

## List of topics in this lecture

- Feynman-Kac formula for the forward equation, path integral  $u(x, t)$ , interpretation of path integral as mass density of the surviving cell population at time  $t$ , governing equation of  $u(x, t)$
- An application of Feynman-Kac formula: reconstructing potential  $V(x)$  from a set of sample paths of particle position; exploring non-equilibrium with an applied force; modeling the effect of applied force as a fatality/growth rate

## Recap

### Feynman-Kac formula for the backward equation

We are back to the time-dependent SDE

$$dX = b(X, t)dt + \sqrt{a(X, t)}dW$$

$u(x, t, T)$  is defined by the Feynman-Kac path integral formula

$$u(x, t, T) = E \left( \exp \left( - \int_t^T \psi(X(s), s) ds \right) f(X(T)) \middle| X(t) = x \right)$$

### Meaning of $u(x, t, T)$

$\psi(x, s)$  = fatality/growth rate of a cell at time  $s$  with  $X(s) = x$ .

$f(x)$  = reward for a cell surviving to time  $T$  with  $X(T) = x$ .

$u(x, t, T)$  = expected reward at final time  $T$  per unit population at time  $t$

**Each cell of the population gets its own reward.** The growth increases the population size and increases the reward for the population.

### Governing equation for $u(x, t, T)$ and the FVP

$$\begin{cases} 0 = u_t + b(x, t)u_x + \frac{1}{2}a(x, t)u_{xx} - \psi(x, t)u \\ u(x, t, T)|_{t=T} = f_0(x) \end{cases}$$

The solution of the FVP is given by the Feynman-Kac path integral formula.

**Feynman-Kac formula for the forward equation (continued)**

$$u(x, t) = E \left( \delta(X(t) - x) \exp \left( - \int_0^t \psi(X(s), s) ds \right) \right)$$

Items of the discussion:

- 1) We need to explain the  $\delta$  function in the average.
- 2) We need to derive the governing equation for  $u(x, t)$ .
- 3) We need to explain the meaning of  $u(x, t)$  and discuss the distribution of  $X(0)$ .

Item #1: Explaining the  $\delta$  function in the average

View 1: 
$$u(x, t) = \lim_{\Delta x \rightarrow 0} \frac{1}{\Delta x} E \left( I_{[x, x+\Delta x]}(X(t)) \exp \left( - \int_0^t \psi(X(s), s) ds \right) \right)$$

View 2: 
$$\int h(x) u(x, t) dx = E \left( h(X(t)) \exp \left( - \int_0^t \psi(X(s), s) ds \right) \right) \quad \text{for any } h(x)$$

(We use this view to derive the equation for  $u(x, t)$ ).

Item #2: Derivation of the governing equation for  $u(x, t)$

We start with View #2 at  $(t+\Delta t)$ . We apply the forward view on  $[0 \rightarrow t+\Delta t]$ .

$[0 \rightarrow t+\Delta t]$  is divided into  $[0 \rightarrow t]$  and  $[t \rightarrow t+\Delta t]$ .

$$\underbrace{\int h(x) u(x, t+dt) dx}_{\text{LHS}} = \underbrace{E \left( h(X(t) + dX) \exp \left( - \int_0^{t+dt} \psi(X(s), s) ds \right) \right)}_{\text{RHS}}$$

$$= E \left( h(X(t) + dX) \exp \left( - \int_t^{t+dt} \psi(X(s), s) ds \right) \times \exp \left( - \int_0^t \psi(X(s), s) ds \right) \right)$$

In the RHS, we expand  $h(X(t) + dX)$  and  $\exp \left( - \int_t^{t+dt} \psi(X(s), s) ds \right)$

$$\begin{aligned} \text{RHS} &= E \left( \left[ h(X(t)) + h'(X(t))dX + \frac{1}{2}h''(X(t))(dX)^2 \right] (1 - \psi(X(t), t)dt) \right. \\ &\quad \left. \times \exp \left( - \int_0^t \psi(X(s), s) ds \right) \right) \\ &= E \left( \left[ h(X(t)) + h'(X(t))dX + \frac{1}{2}h''(X(t))(dX)^2 - h(X(t))\psi(X(t), t)dt \right] \right. \\ &\quad \left. \times \underbrace{\exp \left( - \int_0^t \psi(X(s), s) ds \right)}_{\text{independent of } dX} \right) \end{aligned}$$

In the RHS, the average is over  $\{X(s), 0 \leq s \leq t+\Delta t\}$ . We use the law of total expectation to rewrite is as averaging over  $(dX(t) | X(t))$  and then over  $\{X(s), 0 \leq s \leq t\}$ .

$$E_{\{X(s), 0 \leq s \leq t+\Delta t\}}(Q) = E_{\{X(s), 0 \leq s \leq t\}} \left( E_{dX(t)}(Q | X(t)) \right)$$

We use the moments of  $(dX(t) | X(t))$ :

$$E_{dX(t)}(h'(X(t))dX | X(t)) = h'(X(t))b(X(t), t)dt$$

$$E_{dX(t)}(h''(X(t))(dX)^2 | X(t)) = h''(X(t))a(X(t), t)dt$$

$$\begin{aligned} \text{RHS} = E \left[ \left( h(X(t)) + h'(X(t))b(X(t), t)dt + \frac{1}{2}h''(X(t))a(X(t), t)dt - h(X(t))\psi(X(t), t)dt \right) \right. \\ \left. \times \exp \left( - \int_0^t \psi(X(s), s)ds \right) \right] \end{aligned}$$

View #2 (method of test function) of  $u(x, t)$  gives

$$E \left( g(X(t)) \exp \left( - \int_0^t \psi(X(s), s)ds \right) \right) = \int g(x)u(x, t)dx \quad \text{for any } g(x).$$

In the RHS, we set respectively  $g(x) = h(x)$  for term 1,  $g(x) = h'(x)b(x, t)$  for term 2, ...,

$$\text{RHS} = \int \left( h(x) + h'(x)b(x, t)dt + \frac{1}{2}h''(x)a(x, t)dt - h(x)\psi(x, t)dt \right) u(x, t)dx$$

In the RHS, we integrate by parts; in the LHS, we expand  $u(x, t+dt)$ .

$$\text{RHS} = \int h(x) \left( u(x, t) - \left( b(x, t)u \right)_x dt + \frac{1}{2} \left( a(x, t)u \right)_{xx} dt - \psi(x, t)u dt \right) dx$$

$$\text{LHS} = \int h(x)u(x, t+dt)dx = \int h(x)(u(x, t) + u_t dt)dx$$

Subtracting  $\int h(x) u(x, t)dx$ , dividing by  $dt$ , and taking the limit as  $dt \rightarrow 0$ , we obtain

$$\underbrace{\int h(x)u_t dx}_{\text{LHS}} = \underbrace{\int h(x) \left( - \left( b(x, t)u \right)_x + \frac{1}{2} \left( a(x, t)u \right)_{xx} - \psi(x, t)u \right) dx}_{\text{RHS}}$$

Since LHS = RHS for arbitrary test function  $h(x)$ , we arrive at

$$u_t = - \left( b(x, t)u \right)_x + \frac{1}{2} \left( a(x, t)u \right)_{xx} - \psi(x, t)u$$

This is the governing PDE for  $u(x, t)$ .

It is the forward equation with a fatality/growth term.

Item #3: Meaning of  $u(x, t)$

View #1 of  $u(x, t)$  gives

$$u(x, t) = \lim_{\Delta x \rightarrow 0} \frac{1}{\Delta x} E_{\omega} \left( I_{[x, x+\Delta x]}(X(t, \omega)) \exp \left( - \int_0^t \psi(X(s, \omega), s) ds \right) \right)$$

The average is based on an ensemble  $\{X(s, \omega), \omega \in \Omega\}$ . It is worthwhile to emphasize that the evolution of  $X(s)$  is governed solely by the SDE

$$dX = b(X, t)dt + \sqrt{a(X, t)}dW, \quad \text{independent of } \psi(x, s)$$

In particular,  $X(s)$  does not die or split. In  $u(x, t)$ , the fatality/growth effect of  $\psi(x, s)$  is reflected in the factor  $\exp \left( - \int_0^t \psi(X_j(s), s) ds \right)$ .

To interpret  $u(x, t)$ , we consider **the “cell” population associated with  $X(s)$** . The cell population includes the fatality/growth effect of  $\psi(x, s)$ , in the form of termination/addition of cells.

For a cell at time  $s$  with  $X(s) = z$ , the outcome of the cell in  $[s, s+\Delta s]$  is

$$\begin{aligned} & \text{(Outcome in } [s, s+\Delta s] \mid \text{having survived to } s \text{ with } X(s) = z) \\ &= \begin{cases} \text{fatality with prob} = \psi(z, s)\Delta s & \text{if } \psi(z, s) > 0 \\ \text{split into two with prob} = (-\psi(z, s))\Delta s & \text{if } \psi(z, s) < 0 \end{cases} \end{aligned}$$

We write  $u(x, t)$  in terms of the cell population and associated  $X(s)$ .

$$\begin{aligned} u(x, t) &= \lim_{\Delta x \rightarrow 0} \frac{1}{\Delta x} E_{\omega} \left( \# \text{ of cells surviving to time } t \text{ with } X(t, \omega) \in [x, x+dx] \right) \\ &= \text{mass density in } x \text{ of the surviving cell population at time } t. \end{aligned}$$

The size of surviving population is different from the size of starting population.

Initial condition for  $u(x, t)$

$$u(x, 0) = f_0(x) = \text{mass density in } x \text{ of the starting cell population.}$$

Remarks on the Feynman-Kac formula for the forward equation

- The evolution of  $X(s)$  is governed solely by the SDE, independent of  $\psi(x, s)$ .  $X(s)$  does not die or split.
- In the Feynman-Kac formula, the average is over an ensemble of  $X(s)$ . The fatality/growth effect of  $\psi(x, s)$  is reflected in the factor  $\exp \left( - \int_0^t \psi(X_j(s), s) ds \right)$ .  
The Feynman-Kac formula **is good for calculating  $u(x, t)$  from an ensemble of  $X(s)$** .
- Given a LARGE set of sample paths of  $\{X_j(s), j = 1, 2, \dots, N\}$ , we have

$$u(x, t) \approx \frac{1}{N \cdot \Delta x} \sum_{X_j(t) \in [x, x+\Delta x]} \exp\left(-\int_0^t \psi(X_j(s), s) ds\right)$$

The calculation does not directly involve the cell population.

- To interpret  $u(x, t)$  defined in the Feynman-Kac formula, we follow the surviving cell population associated with  $X(s)$ , which includes the effect of fatality/growth.
- Although we interpret  $\psi(x, s)$  as the fatality/growth rate, the Feynman-Kac formula is well defined for any  $\psi(x, s)$ , not associated with any physical fatality/growth.
- When  $\psi(x, s)$  is not associated with any physical fatality/growth, the cell population exists only in our mathematical imagination (see the application below).

### An application of Feynman-Kac formula

The big picture: When  $b(x, t)$  in the SDE is unknown but a LARGE set of sample paths is available, we can use the Feynman-Kac formula to calculate the unknown  $b(x, t)$ .

Consider a particle diffusing in a potential well.

$X(t)$ : position of particle at time  $t$

$V(x)$ : static potential well

The stochastic motion is governed by the over-damped Langevin equation.

$$dX = -\frac{D}{k_B T} V'(X) dt + \sqrt{2D} dW$$

After non-dimensionalization, we have

$$dX = -V'(X) dt + \sqrt{2} dW$$

What we can measure (data)

A large set of sample paths  $\{X_j(t), j = 1, 2, \dots, N\}$

Goal:

To reconstruct potential  $V(x)$  from the data

Method:

We use Feynman-Kac formula to reconstruct potential  $V(x)$ .

Items of the discussion

- Forward equation and equilibrium of probability density
- Estimating potential from equilibrium measurements
- A practical issue with equilibrium data
- Exploring non-equilibrium with an applied force

E. Steps in constructing potential  $V(x)$

Item A:

Forward equation (Fokker-Planck equation) of  $dX = -V'(X)dt + \sqrt{2} dW$

Let  $\rho(x, t)$  = probability density of  $X$  at time  $t$ .

Note: probability density = mass density of a population of one path.

The time evolution of  $\rho(x, t)$  is governed by the forward equation

$$\rho_t = -\left(b(x)\rho\right)_x + \frac{1}{2}\left(a(x)\rho\right)_{xx}, \quad a(x) = 2, \quad b(x) = -V'(x)$$

$$\Rightarrow \rho_t = \left(V'(x)\rho\right)_x + \rho_{xx}$$

We write the forward equation in conservation form

$$\rho_t = -\frac{\partial}{\partial x} J(x, t), \quad J(x, t) \equiv -\left(V'(x)\rho + \rho_x\right)$$

where  $J(x, t)$  is the probability flux.

Equilibrium distribution

At equilibrium, the probability flux must be identically zero.

$$J(x) = 0, \quad \text{for all } x$$

$$\Rightarrow V'(x)\rho + \rho_x = 0$$

$$\Rightarrow \left(\exp(V(x))\rho(x)\right)' = 0$$

$$\Rightarrow \exp(V(x))\rho(x) = \text{const}$$

$$\Rightarrow \rho^{(\text{eq})}(x) \propto \exp(-V(x))$$

As expected, the equilibrium is the **Maxwell-Boltzmann distribution**.

Caution: A steady state is different from equilibrium.

When  $J(x) = \text{const} \neq 0$  for all  $x$ , we still have  $\rho = \rho(x)$ , independent of  $t$ .

It is called a steady state, which is different from equilibrium.

At equilibrium, we must have  $J(x) = 0$  for all  $x$ .

**In the terminology of deterministic dynamical systems, “steady state” and “equilibrium” are usually not distinguished.**

Item B:

### Estimating potential from equilibrium measurements

Suppose the system is at equilibrium and we measure a large set of sample paths  $\{X_j(t), j = 1, 2, \dots, N\}$ .

The equilibrium density  $\rho^{(eq)}(x)$  can be calculated as

$$\rho^{(eq)}(x) \approx \frac{1}{N \cdot \Delta x} \sum_{X_j(t_k) \in [x, x+\Delta x]} 1 \quad \text{at a particular time level } t_k$$

where  $N$  is the number of sample paths.

To fully utilize the data set, we average  $\rho^{(eq)}(x)$  over all  $t_k$ 's

$$\rho^{(eq)}(x) \approx \frac{1}{K_T} \sum_{k=1}^{K_T} [\rho^{(eq)}(x) \text{ estimated at } t_k]$$

where  $K_T$  is the number of time levels.

Recall that  $\rho^{(eq)}(x) \propto \exp(-V(x))$ . We write potential  $V(x)$  as

$$V(x) = -\log \rho^{(eq)}(x) + \underbrace{C}_{\text{additive constant}}$$

### Item C:

#### A practical issue with equilibrium data

Unfortunately, the approach of using only equilibrium measurements does not work well. It requires an impractically large amount of data.

At equilibrium, a region of high  $V(x)$  value is visited only very infrequently.

$$\rho^{(eq)}(x) \propto \exp(-V(x))$$

Consider a set of discrete sites (intervals of  $x$ ). For a site with probability  $10^{-8}$ , we need to sample  $10^9$  times to get 10 visits to that particular site. It is practically impossible to accurately estimate  $\rho^{(eq)}(x)$  in a region of high  $V(x)$  value.

Remedy: We need to perturb the system to non-equilibrium.

### Item D:

#### Exploring non-equilibrium with an applied force

Let  $F(t)$  be the applied force (non-dimensionalized).

In experiments,  $F(t)$  is controlled. In AFM (Atomic Force Microscopy) experiments, the force is controlled by moving an actuator to stretch an elastic link.

$$F^{(AFM)}(t) = k \int_0^t u(s) ds$$

where  $k$  = spring constant;  $u(s)$  velocity of actuator at time  $s$ .

Stochastic differential equation in the presence of an applied force

The applied force tilts the static potential. At time  $t$ , the tilted potential is

$$H(x, t) = V(x) - \underbrace{F(t) \cdot x}_{\text{Effect of applied force}}$$

Replacing  $V(x)$  with  $H(x, t)$ , we get the new SDE.

$$dX = -H_x(X, t)dt + \sqrt{2} dW$$

In the presence of an applied force, potential  $H(x, t)$  changes with time. As a result, the system is not at equilibrium and the Boltzmann distribution does not apply.

Nevertheless we consider a “hypothetical” density,  $\rho^{(F)}(x, t)$ , defined as

$$\rho^{(F)}(x, t) \equiv \frac{1}{Z} \exp(-H(x, t)) = \frac{1}{Z} \exp(-V(x) + F(t) \cdot x)$$

where  $Z = \int \exp(-V(x)) dx$

Remark:

- We call  $\rho^{(F)}(x, t)$  “hypothetical” density because it is not the mass density of any physical population. In particular,  $\rho^{(F)}(x, t)$  is NOT the probability density of particle position. In the discussion below, we interpret  $\rho^{(F)}(x, t)$  as the mass density of a “hypothetical” cell population, whose existence/fatality/growth is only in our mathematical imagination.

Advantage of working with  $H(x, t)$  and  $\rho^{(F)}(x, t)$

With a properly designed force schedule  $F(t)$ , a region of relatively high  $V(x)$  value becomes a region of relatively low  $H(x, t)$  value in the tilted potential.

In this way, different regions of  $V(x)$  can be very well explored/sampled at different time  $t$  with a time-dependent force schedule  $F(t)$ .

Item E:

Steps in constructing potential  $V(x)$

1. Find the governing PDE for  $\rho^{(F)}(x, t)$
2. Identify the “fatality/growth” term  $\psi(x, s)$  in the PDE
3. Express  $\rho^{(F)}(x, t)$  in the Feynman-Kac formula



4. Use the Feynman-Kac formula to calculate  $\rho^{(F)}(x, t)$  from a set of sample paths.
5. Determine potential  $V(x)$  from  $\rho^{(F)}(x, t)$ .

**Step 1:** Find the governing PDE for  $\rho^{(F)}(x, t)$

Let  $\rho(x, t)$  be the density of particle position in the presence of applied force  $F(t)$ .

$\rho(x, t)$  is NOT the same as  $\rho^{(F)}(x, t)$ .

$$\rho(x, t) \neq \rho^{(F)}(x, t)$$

Stochastic motion of particle is governed by

$$dX = -H_x(X, t)dt + \sqrt{2} dW, \quad H(x, t) \equiv V(x) - F(t) \cdot x$$

The forward equation (Fokker-Planck equation) for  $\rho(x, t)$  is

$$\frac{\partial \rho}{\partial t} = \frac{\partial}{\partial x} \left( H_x(x, t) \rho + \frac{\partial}{\partial x} \rho \right)$$

We write the Fokker-Planck equation in terms of a differential operator.

$$\rho_t = L_{\{H\}}[\rho] \quad (\text{FP\_forced})$$

$$\text{where} \quad L_{\{H\}}[\cdot] = \frac{\partial}{\partial x} \left( H_x(x, t) \cdot + \frac{\partial}{\partial x} \cdot \right)$$

We find the governing equation for the “hypothetical” density  $\rho^{(F)}(x, t)$ .

$$\rho^{(F)}(x, t) \equiv \frac{1}{Z} \exp(-H(x, t)) = \frac{1}{Z} \exp(-V(x) + F(t) \cdot x), \quad Z = \int \exp(-V(x)) dx$$

Substitute  $\rho^{(F)}(x, t)$  into operator  $L_{\{H\}}[\cdot]$  and into operator  $\partial/\partial t$

$$L_{\{H\}}[\rho^{(F)}(x, t)] = \frac{\partial}{\partial x} \left( H_x(x, t) \rho^{(F)}(x, t) + \frac{\partial}{\partial x} \rho^{(F)}(x, t) \right) = 0$$

$$\rho_t^{(F)}(x, t) = F'(t)x \cdot \rho^{(F)}(x, t)$$

It follows that  $\rho^{(F)}(x, t)$  satisfies

$$\rho_t^{(F)} = L_{\{H\}}[\rho^{(F)}] + \underbrace{F'(t)x \cdot \rho^{(F)}}_{\text{fatality/growth}}$$

**Step 2:** Identify the “fatality/growth” term  $\psi(x, s)$  in the PDE

$\rho^{(F)}(x, t)$  satisfies the forward equation with a fatality/growth term

$$\rho_t^{(F)} = L_{\{H\}}[\rho^{(F)}] - \psi(x, t) \cdot \rho^{(F)}, \quad \psi(x, t) = -F'(t)x$$

**Step 3:** Express  $\rho^{(F)}(x, t)$  using the Feynman-Kac formula

$$\rho^{(F)}(x, t) = E \left( \delta(X(t) - x) \exp \left( - \int_0^t \psi(X(s), s) ds \right) \right), \quad \psi(x, s) = -F'(s)x$$

$$\Rightarrow \rho^{(F)}(x, t) = E \left( \delta(X(t) - x) \exp \left( \int_0^t F'(s)X(s) ds \right) \right)$$

**Remark:** The ensemble of  $X(s)$  is good for calculating  $\rho^{(F)}(x, t)$ . For that calculation, we don't need the hypothetical cell population that includes the stochastic addition/termination of cells.

**Step 4:** Calculate  $\rho^{(F)}(x, t)$  from a set of sample paths.

Suppose we apply force  $F(t)$  and measure a set of sample paths  $\{X_j(s), j = 1, 2, \dots, N\}$ .

At each time level  $t_k$ ,  $\rho^{(F)}(x, t_k)$  can be calculated using the Feynman-Kac formula.

$$\rho^{(F)}(x, t_k) \approx \frac{1}{N \cdot \Delta x} \sum_{X_j(t_k) \in [x, x + \Delta x]} \exp \left( \int_0^{t_k} F'(s)X_j(s) ds \right) \quad \text{at each time level } t_k$$

where  $N$  is the number of sample paths.

**Step 5:** Determine potential  $V(x)$  from  $\rho^{(F)}(x, t)$

Note that  $\rho^{(F)}(x, t)$  and  $\rho^{(eq)}(x)$  are related by

$$\begin{aligned} \rho^{(eq)}(x) &= \frac{1}{Z} \exp(-V(x)) \\ \rho^{(F)}(x, t) &= \frac{1}{Z} \exp(-V(x) + F(t)x) \\ \Rightarrow \rho^{(eq)}(x) &= \rho^{(F)}(x, t) \exp(-F(t)x) \end{aligned}$$

Once  $\rho^{(F)}(x, t_k)$  is obtained at a time level  $t_k$ , we use it to calculate a sample version of equilibrium density  $\rho^{(eq)}(x)$ .

$$\rho^{(eq)}(x) = \rho^{(F)}(x, t_k) \exp(-F(t_k)x) \quad \text{at each time level } t_k$$

Then we average the sample versions of  $\rho^{(eq)}(x)$  over all  $t_k$ 's.

$$\rho^{(eq)}(x) \approx \frac{1}{K_T} \sum_{k=1}^{K_T} \rho^{(F)}(x, t_k) \exp(-F(t_k)x)$$

where  $K_T$  is the number of time levels.

Once  $\rho^{(eq)}(x)$  is accurately estimated, we write potential  $V(x)$  as

$$V(x) = -\log \rho^{(eq)}(x) + \underbrace{C}_{\text{additive constant}}$$

### Remarks on constructing potential $V(x)$

1. The “hypothetical” density  $\rho^{(F)}(x, t)$  contains potential  $V(x)$  and applied force  $F(t)$ , which allows us to extract potential  $V(x)$  from  $\rho^{(F)}(x, t)$  once  $\rho^{(F)}(x, t)$  is obtained.
2.  $\rho^{(F)}(x, t)$  satisfied the forward equation with a fatality/growth term  $\psi(x, t) = -F'(t)x$ . This hypothetical fatality/growth term exists only in our mathematical imagination. It does not correspond to the fatality/growth of any physical process.
3. The path integral expression (Feynman-Kac formula) of  $\rho^{(F)}(x, t)$  allows us to calculate  $\rho^{(F)}(x, t)$  from sample paths  $\{X_j(s), j = 1, 2, \dots, N\}$  and applied force  $F(t)$ .
4. Once  $\rho^{(F)}(x, t)$  is obtained, we extract potential  $V(x)$  from  $\rho^{(F)}(x, t)$ .