

Crescita logistica di una singola risorsa (x)

$x(t)$ = densità/biomassa della risorsa

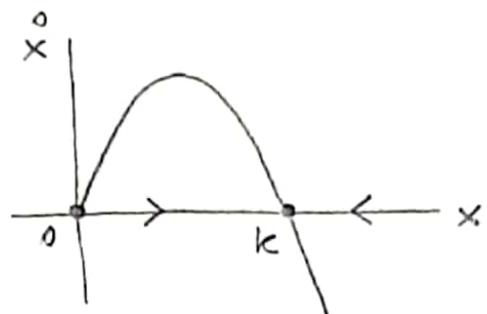
$$\dot{x} = r x \left(1 - \frac{x}{k}\right)$$

$r, k > 0$

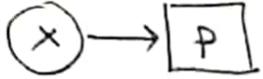
tasso intrinseco di crescita

capacità portante

Equilibri: $\dot{x} = 0$ $\left\{ \begin{array}{l} \bar{x} = 0 \\ \bar{x} = k \end{array} \right.$



Risorsa - Consumatore costante (P) (prede - predatore costante)



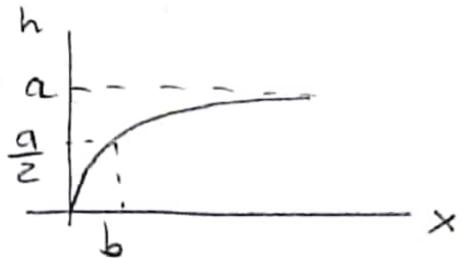
$$\dot{x} = r x \left(1 - \frac{x}{k}\right) - \frac{a x P}{b + x}$$

crescita logistica
mortalità per predazione

$$h(x) = \frac{a x}{b + x}$$

Risposta funzionale di Holling di tipo II

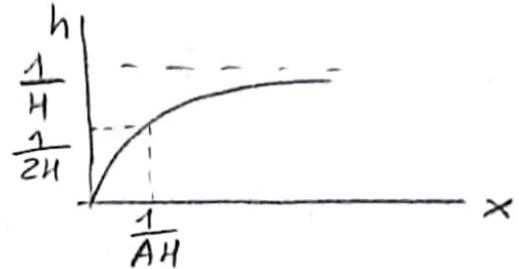
$$[h] = \frac{[x]}{[P][t]}$$



a = massima capacità predatoria
 b = costante di semi-saturazione

In alternativa $h(x) = \frac{A x}{1 + A H x}$

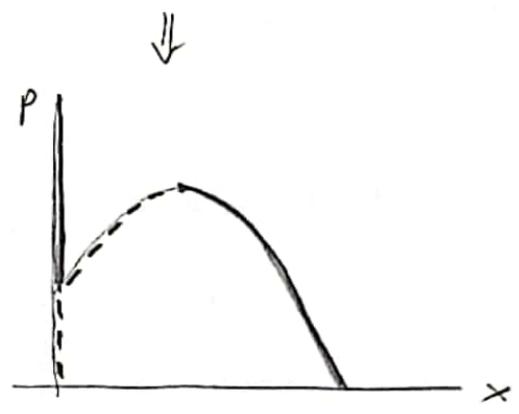
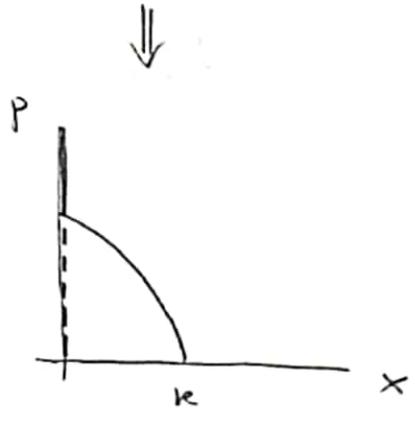
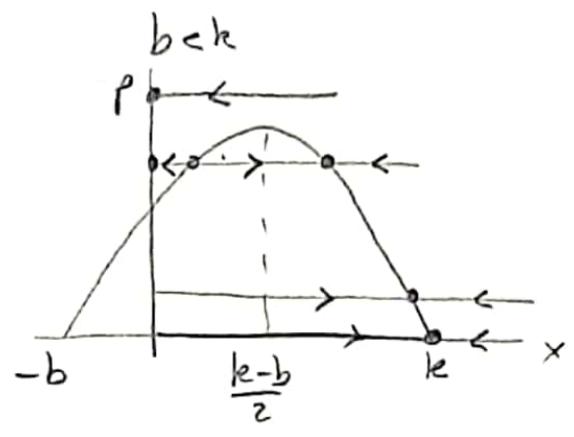
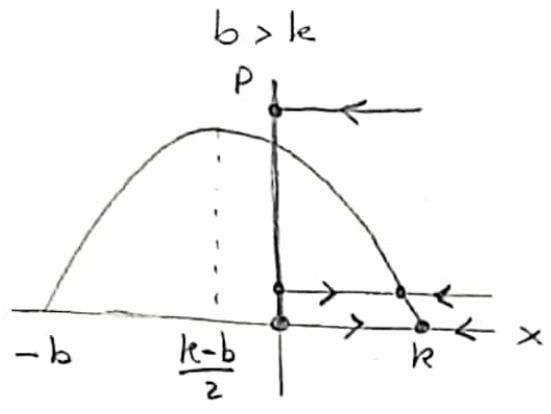
A = attack rate
 H = handling time



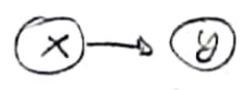
$$a = \frac{1}{H} \quad b = \frac{1}{A H} \quad \rightarrow \quad A = \frac{a}{b} \quad H = \frac{1}{a}$$

Equilibri: $\dot{x} = 0 \begin{cases} x=0 \\ r(1-\frac{x}{k}) - \frac{a}{b+x}P = 0 \end{cases} \downarrow$

$$P = \frac{r}{a} (1 - \frac{x}{k})(b+x)$$



Risorsa - Consumatore
(x) (y)



Preda - Predatore

Modello di Rosenzweig-MacArthur

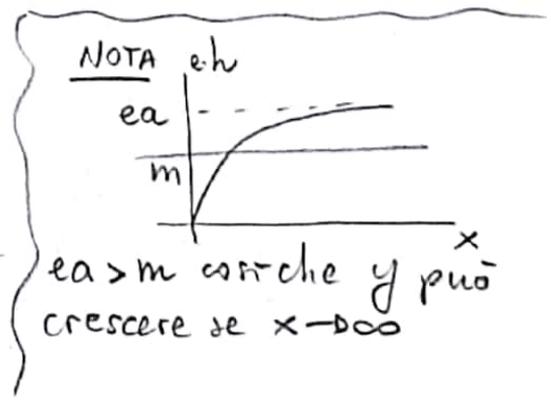
$$\dot{x} = rx(1 - \frac{x}{k}) - \frac{axy}{b+x}$$

crescita logistica mortalità per predazione

$$\dot{y} = e \frac{axy}{b+x} - my$$

efficienza di conversione

$-my$
 mortalità naturale
 tasso di mortalità



Equilibri $\begin{cases} \dot{x} = 0 \\ \dot{y} = 0 \end{cases}$

$\dot{x} = 0 \begin{cases} x = 0 \\ y = \frac{r}{a} \left(1 - \frac{x}{k}\right) (b+x) \end{cases}$

$\dot{y} = 0 \begin{cases} y = 0 \\ x = \frac{mb}{ea-m} \end{cases}$

$E_0 = (0,0)$ $E_k = (k, 0)$ $E = (\bar{x}, \bar{y})$

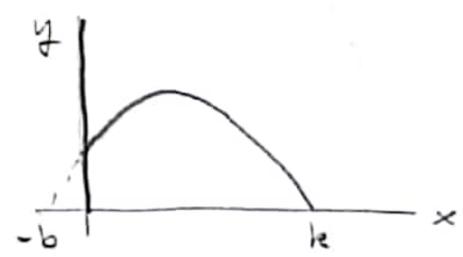
$\frac{mb}{ea-m}$ $\frac{r}{a} \left(1 - \frac{\bar{x}}{k}\right) (b + \bar{x})$

con $1 - \frac{\bar{x}}{k} > 0$

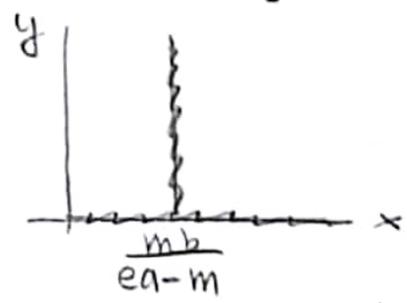
$\Rightarrow \bar{x} < k \rightarrow \frac{mb}{ea-m} < k$

Equilibri \rightarrow Nisocline

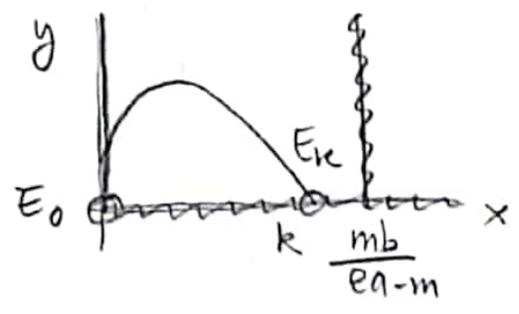
$\dot{x} = 0 \begin{cases} x = 0 \\ y = \frac{r}{a} \left(1 - \frac{x}{k}\right) (b+x) \end{cases}$



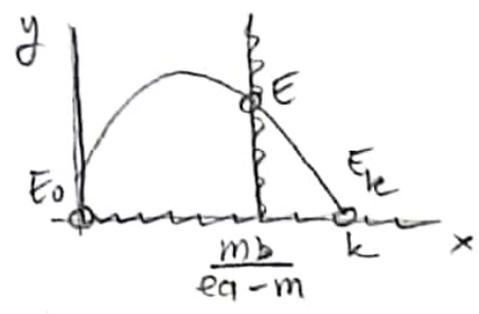
$\dot{y} = 0 \begin{cases} y = 0 \\ x = \frac{mb}{ea-m} \end{cases}$



$\frac{mb}{ea-m} > k$



$\frac{mb}{ea-m} < k$

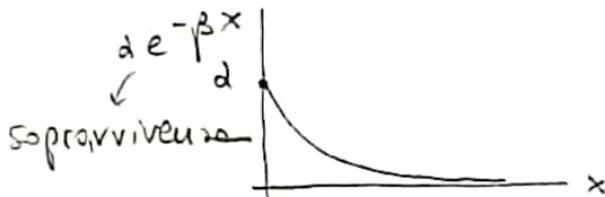


Modello di Ricker → crescita di una singola misura a tempo discreto $x(t)$

Salmanni
dell'Oceano Pacifico
(popolazione semelpara)

$$x(t+1) = d x(t) e^{-\beta x(t)} = f(x(t))$$

NOTA: $d > 1$
 x piccolo $e^{-\beta x} \sim 1$
 $x(t+1) = d x(t)$ e $x(t)$ cresce
se $d < 1$ $x(t)$ decresce

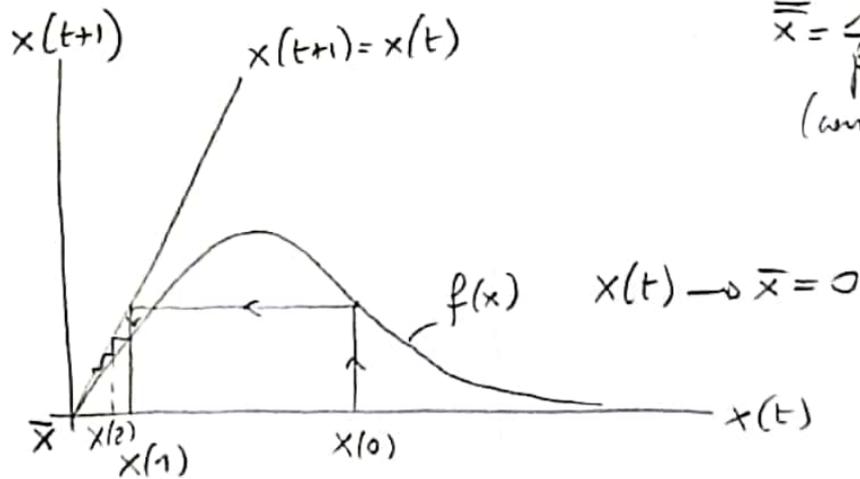


Equilibri: $x(t+1) = x(t) = \bar{x}$

$$\bar{x} = d \bar{x} e^{-\beta \bar{x}} \begin{cases} \bar{x} = 0 \\ 1 = d e^{-\beta \bar{x}} \\ \downarrow \\ \bar{x} = \frac{1}{\beta} \ln d \\ (\text{con } d > 1) \end{cases}$$

Diagramma di Moran

$d < 1$ ($\neq \bar{x}$)
 $(f'(0) = d < 1)$



$d > 1$? Dipende!
 $1 < d < e^2 \Rightarrow x(t) \rightarrow \bar{x}$
 $d > e^2 \Rightarrow x(t) \rightarrow$ qualcosa di più complicato
• ciclo
• caos

Vedi simulatore

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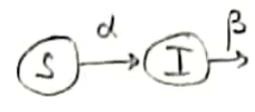
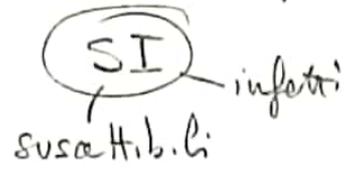
function [stato]=ricker(alfa,beta,x0)
% simulazione modello di Ricker
% [stato]=ricker(alfa,beta,x0)
% [stato]=ricker(0.8,1,0.2); % ---> equilibrio nullo
% [stato]=ricker(6,1,0.2); % ---> equilibrio positivo
% [stato]=ricker(10,1,0.2); % ---> ciclo di periodo 2
% [stato]=ricker(13,1,0.2); % ---> ciclo di periodo 4
% [stato]=ricker(14.5,1,0.2); % ---> ciclo di periodo 8
% [stato]=ricker(15,1,0.2); % ---> caos
% [stato]=ricker(20,1,0.2); % ---> caos
% [stato]=ricker(23,1,0.2); % ---> ciclo di periodo 3
% [stato]=ricker(24,1,0.2); % ---> ciclo di periodo 6

close all;

##### SIMULAZIONE #####
tfin=30; stato=x0;
for ind=2:tfin
    x=stato(ind-1);
    stato=[stato;alfa*x*exp(-beta*x)];
end;
##### MOVIMENTO #####
figure;
plot(stato,'-');
xlabel('tempo')
ylabel('x(t)')
pause;
##### COSTRUZIONE DEL DIAGRAMMA DI MORAN "DINAMICO" #####
moran1=ones(2*tfin,1);
moran2=ones(2*tfin,1);
for h=1:tfin
    moran1(2*h-1)=stato(h);
    moran1(2*h)=stato(h);
end;
moran2=[0;moran1(3:2*tfin);1];
figure;
hold on
massimo=1.2*max(x0,alfa/beta*exp(-1));
x=0:massimo/100:massimo;
y=alfa*x.*exp(-beta*x);
plot(x,y,'r');
plot(x,x,'k');
xlabel('x(t)')
ylabel('x(t+1)')
title('diagramma di Moran con transitorio');
for ind=1:length(moran1)-2
    plot([moran1(ind) moran1(ind+1)],[moran2(ind) moran2(ind+1)],'b-');
    pause(0.2)
end;
pause;
% Diagramma di Moran a transitorio esaurito
tfin=200;
stato=x0;
for ind=2:tfin
    x=stato(ind-1);
    stato=[stato;alfa*x*exp(-beta*x)];
end;
stato=stato(length(stato));
for ind=2:tfin
    x=stato(ind-1);
    stato=[stato;alfa*x*exp(-beta*x)];
end;
figure;
plot(stato(1:length(stato)-1),stato(2:length(stato)),'*');
hold on;
massimo=1.2*max(x0,alfa/beta*exp(-1));
x=0:massimo/100:massimo;
y=alfa*x.*exp(-beta*x);
plot(x,y,'r');
xlabel('x(t)')
ylabel('x(t+1)')
title('diagramma di Moran su attrattore (transitorio esaurito)');
moran1=ones(2*tfin,1);
moran2=ones(2*tfin,1);
for h=1:tfin
    moran1(2*h-1)=stato(h);
    moran1(2*h)=stato(h);
end;
moran2=[0;moran1(3:2*tfin);1];
plot(moran1(1:length(moran1)-1),moran2(1:length(moran1)-1),'-.');
plot(x,x,'k');
pause;
% -----
figure;
hold on;
axis([0 massimo -1 1]);
plot([0 massimo],[0 0],'k');
axis('off');
text(massimo,-0.1,'x(t)');
text(0,-0.1,'0');
title('attrattore');
plot(stato(1),0,'ro','markersize',5);
pause;
for ind=1:length(stato)
    plot(stato(ind),0,'b*','markersize',5);
    pause(0.1)
end;

```

Modelli di epidemie



Immunizzazione permanente
(morbilli/rosha/rar:cella)

$$\dot{S} = -\alpha SI$$

contagio

$$\dot{I} = \alpha SI - \beta I$$

guarigione

$\alpha \downarrow$ con mascherine/raccini

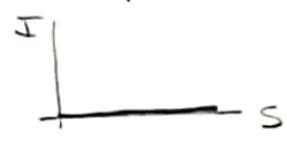
$\beta \uparrow$ con cure

Equilibri

$$\dot{S} = 0 \begin{cases} S = 0 \\ I = 0 \end{cases}$$

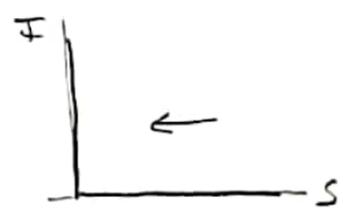
$$\dot{I} = 0 \begin{cases} I = 0 \\ S = \frac{\beta}{\alpha} \end{cases}$$

$\Rightarrow \exists \infty$ equilibri $(\bar{S}, 0)$

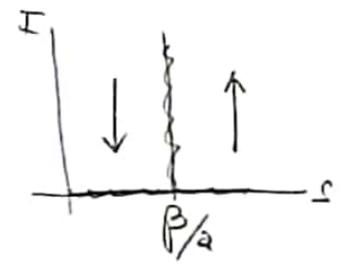


Isocline

$\dot{S} = 0$

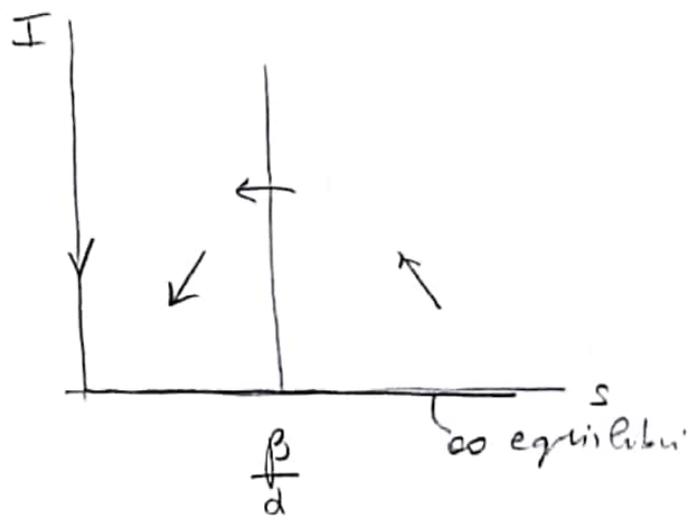


$\dot{I} = 0$



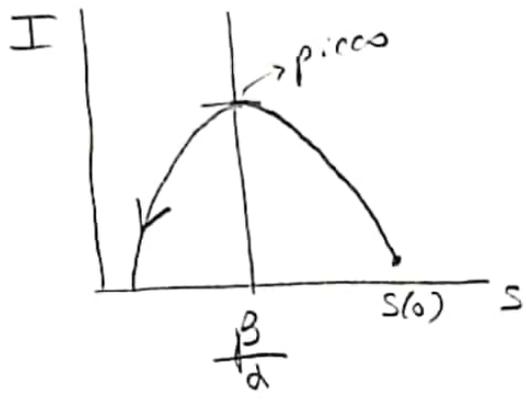
Equilibri \rightarrow Isocline $\rightarrow I = 0 \forall S$

Vettore tangente

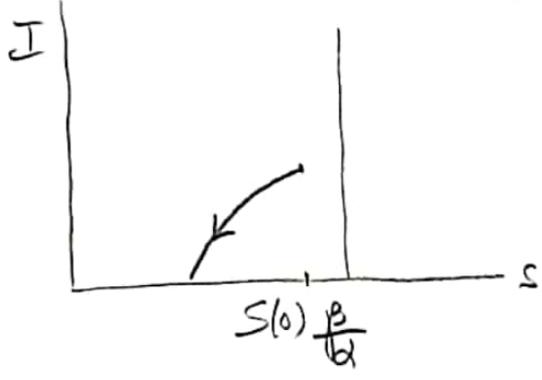


$S(0) > \frac{\beta}{\alpha} \rightarrow$ L'epidemia "parte"

$I(0) > 0$



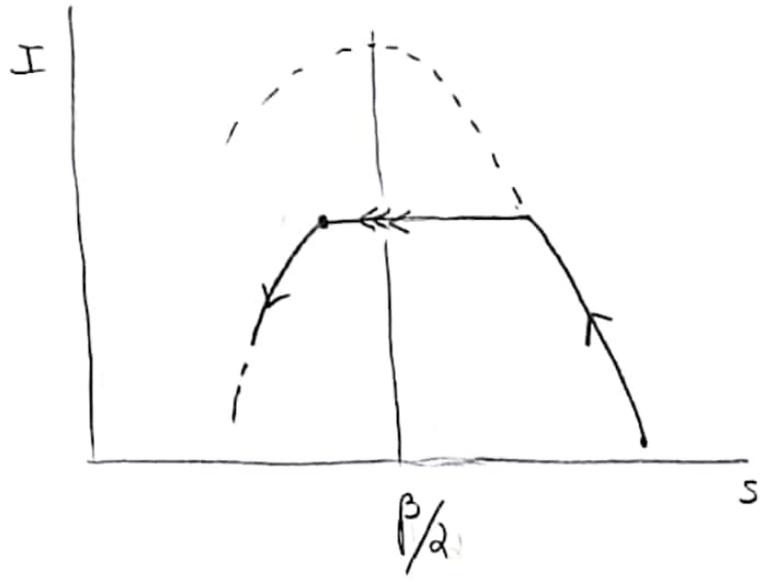
$S(0) < \frac{\beta}{\alpha} \rightarrow$ L'epidemia "non parte"



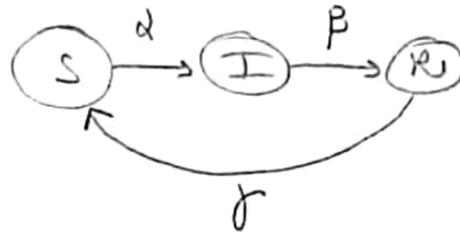
Per fare in modo che $S(0)$ sia $< \frac{\beta}{\alpha} \rightarrow$ β alto (cure)
 α piccolo (mascherine)
 Vaccini

lockdown?

$S \downarrow$ sotto $\frac{\beta}{\alpha}$



$\textcircled{\text{SIR}}$ ^{infezz.}
 rimossi
 suscettibili



Immunità associata
 temporanea
 (influenza/pertosse...)

$$\begin{aligned}
 \dot{S} &= -\alpha SI + \gamma R \\
 \dot{I} &= \alpha SI - \beta I \\
 \dot{R} &= \beta I - \gamma R
 \end{aligned}$$

\neq morti (opp. trascurabili)
 $S + I + R = N \quad \forall t$
 \downarrow
 popolazione totale

Equilibrio

$$\begin{aligned}
 -\alpha SI + \gamma R &= 0 \\
 \alpha SI - \beta I &= 0 \rightarrow I = 0 \xrightarrow{\gamma S} \\
 \beta I - \gamma R &= 0 \xrightarrow{\gamma R}
 \end{aligned}
 \Rightarrow \bar{E} = (\bar{S}, 0, 0)$$

oppure $S = \frac{\beta}{\alpha} \quad R = \frac{\beta}{\gamma} I$

$$\frac{\beta}{\alpha} + I + \frac{\beta}{\gamma} I = N \rightarrow I = \frac{N - \frac{\beta}{\alpha}}{1 + \frac{\beta}{\gamma}} = \frac{\gamma}{\alpha} \frac{\alpha N - \beta}{\gamma + \beta}$$

$$\Rightarrow \bar{E} = \left(\frac{\beta}{\alpha}, \frac{\gamma}{\alpha} \frac{\alpha N - \beta}{\gamma + \beta}, \frac{\beta}{\alpha} \frac{\alpha N - \beta}{\gamma + \beta} \right)$$

$\textcircled{\text{SIRV}}$

