt)^{\Delta N_i(t)} \\ &= \underbrace{\exp\left(-\int_0^\infty Y^i(t) \mu_t dt\right)}_{P[\text{Survival}]} \underbrace{(Y^i(b_i) \mu_{b_i} dt)^{\Delta N_i(b_i)}}_{P[\text{Death}]} \end{aligned}$$

MACRO: Universal Poisson-type likelihood

Poisson-type Models I: Survival Models

M lives, lifetimes T_1, T_2, \dots, T_M , life i observed on $(a_i, b_i]$.

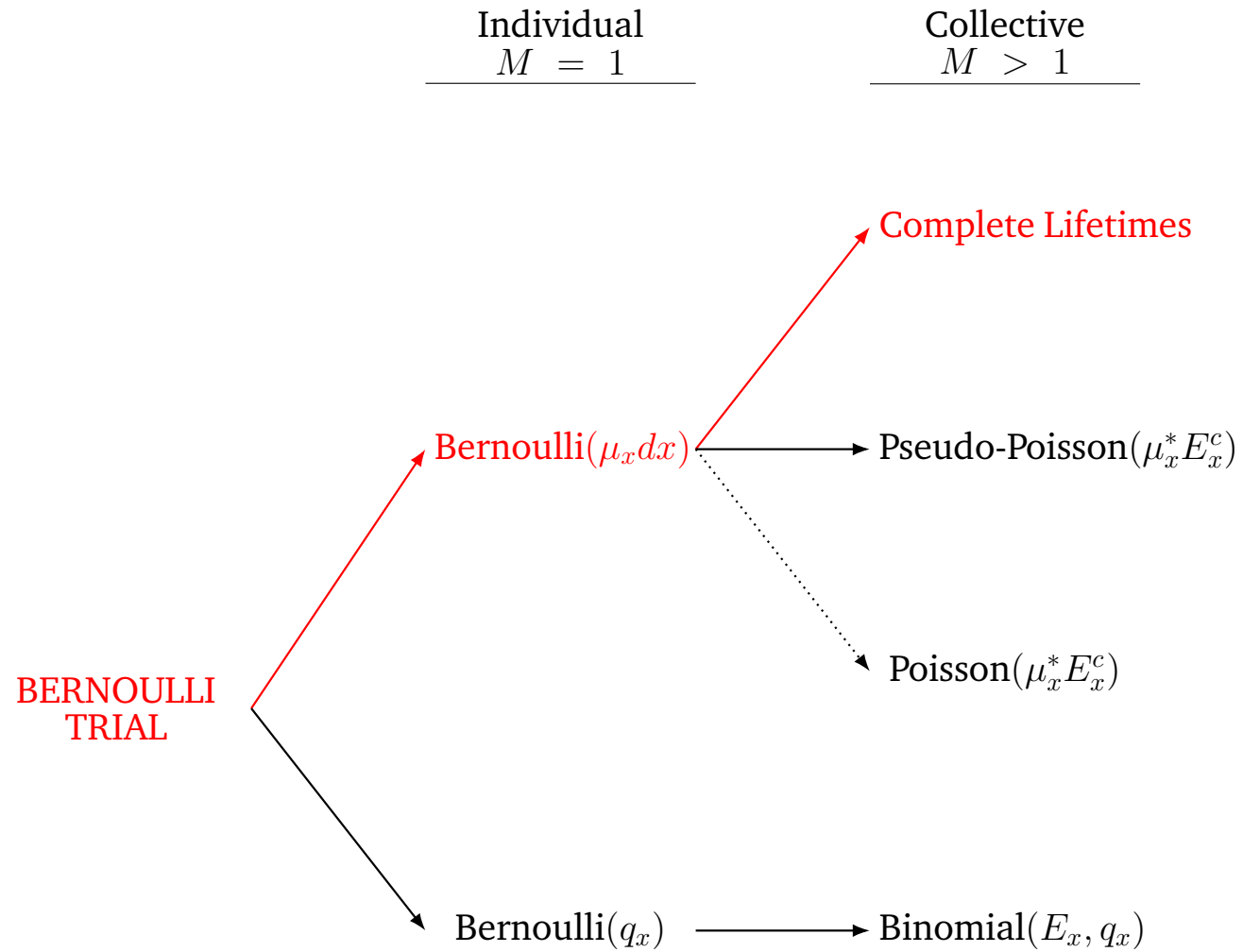
$\mu_x^\theta =$ Parametric hazard rate on $[0, \infty)$.

$Y^i(t) = I_{\{\text{Life } i \text{ alive and under observation}\}}$.

$$\begin{aligned} L &= \prod_i L_i = \prod_i \prod_{[0, \infty)} (1 - Y^i(t) \mu_t^\theta dt)^{(1 - \Delta N_i(t))} (Y^i(t) \mu_t^\theta dt)^{\Delta N_i(t)} \\ &= \prod_i \exp \left(- \int_{a_i}^{b_i} Y^i(t) \mu_t^\theta dt \right) (Y^i(b_i) \mu_{b_i}^\theta dt)^{\Delta N_i(b_i)}. \end{aligned}$$

INDIVIDUAL DATA/COMPLETE OBSERVED LIFETIMES/SURVIVAL MODEL

Bernoulli Family Tree



Poisson-type Models II: Pseudo-Poisson Models

M lives under observation, life i on $(a_i, b_i]$.

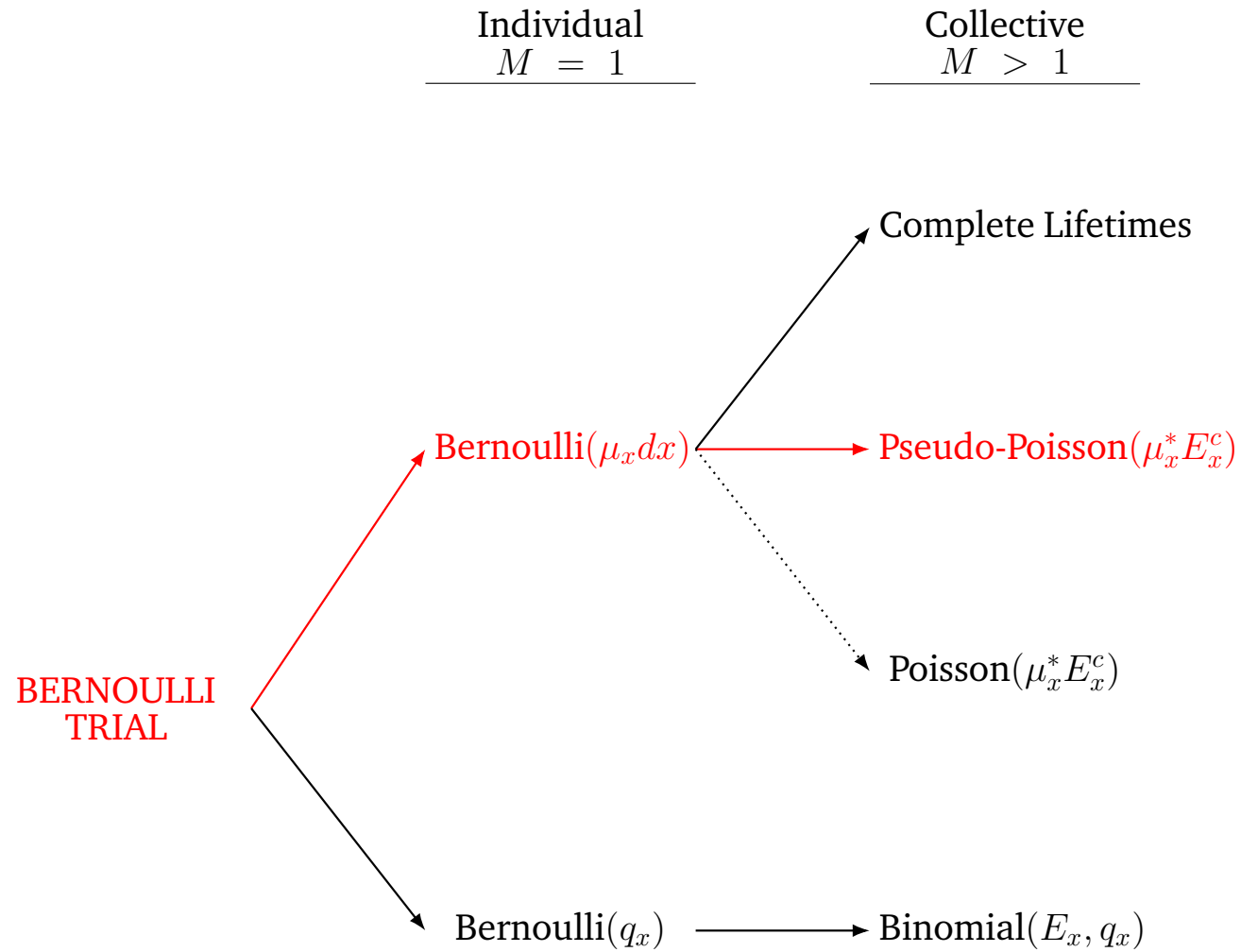
μ_x^* = Constant hazard rate on $(x, x + 1]$.

$Y_x^i(t) = I_{\{\text{Life } i \text{ alive and recorded as 'active' on } (a_i, b_i] \cap (x, x + 1]\}}$.

$$\begin{aligned}
 L &= \prod_x \prod_i L_{x,i}^* = \prod_x \prod_i \prod_{[0, \infty)} (1 - Y_x^i(t) \mu_x^* dt)^{(1 - \Delta N_i(t))} (Y_x^i(t) \mu_x^* dt)^{\Delta N_i(t)} \\
 &= \prod_x \prod_i \exp\left(-\int_x^{x+1} Y_x^i(t) \mu_x^* dt\right) \prod_{[0, \infty)} (Y_x^i(t) \mu_x^* dt)^{\Delta N_i(t)} \\
 &= \prod_x \exp(E_x^c) (\mu_x^*)^{D_x}.
 \end{aligned}$$

M known, E_x^c random variable \Rightarrow GROUPED DATA/PSEUDO-POISSON

Bernoulli Family Tree



Poisson-type Models III: True Poisson Models

Random M lives under observation, life i on $(a_i, b_i]$.

μ_x^* = Constant hazard rate on $(x, x + 1]$.

$\tilde{Y}_x^i(t) = I_{\{\text{Life } i \text{ alive and recorded as 'active' on } (a_i, b_i] \cap (x, x + 1]\}}$ constrained so that E_x^c is a pre-determined constant.

$$\begin{aligned}
 L &= \prod_x \prod_i L_{x,i}^* \propto \prod_x \prod_i \prod_{[0, \infty)} (1 - \tilde{Y}_x^i(t) \mu_x^* dt)^{(1 - \Delta N_i(t))} (\tilde{Y}_x^i(t) \mu_x^* dt)^{\Delta N_i(t)} \\
 &= \prod_x \prod_i \exp \left(- \int_x^{x+1} \tilde{Y}_x^i(t) \mu_x^* dt \right) \prod_{[0, \infty)} (\tilde{Y}_x^i(t) \mu_x^* dt)^{\Delta N_i(t)} \\
 &= \prod_x \exp (E_x^c) (\mu_x^*)^{D_x}.
 \end{aligned}$$

M random variable, E_x^c known \Rightarrow GROUPED DATA/TRUE POISSON

Bernoulli Family Tree

