### Markov chains with rewards

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### Lifetime accumulation

- imagine accumulating something over lifetime
- call it a "reward"



(much like kaiten sushi)



### Individual lives

- trajectories through the life cycle
- probabilities given by transition matrix

$$\mathbf{P} = \left(\begin{array}{c|c} \mathbf{U} & \mathbf{0} \\ \mathbf{M} & \mathbf{I} \end{array}\right)$$

• trajectories (lives) are stochastic



# Examples of rewards

reward	lifetime accumulation
a year of life	longevity
a year of healthy life	healthy longevity
an offspring	lifetime reproductive output
income	lifetime income



### **Analysis**

transition matrix

$$\mathbf{P} = \left(\begin{array}{c|c} \mathbf{U} & \mathbf{0} \\ \hline \mathbf{M} & \mathbf{I} \end{array}\right)$$

define a reward matrix

$$\mathbf{R}(i,j)$$
 = reward for transition  $j \rightarrow i$ 

but rewards are random, so define the moments of reward

$$\mathbf{R}_m(i,j) = m$$
th moment of reward  $m = 1, 2, 3, ...$ 

assume no rewards for the dead

$$\mathbf{R}_m(i,j) = 0$$
 if  $j$  an absorbing state



### **Analysis**

$$\begin{array}{rcl} E(\widetilde{\rho}) & = & \widetilde{\rho}_1 \\ V(\widetilde{\rho}) & = & \widetilde{\rho}_2 - \widetilde{\rho}_1 \circ \widetilde{\rho}_1 \\ SD(\widetilde{\rho}) & = & \sqrt{V(\widetilde{\rho})} \end{array}$$

(and skewness, kurtosis, etc.)



### Analysis<sup>1</sup>

 $\widetilde{\rho}_i$  = vector of *i*th moments of lifetime rewards

$$\begin{split} \widetilde{\rho}_1 &= \mathbf{N}^\mathsf{T} \mathbf{Z} (\mathbf{P} \circ \mathbf{R}_1)^\mathsf{T} \mathbf{1}_{s+1} \\ \widetilde{\rho}_2 &= \mathbf{N}^\mathsf{T} \left[ \mathbf{Z} (\mathbf{P} \circ \mathbf{R}_2)^\mathsf{T} \mathbf{1}_{s+1} + 2 (\mathbf{U} \circ \widetilde{\mathbf{R}}_1)^\mathsf{T} \widetilde{\rho}_1 \right] \\ \widetilde{\rho}_3 &= \mathbf{N}^\mathsf{T} \left[ \mathbf{Z} (\mathbf{P} \circ \mathbf{R}_3)^\mathsf{T} \mathbf{1}_{s+1} + 3 (\mathbf{U} \circ \widetilde{\mathbf{R}}_2)^\mathsf{T} \widetilde{\rho}_1 + 3 (\mathbf{U} \circ \widetilde{\mathbf{R}}_1)^\mathsf{T} \widetilde{\rho}_2 \right] \\ \text{where} \\ & \mathbf{N} &= (\mathbf{I} - \mathbf{U})^{-1} \\ & \mathbf{Z} &= \left( \mathbf{I}_{\tau \times \tau} \mid \mathbf{0}_{\tau \times \alpha} \right) \\ & \widetilde{\mathbf{R}}_m &= \mathbf{R}_m (1 : \tau, 1 : \tau) \end{split}$$

### Healthy longevity: prevalence<sup>2</sup>

- prevalence is probability of having condition of interest
- prevalence measured from cross-sectional data
- health expectancy (HE), via Sullivan method
- Sullivan method for health expectancy

$$\ell(x) = P[\text{survival to age } x]$$

$$v(x)$$
 = prevalence of health at age  $x$ 

$$HE(a) = \frac{1}{\ell(a)} \int_{a}^{\infty} \ell(x) v(x) dx$$

= E (healthy years remaining at age a)

treat health condition as a reward

¹van Daalen and Caswell 2017, Theoretical Ecology, Thm. 1. 2 > 4 2 > 2 < 9 < @

<sup>&</sup>lt;sup>2</sup>Caswell and Zarulli 2018, Population Health Metrics

### Reward matrices for prevalence

$$v_j$$
 = prevalence in age class  $j$ 

then

reward 
$$j \rightarrow i = \begin{cases} 1 & \text{with probability } v_j \\ 0 & \text{with probability } 1 - v_j \end{cases}$$

so

$$\mathbf{R}_1 = \left(egin{array}{ccc|c} v_1 & \cdots & v_\omega & 0 \ dots & dots & dots \ v_1 & \cdots & v_\omega & 0 \ \hline v_1/2 & \cdots & v_\omega/2 & 0 \end{array}
ight)$$

and

$$\mathbf{R}_2 = \mathbf{R}_3 = \cdots = \mathbf{R}_1$$

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### Types of health rewards

- binary (e.g., disability-free)
  - Bernoulli distribution
- counts (e.g., number of hospitalizations)
  - maybe Poisson distribution
- quantitative measures (e.g., grip strength)
  - empirical measurement of moments
- amount of life lost (e.g., in DALY)
  - · moments calculated from Markov chain model



# Stochasticity in healthy longevity: the missing ingredient

- stochasticity in trajectories
- stochasticity in health outcomes (prevalence)
- variance among individuals in lifetime experience



### SHARE survey data

- Survey of Health, Ageing and Retirement in Europe
- wave 4, 2011, data for Germany, Sweden, France, Denmark, Switzerland, Belgium, Czech Republic, Portugal, and Estonia
- prevalence of disability (binary): moments calculated from Bernoulli distribution
- grip strength<sup>3</sup> (continuous): moments calculated from empirical individual-level measures
- measurements only reliable to age 90

<sup>&</sup>lt;sup>3</sup>Affects mortality, myocardial infarction, stroke, cardiovascular mortality 📱 🗸 a c

# An example: healthy longevity from SHARE data<sup>4</sup>

 $v_j$  = prevalence of disability-freedom in age class j

Rewards

$$r_{ij} = \begin{cases} 1 & \text{with probability } v_j \\ 0 & \text{with probability } 1 - v_j \end{cases}$$

# 

### Disability-free longevity (Belgium)

		M	en	Wo	Women		
		55	75	55	75		
mean	L	24.5	10.1	27.9	11.8		
	HL	20.6	7.6	22.2	7.9		
SD	L	9.4	5.1	8.5	5.1		
	HL	7.6	3.8	6.4	3.2		
CV	L	0.38	0.50	0.31	0.40		
	HL	0.37	0.50	0.29	0.41		
Sk	L	-0.75	-0.38	-1.34	-0.89		
	HL	-0.80	-0.28	-1.35	-0.51		



# See the paper for more results



<sup>&</sup>lt;sup>4</sup>Caswell and Zarulli 2018, Population Health Metrics

### Healthy longevity: incidence models<sup>5</sup>

- incidence refers to transitions among health states
- · requires longitudinal data
- age-specific health transition matrices ...
- lead to age-stage multistate Markov chains
- health is part of the i-state, not just a prevalence
- basic questions
  - occupancy
  - longevity (total, healthy)
  - eventual fate
  - accumulation of rewards
  - life lost due to causes
  - · sensitivity analysis

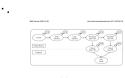
### Multistate model: ages and health stages



x = 1



x = 2



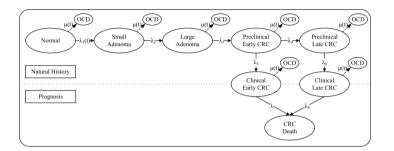
 $\mathbf{X} = \omega$ 



### An example: a model for colorectal cancer<sup>6</sup>

BMC Cancer 2006. 6:136

http://www.biomedcentral.com/1471-2407/6/136



<sup>6</sup>Wu et al. 2006, BMC Cancer



### How to measure "healthy longevity"

- 1. **Occupancy of health states.** How much life spent in a specified health condition, including combinations of ages and health stages?
- 2. Transitions among health states.
- 3. The "value" of occupancy of health states.
- 4. The "value" of transitions among states.



<sup>&</sup>lt;sup>5</sup>Caswell and van Daalen 2021, Demographic Research → ⟨⟨⟨⟨⟨⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩ ⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩ ⟨⟨⟨⟩⟩⟩⟩⟩

### Constructing age × stage matrix models<sup>7</sup>

 $\mathbf{U}_i$  = age-specific state transitions

 $\mathbb{U} = \mathsf{block} \, \mathsf{diagonal} \, \mathsf{matrix}$ 

 $\mathbf{D}_{j}$  = stage-specific age transitions

 $\mathbb{D}$  = block diagonal matrix

**K** = vec-permutation matrix

Then

$$\widetilde{\mathbf{U}} = \mathbf{K}^{\mathsf{T}} \mathbb{D} \mathbf{K} \mathbb{U}$$

$$\widetilde{\mathbf{P}} = \left( \begin{array}{c|c} \widetilde{\mathbf{U}} & \mathbf{0} \\ \hline \mathbf{M} & \mathbf{I} \end{array} \right)$$

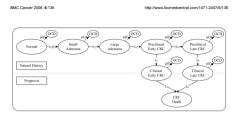
### Longevity by age and stage: the health matrix

Create an array  ${\bf H}$  to specify age-stage combinations of interest

Stages
healthy
small adenoma
large adenoma
preclinical early CRC
preclinical late CRC
clinical early CRC
clinical late CRC

1	2	Age 3	es 4	 ω

### Define "healthy" longevity in the CRC model



### For example:

- 1. **Cancer-free longevity.** Time spent in any of the cancer-free stages (1–3) over a lifetime.
- 2. **Old age clinical cancer.** The time spent in any of the clinical cancer stages (6–7) after a specified age (65 years)



### e.g., Cancer-free longevity

Put a 1 in cells that count as "healthy"

	Ages					
Stage	1	2	3	4		$\omega$
normal cells	1	1	1	1	1	1
small adenoma	1	1	1	1	1	1
large adenoma	1	1	1	1	1	1
preclinical early CRC						
preclinical late CRC						
clinical early CRC						
clinical late CRC						

(zeros elsewhere)



### Old age clinical cancer

Stages healthy small adenoma large adenoma preclinical early CRC preclinical late CRC clinical early CRC clinical late CRC

	$egin{array}{cccccccccccccccccccccccccccccccccccc$						
,	1	2	3	4		$\omega$	
L							
;							
				1	• • •	1	
				1	• • •	1	

(zeros elsewhere)



### How many definitions of health are there?

7 cancer stages  $\times$  50 ages = 350 combinations

Number of definitions of health  $= 2^{350} \approx 10^{105}$ 

Compare this to  $10^{24}$  stars in the visible universe

### How many definitions of health are there?





### **Reward matrices**

Including partial occupancy when transitions into or out of states occur

Define

 $\mathbf{r} = \mathsf{vec}\,\mathbf{X}$ 

Then

 $\widetilde{\mathbf{R}}_1 = \mathbf{r}\mathbf{r}^{\mathsf{T}} + 0.5 \, (\sim \mathbf{r}) \, (\mathbf{r}^{\mathsf{T}}) + 0.5 \, (\mathbf{r}) \, (\sim \mathbf{r}^{\mathsf{T}})$ 

and

$$\begin{array}{rcl} \widetilde{R}_2 & = & \widetilde{R}_1 \circ \widetilde{R}_1 \\ \widetilde{R}_3 & = & \widetilde{R}_1 \circ \widetilde{R}_1 \circ \widetilde{R}_1 \end{array}$$

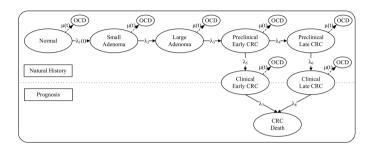
$$\widetilde{\mathbf{R}}_3 = \widetilde{\mathbf{R}}_1 \circ \widetilde{\mathbf{R}}_1 \circ \widetilde{\mathbf{R}}$$



### Colorectal cancer

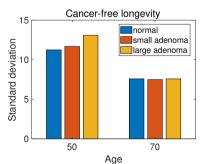
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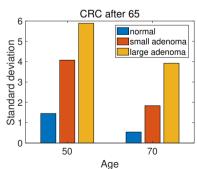
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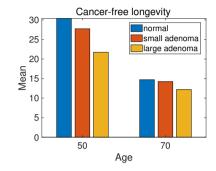
# Standard deviation healthy longevity

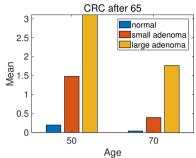






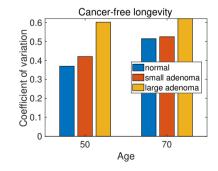
### Mean healthy longevity

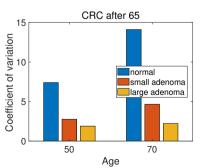






# Coefficient of variation of healthy longevity







# Concluding thoughts: MCWR

- very powerful tool for analyzing variation in lifetime experiences
- a fixation on expectancies neglects important aspects related to risk
- Markov chains with rewards give all the moments of lifetime accumulation
- in the case of health, provides limitless defintion of "healthy" longevity: any combination of health stages and ages
- extensions 

   value or cost of occupancy, transitions into
   or out of stages, components of variance, decomposition of
   differences

