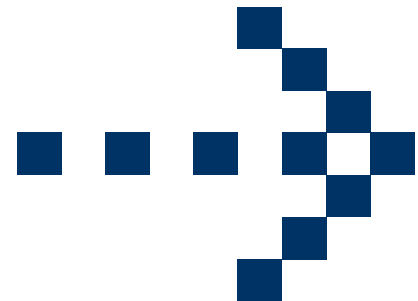


Use of discrete event simulation in health service planning: a luxury or necessity?

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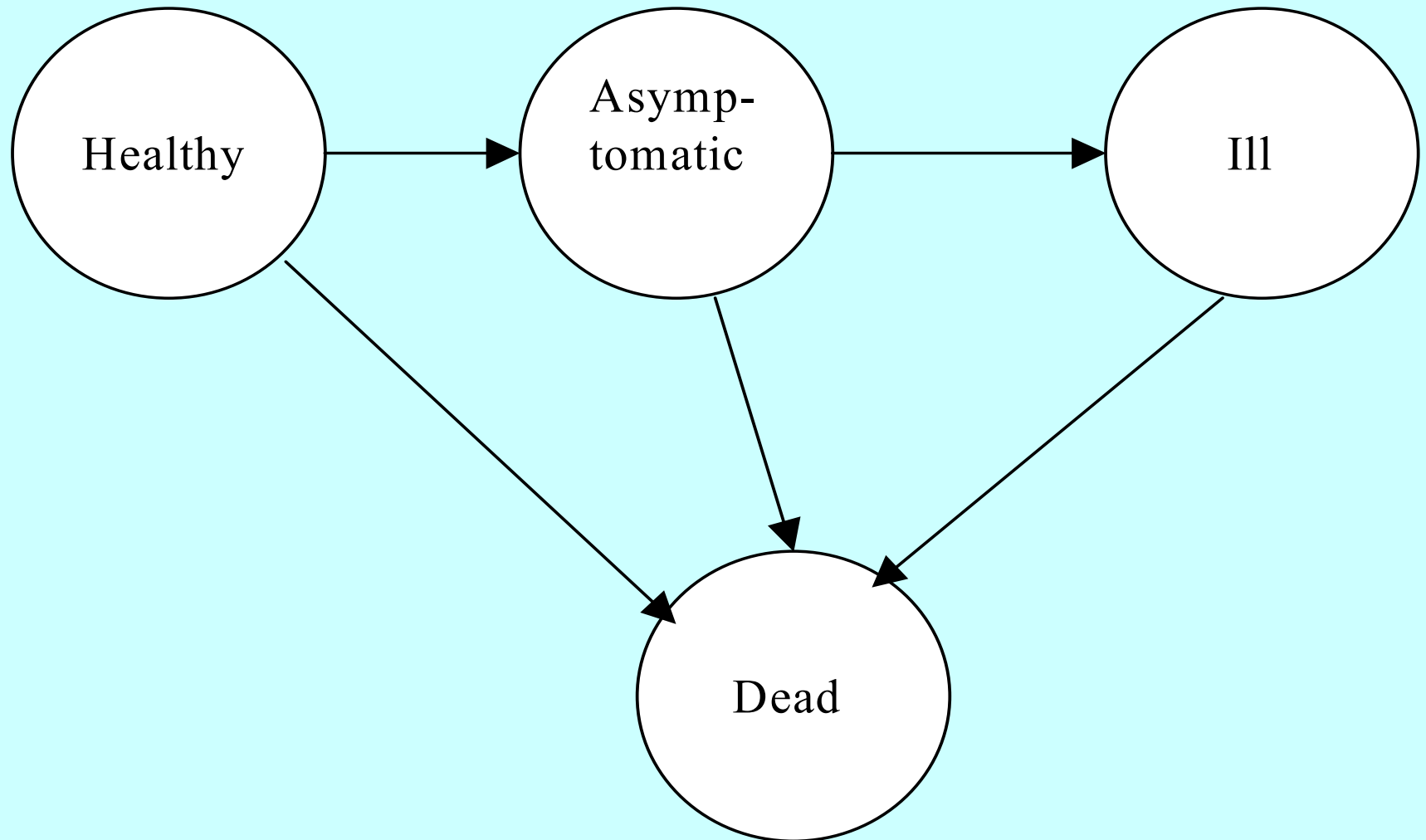
Health service planning problems

- ▣▣▣ What services should be plan for patients with end-stage renal failure?
- ▣▣▣ Is it cost-effective to give CHD patients statins?
- ▣▣▣ Should we screen for diabetic retinopathy or breast cancer?

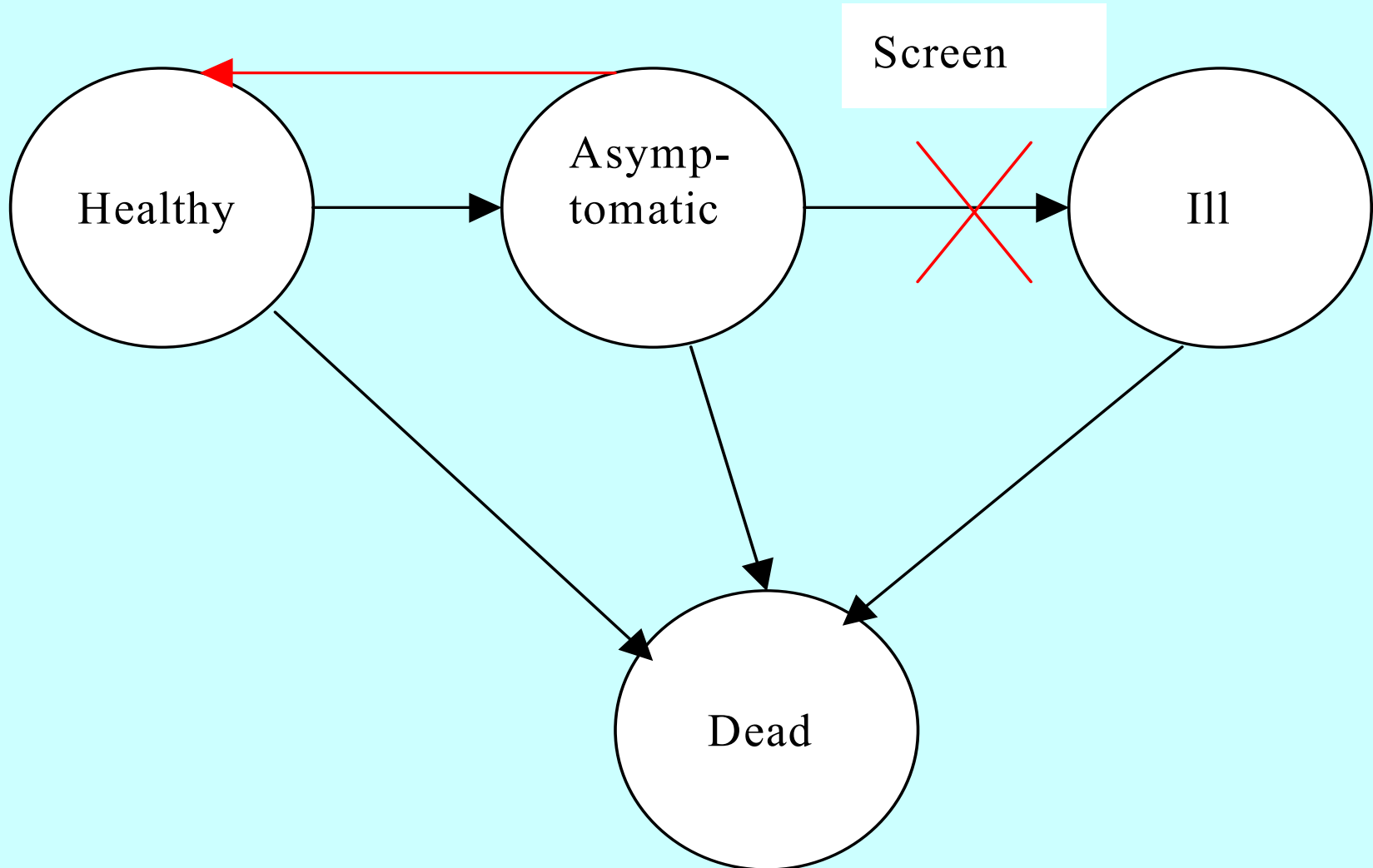
General characteristics

- ▣▣▣ Individuals, often patients, move from state to state;
- ▣▣▣ The states may be different states of health or opportunities for treatment;
- ▣▣▣ Patient characteristics e.g. age are important;
- ▣▣▣ The populations are large and the difference in outcome may be small and occur over a long period of time.

How many remain alive in each state in each year?



Screening – what benefits are there? More people? More quality of life? More cost?



Output measures

- ▣ Cost
- ▣ Deaths prevented
- ▣ Life years saved
- ▣ QALYS (quality adjusted life years) gained
- ▣ Cost per life year saved
- ▣ Cost per QALY gained

Markov models

- ❑❑❑ Classifies system into states
- ❑❑❑ Need to know transition probabilities from state to state
- ❑❑❑ Assumes population is homogeneous
- ❑❑❑ Assumes probability of moving from one state to the next is independent of time spent in that state – Markov assumption.

Probabilities of moving on in a fixed time period
Death is an absorbing state

States	Healthy	Asymptomatic	Ill	Dead
Healthy	p_{11}	p_{12}	0	p_{14}
Asymptomatic	0	p_{22}	p_{23}	p_{24}
Ill	0	0	p_{33}	p_{34}
Dead	0	0	0	1

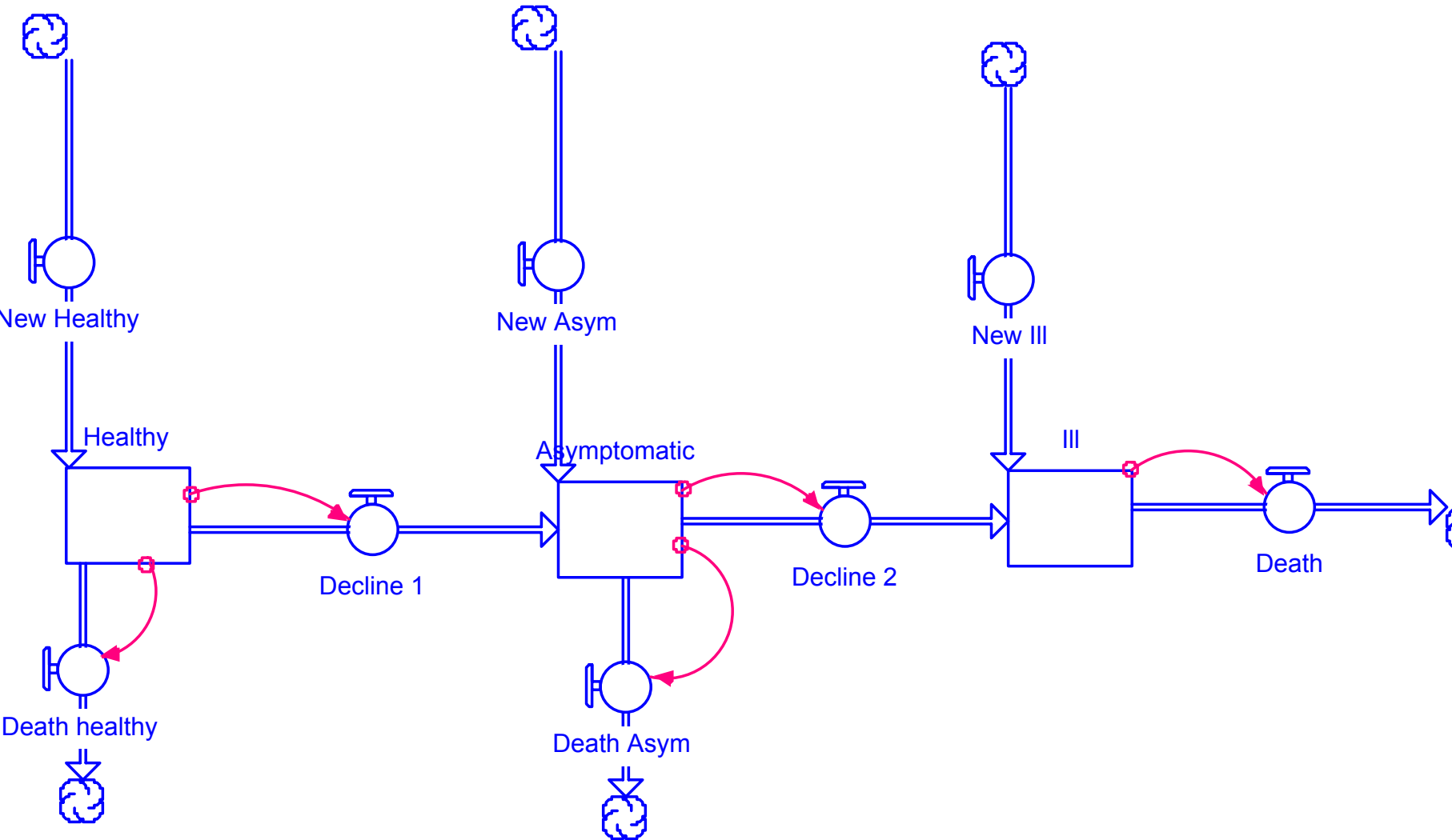
$$N(t) = N(t-1) P + M(t) I$$

Where N is the vector of the number in each state at time t ,

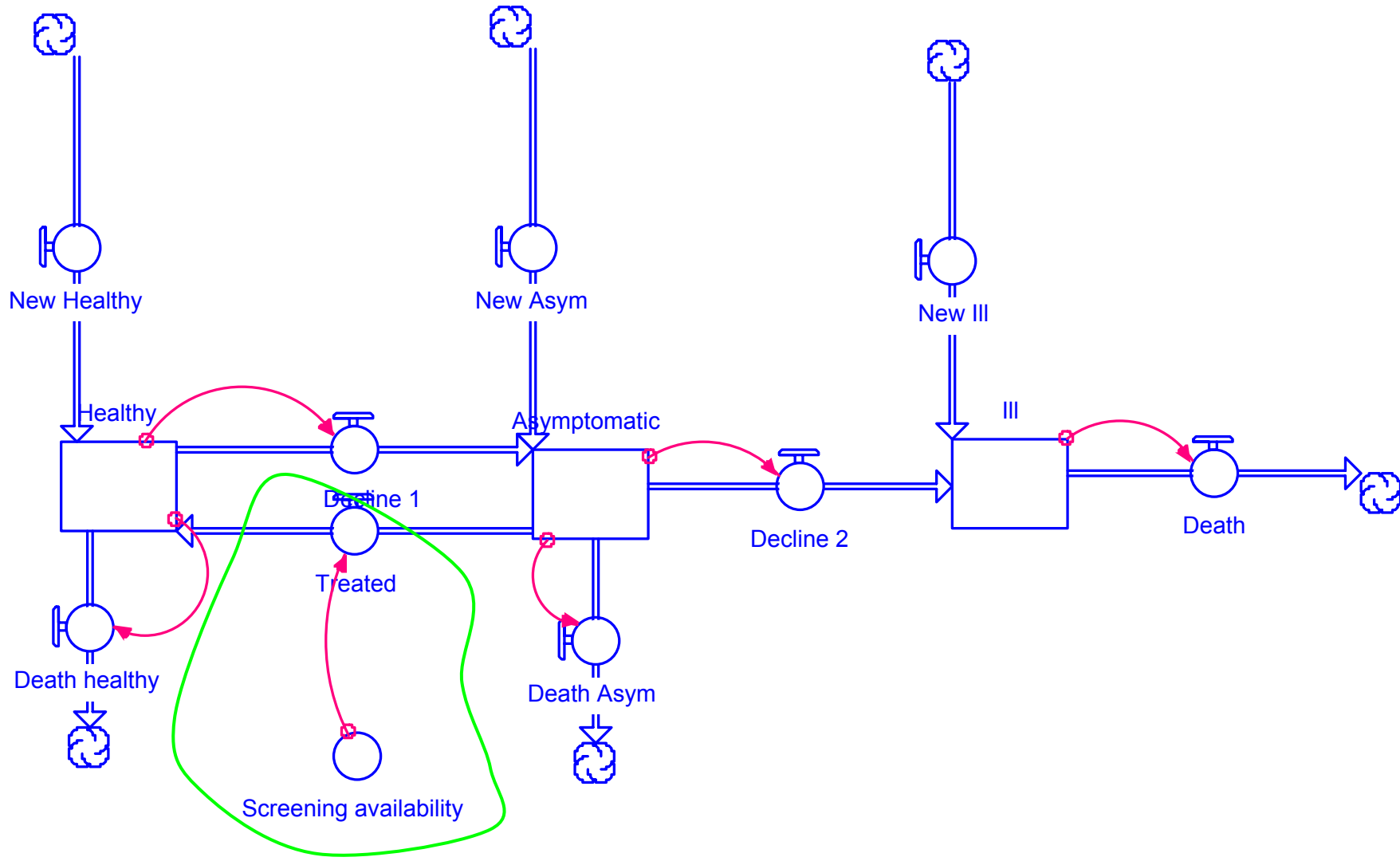
P is the matrix of probabilities,

$M(t)$ are the new people entering each state in each time period and I is the identity matrix.

Deterministic system dynamics model of the same system



It is possible to include constraints – but limited



Other software for aggregate transition models

- ❑❑❑ **Spreadsheets** may describe the transitions of proportions of patients in fixed time periods.
 - ❑❑❑ Deal with *average* numbers and are deterministic.
 - ❑❑❑ May incorporate exogenous influences e.g. screening at exact time periods.
- ❑❑❑ Another possibility is to use commercial software like **Treeage**.

Sensitivity Analysis

It is possible to do sensitivity analysis on input parameters quite quickly with SD, commercial or spreadsheet software.

Limitations

- Time intervals are discrete and equal
- Markovian assumption implies negative exponential decay from each state
- Homogeneous populations
- Rule of thumb – time interval should be at less than half the average activity time of the shortest activity
- States can be split to overcome this problem – to give Gamma distributions
- Replicate all states for each risk group

What are the potential problems?

- ▣ Assumption system is deterministic
- ▣ Subdivide states to take account of risk groups
- ▣ Subdivide states to model distributions
- ▣ Explosion of states
- ▣ Over-simplification of model
- ▣ Limited view of output

Suppose we are interested in the stochastic aspects of the system?

Use Monte Carlo simulation of a Markov process.

- ▣▣▣ Sample to decide when each individual goes from one state to the next;
- ▣▣▣ Like tossing a coin, throwing a die;
- ▣▣▣ Have to do it many times to get reliable averages for each time period;
- ▣▣▣ Can measure uncertainty.

G

Probability of going from healthy to asymptomatic in one year

0.3

Probability of going from asymptomatic to ill in one year

0.2

Year

Patients

1

2

3

4

5

6

1

1

C6

2

3

3

3

3

2

1

1

1

2

2

2

3

1

1

1

2

2

2

4

1

1

1

1

1

2

5

1

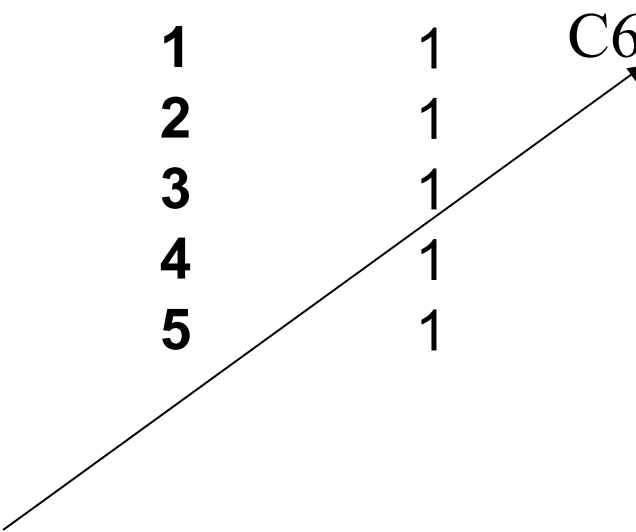
1

2

2

2

2



```
=IF(B6=1,IF(RAND()>$G$2,1,2),IF(B6=2,IF(RAND()>$G$3,2,3),3))
```

Output

```
=COUNTIF($K$6:$K$25,3)
```

Monte Carlo simulation 1

- ⌘ With @Risk or Crystal Ball can simulate just one individual over and over again
- ⌘ Eg. Run scenario with screening and without and compare output
- ⌘ Possible to use Treeage

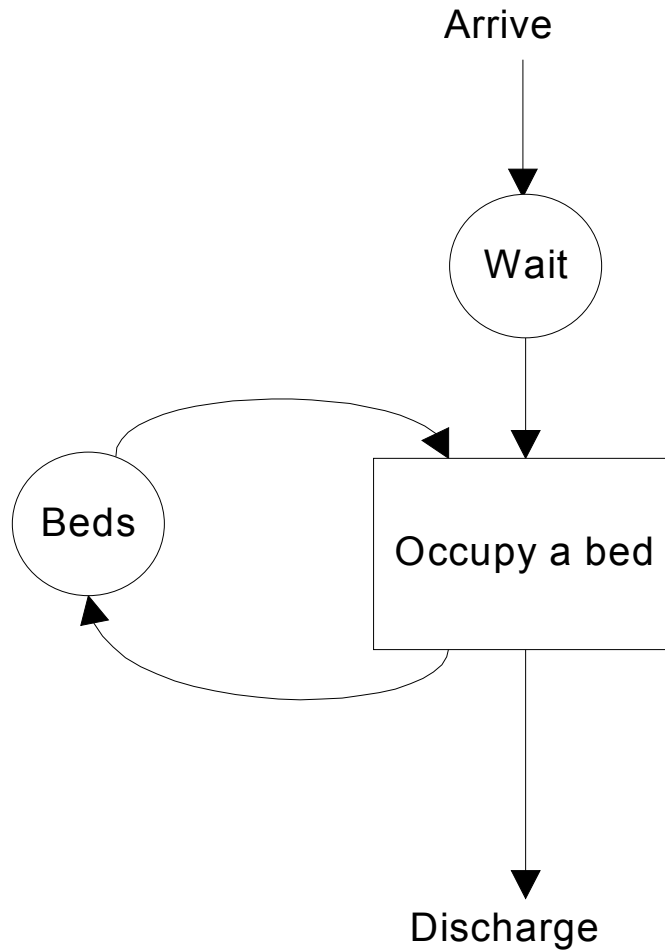
Monte Carlo simulation 2

- ▣▣▣ Individuals can be given random *characteristics* which affect their progress
- ▣▣▣ Have to finish with one patient before starting with next.
- ▣▣▣ Can use @Risk – very fast and can process lots of people.

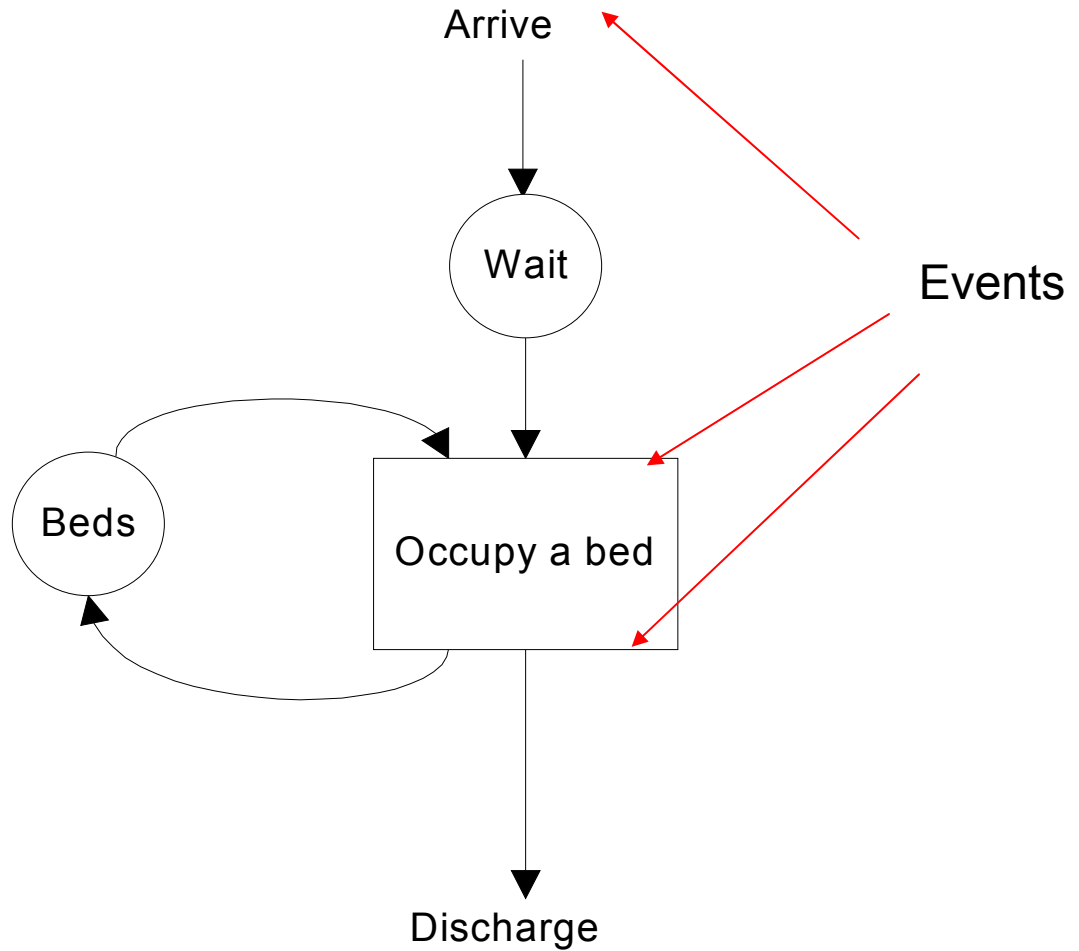
What is Discrete Event Simulation?

- ▣ The system is represented by a series of events.
- ▣ Individual, called entities, take part in these events.
- ▣ Simulation moves from one event to the next in continuous time.
- ▣ Future events are kept in a list and performed in time order.

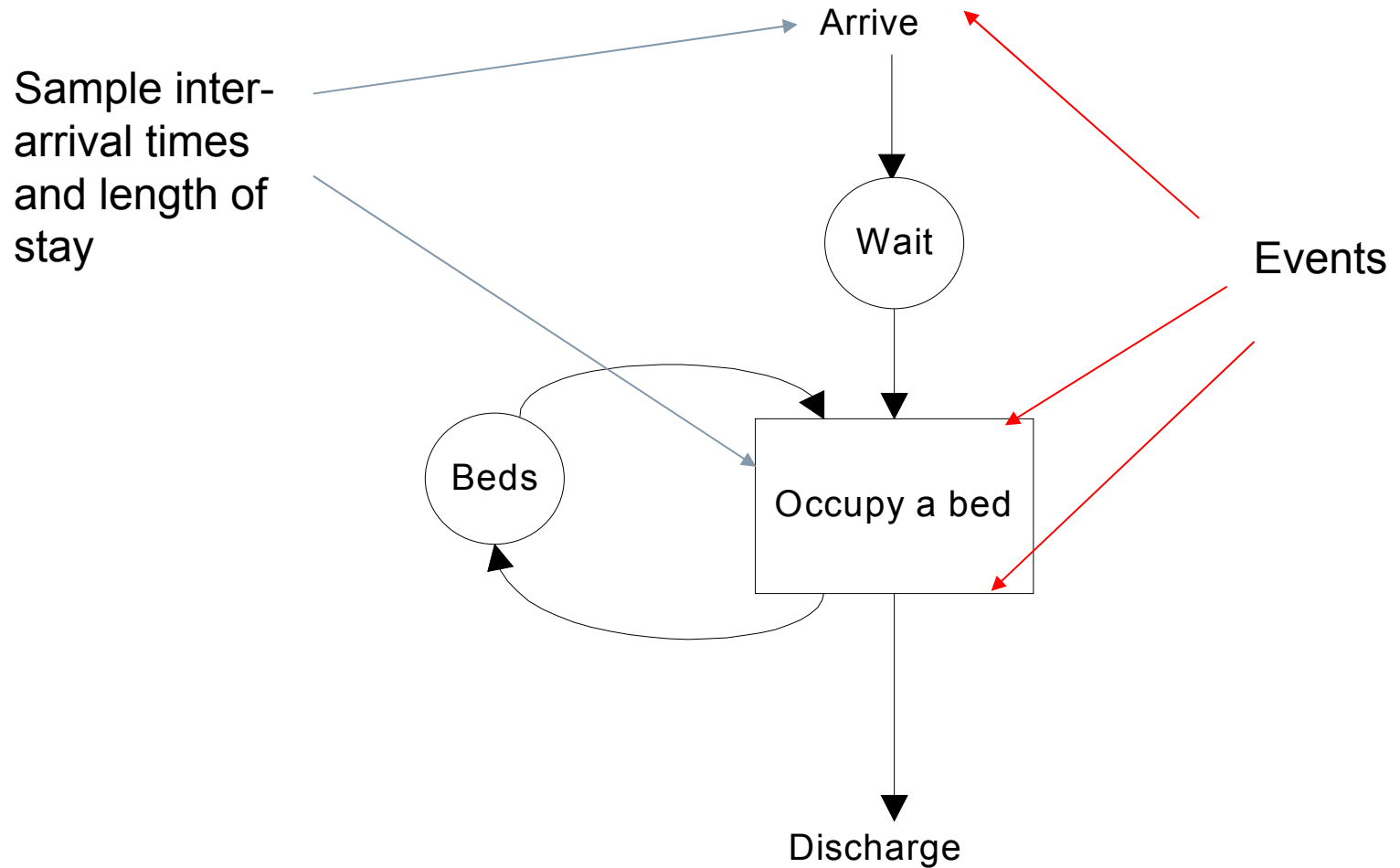
Simple DES structure

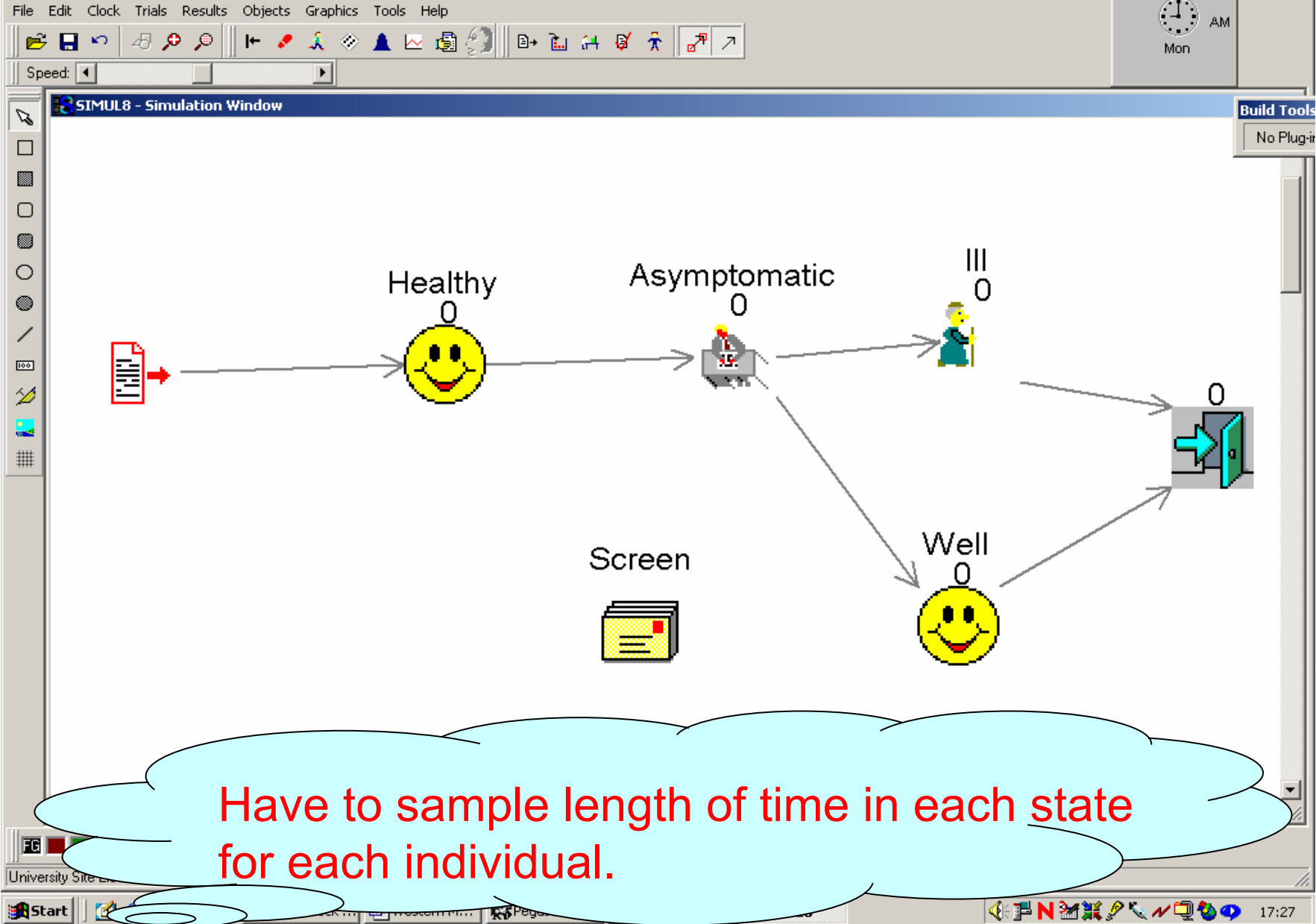


Simple DES structure



Simple DES structure





Advantages of Discrete Event Simulation

- ▣▣▣ Characteristics influence progression – No explosion of states
- ▣▣▣ Time advances in continuous time, as necessary
- ▣▣▣ Use of any activity time distribution
- ▣▣▣ Can incorporate more complex logic
- ▣▣▣ Can incorporate delays or interactions
- ▣▣▣ Can incorporate constraints

Disadvantages of Discrete Event Simulation

- ❑ Lots of software available but most not specifically designed for patient modelling
 - ❑ Simul8 – generic DES software
 - ❑ Program in high level language eg POST
 - ❑ ModGen
- ❑ Temptation to make model complex
- ❑ Time consuming to get statistically significant results
- ❑ Even more time consuming to do sensitivity analyses.

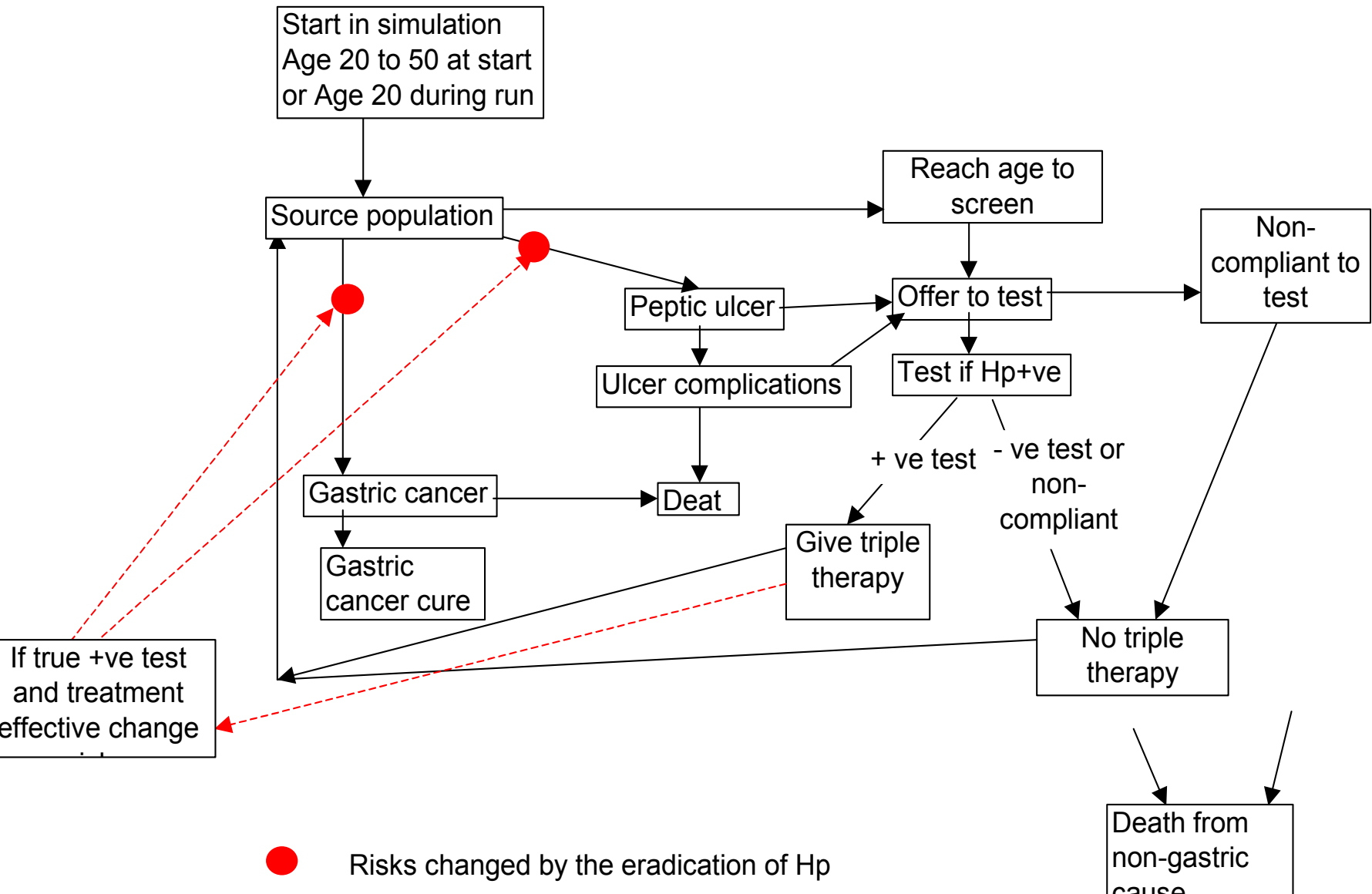
CASE STUDY Helicobacter Pylori

- ❏ Bacteria which populate the stomach
- ❏ Affects approximately 50% of those now 60 years old but only 15% or fewer of 20 year olds
- ❏ Is implicated in gastric and duodenal ulcers
- ❏ Is thought to be responsible for much gastric cancer

Should we screen and eliminate Hp?

- Screening for Hp fulfils some of the criteria for screening programs:
 - burden of disease (gastric cancer, peptic ulcer)
 - test available – blood test or saliva test
 - effective treatment – two antibiotics and proton pump inhibitor for two weeks.

Flow chart showing the effects of a screening program



Characteristics of the model

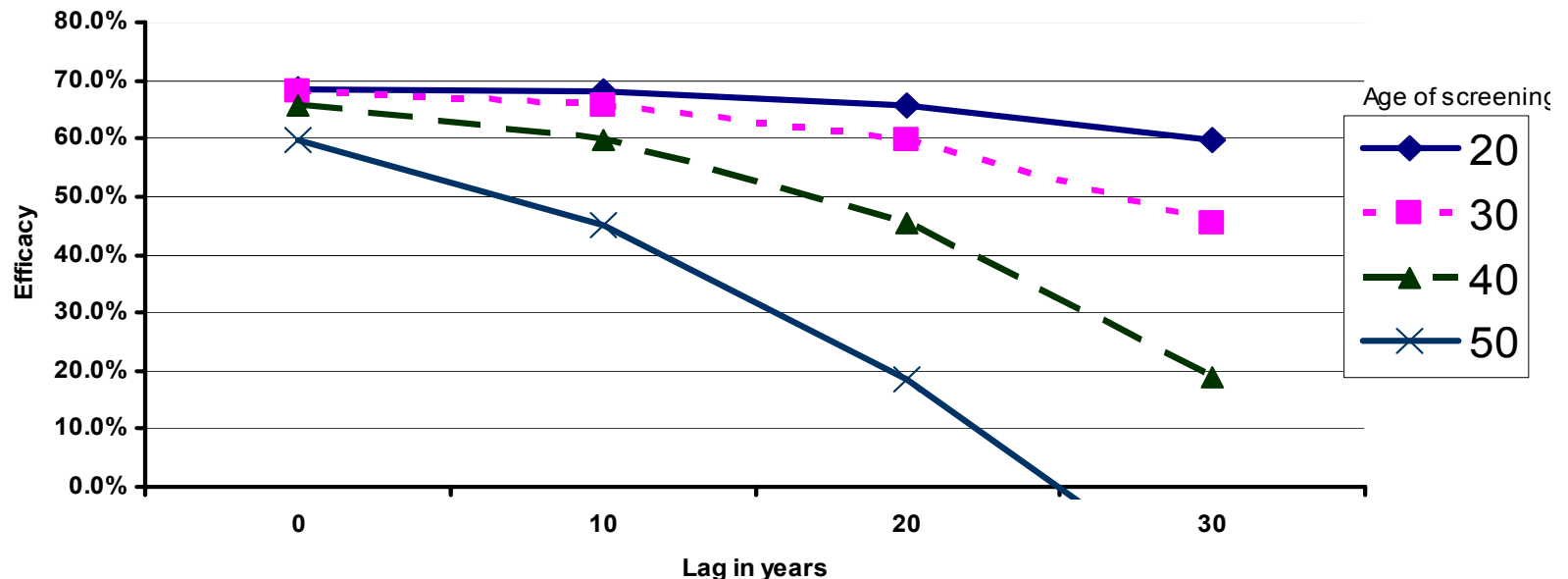
Ulcers

- Ulcer – some to hospital – some of those to death;
- Test, if Hp+, treat, if eradicated, reduce relative risk of next ulcer and of cancer.

Characteristics of the model

Cancer

- If patient goes from Hp+ to Hp- then increase time to cancer, subject to a lag.
- Lag ensures that those about to get cancer do not benefit.



Is it cost-effective to screen?

But.....

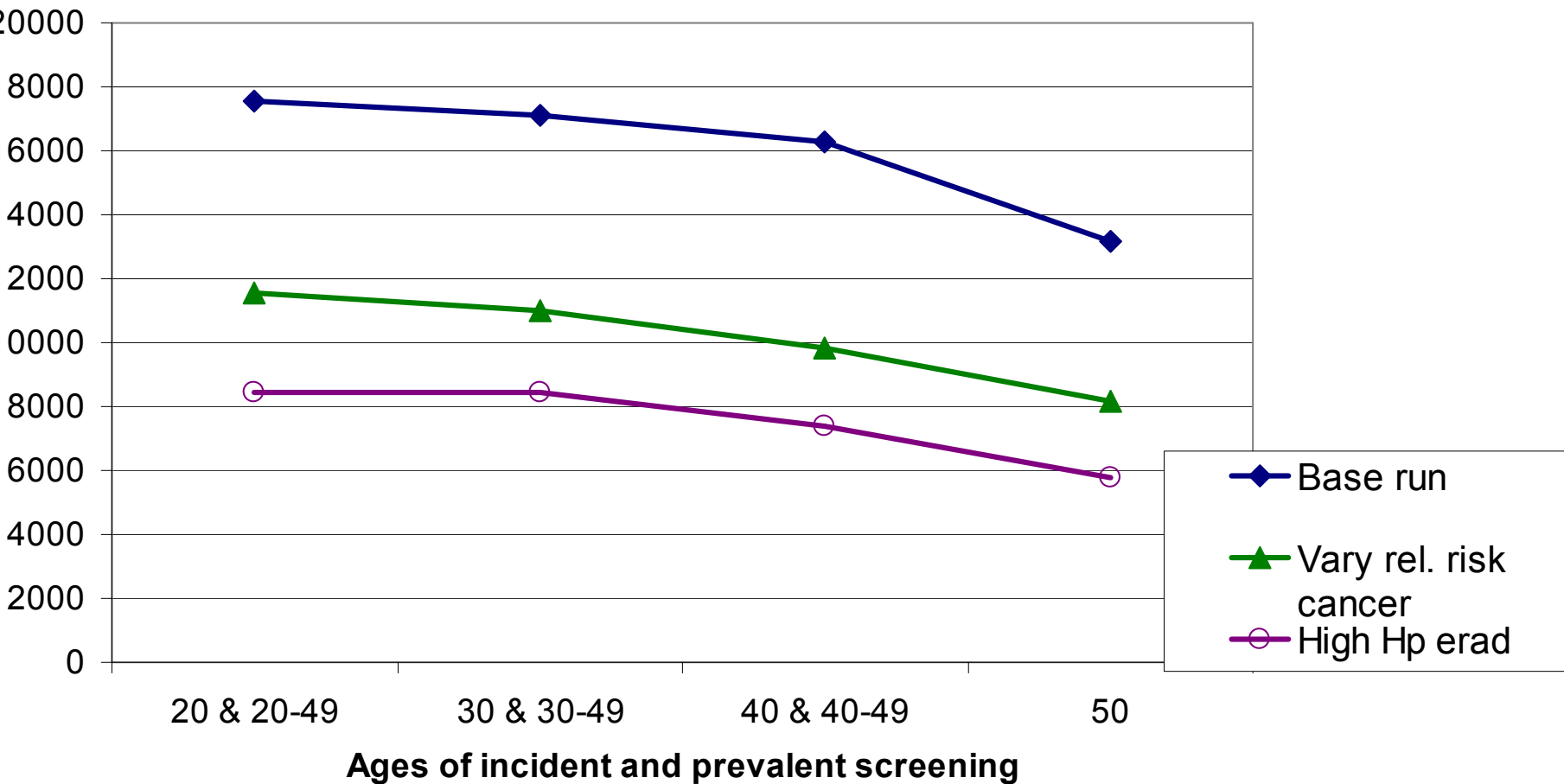
At what age?

What do we do with the prevalent unscreened population?

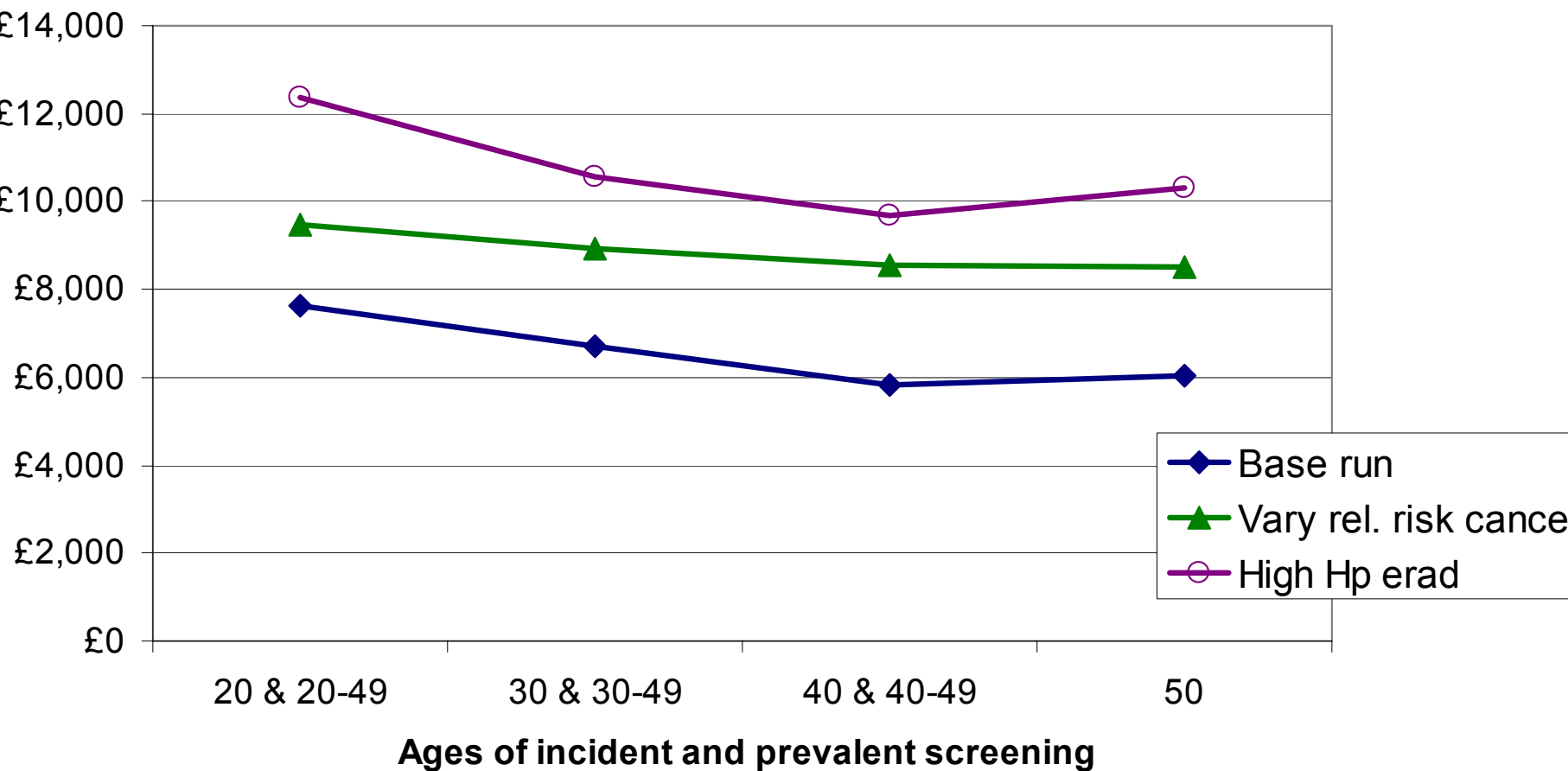
What assumptions have we made about the opportunistic treatment?

What will it cost and how soon will we have pay-back?

Deaths prevented to age 75 for population of England and Wales

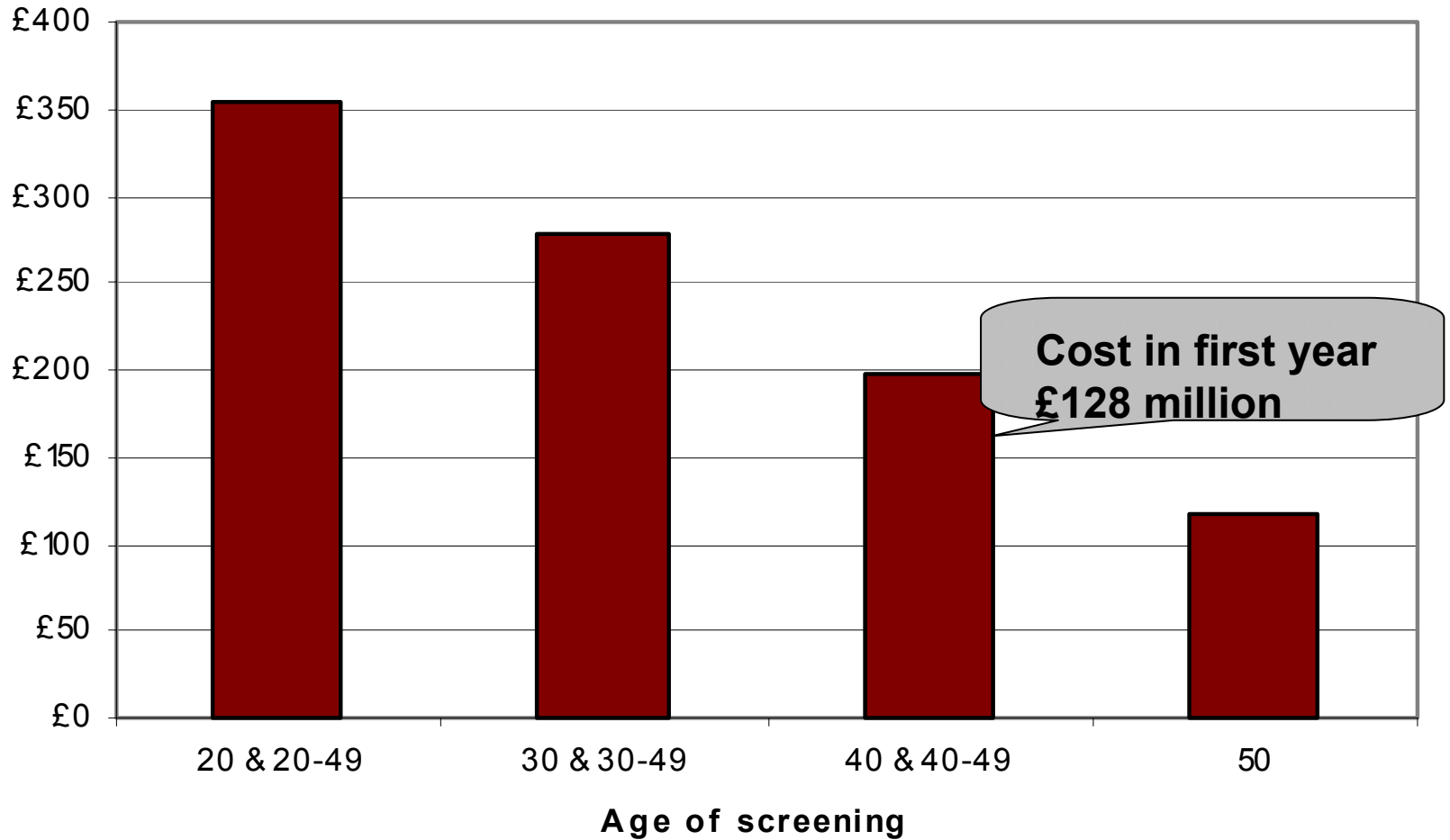


Cost per life year saved, all ages, discount rate 6%



Costs for England and Wales

Present value of costs - discount rate 6%



Variability

Needed 3 million individuals, aged 20 to 50 for statistical significance between no-screening and screening scenario

Screening at 40 years -170 deaths prevented in a population of 500,000 (population of Sheffield) over 80 years. Most after 20 years. Many years no deaths prevented.

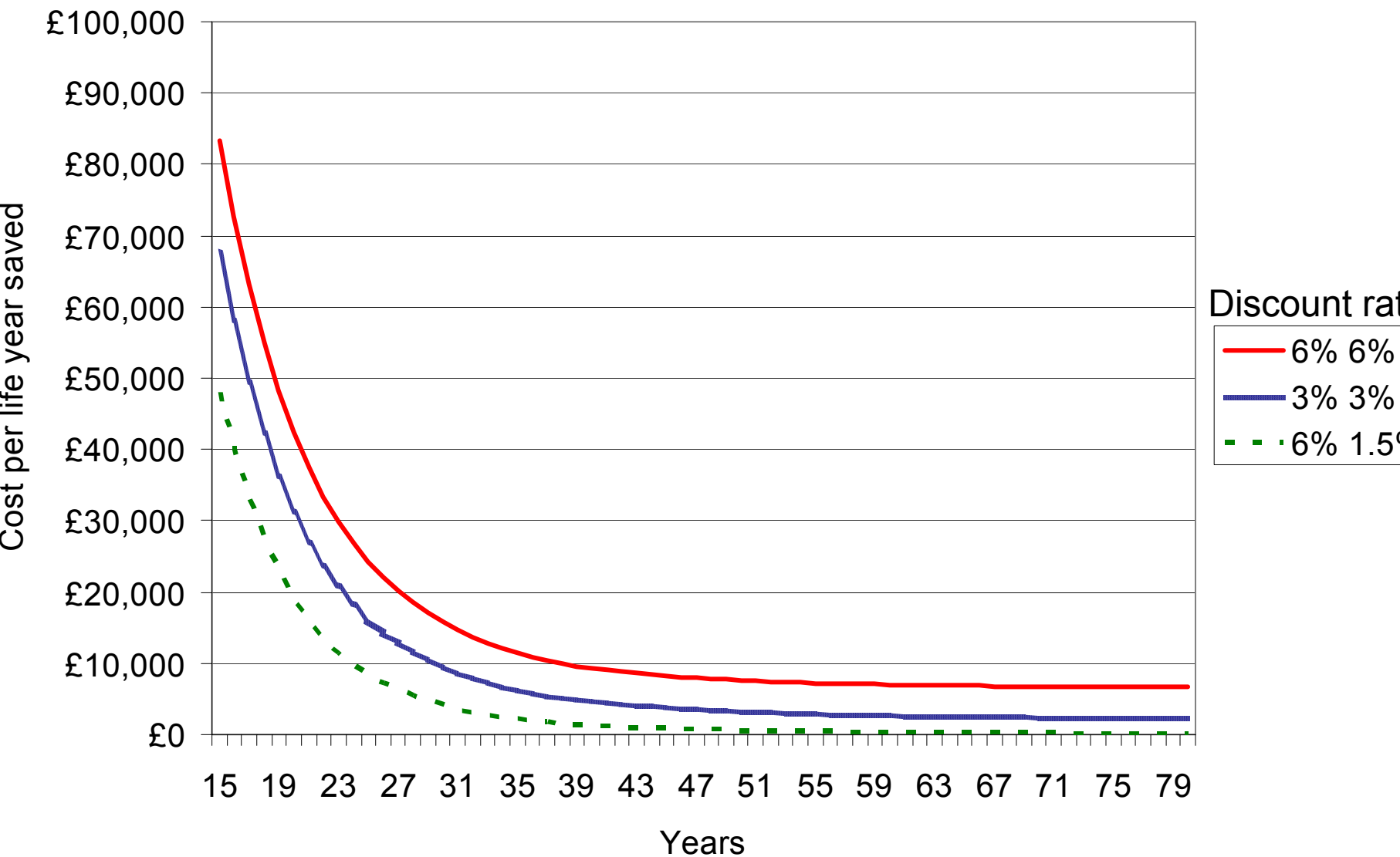
CPLYS – screening at 40 years

Upper age limit	Deaths prevented	Life years saved	CLYS no averted costs, no disc	CLYS with averted costs, no disc	CLYS no averted costs, disc 6%.	CLYS with averted costs, disc 6%.
75	16,263	130,315	£4,402	£1,593	£15,553	£10,924
85	27,684	270,949	£2,117	£700	£10,155	£6,834
None	34,456	368,045	£1,559	£334	£8,799	£5,866

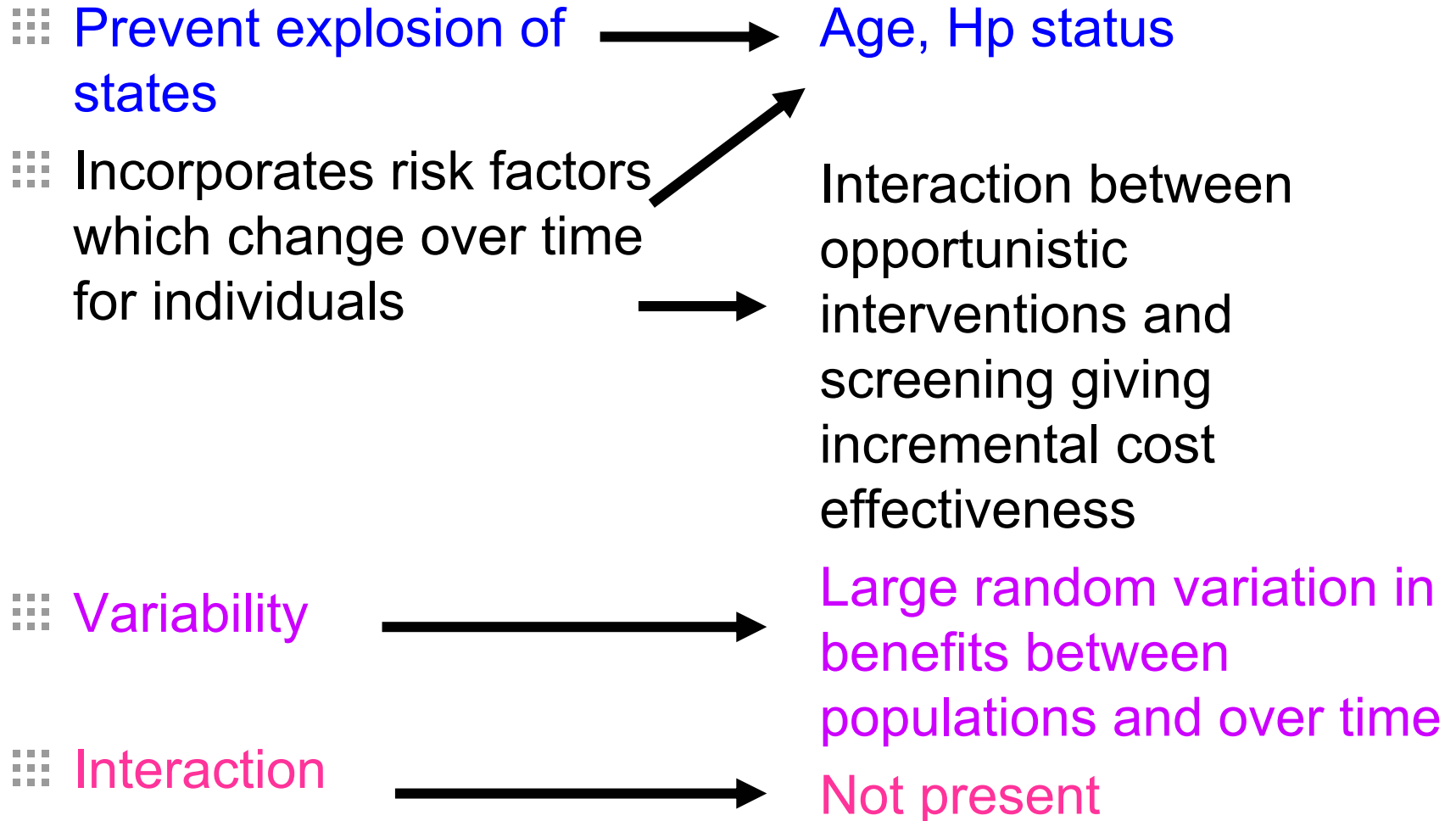
CPLY maximum 50,000 in sensitivity analysis of parameter input

Accumulated cost per life years saved over different numbers of years

Screening at 40 years



Why simulation?



Why simulation? (cont)

- ▣ More complex modelling → Use of time delay in onset of cancer gives overall effect of screening and describes age dependent efficacy of treatment
- ▣ Time advance over many years → Models lifetime disease, treatment, costs and CE
→ Shows changes in cumulative CE over time.
- ▣ Can explore a range of output → Proved to be valuable.

Simulation run times

Simulation lengthy to run

- need to compare difference from base case scenario;

- used variance reduction (common random numbers);

- possible to model patient by patient if no interaction.

Need many runs to

- investigate sensitivity of model to uncertainty in input parameters (probabilistic sensitivity analysis)

- derive “best” scenario

Analysis of output in various ways showed CLYS to be dependent on:

- ▣▣▣ age of screening,
- ▣▣▣ opportunistic activity,
- ▣▣▣ random effects – population by population,
- ▣▣▣ upper age for life years saved,
- ▣▣▣ discount rates,
- ▣▣▣ time horizon for calculations.

ARRIVES AT DIFFERENT CONCLUSIONS FROM OTHER STUDIES.

Need for Research

- ⌘⌘⌘ Appropriate model for particular problem
- ⌘⌘⌘ Development of user friendly, efficient simulation software for this type of problem
- ⌘⌘⌘ Range of output necessary/ desirable for evaluating screening and treatment for policy decisions
- ⌘⌘⌘ Development of easy-to-use meta-models

References

❏ Paper

- ❏ Davies R, Roderick P, Raftery J. The evaluation of disease prevention and treatment using simulation models. *European Journal of OR*. 2003, 150:1, 53-66.

❏ Book

- ❏ Robinson S. Simulation. The practice of model development and use. 2004, Wiley, Chichester