

# Numerical results for trophical transmission model with multiple infections

```
In[486]:= SetDirectory [NotebookDirectory []]
          << Tools`
          ParallelNeeds ["Tools`"](* Import package Tools that define useful functions *)
Out[486]:= /Users/phuongnguyen/Work/multipleinfections/code
```

## ODEs

Dynamics of intermediate hosts

```
In[489]:= dIsdt = R[Iw, Is, Iww] - d Is -  $\Pi_s$ [Ds, Dw, Dww] Is -  $\eta$  Is; (*Susceptible host *)
          dIwdt = (1 - p)  $\eta$  Is - (d +  $\alpha_w$ ) Iw -  $\Pi_w$ [Ds, Dw, Dww,  $\beta_w$ ] Iw;
          (* Host infected with one parasite *)
          dIwwdt = p  $\eta$  Is - (d +  $\alpha_{ww}$ ) Iww -  $\Pi_{ww}$ [Ds, Dw, Dww,  $\beta_{ww}$ ] Iww;
          (* Host infected with two parasites *)
```

Dynamics of definitive hosts

```
In[7]:= dDsdt = B[Ds, Dw, Dww, Is, Iw, Iww] -  $\mu$  Ds - ( $\lambda_{ww}$  +  $\lambda_w$ ) Ds; (*Susceptible host*)
          dDwdt = ( $\lambda_w$  + (1 - q)  $\lambda_{ww}$ ) Ds - ( $\mu$  +  $\sigma_w$ ) Dw - ((1 - q)  $\lambda_{ww}$  +  $\lambda_w$ ) Dw;
          (*Host infected with one parasite*)
          dDwwdt = q  $\lambda_{ww}$  Ds + ((1 - q)  $\lambda_{ww}$  +  $\lambda_w$ ) Dw - ( $\mu$  +  $\sigma_{ww}$ ) Dww;
          (* Host infected with two parasites*)
```

Dynamics of free-living parasites

```
In[10]:= dWdt =  $f_w$  Dw +  $f_{ww}$  Dww -  $\delta$  W -  $\eta$  Is;
```

Setting force of infection, and list of parameters for solving the odes

```
In[11]:= forceInf = { $\eta$  ->  $\gamma$  W,  $\lambda_w$  -> h ( $\rho$  +  $\beta_w$ ) Iw,  $\lambda_{ww}$  -> h ( $\rho$  +  $\beta_{ww}$ ) Iww};
          odesRes = {dIsdt, dIwdt, dIwwdt, dDsdt, dDwdt, dDwwdt, dWdt};
          varRes = {Is, Iw, Iww, Ds, Dw, Dww, W};
          vartRes = {Is[t], Iw[t], Iww[t], Ds[t], Dw[t], Dww[t], W[t]};
```

Testing that the total dynamics of definitive host does not involve transmission

```
In[15]:= dDsdt + dDwdt + dDwwdt //FullSimplify
```

```
Out[15]:= -Ds  $\mu$  - Dw ( $\mu$  +  $\sigma_w$ ) - Dww ( $\mu$  +  $\sigma_{ww}$ ) + B[Ds, Dw, Dww, Is, Iw, Iww]
```

## Reproduction ratio R0

This is the result from the file *tropicaltransmission\_analytical\_git.nb*

$$\begin{aligned} \text{In[16]:= } R_0 = & \gamma_{ls} \frac{p q h (\beta_{ww} + \rho)}{\alpha_{ww} + d + \Pi_{ww} [D_s, D_w, D_{ww}, \beta_{ww}]} \frac{D_s}{\mu + \sigma_{ww}} \frac{f_{ww}}{\delta + \gamma_{ls}} + \\ & \gamma_{ls} \left( \frac{(1-p) (\beta_w + \rho) h}{\alpha_w + d + \Pi_w [D_s, D_w, D_{ww}, \beta_w]} + \frac{p (1-q) (\beta_{ww} + \rho) h}{\alpha_{ww} + d + \Pi_{ww} [D_s, D_w, D_{ww}, \beta_{ww}]} \right) \\ & \frac{D_s}{\mu + \sigma_w} \frac{f_w}{\delta + \gamma_{ls}}; \end{aligned}$$

## Graph format and parallel computation

```

In[17]:= includeFrame = True;
imageSize = 600;
frameStyle = Directive [Black, Thickness [0.003]];
labelStyle = {Black, FontSize -> 24};
colorlist = ColorData [97, "ColorList" ]
colorstable = ColorData [97, "ColorList" ][[{1, 2, -2}]]
ihostCol = ColorData [97, "ColorList" ][[1]];
dhostCol = ColorData [97, "ColorList" ][[4]];
fCol1 = ColorData [97, "ColorList" ][[3]];
fCol2 = ColorData [97, "ColorList" ][[{-1}]];
pointsize = 0.02;
figResolution = 300;
colbifur = {Black, Black, Black, Black };
colbifurfilling =
  {Lighter [Gray, 0.1 ], Lighter [Gray, 0.3 ], Lighter [Gray, 0.5 ], Lighter [Gray, 0.7 ]}
panellabel = 250;
thick = 0.003;
coldat = "Pastel";
intorder = 5;

```

```
Out[21]= {Blue, Orange, Green, Red, Purple, Brown, Teal, Yellow, Magenta, Olive, Coral, Blue, Orange, Purple, Green}
```

```
Out[22]= {Blue, Orange, Purple}
```

```
Out[30]= {Gray, Gray, Gray, Gray}
```

# Linear birth function for intermediate host

Setting birth function

```
In[35]:= func0 = {R[Iw, Is, Iww] → r (Is + Iw + Iww), Πs[Ds, Dw, Dww] → ρ (Ds + Dw + Dww),
  Πw[Ds, Dw, Dww, βw] → βw (Ds + Dw + Dww),
  Πww[Ds, Dw, Dww, βww] → βww (Ds + Dw + Dww),
  B[Ds, Dw, Dww, Is, Iw, Iww] → ρ c (Ds + Dw + Dww) (Is + Iw + Iww)};
sysfunc0 = odesRes /. forceInf /. func0;
sysNDSolvefunc0 = MakeSystem [varRes, t, sysfunc0];
```

Solving the odes with linear birth function

```
In[38]:= sollds = Solve[Thread[sysfunc0 == 0] /.
  {Iw -> 0, Iww -> 0, Dw -> 0, Dww -> 0, W -> 0}, varRes][[2]]
```

 **Solve:** Equations may not give solutions for all "solve" variables.

```
Out[38]:= {Is →  $\frac{\mu}{c \rho}$ , Ds →  $-\frac{d - r}{\rho}$ }
```

## Jacobian matrix

```
In[39]:= Jmatfunc0 = D[sysfunc0, {varRes}];
```

## Ecological trajectories (supplementary figure)

(\*Set parameters \*)

```
prEco0 = {ρ → 1.2, d → 0.9, r → 2.5, γ → 2.9, αw → 0, αww -> 0, βw → 1.5,
  βww -> 1.5, p -> 0.1, c → 1.4, μ → 0.9, σw → 0, σww -> 0,
  q -> 0.01, fw → 6.5, fww -> 7.5, δ → 0.9, h -> 0.8};
maxt = 100;
```

(\*Set initial conditions for population dynamics \*)

```
init0 = {Is[0] == 1, Iw[0] == 1, Iww[0] == 0.3,
  Ds[0] == 1.5, Dw[0] == 1.7, Dww[0] == 1, W[0] == 10};
```

(\*Solving the equilibrium \*)

```
sols0 = NSolve [Thread [(odesRes /. func0 /. forceInf /. prEco0) == 0], varRes];
```

(\*Solving the dynamics of the odes \*)

```
nds0 = NDSolve [Join [sysNDSolvefunc0 /. prEco0, init0], varRes, {t, 0, maxt}];
```

(\*Set the range for the graph \*)

```
range = {{-1, 20}, {-1, 20}};
```

(\* Annotation for the different lines \*)

```

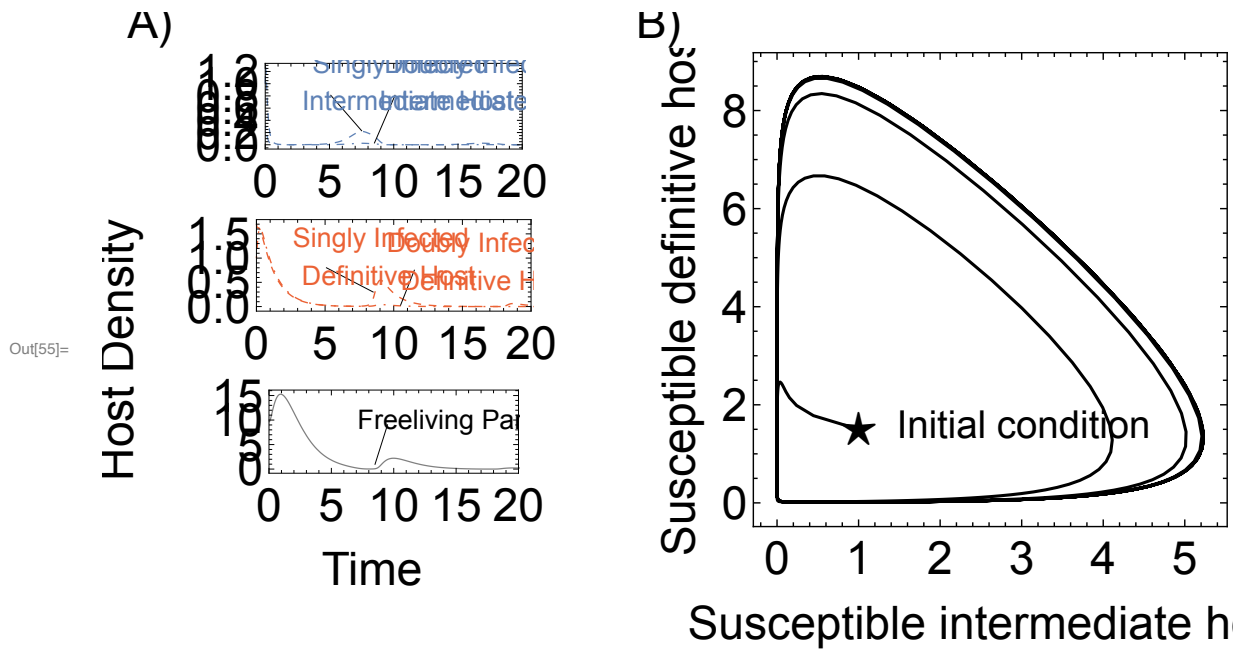
epilogParasite01 =
  {{Black, Line [{{8.5, 0.03 }, {10.1, 0.81 }]}], {Black, Line [{{7.5, 0.23 }, {5.1, 0.81 }]}],
   {ihostCol, Text [Style ["Doubly Infected \n Intermediate Host", FontSize → 14],
    {10., 1. }, {-0.8, 0}]], {ihostCol, Text [Style [
    "Singly Infected \n Intermediate Host", FontSize → 14], {4., 1. }, {-0.8, 0}]]];
epilogParasite02 = {{Black, Line [{{10.5, 0.03 }, {11.5, 0.75 }]}],
  {Black, Line [{{8.5, 0.3 }, {5.1, 0.8 }]}],
  {dhostCol, Text [Style ["Doubly Infected \n Definitive Host", FontSize → 14],
   {10.9, 0.9 }, {-0.8, 0}]], {dhostCol, Text [
   Style ["Singly Infected \n Definitive Host", FontSize → 14], {4., 1. }, {-0.8, 0}]]];
epilogParasite03 = {{Black, Line [{{9.5, 9.03 }, {8.5, 0.98 }]}],
  {Black, Text [Style ["Freeliving Parasites", FontSize → 14], {9., 10. }, {-0.8, 0}]]];

(*Plot the dynamics with time for infected intermediate hosts *)
p11 = Plot[Evaluate[{lw[t], lww[t]}/. ndsol0],
  {t, 0, maxt }, PlotRange → {{0, 20}, All}, AspectRatio → 1/3,
  ImageSize → imageSize /3, Frame → True, FrameStyle → frameStyle,
  LabelStyle → labelStyle, PlotStyle → {{ihostCol, Dashed, Thickness [0.005 ]},
   {ihostCol, DotDashed, Thickness [0.005 ]}}, Epilog → epilogParasite01 ];
(* Plot the dynamics with time for infected definitive hosts *)
p12 = Plot[Evaluate[{Dw[t], Dww[t]}/. ndsol0],
  {t, 0, maxt }, PlotRange → {{0, 20}, All}, AspectRatio → 1/3,
  ImageSize → imageSize /3, Frame → True, FrameStyle → frameStyle,
  LabelStyle → labelStyle, PlotStyle → {{dhostCol, Dashed, Thickness [0.005 ]},
   {dhostCol, DotDashed, Thickness [0.005 ]}}, Epilog → epilogParasite02 ];
(*Plot the dynamics with time for free -living parasites *)
p13 = Plot[Evaluate[{W[t]}/. ndsol0], {t, 0, maxt },
  PlotRange → {{0, 20}, All}, AspectRatio → 1/3, ImageSize → imageSize /3,
  Frame → True, FrameStyle → frameStyle, LabelStyle → labelStyle,
  PlotStyle → {Gray, Thickness [0.005 ]}, Epilog → epilogParasite03 ];
(* Combine the graphs of dynamics with respect to time *)
p1 = GraphicsColumn[{p11, p12, p13 }, Spacings → 0, ImageSize → {900, 1000 },
  Epilog → {Text [Style ["Time", Black, 24 ], {Center, -215}],
   Rotate [Text [Style ["Host Density", Black, 24 ], {0.5, Center }], 90 Degree ]];

(*Annotation for initial condition and equilibrium *)
epilogHost0 = {{Black, Text [Style [★, FontSize → 24], {1., 1.5 }, {0, 0}]},
  {Black, Text [Style ["Initial condition", FontSize → 20], {1., 1.5 }, {-1.3, -0.2}]]];

(*Phase plane of susceptible intermediate and definitive hosts *)
p2 = ParametricPlot [{Evaluate[{Is[t], Ds[t]}/. ndsol0]}, {t, 0, maxt }, AspectRatio → 1,
  PlotRange → All, Frame → includeFrame, PlotStyle → Black, FrameStyle → frameStyle,
  FrameLabel → {Style ["Susceptible intermediate host", Black, 24 ],
   Style ["Susceptible definitive host", Black, 24 ]},
  ImageSize → imageSize * 1.5, LabelStyle → labelStyle, Epilog → epilogHost0 ];
(*Combine graphs of infected and susceptible hosts *)
GraphicsRow[{p1, p2}, Spacings → 0, Epilog →
  {Text [Style ["A", Black, 24 ], {100, -10}], Text [Style ["B", Black, 24 ], {900, -10}]]
(*Export["diseasefree_linear.pdf", %, ImageResolution → figResolution]*)

```



## Nonlinear birth function for intermediate host

Setting birth function

```
In[56]:= func1 = {R[lw, ls, lww] -> r (1 - k (ls + lw + lww)) (ls + lw + lww),  $\Pi_s[D_s, D_w, D_{ww}]$  ->  $\rho (D_s + D_w + D_{ww})$ ,  $\Pi_w[D_s, D_w, D_{ww}, \beta_w]$  ->  $(\rho + \beta_w) (D_s + D_w + D_{ww})$ ,  $\Pi_{ww}[D_s, D_w, D_{ww}, \beta_{ww}]$  ->  $(\rho + \beta_{ww}) (D_s + D_w + D_{ww})$ , B[D_s, D_w, D_{ww}, ls, lw, lww] ->  $\rho c (D_s + D_w + D_{ww}) (ls + lw + lww)}$ ;
```

```
sysfunc1 = odesRes /. forceInf /. func1;
sysNDSolvefunc1 = MakeSystem [varRes, t, sysfunc1];
Thread [sysfunc1 == 0] /. {lw -> 0, lww -> 0, D_w -> 0, D_{ww} -> 0, W -> 0};
```

Equilibrium of susceptible intermediate and definitive host at disease free scenario

```
In[60]:= diseasefree = Solve[%, {Is, Ds}][[3]];
```

Reproduction ratio

```
In[61]:= R0func1 = R0 /. func1 /. D_w -> 0 /. D_{ww} -> 0 /. lw -> 0 /. lww -> 0 // Simplify;
```

## Jacobian matrix

```
In[62]:= Jmatfunc1 = D[sysfunc1, {varRes}];
```

## Ecological trajectories (Figure 3)

## Disease free equilibrium

```

In[65]:= (*Set parameters *)
prEcoNL = { $\rho \rightarrow 1.2$ ,  $d \rightarrow 0.9$ ,  $r \rightarrow 2.5$ ,  $\gamma \rightarrow 2.9$ ,  $\alpha_w \rightarrow 0$ ,  $\alpha_{ww} \rightarrow 0$ ,
 $\beta_w \rightarrow 1.5$ ,  $\beta_{ww} \rightarrow 1.5$ ,  $p \rightarrow 0.05$ ,  $c \rightarrow 1.4$ ,  $\mu \rightarrow 3.9$ ,  $\sigma_w \rightarrow 0$ ,  $\sigma_{ww} \rightarrow 0$ ,
 $q \rightarrow 0.05$ ,  $fw \rightarrow 7.5$ ,  $fww \rightarrow 7.5$ ,  $\delta \rightarrow 0.9$ ,  $k \rightarrow 0.26$ ,  $h \rightarrow 0.8$ };
maxt = 100;

(*Solving equilibrium *)
solsNL = NSolve[Thread[(sysfunc1 /. prEcoNL) == 0], varRes];
(*Set range for the graph *)
range1 = {{-0.05, 2.55}, {-0.05, 1.05}};

(*Set different initial conditions for population dynamics *)
ics = {
  {Is[0] == 0.6, Iw[0] == 0.2, Iww[0] == 0.3,
   Ds[0] == 0.25, Dw[0] == 0.7, Dww[0] == 0.1, W[0] == 0.8},
  {Is[0] == 1.2, Iw[0] == 0.2, Iww[0] == 0.3, Ds[0] == 0.5,
   Dw[0] == 0.7, Dww[0] == 0.1, W[0] == 0.8},
  {Is[0] == 1.8, Iw[0] == 0.2, Iww[0] == 0.3, Ds[0] == 0.75,
   Dw[0] == 0.7, Dww[0] == 0.1, W[0] == 0.8}};

(*Solving the odes at different initial conditions *)
ndsolsNL = Table[NDSolve[Join[sysNDSolvefunc1 /. prEcoNL, ics[[i]]],
  varRes, {t, 0, maxt}], {i, 1, Length[ics], 1}];

(*Population dynamics at equilibrium 1 *)
ieq1 = Evaluate[Is[maxt] /. ndsolsNL][[1]];
deq1 = Evaluate[Ds[maxt] /. ndsolsNL][[1]];

(*Annotation for initial equilibrium and starting condition 1 *)
epilogHost1 = {{Black, PointSize[pointsize], Point[{ieq1, deq1}]},
  {Black, Text[Style[ $\star$ , FontSize  $\rightarrow$  24], {1.2, 0.5}, {0, 0}]},
  {Black, Text[Style["Initial condition", FontSize  $\rightarrow$  14], {1.2, 0.5}, {-1.2, -0.2}]},
  {Black, Text[Style["Equilibrium", FontSize  $\rightarrow$  14], {ieq1, deq1}, {1.2, -0.2}]}];

(*Phase plane for susceptible intermediate and definitive hosts *)
ppHost1 = ParametricPlot[Evaluate[{Is[t], Ds[t]} /. ndsolsNL], {t, 0, maxt}, AspectRatio  $\rightarrow$  1,
  Frame  $\rightarrow$  includeFrame, PlotStyle  $\rightarrow$  Black, FrameStyle  $\rightarrow$  frameStyle,
  PlotRange  $\rightarrow$  range1, FrameLabel  $\rightarrow$  {"Density of \n Susceptible Intermediate Host",
    "Density of \n Susceptible Definitive Host"},
  LabelStyle  $\rightarrow$  labelStyle, ImageSize  $\rightarrow$  imageSize / 1.1, Epilog  $\rightarrow$  epilogHost1];

(*Annotation for different lines *)
epilogParas11 = {{ihostCol, Text[Style["Singly Infected \n Intermediate Hosts",
  FontSize  $\rightarrow$  14], {2, 0.6}, {-1.05, 0}]},
  {ihostCol, Text[Style["Doubly Infected \n Intermediate Hosts", FontSize  $\rightarrow$  14],
    {7, 0.2}, {-1.05, 0}]}];

```

```

epilogParas12 = {{dhostCol, Text [Style["Doubly Infected \n Definitive Hosts",
    FontSize → 14], {7, 0.2}, {-1.05, 0}]},
    {dhostCol, Text [Style["Singly Infected \n Definitive Hosts", FontSize → 14],
    {0.2, 0.6}, {-1.05, 0}]}];
epilogParas13 = {{Black, Text [Style["Freeliving Parasites", FontSize → 14],
    {0.75, 0.8}, {-1.05, 0}]}];

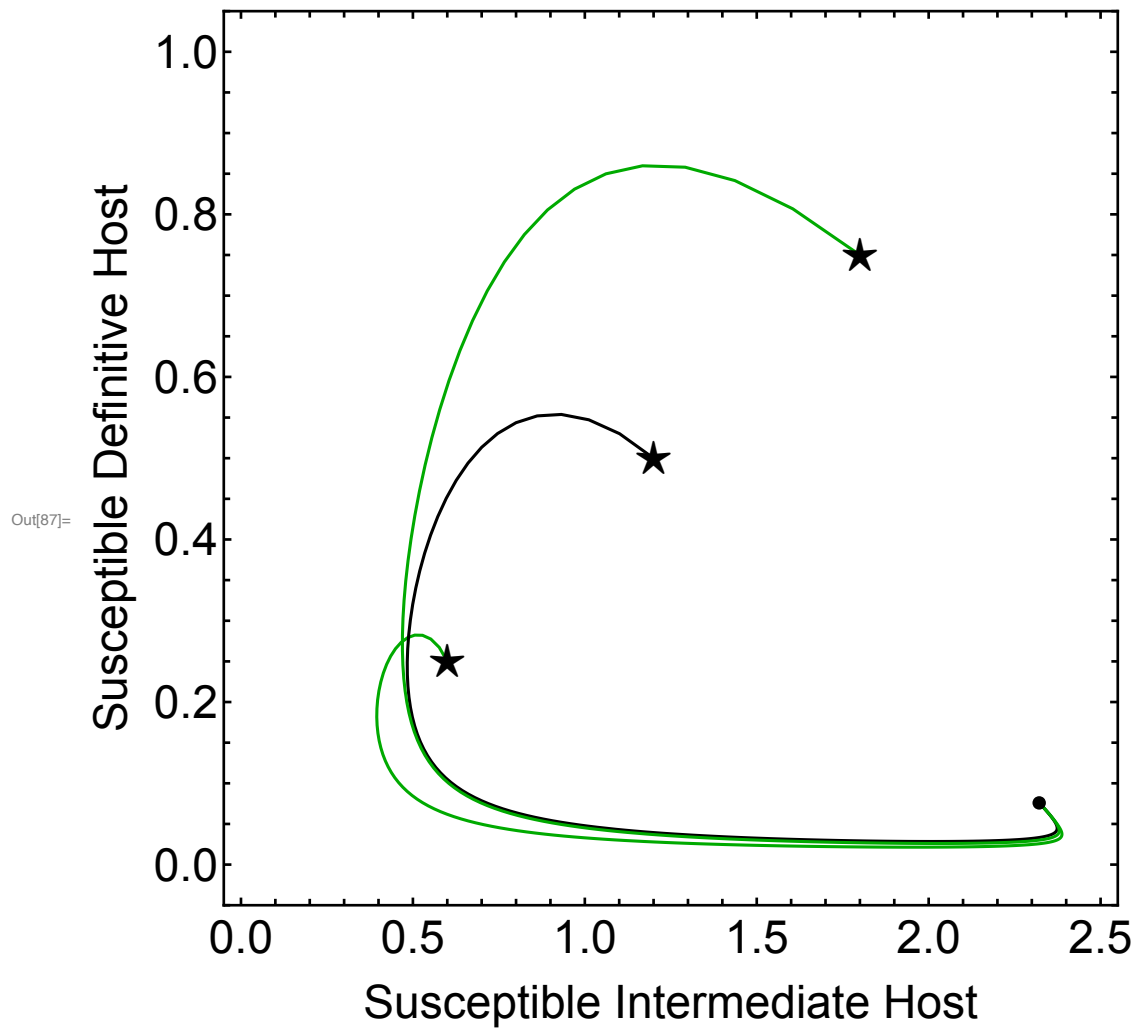
(*Plot dynamics with respect to time for intermediate host,
definitive host and free -living parasite *)
spcl = ndsolNL[[2]];
ppParasite11 = Plot[Evaluate[{lw[t], lww[t]}/. spcl],
    {t, 0, maxt}, AspectRatio → 1/3, PlotRange → {{0, 10}, {-0.1, 1.2}},
    PlotStyle → {{ihostCol, Dashed}, {ihostCol, DotDashed}}, Frame → includeFrame,
    ImageSize → imageSize, FrameStyle → frameStyle, Epilog → epilogParas11];
ppParasite12 = Plot[Evaluate[{Dw[t], Dww[t]}/. spcl], {t, 0, maxt},
    AspectRatio → 1/3, PlotRange → {{0, 10}, {-0.1, 1.2}},
    PlotStyle → {{dhostCol, Dashed}, {dhostCol, DotDashed}}, Frame → includeFrame,
    ImageSize → imageSize, FrameStyle → frameStyle, Epilog → epilogParas12];
ppParasite13 = Plot[Evaluate[{W[t]}/. spcl], {t, 0, maxt}, AspectRatio → 1/3,
    PlotRange → {{0, 10}, {-0.1, 1.2}}, PlotStyle → Gray, Frame → includeFrame,
    ImageSize → imageSize, FrameStyle → frameStyle, Epilog → epilogParas13];
ppParasite1 = GraphicsColumn[{ppParasite11, ppParasite12, ppParasite13},
    Spacings → 0, ImageSize → {600, 600}];

```

```

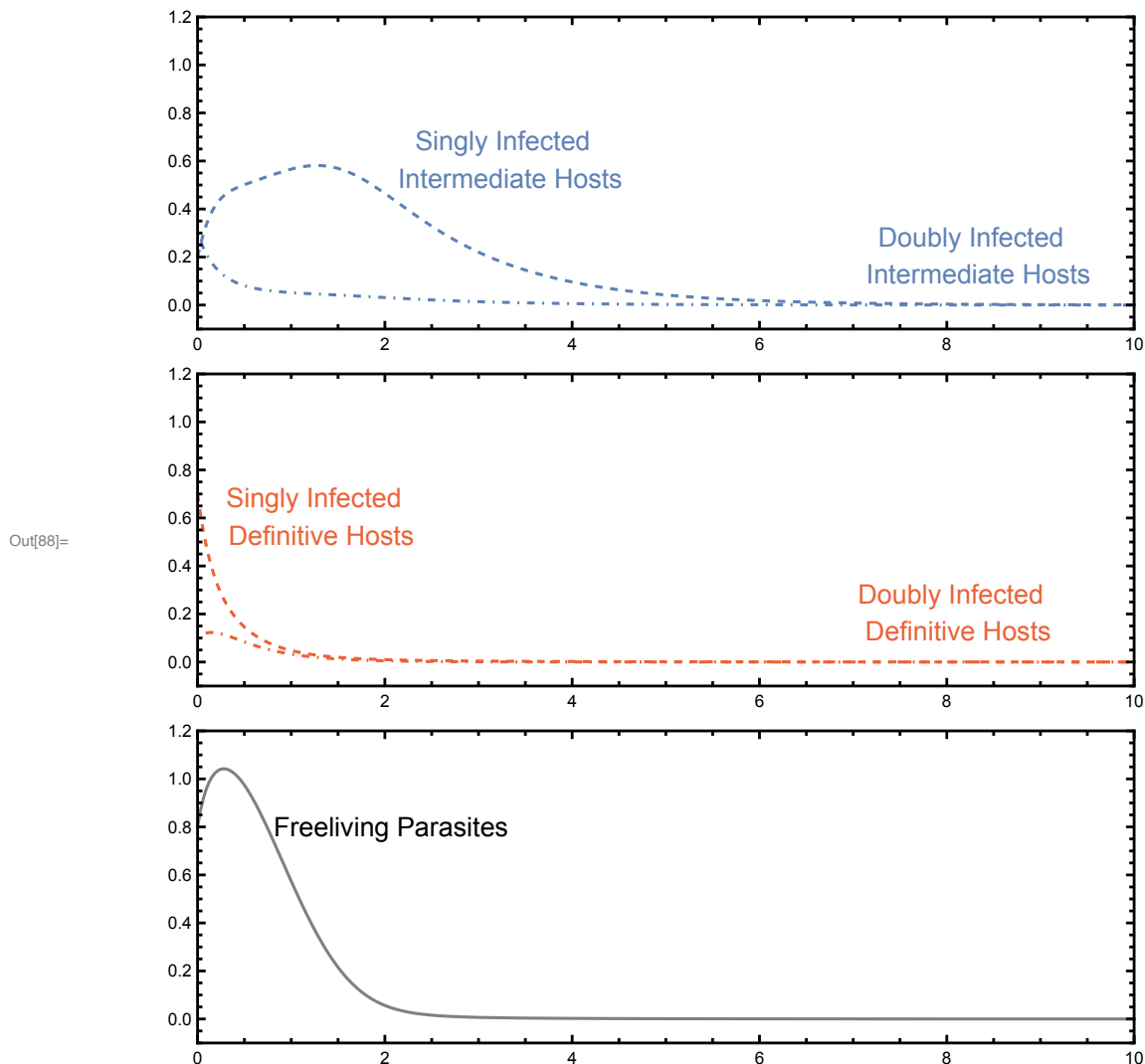
In[86]:= epilogHost1 =
    {{PointSize[Large], Point[Partition[Riffle[ieq1, deq1], 2]]}, Table[
        {Black, Text[Style[*, FontSize → 24], {ics[[i, 1, 2]], ics[[i, 4, 2]]}, {0, 0}]},
        {i, 1, Length[ics], 1}];
ppHost1 = ParametricPlot[{Evaluate[{Is[t], Ds[t]}/. ndsolNL[[1]],
    Evaluate[{Is[t], Ds[t]}/. ndsolNL[[2]],
    Evaluate[{Is[t], Ds[t]}/. ndsolNL[[3]]}], {t, 0, maxt}, AspectRatio → 1,
    Frame → includeFrame, PlotStyle → {Darker[Green], Black, Darker[Green]},
    FrameStyle → frameStyle, PlotRange → range1, FrameLabel →
        {"Susceptible Intermediate Host", "Susceptible Definitive Host"},
    LabelStyle → labelStyle, ImageSize → imageSize / 1.1, Epilog → epilogHost1]

```





In[88]:= ppParasite1



## Disease stable equilibrium

In[89]:= (\*Set parameters \*)

```
prEcoDSNL = { $\rho \rightarrow 1.2$ ,  $d \rightarrow 0.9$ ,  $r \rightarrow 2.5$ ,  $\gamma \rightarrow 2.9$ ,  $\alpha \rightarrow 0$ ,  $\beta_w \rightarrow 1.5$ ,  $\beta_{ww} \rightarrow 1.5$ ,  
p  $\rightarrow 0.05$ ,  $c \rightarrow 1.4$ ,  $\mu \rightarrow 3.9$ ,  $\sigma \rightarrow 0$ ,  $q \rightarrow 0.05$ ,  
fw  $\rightarrow 45$ ,  $f_{ww} \rightarrow 45$ ,  $\delta \rightarrow 0.9$ ,  $k \rightarrow 0.26$ ,  $h \rightarrow 0.6$ };  
condsimplifyNL = { $\alpha_w \rightarrow \alpha$ ,  $\alpha_{ww} \rightarrow \alpha$ ,  $\sigma_w \rightarrow \sigma$ ,  $\sigma_{ww} \rightarrow \sigma$ };  
maxt = 80;
```

(\*Set range for the plot and initial values \*)

```
range2 = Full;
```

```
inits =
```

```
{Is[0] == 0.9, lw[0] == 0.27287415391003156`, lww[0] == 0.46321506893225517`,  
Ds[0] == 0.9, Dw[0] == 0.214956687585793`, Dww[0] == 0.018729116483236274`,  
W[0] == 0.025702772691754472`, Is[0] == 0.4588921303535818`},
```

```

lw[0] == 0.27287415391003156`, lww [0] == 0.46321506893225517`,
Ds[0] == 0.4460456255182149`, Dw [0] == 0.214956687585793`,
Dww[0] == 0.018729116483236274`, W [0] == 0.025702772691754472` },
{Is[0] == 1.45, lw [0] == 0.27287415391003156`, lww [0] == 0.46321506893225517`,
Ds[0] == 1.2, Dw [0] == 0.214956687585793`,
Dww[0] == 0.018729116483236274`, W [0] == 0.025702772691754472` });
(*Solving the ode for different initial values *)
ndsolsDS =
Table [NDSolve [Join [sysNDSolvefunc1 /. condsimplifyNL /. prEcoDSNL, inits [i]],
varRes, {t, 0, maxt}], {i, 1, Length [inits], 1}];

(*Get the dynamics of initial condition 2 *)
ieq2 = Evaluate [Is[maxt] /. ndsolsDS][[1]];
deq2 = Evaluate [Ds[maxt] /. ndsolsDS][[1]];

(*Annotation for different lines *)
epilogParas21 = {{ihostCol, Text [Style ["Singly Infected \n Intermediate Hosts",
FontSize → 14], {30, -1.}, {-1.05, 0}]},
{ihostCol, Text [Style ["Doubly Infected \n Intermediate Hosts", FontSize → 14],
{30, -4.2}, {-1.05, 0}]}];
epilogParas22 = {{dhostCol, Text [Style ["Singly Infected \n Definitive Hosts",
FontSize → 14], {30, -4.1}, {-1.05, 0}]},
{dhostCol, Text [Style ["Doubly Infected \n Definitive Hosts", FontSize → 14],
{30, -7.5}, {-1.05, 0}]}];
epilogParas23 = {{Black, Text [Style ["Freeliving Parasites", FontSize → 14],
{30, -3.2}, {-1.08, 0}]}];
trajectStyle = {{Gray, Thick }};
spcl = ndsolsDS[[2]];

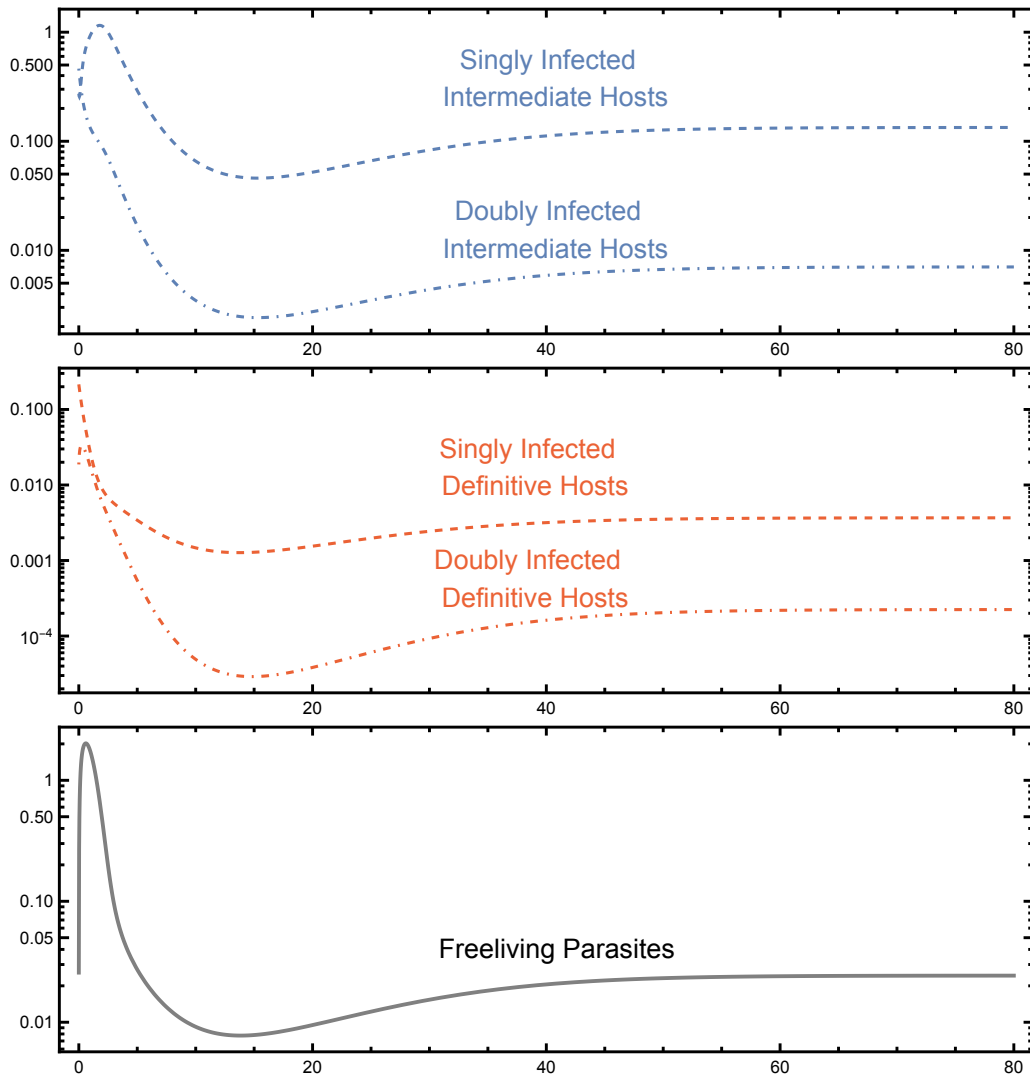
(*Plot trajectories of infected intermediate hosts,
definitive hosts, and free -living parasite with respect to time *)
ppParasite21 = LogPlot [Evaluate [{lw[t], lww[t]} /. spcl], {t, 0, maxt },
AspectRatio → 1/3, PlotStyle → {{ihostCol, Dashed }, {ihostCol, DotDashed }},
PlotRange → All, Frame → includeFrame, FrameStyle → frameStyle,
ImageSize → imageSize, Epilog → epilogParas21 ];
ppParasite22 = LogPlot [Evaluate [{Dw[t], Dww[t]} /. spcl], {t, 0, maxt },
AspectRatio → 1/3, PlotStyle → {{dhostCol, Dashed }, {dhostCol, DotDashed }},
PlotRange → All, Frame → includeFrame, FrameStyle → frameStyle,
ImageSize → imageSize, Epilog → epilogParas22 ];
ppParasite23 = LogPlot [Evaluate [{W[t]} /. spcl], {t, 0, maxt }, AspectRatio → 1/3,
PlotStyle → trajectStyle, PlotRange → All, Frame → includeFrame,
FrameStyle → frameStyle, ImageSize → imageSize, Epilog → epilogParas23 ];

ppParasite2 = GraphicsColumn [{ppParasite21, ppParasite22, ppParasite23 },
Spacings → 0, ImageSize → {600, 600}];

```

In[106]:= **ppParasite2**

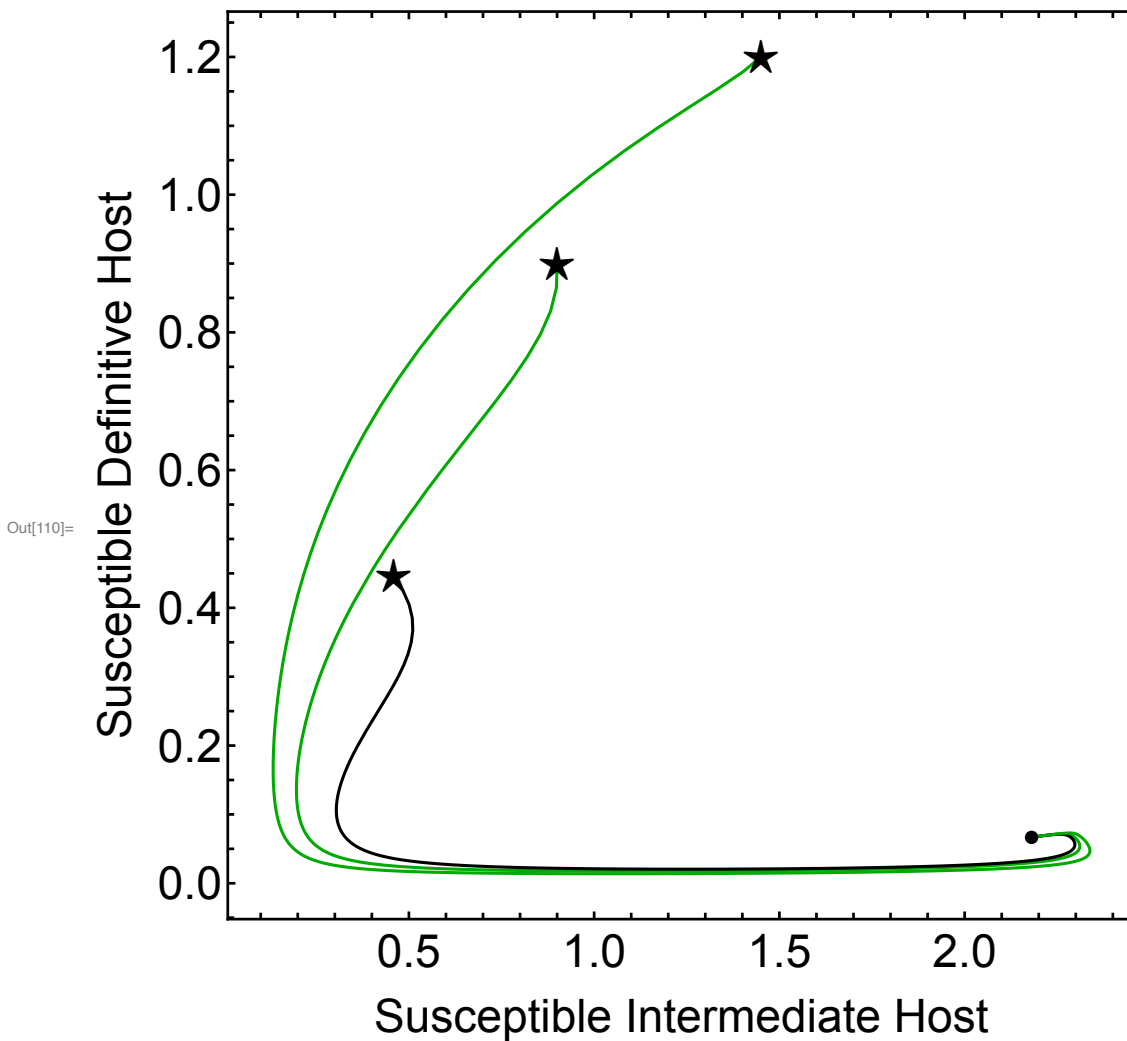
Out[106]=



```

In[109]:= (*Plot phase plane of susceptible intermediate and definitive hosts*)
epilogHost2 = {{PointSize[Large], Point[Partition[Riffle[ieq2, deq2], 2]]},
  Table[{Black, Text[Style[*, FontSize → 24],
    {inits[[i, 1, 2]], inits[[i, 4, 2]]}, {0, 0}]], {i, 1, Length[inits], 1}}];
ppHost2 = ParametricPlot[{Evaluate[{Is[t], Ds[t]} /. ndsolDS[[1]],
  Evaluate[{Is[t], Ds[t]} /. ndsolDS[[2]],
  Evaluate[{Is[t], Ds[t]} /. ndsolDS[[3]]}], {t, 0, maxt}, AspectRatio → 1,
Frame → includeFrame, PlotStyle → {Darker[Green], Black, Darker[Green]},
FrameStyle -> frameStyle, PlotRange → range2, FrameLabel →
{"Susceptible Intermediate Host", "Susceptible Definitive Host"},
LabelStyle → labelStyle, ImageSize → imageSize / 1.1, Epilog → epilogHost2]

```

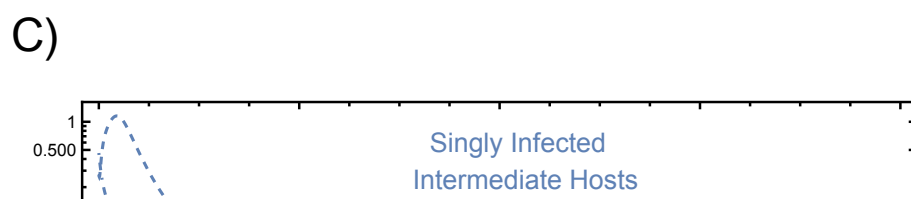
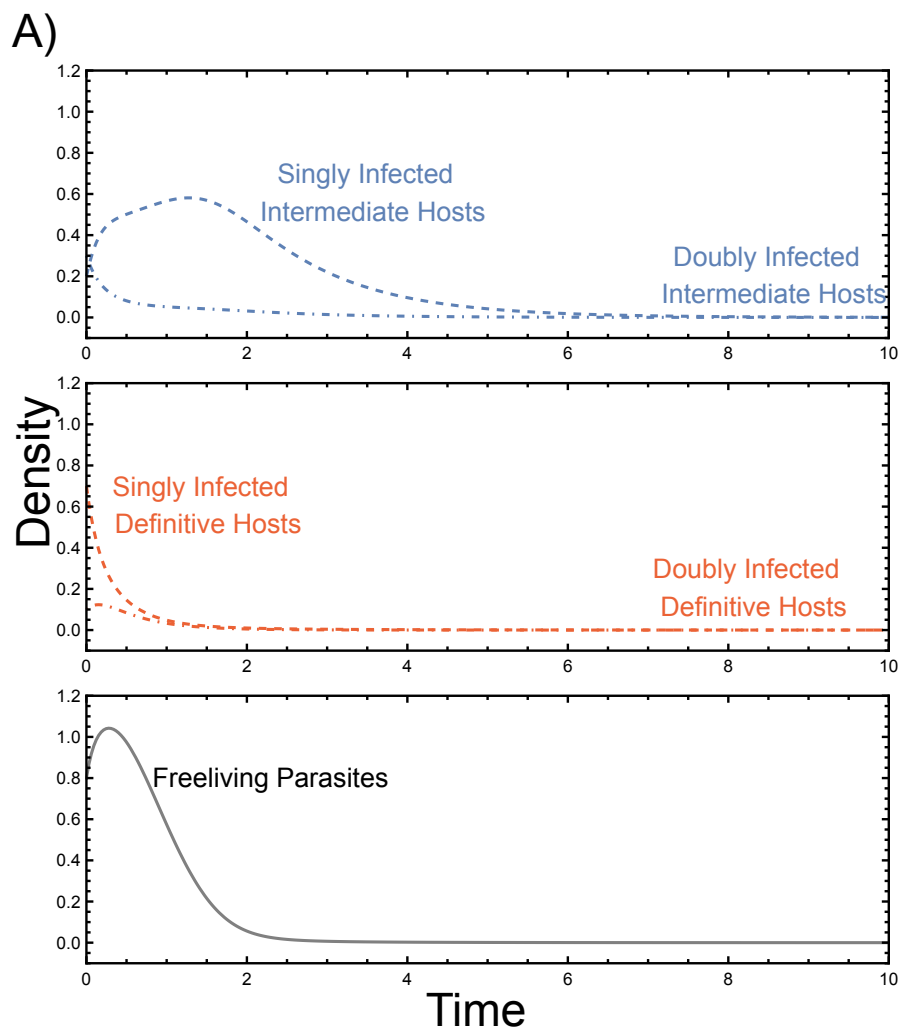


## Combined disease free and disease stable ecology plot (Figure 3)

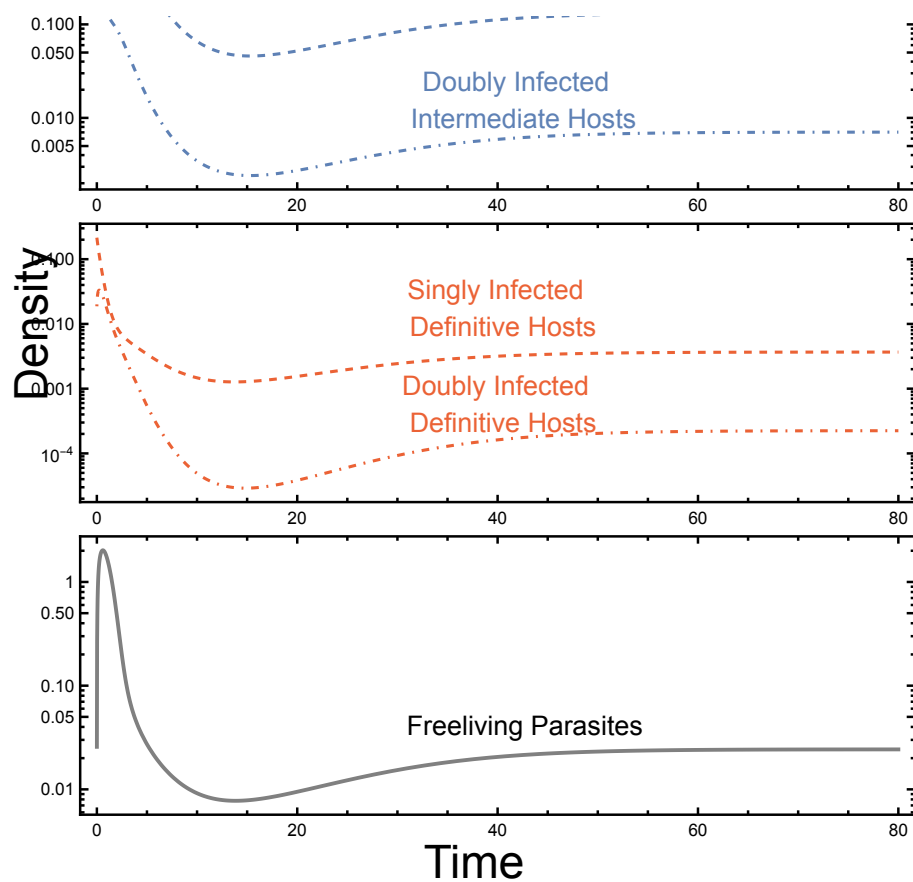
```

In[111]:= epilogAxesCombiPanel = {Text [Style["A)", Black, 24 ], {100, -10}],
  Text [Style["Time", Black, 24 ], {330, -520}],
  Rotate [Text [Style["Density", Black, 24 ], {100, -240}], 90 Degree ],
  Text [Style["B)", Black, 24 ], {660, -10}], Text [Style["C)", Black, 24 ], {100, -590}],
  Text [Style["Time", Black, 24 ], {330, -1120}],
  Rotate [Text [Style["Density", Black, 24 ], {100, -840}], 90 Degree ],
  Text [Style["D)", Black, 24 ], {660, -590}];
GraphicsGrid [{{ppParasite1, ppHost1 }, {ppParasite2, ppHost2 }},
  ImageSize → {1200, 1200 }, Spacings → 0, Epilog → epilogAxesCombiPanel ]
(*Export ["ecotraject _nonlinear.pdf", %, ImageResolution → figResolution]*)

```



Out[112]=



Bifurcation for reproduction in single infection  $f_w$  ( $f_{ww} = \epsilon f_w$ )  
(Figure 5)

```
In[113]:= fwstart = 33;
          fwend   = 45;
          fwstep  = 0.1;
          fwrang  = Range[fwstart, fwend, fwstep];
```

Parameter set 1 (bistability) Figure 5 Panel B-C

CALCULATE BIFURCATION

```

In[117]:= parfw1 = { $\rho \rightarrow 1.2$ ,  $d \rightarrow 0.9$ ,  $r \rightarrow 2.5$ ,  $\gamma \rightarrow 2.9$ ,  $\alpha_w \rightarrow 0$ ,
                $\alpha_{ww} \rightarrow 0$ ,  $\beta_w \rightarrow 1.5$ ,  $\beta_{ww} \rightarrow 1.5$ ,  $p \rightarrow 0.05$ ,  $c \rightarrow 1.4$ ,  $\mu \rightarrow 3.9$ ,
                $\sigma_w \rightarrow 0$ ,  $\sigma_{ww} \rightarrow 0$ ,  $q \rightarrow 0.05$ ,  $\delta \rightarrow 0.9$ ,  $k \rightarrow 0.26$ ,  $\epsilon \rightarrow 4$ ,  $h \rightarrow 0.6$ };
(*Find all solutions of the system with the given parameter values *)
solsfwAll1 =
  NSolve[Thread[(odesRes /. func1 /. forceInf /. fww ->  $\epsilon$ fw /. parfw1 /. fw -> 36) == 0],
    varRes, Reals];

(*Select disease circulation solutions *)
fwAllPositive1 = ParallelTable[NSolvePositive[sysfunc1 /. fww ->  $\epsilon$ fw,
  parfw1, fw -> i, varRes, eq], {i, fwstart, fwend, fwstep}];
(*Select disease free solutions *)
solsfwzero1 =
  Select[solsfwAll1, (Is /. #) > 0 && (lw /. #) == 0 && (lww /. #) == 0 && (Ds /. #) > 0 &&
    (Dw /. #) == 0 && (Dww /. #) == 0 && (W /. #) == 0 &][[1]];

CALCULATE DYNAMICS

In[121]:= maxt1 = 300;
init1 = {Is[0] == 0.1588921303535818`,
  lw[0] == 0.27287415391003156`, lww [0] == 0.46321506893225517`,
  Ds[0] == 0.4460456255182149`, Dw [0] == 0.214956687585793`,
  Dww[0] == 0.018729116483236274`, W [0] == 0.025702772691754472` };
solsbistable1 = NDSolve[Join[sysNDSolvefunc1 /. fww ->  $\epsilon$ fw /. parfw1 /. fw -> 39 /.
   $\epsilon \rightarrow 4$ , init1], varRes, {t, 0, maxt1}];
maxt2 = 300;
init2 = {Is[0] == 1.1588921303535818`,
  lw[0] == 0.27287415391003156`, lww [0] == 0.46321506893225517`,
  Ds[0] == 0.4460456255182149`, Dw [0] == 0.214956687585793`,
  Dww[0] == 0.018729116483236274`, W [0] == 0.025702772691754472` };
solsbistable2 = NDSolve[Join[sysNDSolvefunc1 /. fww ->  $\epsilon$ fw /. parfw1 /. fw -> 39 /.
   $\epsilon \rightarrow 4$ , init2], varRes, {t, 0, maxt2}];

PLOT

```

```

In[127]:= fwlist = fw /. Flatten [fwAllPositive1, 1 ];
eqfwlist = eq /. Flatten [fwAllPositive1, 1 ];
eqfwzero = ConstantArray [solswzero1, Length [fwrangle]];
col1 = ColorData ["GeologicAges", "ColorList" ][[45]];
colLine1 = ColorData ["GeologicAges", "ColorList" ][[27]]
colLine2 = ColorData ["GeologicAges", "ColorList" ][[25]]
thickness = 0.005;
rangelw = {{fwstart, fwend }, {-0.1, 1. }};

lwfwlist = lw /. eqfwlist;


sdat1 = MakeListPlotData [fwlist, lwfwlist ];
markers1 = ListStableMark [Jmatfunc1 /. fww ->  $\epsilon$ fw,
    parfw1, fw, fwlist, eqfwlist, {"★", "○", "●"}];


zdat1 = MakeListPlotData [fwrangle, lw /. eqfwzero ];
markerz1 = ListStableMark [Jmatfunc1 /. fww ->  $\epsilon$ fw,
    parfw1, fw, fwrangle, eqfwzero, {"★", "○", "●"}];

plw = ListPlot [Join [sdat1, zdat1 ], PlotMarkers -> Join [markers1, markerz1 ],
    PlotStyle -> col1, PlotRange -> rangelw, ImageSize -> imageSize,
    Frame -> includeFrame, FrameLabel -> {"Reproduction in single infection ( $f_w$ )",
        "Equilibrium of density of \n singly infected intermediate host" },
    LabelStyle -> labelStyle, FrameStyle -> frameStyle ]
Export ["reproduction _bifurcationB.pdf", %, ImageResolution -> figResolution ]

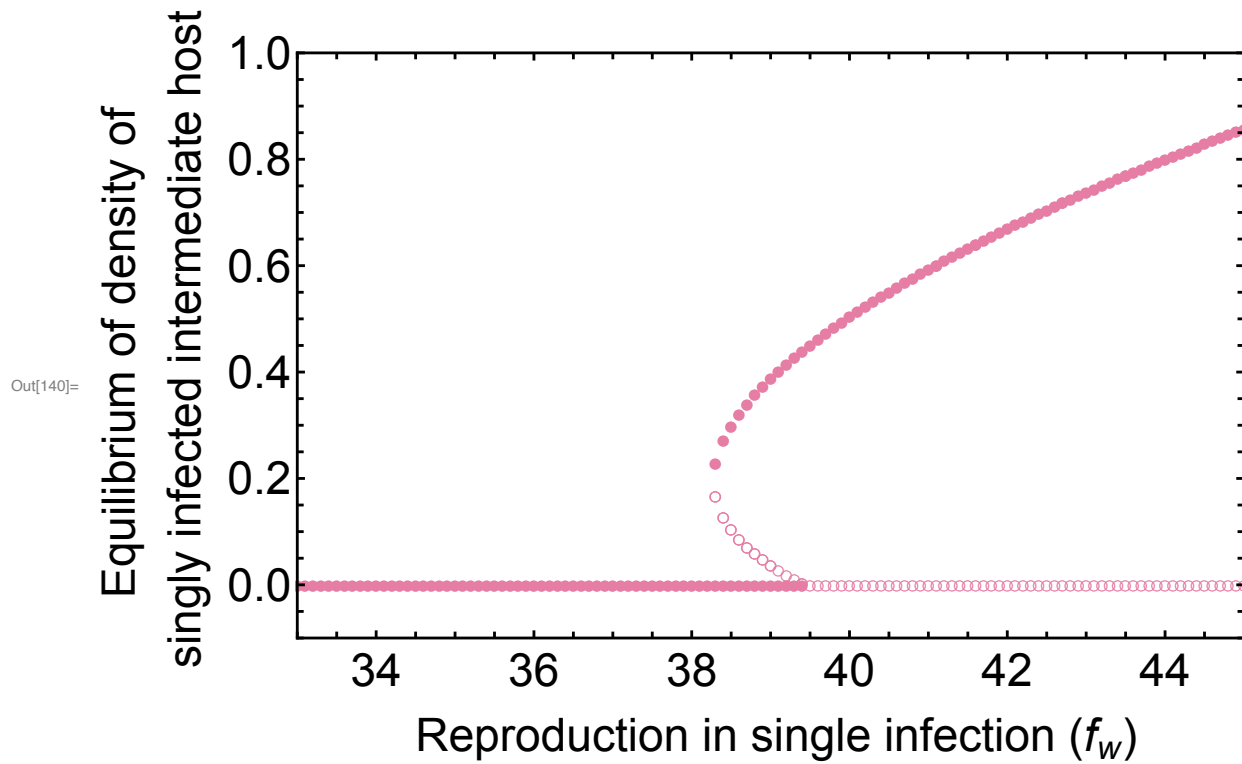
parfw1Dynamics =
LogLogPlot [{Evaluate [lw[t]] /. solsbistable1, Evaluate [lww[t]] /. solsbistable1,
    Evaluate [lw[t]] /. solsbistable2, Evaluate [lww[t]] /. solsbistable2 },
    {t, 0, maxt1 }, PlotStyle -> {{colLine1, Dashed, Thickness [thickness * 2]},
        {colLine1, DotDashed, Thickness [thickness * 2]}, {colLine2, Dashed,
            Thickness [thickness ]}, {colLine2, DotDashed, Thickness [thickness ]}},
    PlotRange -> {{0, maxt1 }, {-0.1, 1.9 }}, Frame -> includeFrame,
    FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    ImageSize -> imageSize, FrameTicksStyle -> Directive [Black, 20 ],
    FrameLabel -> {"Time", "Dynamics of singly and doubly \n infected intermediate host" }]
Export ["reproduction _bifurcationD.pdf", %, ImageResolution -> figResolution ]

```

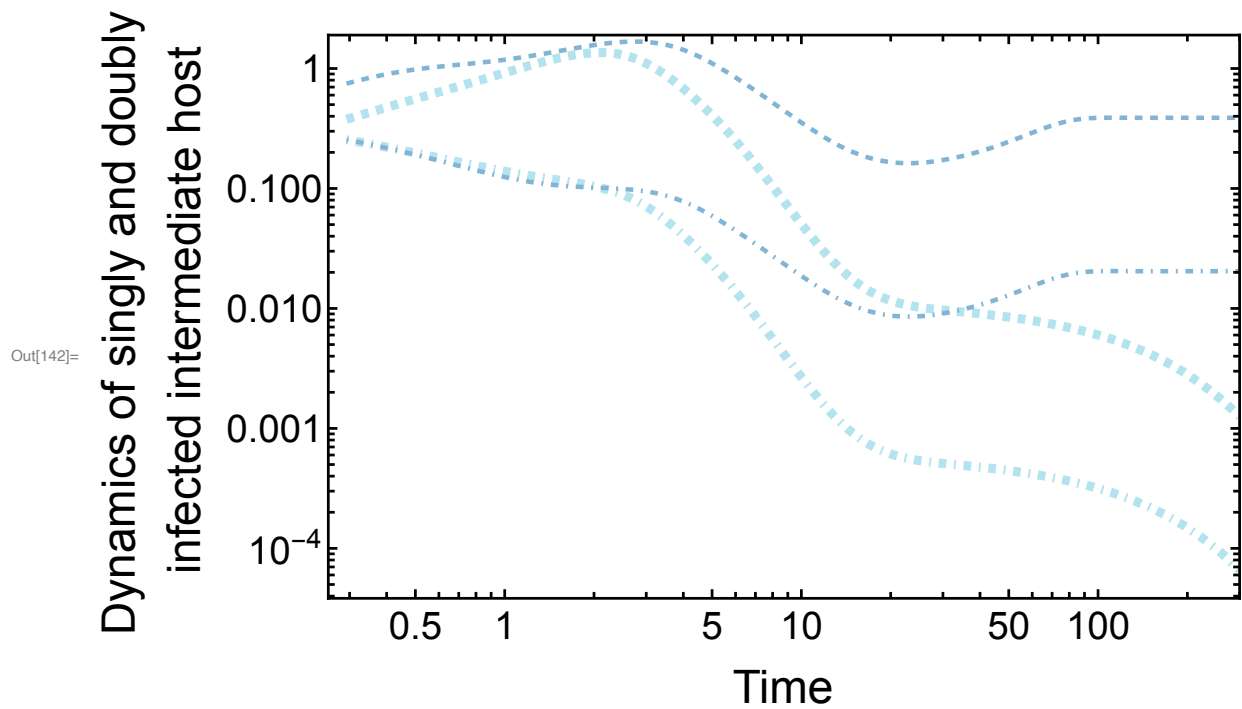
Out[131]= 

Out[132]= 





Out[141]= reproduction\_bifurcationB.pdf



Out[143]= reproduction\_bifurcationD.pdf

## Parameter set 2 (single equilibrium) Figure 5 Panel B

CALCULATION

```

In[144]:= parfw2 = { $\rho \rightarrow 1.2$ ,  $d \rightarrow 0.9$ ,  $r \rightarrow 2.5$ ,  $\gamma \rightarrow 2.9$ ,  $\alpha_w \rightarrow 0$ ,
   $\alpha_{ww} \rightarrow 0$ ,  $\beta_w \rightarrow 1.5$ ,  $\beta_{ww} \rightarrow 1.5$ ,  $p \rightarrow 0.05$ ,  $c \rightarrow 1.4$ ,  $\mu \rightarrow 3.9$ ,
   $\sigma_w \rightarrow 0$ ,  $\sigma_{ww} \rightarrow 0$ ,  $q \rightarrow 0.05$ ,  $\delta \rightarrow 0.9$ ,  $k \rightarrow 0.26$ ,  $\epsilon \rightarrow 1$ ,  $h \rightarrow 0.6$ };
(*Find all solutions of the system with the given parameter values *)
solsfwAll2 =
  NSolve[Thread[(odesRes /. func1 /. forceInf /. fww ->  $\epsilon$ fw /. parfw2 /. fw -> 36) == 0],
    varRes, Reals];

(*Select disease circulation solutions *)
solsfw2 =
  Select[solsfwAll2, (ls /. #) > 0 && (lw /. #) > 0 && (lww /. #) > 0 && (W /. #) > 0 &][[1]];

(*Select disease free solutions *)
solsfwzero2 =
  Select[solsfwAll2, (ls /. #) > 0 && (lw /. #) == 0 && (lww /. #) == 0 && (Ds /. #) > 0 &&
    (Dw /. #) == 0 && (Dww /. #) == 0 && (W /. #) == 0 &][[1]];

(*Select disease circulation solutions *)
fwAllPositive2 = ParallelTable[NSolvePositive[sysfunc1 /. fww ->  $\epsilon$ fw,
  parfw2, fw -> i, varRes, eq], {i, fwstart, fwend, fwstep}];

```

Part: Part 1 of {} does not exist.

## PLOTTING

```

In[149]:= rangelw = {{fwstart, fwend}, {-0.01, 0.2}};
fwlist = fw /. Flatten[fwAllPositive2, 1];
eqfwlist = eq /. Flatten[fwAllPositive2, 1];
eqfwzero = ConstantArray[solsfwzero2, Length[fwrange]];
col2 = ColorData["GeologicAges", "ColorList"][[35]];

lwfwlist = lw /. eqfwlist;

sdat3 = MakeListPlotData[fwlist, lwfwlist];
markers3 = ListStableMark[Jmatfunc1 /. fww ->  $\epsilon$ fw,
  parfw2, fw, fwlist, eqfwlist, {"★", "○", "●"}];
zdat3 = MakeListPlotData[fwrange, lw /. eqfwzero];
markerz3 = ListStableMark[Jmatfunc1 /. fww ->  $\epsilon$ fw,
  parfw2, fw, fwrange, eqfwzero, {"★", "○", "●"}];

pplw = ListPlot[Join[sdat3, zdat3], PlotMarkers -> Join[markers3, markerz3],
  PlotStyle -> col2, PlotRange -> rangelw, Frame -> includeFrame,
  ImageSize -> imageSize, FrameLabel -> {"Reproduction in single infection ( $f_w$ )",
    "Equilibrium of density of \n singly infected intermediate host"},
  LabelStyle -> labelStyle, FrameStyle -> frameStyle (*,
  PlotLabel -> Pane["B)  $f_{ww} = f_w$ ", Alignment -> Left, ImageSize -> panellabel *]);
Export["reproduction_bifurcationC.pdf", %, ImageResolution -> figResolution]

Out[160]= reproduction_bifurcationC.pdf

```

## Bifurcation $\epsilon$ and $f_w$ ( $f_{ww} = \epsilon f_w$ ) Figure 5 Panel A

### CALCULATE

```

par $\epsilon$ fw = { $\rho \rightarrow 1.2$ ,  $d \rightarrow 0.9$ ,  $r \rightarrow 2.5$ ,  $\gamma \rightarrow 2.9$ ,  $\alpha_w \rightarrow 0$ ,
 $\alpha_{ww} \rightarrow 0$ ,  $\beta_w \rightarrow 1.5$ ,  $\beta_{ww} \rightarrow 1.5$ ,  $p \rightarrow 0.05$ ,  $c \rightarrow 1.4$ ,  $\mu \rightarrow 3.9$ ,
 $\sigma_w \rightarrow 0$ ,  $\sigma_{ww} \rightarrow 0$ ,  $q \rightarrow 0.05$ ,  $\delta \rightarrow 0.9$ ,  $k \rightarrow 0.26$ ,  $h \rightarrow 0.6$ };
 $\epsilon$ start = 0;
 $\epsilon$ end = 6;
 $\epsilon$ interval = 0.025;
fwstart = 33;
fwend = 45;
fwinterval = 0.25;

```

```

 $\epsilon$ fwAllresults = ParallelTable [NSolveCodim2Positive [
  sysfunc1 /. fww  $\rightarrow$   $\epsilon$ fw, par $\epsilon$ fw, {fw  $\rightarrow$  i}, { $\epsilon \rightarrow$  j},  $\epsilon$ fw, eq, varRes, 12 ],
  {i, fwstart, fwend, fwinterval }, {j,  $\epsilon$ start,  $\epsilon$ end,  $\epsilon$ interval}];

```

(kernel 5) NSolve::sfail : Subsystem could not be solved for

$$\frac{172209 Ds}{183067} - \frac{117425 Dw}{183067} - \frac{116690 Dww}{183067} - \frac{138285 Is}{183067} + \frac{169206 Iw}{183067} - \frac{151916 Iww}{183067} - \frac{185189 W}{183067}$$

at value -1.8595047574076894. The likely cause is failure to detect zero due to low precision. The likely effect is the loss of one or more solutions. Increasing WorkingPrecision might prevent some solutions from being lost.

(kernel 3) NSolve::sfail : Subsystem could not be solved for

$$\frac{172209 Ds}{183067} - \frac{117425 Dw}{183067} - \frac{116690 Dww}{183067} - \frac{138285 Is}{183067} + \frac{169206 Iw}{183067} - \frac{151916 Iww}{183067} - \frac{185189 W}{183067}$$

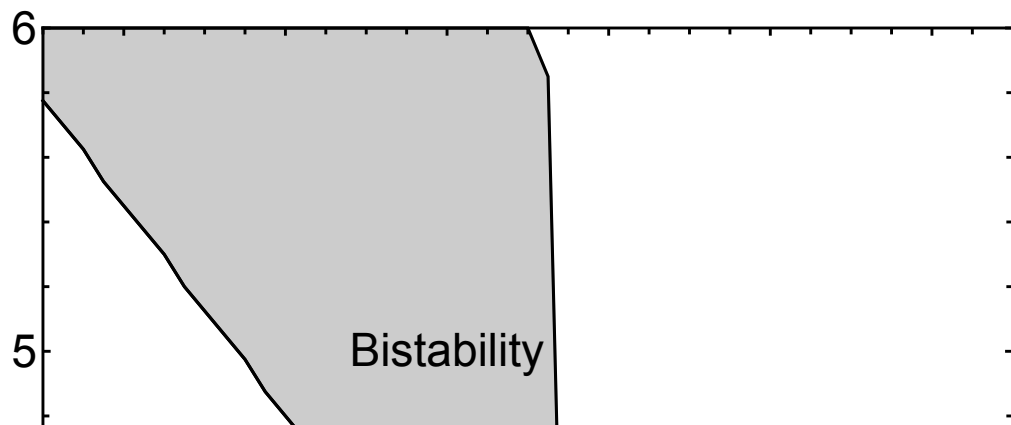
at value -1.8594515089778949. The likely cause is failure to detect zero due to low precision. The likely effect is the loss of one or more solutions. Increasing WorkingPrecision might prevent some solutions from being lost.

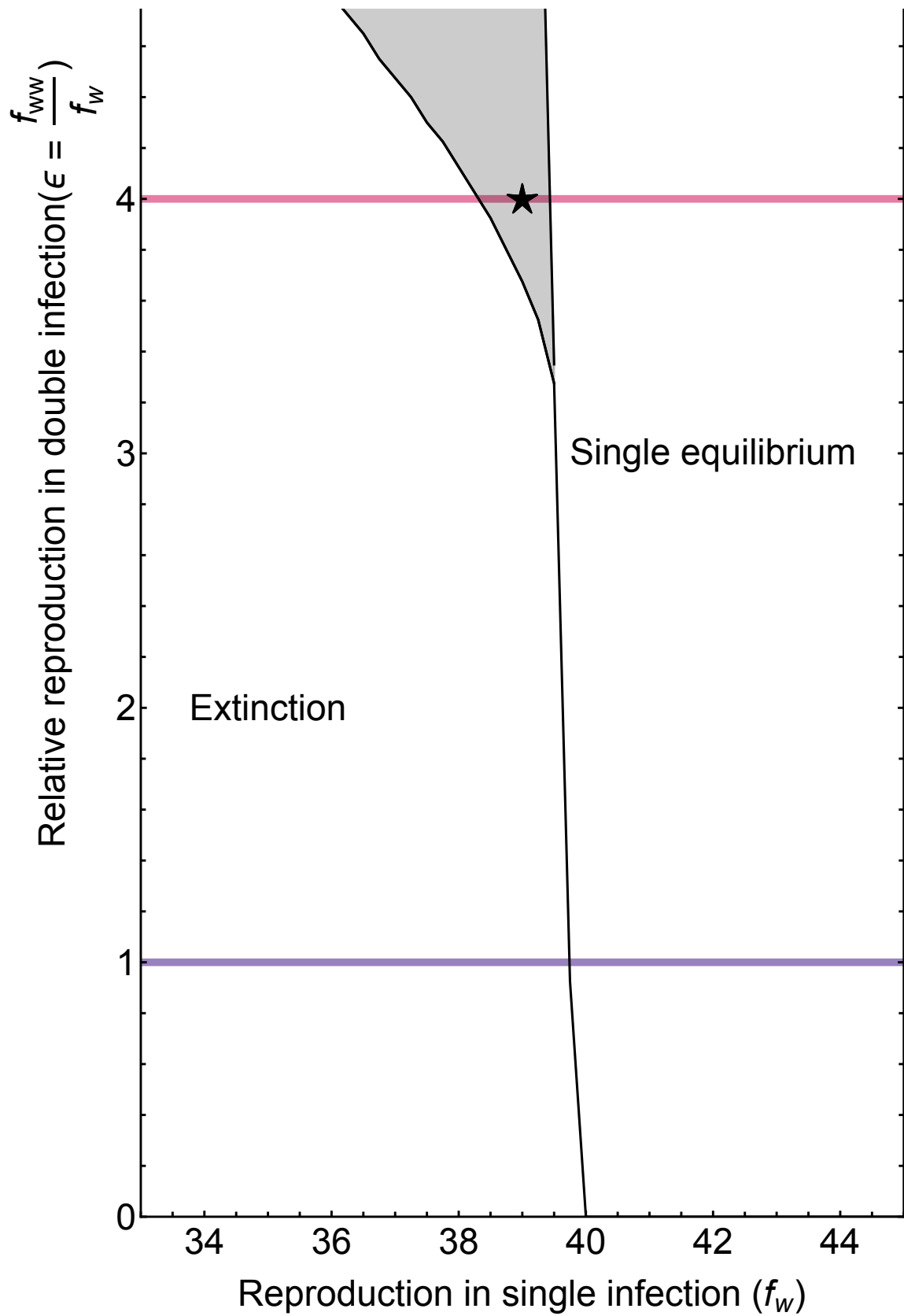
```

In[169]:= eqfww = eq /. Flatten [efwAllresults, 2 ];
efwlist = efww /. Flatten [efwAllresults, 2 ];
rangeefw = {{fwwstart, fwwend }, {ϵstart, ϵend}};

marklist = ListStableMarkTwoParameters [
  Jmatfunc1 /. fww -> efww, par efww, {ϵ, fww}, efwlist, eq efww, colorstable, True ];
MakeListPlotData [efwlist [[All, 1]], efwlist [[All, 2]]];
p0 =
  ListPlot [% , PlotStyle -> marklist, PlotRange -> rangeefw, ImageSize -> imageSize,
    Frame -> includeFrame, FrameLabel -> {"ϵ", "Reproduction in single infection fw"},
    GridLines -> {{1}, {}}, GridLinesStyle -> Directive [Thick,
      Red], LabelStyle -> labelStyle, FrameStyle -> frameStyle, AspectRatio -> 1];
GetBoundaryLineBiStable [efwlist ];
p1 = ListLinePlot [% , PlotRange -> rangeefw, Frame -> includeFrame,
  FrameStyle -> frameStyle, FrameLabel -> {"Reproduction in single infection (fw)",
    "Relative reproduction in double infection (ϵ =  $\frac{f_{ww}}{f_w}$ )"},
  LabelStyle -> labelStyle, GridLines -> {{}, {{1, col2}, {4, col1}}},
  GridLinesStyle -> Thickness [0.01], PlotStyle -> Black, Filling -> {1 -> {2}}];
GetBoundaryLineSingle [efwlist ];
p2 = ListLinePlot [% , PlotRange -> rangeefw,
  Frame -> includeFrame, FrameStyle -> frameStyle, PlotStyle -> Black];
ip = ip = ListLinePlot [{39, 4}], PlotRange -> rangeefw, PlotMarkers -> {"★", 28},
  Frame -> includeFrame, FrameStyle -> frameStyle,
  PlotStyle -> Black, ImageSize -> imageSize, LabelStyle -> labelStyle];
GraphicsGrid [{p0, Show [p1, p2]}];
pefw = Show [p1, p2, ip], ImageSize -> {imageSize, imageSize * 2},
  Epilog -> {Text [Style ["Extinction", FontSize -> 24], {35, 2}],
    Text [Style ["Single equilibrium", FontSize -> 24], {42, 3}],
    Text [Style ["Bistability", FontSize -> 24], {38, 5.}],
  AspectRatio -> 2/1, LabelStyle -> labelStyle,
  ImagePadding -> {{90(*left*), 5(*right*)}, {65(*bottom*), 10(*top*)}}];
Export ["reproduction _bifurcationA.pdf", %, ImageResolution -> figResolution ]

```





Out[181]=

Out[182]= reproduction\_bifurcationA.pdf

## Contour plot (Figure 4)

### Manipulation ratio representation

```
In[222]:=  $\beta_{wwstart}$  = 0.1;
 $\beta_{wwend}$  = 3;
 $\beta_{wwstep}$  = 0.03;
 $\epsilon_{start}$  = 0.5;
 $\epsilon_{end}$  = 12;
 $\epsilon_{step}$  = 0.2;
alpha = 0.6;
```

## Changing reproduction in single infection

$fw = 37, \beta_w = 1.65$  (Figure 6 Panel D)

### CALCULATION

```
In[245]:= (*Set parameters *)
parmanip  $\beta\epsilon$  = { $\rho \rightarrow 1.2$ ,  $d \rightarrow 0.9$ ,  $r \rightarrow 2.5$ ,  $\gamma \rightarrow 2.9$ ,  $\alpha_w \rightarrow 0$ ,  $\alpha_{ww} \rightarrow 0$ ,
   $p \rightarrow 0.05$ ,  $c \rightarrow 1.4$ ,  $\mu \rightarrow 3.9$ ,  $\sigma_w \rightarrow 0$ ,  $\sigma_{ww} \rightarrow 0$ ,  $q \rightarrow 0.05$ ,
   $\delta \rightarrow 0.9$ ,  $k \rightarrow 0.26$ ,  $fw \rightarrow 37$ ,  $h \rightarrow 0.6$ ,  $\beta_w \rightarrow 1.65$ };
range1 = {{ $\beta_{wwstart} / (\beta_w /. \text{parmanip } \beta\epsilon)$ ,  $\beta_{wwend} / (\beta_w /. \text{parmanip } \beta\epsilon)$ },
  { $\epsilon_{start}$ ,  $\epsilon_{end} + 0.1$ }}
Out[246]:= {{0.0606061, 1.81818}, {0.5, 12.1}}

(*Bifurcation value*)
manip $\beta\epsilon$ AllResults = ParallelTable[NSolveCodim2Positive[sysfunc1 /.  $fw \rightarrow \epsilon fw$ ,
  parmanip $\beta\epsilon$ , { $\beta_{ww} \rightarrow x$ }, { $\epsilon \rightarrow y$ },  $\beta\epsilon$ , eq, varRes, 15],
  {x,  $\beta_{wwstart}$ ,  $\beta_{wwend}$ ,  $\beta_{wwstep}$ }, {y,  $\epsilon_{start}$ ,  $\epsilon_{end}$ ,  $\epsilon_{step}$ }]];
```

### PLOT

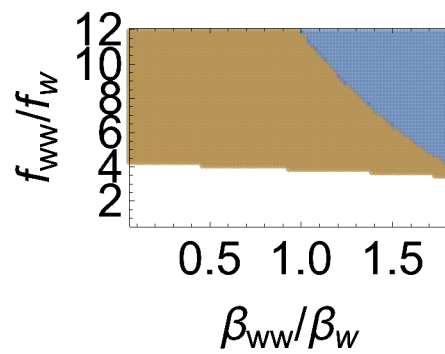
```

equilibrium = eq /. Flatten [manip  $\beta\epsilon$ AllResults, 2];
parslist =  $\beta\epsilon$  /. Flatten [manip  $\beta\epsilon$ AllResults, 2];

(*Create markers for equilibrium *)
marklist = ListStableMarkTwoParameters [Jmatfunc1 /. fww ->  $\epsilon$ fww,
    parmanip  $\beta\epsilon$ , { $\beta_{ww}$ ,  $\epsilon$ }, parslist, equilibrium, colorstable, True];
MakeListPlotData [parslist [[All, 1]]/( $\beta_w$  /. parmanip  $\beta\epsilon$ ), parslist [[All, 2]]];
(*Plot equilibrium as points with blue as stable points and orange as unstable points *)
p $\beta\epsilon$  = ListPlot [% , PlotStyle -> marklist,
    PlotRange -> range1, ImageSize -> imageSize, Frame -> includeFrame,
    FrameLabel -> {" $\beta_{ww}/\beta_w$ ", " $f_{ww}/f_w$ "}, LabelStyle -> labelStyle];
(*MapThread to get ratio as xaxis *)
(*Connect points at boundary *)
MapThread [{#1/#2 &,
    {GetBoundaryLineBiStable [parslist], {ConstantArray [{ $\beta_w$  /. parmanip  $\beta\epsilon$ , 1},
        Length @GetBoundaryLineBiStable [parslist] [[1, All]]}, ConstantArray [
            { $\beta_w$  /. parmanip  $\beta\epsilon$ , 1}, Length @GetBoundaryLineBiStable [parslist] [[2, All]]}}];
(*Draw upper boundary *)lp $\beta\epsilon$ 1 = ListLinePlot [% , PlotRange -> range1, Frame ->
    includeFrame, FrameStyle -> frameStyle, LabelStyle -> labelStyle, PlotStyle ->
    {{Thick, Thickness [thick], colbifur [[3]]}, {Thick, Thickness [thick], colbifur [[3]]}},
    Filling -> {1 -> {{2}, Directive [Opacity [alpha], colbifurfilling [[3]]]}},
    ImageSize -> imageSize];
GetBoundaryLineSingle [parslist]/ConstantArray [{ $\beta_w$  /. parmanip  $\beta\epsilon$ , 1},
    Length @GetBoundaryLineSingle [parslist]];
(*Draw lower boundary *)lp $\beta\epsilon$ 2 = ListLinePlot [% , PlotRange -> range1,
    Frame -> includeFrame, FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    PlotStyle -> {Thick, Thickness [thick], colbifur [[3]]},
    FrameLabel -> {"Relative manipulation in double infections ( $\frac{\beta_{ww}}{\beta_w}$ )",
        "Relative reproduction \n in double infections ( $\frac{f_{ww}}{f_w}$ )"},
    Filling -> Bottom, FillingStyle -> Directive [LightGray, HatchFilling []]];
lp $\beta\epsilon$  = Show [{lp $\beta\epsilon$ 2, lp $\beta\epsilon$ 1}];
GraphicsGrid [{p $\beta\epsilon$ , lp $\beta\epsilon$ }]

```

Out[257]=



double infections( $\frac{f_{ww}}{f_w}$ )

e. Manip

$f_w = 38, \beta_w = 1.65$  (Figure 6 Panel B)

(\*Set parameters \*)

parmanip  $\beta \in 1 = \{\rho \rightarrow 1.2, d \rightarrow 0.9, r \rightarrow 2.5, \gamma \rightarrow 2.9, \alpha_w \rightarrow 0, \alpha_{ww} \rightarrow 0,$

$p \rightarrow 0.05, c \rightarrow 1.4, \mu \rightarrow 3.9, \sigma_w \rightarrow 0, \sigma_{ww} \rightarrow 0, q \rightarrow 0.05,$

$\delta \rightarrow 0.9, k \rightarrow 0.26, f_w \rightarrow 38, h \rightarrow 0.6, \beta_w \rightarrow 1.65\}$ ;

(\*Equilibrium values \*)manip  $\beta \in 1$  AllResults = ParallelTable [NSolveCodim2Positive [

sysfunc1 /. fww  $\rightarrow \epsilon f_w$ , parmanip  $\beta \in 1, \{\beta_{ww} \rightarrow x\}, \{\epsilon \rightarrow y\}, \beta \in$ , eq, varRes, 15 ],

$\{x, \beta_{wwstart}, \beta_{wwend}, \beta_{wwstep}\}, \{y, \epsilonstart, \epsilonend, \epsilonstep\}\}$ ];



```

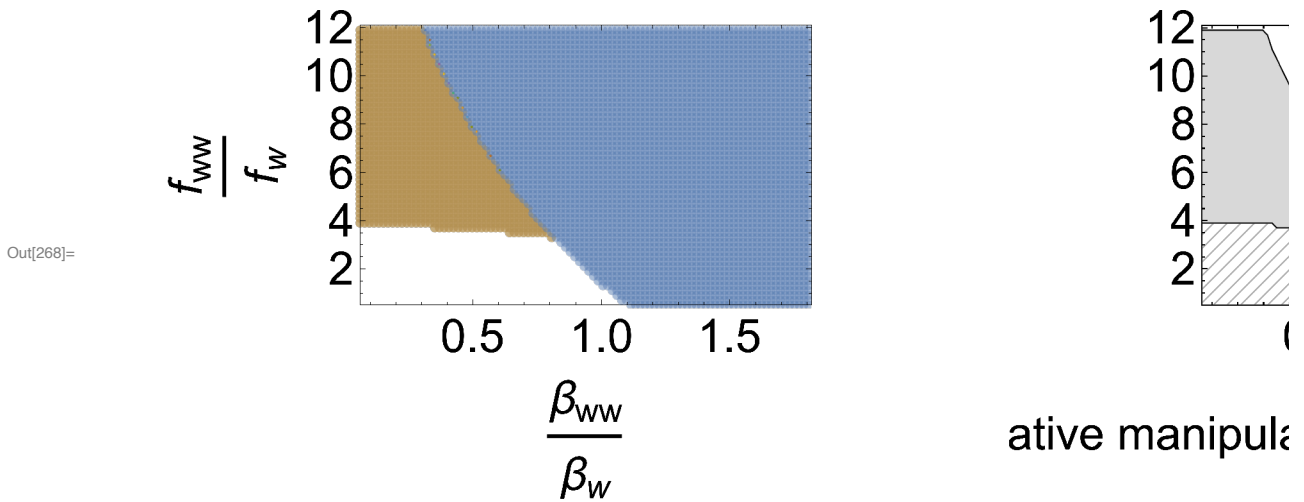
equilibrium = eq /. Flatten [manip  $\beta\epsilon 1$  AllResults, 2 ];
parslist =  $\beta\epsilon$  /. Flatten [manip  $\beta\epsilon 1$  AllResults, 2 ];

(*Mark equilibrium as stable and unstable *)
marklist = ListStableMarkTwoParameters [Jmatfunc1 /. fww ->  $\epsilon$ fw,
    parmanip  $\beta\epsilon 1$ , { $\beta_{ww}$ ,  $\epsilon$ }, parslist, equilibrium, colorstable, True ];
(*Plot unstable and stable equilibrium *)
MakeListPlotData [parslist [[All, 1]]/( $\beta_w$  /. parmanip  $\beta\epsilon 1$ ), parslist [[All, 2]]];
p $\beta\epsilon 1$  = ListPlot [% , PlotStyle -> marklist, PlotRange -> range1, ImageSize -> imageSize,

    Frame -> includeFrame, FrameLabel -> { " $\frac{\beta_{ww}}{\beta_w}$ ", " $\frac{f_{ww}}{f_w}$ " }, LabelStyle -> labelStyle ];

(*Connect boundary points *)
(*MapThread to get ratio as xaxis *)
MapThread [{#1/#2 &,
    {GetBoundaryLineBiStable [parslist ], { ConstantArray [{ $\beta_w$  /. parmanip  $\beta\epsilon 1$ , 1},
        Length @GetBoundaryLineBiStable [parslist ][[1, All]], ConstantArray [
            { $\beta_w$  /. parmanip  $\beta\epsilon 1$ , 1}, Length @GetBoundaryLineBiStable [parslist ][[2, All]]}}];
(*Upper boundary *)lp $\beta\epsilon 1$  = ListLinePlot [% , PlotRange -> range1, Frame -> includeFrame,
    FrameStyle -> frameStyle, LabelStyle -> labelStyle, FrameLabel ->
    { "Relative manipulation in double infections ( $\frac{\beta_{ww}}{\beta_w}$ )", " \n " }, PlotStyle ->
    {{Thick, Thickness [thick ], colbifur [[3]]}, {Thick, Thickness [thick ], colbifur [[3]]}},
    Filling -> {1 -> {2}, Directive [Opacity [alpha ], colbifurfilling [[3]]]},
    ImageSize -> imageSize ];
(*Lower boundary *)
GetBoundaryLineSingle [parslist ]/
    ConstantArray [{ $\beta_w$  /. parmanip  $\beta\epsilon 1$ , 1}, Length @GetBoundaryLineSingle [parslist ]];
lp $\beta\epsilon 2$  = ListLinePlot [% , PlotRange -> range1, Frame -> includeFrame,
    FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    PlotStyle -> {Thick, Thickness [thick ], colbifur [[3]]},
    Filling -> Bottom, FillingStyle -> Directive [LightGray, HatchFilling []];
l $\beta\epsilon 1$  = Show [{lp $\beta\epsilon 1$ , lp $\beta\epsilon 2$ ];
GraphicsGrid [{p $\beta\epsilon 1$ , l $\beta\epsilon 1$ }]

```



$f_w = 37, \beta_w = 1.60$  (Figure 6 Panel C)

```
(*Set parameters *)
parmanip  $\beta\epsilon 2$  = { $\rho \rightarrow 1.2$ ,  $d \rightarrow 0.9$ ,  $r \rightarrow 2.5$ ,  $\gamma \rightarrow 2.9$ ,  $\alpha_w \rightarrow 0$ ,  $\alpha_{ww} \rightarrow 0$ ,
   $p \rightarrow 0.05$ ,  $c \rightarrow 1.4$ ,  $\mu \rightarrow 3.9$ ,  $\sigma_w \rightarrow 0$ ,  $\sigma_{ww} \rightarrow 0$ ,  $q \rightarrow 0.05$ ,
   $\delta \rightarrow 0.9$ ,  $k \rightarrow 0.26$ ,  $f_w \rightarrow 37.$ ,  $h \rightarrow 0.6$ ,  $\beta_w \rightarrow 1.60$ };

(*Calculate equilibrium *)manip $\beta\epsilon 2$ AllResults = ParallelTable [NSolveCodim2Positive [
  sysfunc1 /.  $f_{ww} \rightarrow \epsilon f_w$ , parmanip  $\beta\epsilon 2$ , { $\beta_{ww} \rightarrow x$ }, { $\epsilon \rightarrow y$ },  $\beta\epsilon$ , eq, varRes, 15 ],
  {x,  $\beta_{ww}start$ ,  $\beta_{ww}end$ ,  $\beta_{ww}step$ }, {y,  $\epsilon start$ ,  $\epsilon end$ ,  $\epsilon step$ });
range2 = {{ $\beta_{ww}start / (\beta_w /. parmanip \beta\epsilon 2)$ ,  $\beta_{ww}end / (\beta_w /. parmanip \beta\epsilon 2)$ },
  { $\epsilon start$ ,  $\epsilon end + 0.1$ }};
```

```

equilibrium = eq /. Flatten [manip  $\beta \in 2$  AllResults, 2 ];
parslist =  $\beta \in$  /. Flatten [manip  $\beta \in 2$  AllResults, 2 ];

(*Mark stable and unstable equilibrium *)
marklist = ListStableMarkTwoParameters [Jmatfunc1 /. fww ->  $\epsilon$ fw,
    parmanip  $\beta \in 2$ , { $\beta_{ww}$ ,  $\epsilon$ }, parslist, equilibrium, colorstable, True ];
MakeListPlotData [parslist [[All, 1]]/( $\beta_w$  /. parmanip  $\beta \in 2$ ), parslist [[All, 2]]];
(*Plot stable and unstable equilibrium *)
p $\beta \in 2$  = ListPlot [% , PlotStyle -> marklist, PlotRange -> range1, ImageSize -> imageSize,

    Frame -> includeFrame, FrameLabel -> { " $\frac{\beta_{ww}}{\beta_w}$ ", " $\frac{f_{ww}}{f_w}$ " }, LabelStyle -> labelStyle ];

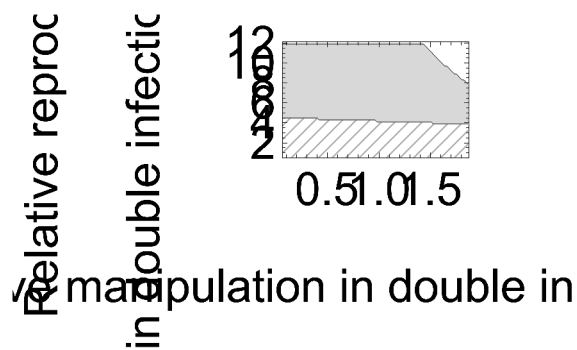
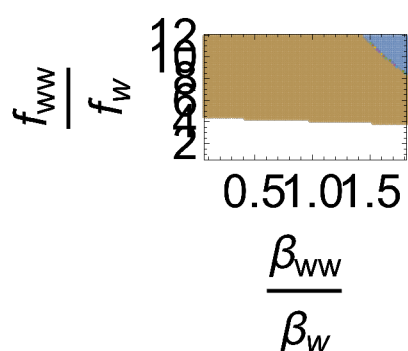
(*Connect boundary points *)
(*MapThread to get ratio as xaxis *)
MapThread [#1/#2 &,
    {GetBoundaryLineBiStable [parslist ], {ConstantArray [{ $\beta_w$  /. parmanip  $\beta \in 2$ , 1},
        Length @GetBoundaryLineBiStable [parslist ][[1, All]]}, ConstantArray [
            { $\beta_w$  /. parmanip  $\beta \in 2$ , 1}, Length @GetBoundaryLineBiStable [parslist ][[2, All]]}}];

(*Upper boundary *)lp $\beta \in 1$  = ListLinePlot [% , PlotRange -> range2,
    Frame -> includeFrame, FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    FrameLabel -> { "Relative manipulation in double infections ( $\frac{\beta_{ww}}{\beta_w}$ )",
        " Relative reproduction \n in double infections ( $\frac{f_{ww}}{f_w}$ ) " },
    PlotStyle -> {{Thickness [thick ], colbifur [[3]]}, {Thickness [thick ], colbifur [[3]]}},
    Filling -> {1 -> {2}}, FillingStyle -> Directive [Opacity [alpha ], colbifurfilling [[3]],
    ImageSize -> imageSize ];

(*Lower boundary *)
GetBoundaryLineSingle [parslist ]/
    ConstantArray [{ $\beta_w$  /. parmanip  $\beta \in 2$ , 1}, Length @GetBoundaryLineSingle [parslist ]];
lp $\beta \in 2$  = ListLinePlot [% , PlotRange -> range2, Frame -> includeFrame,
    FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    PlotStyle -> {Thickness [thick ], colbifur [[3]]}, Filling -> Bottom,
    FillingStyle -> Directive [LightGray, HatchFilling []];
l $\beta \in 2$  = Show [{lp $\beta \in 1$ , lp $\beta \in 2$ };
GraphicsGrid [{p $\beta \in 2$ , l $\beta \in 2$ }]

```

Out[295]=



$f_w = 38, \beta_w = 1.60$  (Figure 6 Panel A)

(\*Set parameters \*)

```
parmanip  $\beta\epsilon 3 = \{\rho \rightarrow 1.2, d \rightarrow 0.9, r \rightarrow 2.5, \gamma \rightarrow 2.9, \alpha_w \rightarrow 0, \alpha_{ww} \rightarrow 0,$   

 $p \rightarrow 0.05, c \rightarrow 1.4, \mu \rightarrow 3.9, \sigma_w \rightarrow 0, \sigma_{ww} \rightarrow 0, q \rightarrow 0.05,$   

 $\delta \rightarrow 0.9, k \rightarrow 0.26, f_w \rightarrow 38, h \rightarrow 0.6, \beta_w \rightarrow 1.60\};$ 
```

```
(*Calculate equilibrium *)manip  $\beta\epsilon 3$ AllResults = ParallelTable [NSolveCodim2Positive [  

  sysfunc1 /. fww  $\rightarrow \epsilon f_w$ , parmanip  $\beta\epsilon 3, \{\beta_{ww} \rightarrow x\}, \{\epsilon \rightarrow y\}, \beta\epsilon$ , eq, varRes, 15 ],  

  {x,  $\beta_{ww}$ start,  $\beta_{ww}$ end,  $\beta_{ww}$ step }, {y,  $\epsilon$ start,  $\epsilon$ end,  $\epsilon$ step }];
```

```

equilibrium = eq /. Flatten [manip  $\beta \in 3$  AllResults, 2 ];
parslist =  $\beta \in$  /. Flatten [manip  $\beta \in 3$  AllResults, 2 ];

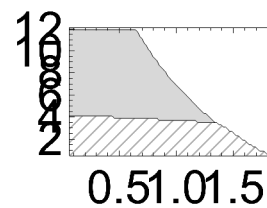
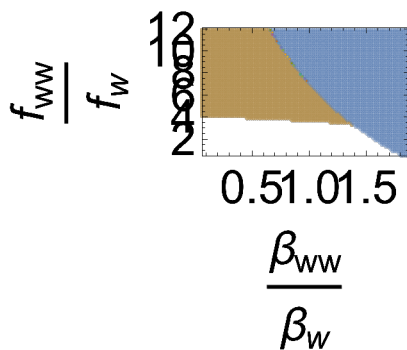
(*Mark stable and unstable equilibrium *)
marklist = ListStableMarkTwoParameters [Jmatfunc1 /. fww ->  $\epsilon$  fw,
    parmanip  $\beta \in 3$ , { $\beta_{ww}$ ,  $\epsilon$ }, parslist, equilibrium, colorstable, True ];
MakeListPlotData [parslist [[All, 1]]/( $\beta_w$  /. parmanip  $\beta \in 3$ ), parslist [[All, 2]]];
p $\beta \in 3$  = ListPlot [% , PlotStyle -> marklist, PlotRange -> range2, ImageSize -> imageSize,

    Frame -> includeFrame, FrameLabel -> { " $\frac{\beta_{ww}}{\beta_w}$ ", " $\frac{f_{ww}}{f_w}$ " }, LabelStyle -> labelStyle ];

(*Connect boundary points *)
(*MapThread to get ratio as xaxis *)
MapThread [#1/#2 &,
    {GetBoundaryLineBiStable [parslist ], {ConstantArray [{ $\beta_w$  /. parmanip  $\beta \in 3$ , 1},
        Length @GetBoundaryLineBiStable [parslist ][[1, All]]], ConstantArray [
            { $\beta_w$  /. parmanip  $\beta \in 3$ , 1}, Length @GetBoundaryLineBiStable [parslist ][[2, All]]}]}];
(*Upper boundary *)lp $\beta \in 1$  = ListLinePlot [% , PlotRange -> range2,
    Frame -> includeFrame, FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    FrameLabel -> { "Relative manipulation in double infections ( $\frac{\beta_{ww}}{\beta_w}$ )", "\n" },
    PlotStyle -> {{Thickness [thick ], colbifur [[3]]}, {Thickness [thick ], colbifur [[3]]}},
    Filling -> {1 -> {2}}, FillingStyle -> Directive [Opacity [alpha], colbifurfilling [[3]]],
    ImageSize -> imageSize ];
(*Lower boundary *)
GetBoundaryLineSingle [parslist ]/
    ConstantArray [{ $\beta_w$  /. parmanip  $\beta \in 3$ , 1}, Length @GetBoundaryLineSingle [parslist ]];
lp $\beta \in 2$  = ListLinePlot [% , PlotRange -> range2, Frame -> includeFrame,
    FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    PlotStyle -> {Thickness [thick ], colbifur [[3]]}, Filling -> Bottom,
    FillingStyle -> Directive [LightGray, HatchFilling []]];
lp $\beta \in 3$  = Show [{lp $\beta \in 1$ , lp $\beta \in 2$ };
GraphicsGrid [{p $\beta \in 3$ , lp $\beta \in 3$ }]

```

Out[306]=



manipulation in double

## Combined graph (Figure 6)

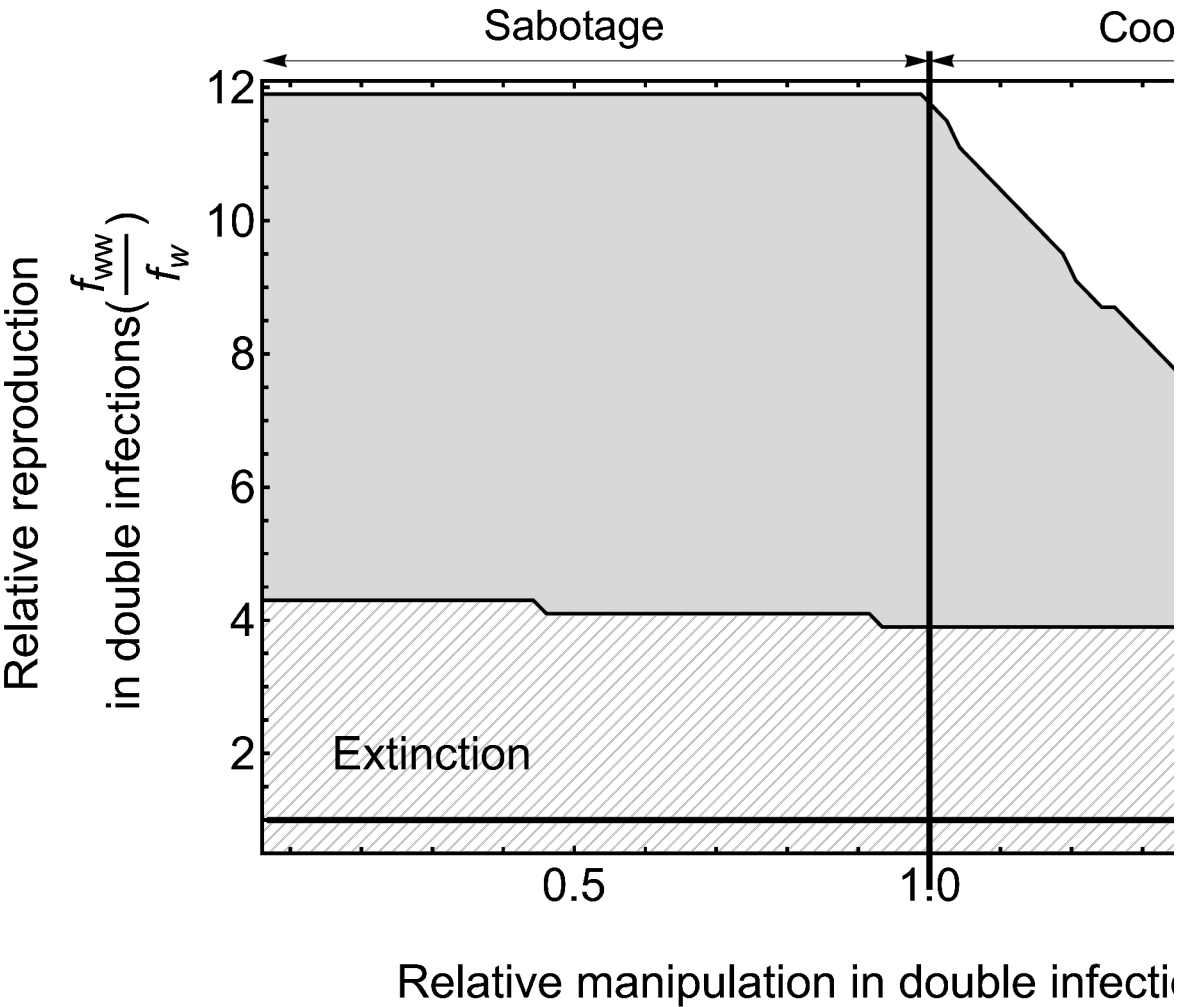
```

In[361]:= line = {Graphics[{Thickness[0.005], Black, Line[{1, range[[2]][[1]], {1, 12.5}}]},
  Graphics[{Thickness[0.005], Black, Line[{range[[1]][[1]], 1, {2.25, 1}}]}];
arrowsize = {-0.02, 0.02}
arrow1 = Graphics[{Arrowheads[arrowsize], Arrow[{1, 12.4}, {range1[[1]][[1]], 12.4}],
  Text[Style["Sabotage", FontSize -> 22], {0.5, 12.9}]}];
arrow2 = Graphics[{Arrowheads[arrowsize], Arrow[{1, 12.4}, {range1[[1]][[2]], 12.4}],
  Text[Style["Cooperation", FontSize -> 22], {1.4, 12.9}]}];
arrow3 = Graphics[{Arrowheads[arrowsize], Arrow[{1.9, 1}, {1.9, range1[[2]][[2]]}],
  Rotate[Text[Style["Enhancement", FontSize -> 22], {1.96, 6.7}], -90 Degree]}];
arrow4 = Graphics[{Text[Style["Suppression", FontSize -> 22], {2.01, 0.55}]}];
plot1βfull1 = Show[{1βϵ, line, arrow1, arrow2, arrow3, arrow4}, PlotRangePadding -> 0,
  PlotRangeClipping -> False, ImagePadding -> {{150, 200}, {100, 45}},
  Epilog -> {Text[Style["Extinction", FontSize -> 24], {0.3, 2.}],
  Text[Style["Single\n equilibrium", FontSize -> 24], {1.6, 9.5}]},
  ImageSize -> {1000, 650}]
Export["ratio_reproduction_manipulationA.pdf", %, ImageResolution -> figResolution]
plot1βfull1 = Show[{1βϵ1, line, arrow1, arrow2, arrow3, arrow4}, PlotRangePadding -> 0,
  PlotRangeClipping -> False, ImagePadding -> {{110, 200}, {100, 45}},
  Epilog -> {Text[Style["Extinction", FontSize -> 24], {0.3, 2.}],
  Text[Style["Single\n equilibrium", FontSize -> 24], {1.6, 9.5}]},
  ImageSize -> {1000, 650}]
Export["ratio_reproduction_manipulationB.pdf", %, ImageResolution -> figResolution]

arrow1 = Graphics[{Arrowheads[arrowsize], Arrow[{1, 12.4}, {range2[[1]][[1]], 12.4}],
  Text[Style["Sabotage", FontSize -> 22], {0.5, 12.9}]}];
arrow2 = Graphics[{Arrowheads[arrowsize], Arrow[{1, 12.4}, {range2[[1]][[2]], 12.4}],
  Text[Style["Cooperation", FontSize -> 22], {1.4, 12.9}]}];
arrow3 = Graphics[{Arrowheads[arrowsize], Arrow[{1.96, 1}, {1.96, range2[[2]][[2]]}],
  Rotate[Text[Style["Enhancement", FontSize -> 22], {2.02, 6.7}], -90 Degree]}];
arrow4 = Graphics[{Text[Style["Suppression", FontSize -> 22], {2.07, 0.55}]}];
plot1βfull1 = Show[{1βϵ2, line, arrow1, arrow2, arrow3, arrow4}, PlotRangePadding -> 0,
  PlotRangeClipping -> False, ImagePadding -> {{150, 200}, {100, 45}},
  Epilog -> {Text[Style["Extinction", FontSize -> 24], {0.3, 2.}],
  Text[Style["Single\n equilibrium", FontSize -> 24], {1.7, 11.2}]},
  ImageSize -> {1000, 650}]
Export["ratio_reproduction_manipulationC.pdf", %, ImageResolution -> figResolution]
plot1βfull1 = Show[{1βϵ3, line, arrow1, arrow2, arrow3, arrow4}, PlotRangePadding -> 0,
  PlotRangeClipping -> False, ImagePadding -> {{110, 200}, {100, 45}},
  Epilog -> {Text[Style["Extinction", FontSize -> 24], {0.3, 2.}],
  Text[Style["Single\n equilibrium", FontSize -> 24], {1.6, 9.5}]},
  ImageSize -> {1000, 650}]
Export["ratio_reproduction_manipulationD.pdf", %, ImageResolution -> figResolution]

Out[362]= {-0.02, 0.02}

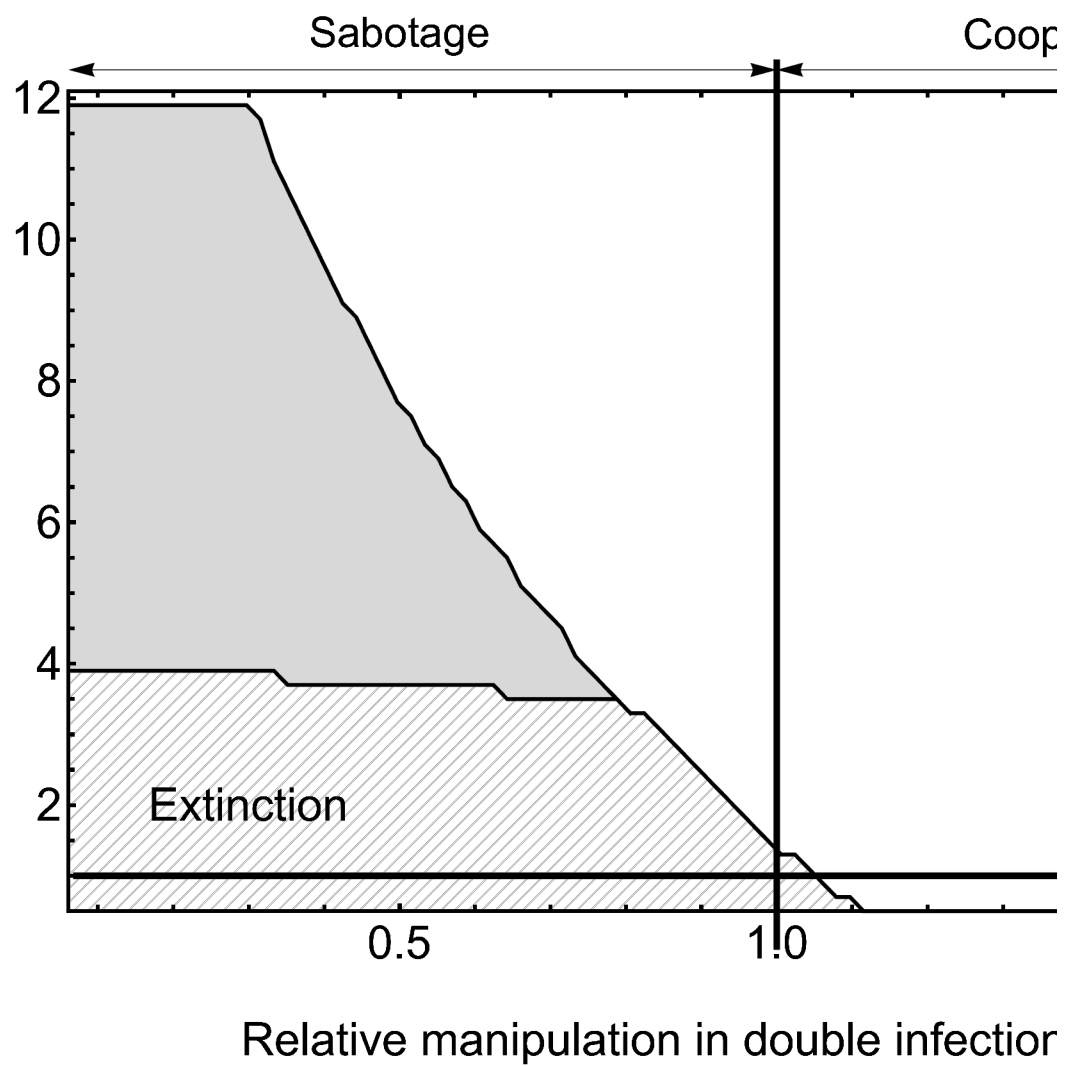
```



Out[367]=

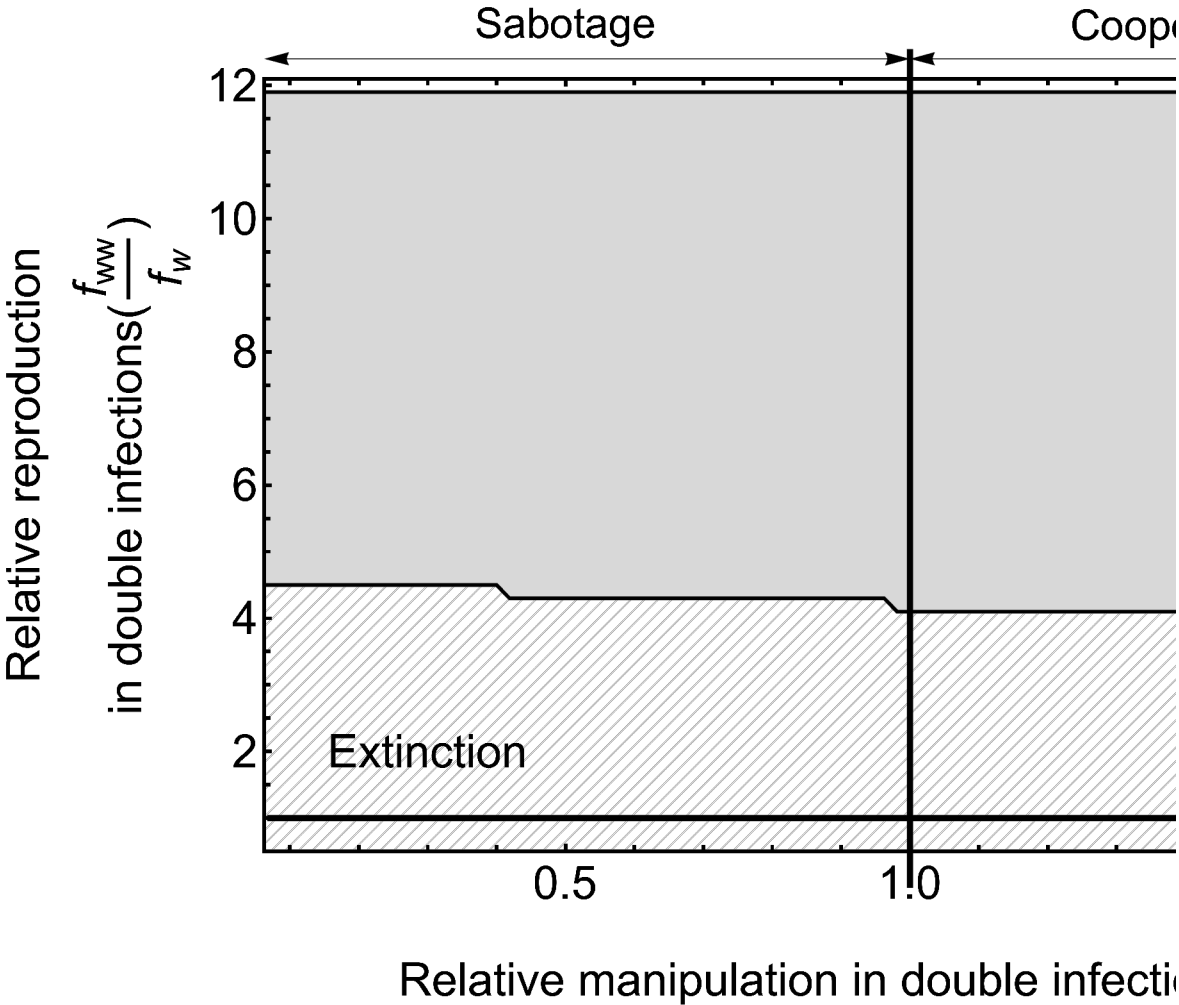
Out[368]= ratio\_reproduction\_manipulationA.pdf

Out[369]=



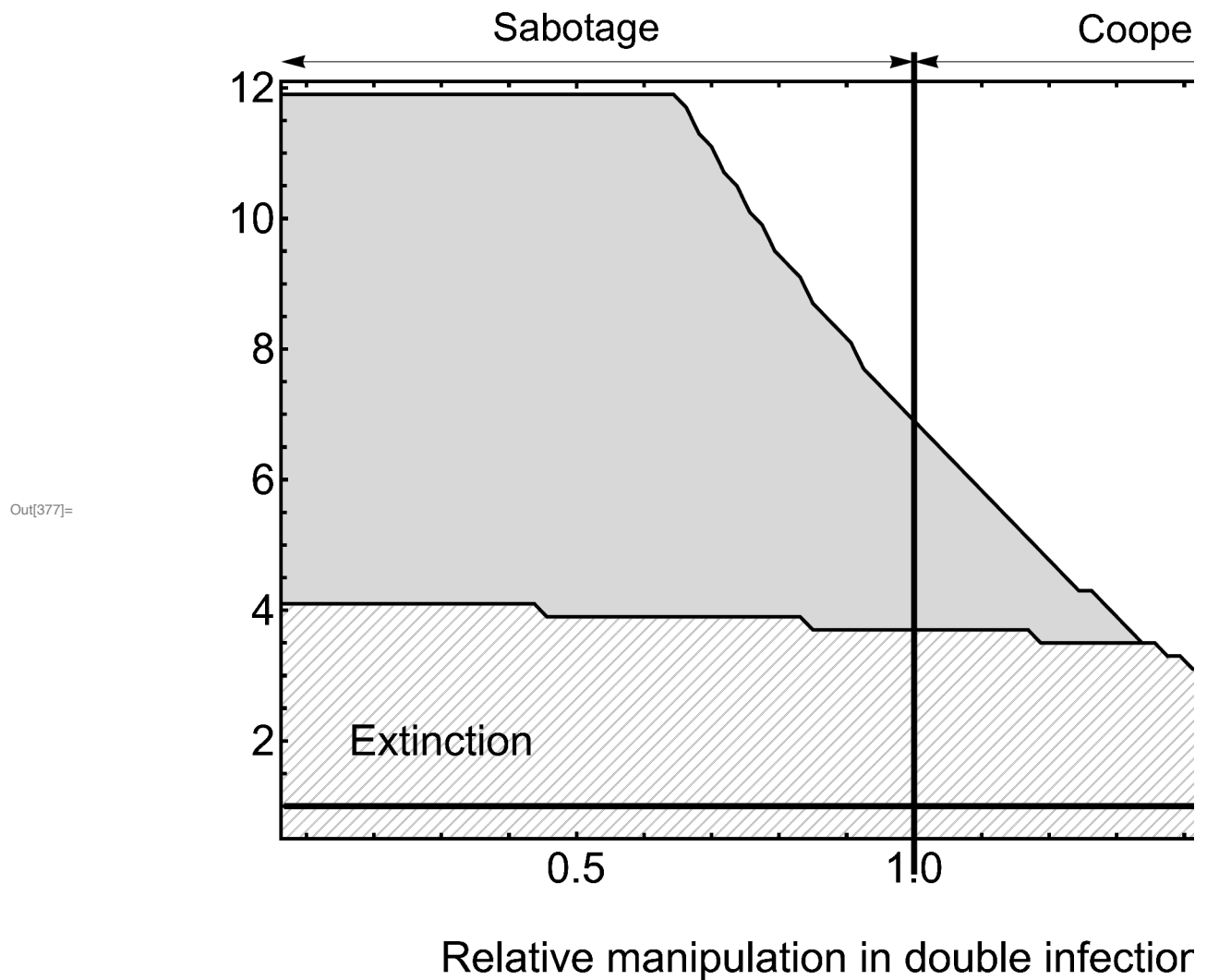
Out[370]= ratio\_reproduction\_manipulationB.pdf





Out[375]=

Out[376]= ratio\_reproduction\_manipulationC.pdf



Out[378]= ratio\_reproduction\_manipulationD.pdf

## Changing p (Figure 7 Panel A)

```
 $\beta_{wwstart}$  = 0.1;
 $\beta_{wwend}$  = 6.8;
 $\beta_{wwstep}$  = 0.05;
coldat = "Rainbow";
```

Out[379]= 0.05

**CALCULATION**

```

parmanip  $\beta_p = \{\rho \rightarrow 1.2, d \rightarrow 0.9, r \rightarrow 2.5, \gamma \rightarrow 2.9, \alpha_w \rightarrow 0,
\alpha_{ww} \rightarrow 0, c \rightarrow 1.4, \mu \rightarrow 3.9, \sigma_w \rightarrow 0, \sigma_{ww} \rightarrow 0, q \rightarrow 0.05,
\delta \rightarrow 0.9, k \rightarrow 0.26, fw \rightarrow 38, h \rightarrow 0.6, \beta_w \rightarrow 1.4, \epsilon \rightarrow 4.5\};
manip\beta_pAllResults = ParallelTable [NSolveCodim2Positive [sysfunc1 /. fww ->  $\epsilon$ fw,
parmanip  $\beta_p$ , { $\beta_{ww} \rightarrow x$ }, { $p \rightarrow y$ },  $\beta_p$ , eq, varRes, 15 ],
{x,  $\beta_{ww}$ start,  $\beta_{ww}$ end,  $\beta_{ww}$ step }, {y, 0, 1, 0.007 }];$ 
```

## PLOT

```

In[448]:= equilibrium = eq /. Flatten [manip\beta_pAllResults, 2 ];
parslist =  $\beta_p$  /. Flatten [manip\beta_pAllResults, 2 ];
range = {{ $\beta_{ww}$ start /( $\beta_w$  /. parmanip  $\beta_p$ ),  $\beta_{ww}$ end /( $\beta_w$  /. parmanip  $\beta_p$ )}, {0, 1}};
marklist = ListStableMarkTwoParameters [Jmatfunc1 /. fww ->  $\epsilon$ fw,
parmanip  $\beta_p$ , { $\beta_{ww}$ , p }, parslist, equilibrium, colorstable, True ];
MakeListPlotData [parslist [[All, 1]]/( $\beta_w$  /. parmanip  $\beta_p$ ), parslist [[All, 2]]];
p $\beta_p$  = ListPlot [% , PlotStyle -> marklist, PlotRange -> range, ImageSize -> imageSize,

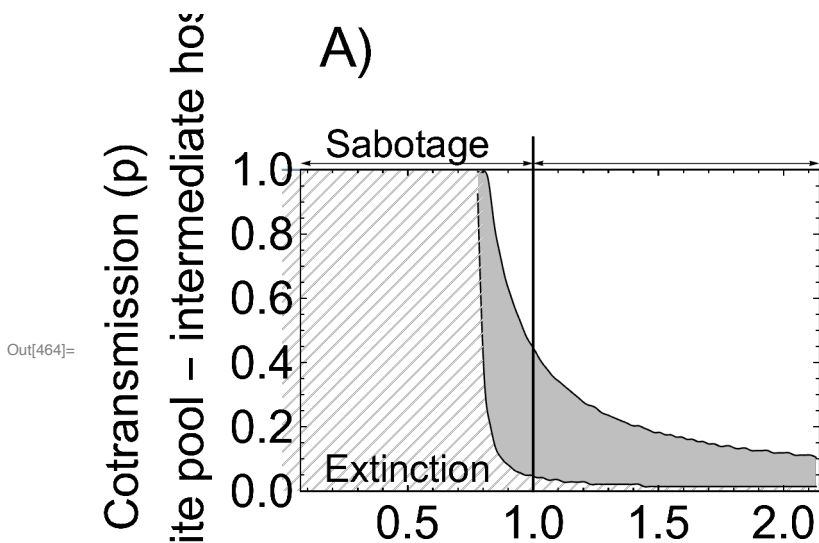
Frame -> includeFrame, FrameLabel -> { " $\frac{\beta_{ww}}{\beta_w}$ ", "p" }, LabelStyle -> labelStyle ];

(*MapThread to get ratio as xaxis *)
MapThread [#1/#2 &,
{GetBoundaryLineBiStable [parslist ], {ConstantArray [{ $\beta_w$  /. parmanip  $\beta_p$ , 1},
Length @GetBoundaryLineBiStable [parslist ][[1, All ]], ConstantArray [
{ $\beta_w$  /. parmanip  $\beta_p$ , 1}, Length @GetBoundaryLineBiStable [parslist ][[2, All ]]]}];
lp $\beta_p$ 1 = ListLinePlot [% , PlotRange -> range, Frame -> includeFrame,
FrameStyle -> frameStyle, LabelStyle -> labelStyle,
FrameLabel -> { "Relative manipulation in double infection ( $\frac{\beta_{ww}}{\beta_w}$ )",
"Cotransmission (p) \n parasite pool - intermediate host" },
PlotStyle -> {{Thickness [thick ], Black }, {Thickness [thick ], Black }},
Filling -> {1 -> {2}}, FillingStyle -> colbifurfilling [[3]], ImageSize -> imageSize,
PlotLabel -> Pane ["A"], Alignment -> Left, ImageSize -> panellabel ],
InterpolationOrder -> intorder ];
GetBoundaryLineSingle [parslist ]/ConstantArray [{ $\beta_w$  /. parmanip  $\beta_p$ , 1},
Length @GetBoundaryLineSingle [parslist ]];
lp $\beta_p$ 2 = ListLinePlot [% , PlotRange -> range, Frame -> includeFrame,
FrameStyle -> frameStyle, LabelStyle -> labelStyle,
PlotStyle -> {Thickness [thick ], Black }, InterpolationOrder -> intorder,
Filling -> Bottom, FillingStyle -> Directive [LightGray, HatchFilling []];
line = {Graphics [{Thickness [0.005 ], Black, Line [{1, 0}, {1, 1.1}]]];
arrow1 = Graphics [
{Arrowheads [arrowsize ], Arrow [{1., 1.02 }, { $\beta_{ww}$ start /( $\beta_w$  /. parmanip  $\beta_p$ ), 1.02 }],
Text [Style ["Sabotage", FontSize -> 20], {0.5, 1.08 }]];
arrow2 = Graphics [{Arrowheads [arrowsize ],
Arrow [{1., 1.02 }, { $\beta_{ww}$ end /( $\beta_w$  /. parmanip  $\beta_p$ ), 1.02 }],
Text [Style ["Cooperation", FontSize -> 20], {2.9, 1.08 }]}];

```

```
lv = ListLinePlot [{{{0., 1}, {0.8, 1}}}, PlotRange -> range, Frame -> includeFrame,
  FrameStyle -> frameStyle, LabelStyle -> labelStyle,
  PlotStyle -> {Thickness [thick], Black}, Filling -> Bottom,
  FillingStyle -> Directive [LightGray, HatchFilling [ ]]];
lp = Show [{lp $\beta$ p1, lp $\beta$ p2, line, lv, arrow1, arrow2 },
  Epilog -> {Text [Style ["Extinction", FontSize -> 20], {0.5, 0.07}],
    Text [Style ["Single \n equilibrium", FontSize -> 20], {3.3, 0.6 }]},
  PlotRangePadding -> 0, PlotRangeClipping -> False,
  ImagePadding -> {{110, 20}, {100, 45}}, ImageSize -> imageSize /1.5];
GraphicsGrid [{{p $\beta$ p, lp}}];
```

In[464]:= lp



Relative manipulation in double infection

## Changing q (Figure 7 Panel B)

### CALCULATION

```
In[379]:= parmanip $\beta$ q = { $\rho$  -> 1.2, d -> 0.9, r -> 2.5,  $\gamma$  -> 2.9,  $\alpha$ w -> 0,
   $\alpha$ ww -> 0, p -> 0.05, c -> 1.4,  $\mu$  -> 3.9,  $\sigma$ w -> 0,  $\sigma$ ww -> 0,
   $\delta$  -> 0.9, k -> 0.26, fw -> 38, h -> 0.6,  $\beta$ w -> 1.45,  $\epsilon$  -> 4.5};
manip $\beta$ qAllResults = ParallelTable [NSolveCodim2Positive [sysfunc1 /. fw ->  $\epsilon$ fw,
  parmanip $\beta$ q, { $\beta$ ww -> x}, {q -> y},  $\beta$ q, eq, varRes, 15 ],
  {x,  $\beta$ wwstart,  $\beta$ wwend,  $\beta$ wwstep }, {y, 0, 1, 0.007 }];
```

### PLOT

```

equilibrium = eq /. Flatten [manip  $\beta$ qAllResults, 2 ];
parslist =  $\beta$ q /. Flatten [manip  $\beta$ qAllResults, 2 ];
range = {{ $\beta$ wwstart / ( $\beta$ w /. parmanip  $\beta$ q),  $\beta$ wwend / ( $\beta$ w /. parmanip  $\beta$ q)}, {0, 1}};
arrowsize = {-0.02, 0.02 };

marklist = ListStableMarkTwoParameters [Jmatfunc1 /. fww ->  $\epsilon$ fw,
    parmanip  $\beta$ q, { $\beta$ ww, q}, parslist, equilibrium, colorstable, True ];
MakeListPlotData [parslist [[All, 1]]/( $\beta$ w /. parmanip  $\beta$ q), parslist [[All, 2 ]]];
p $\beta$ q = ListPlot [% , PlotStyle -> marklist, PlotRange -> range, ImageSize -> imageSize,

    Frame -> includeFrame, FrameLabel -> { " $\frac{\beta_{ww}}{\beta_w}$ ", "q" }, LabelStyle -> labelStyle ];

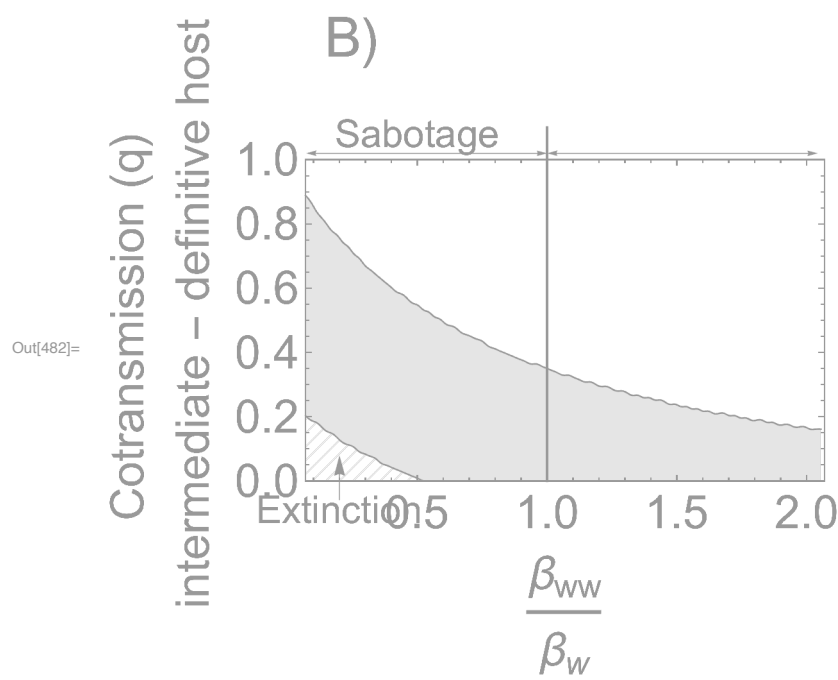
Join [Thread @ {parslist [[All, 1]]/( $\beta$ w /. parmanip  $\beta$ q), parslist [[All, 2 ]], Idensity, 2 ];
MapThread [#1/#2 &,
    {GetBoundaryLineBiStable [parslist ], {ConstantArray [{ $\beta$ w /. parmanip  $\beta$ q, 1},
        Length @GetBoundaryLineBiStable [parslist ][[1, All ]], ConstantArray [
            { $\beta$ w /. parmanip  $\beta$ q, 1}, Length @GetBoundaryLineBiStable [parslist ][[2, All ]]]}}];
lp $\beta$ q1 = ListLinePlot [% , PlotRange -> range, Frame -> includeFrame,
    FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    FrameLabel -> { " $\frac{\beta_{ww}}{\beta_w}$ ", "Cotransmission (q)\n intermediate - definitive host" },
    PlotStyle -> {{Thickness [thick ], Black }, {Thickness [thick ], Black }},
    Filling -> {1 -> {2}}, FillingStyle -> colbifurfilling [[3]], ImageSize -> imageSize,
    PlotLabel -> Pane ["B"], Alignment -> Left, ImageSize -> panellabel ],
    InterpolationOrder -> intorder ];

GetBoundaryLineSingle [parslist ]/ConstantArray [{ $\beta$ w /. parmanip  $\beta$ q, 1},
    Length @GetBoundaryLineSingle [parslist ]];
lp $\beta$ q2 = ListLinePlot [% , PlotRange -> range, Frame -> includeFrame,
    FrameStyle -> frameStyle, LabelStyle -> labelStyle,
    PlotStyle -> {Thickness [thick ], Black }, InterpolationOrder -> intorder,
    Filling -> Bottom, FillingStyle -> Directive [LightGray, HatchFilling []]];
line = {Graphics [{Thickness [0.005 ], Black, Line [{1, 0}, {1, 1.1}]]];
arrow1 = Graphics [
    {Arrowheads [arrowsize ], Arrow [{1., 1.02 }, {  $\beta$ wwstart / ( $\beta$ w /. parmanip  $\beta$ p), 1.02 }]},
    Text [Style ["Sabotage", FontSize -> 20], {0.5, 1.08 }]];
arrow2 = Graphics [{Arrowheads [arrowsize ],
    Arrow [{1., 1.02 }, {  $\beta$ wwend / ( $\beta$ w /. parmanip  $\beta$ p) - 0.1, 1.02 }]},
    Text [Style ["Cooperation", FontSize -> 20], {2.9, 1.08 }]];
arrow3 = Graphics [{Arrow [{0.2, -0.06 }, {0.2, 0.08 }]]];
l $\beta$ q = Show [lp $\beta$ q1, lp $\beta$ q2, line, arrow1, arrow2, arrow3 ],
    Epilog -> {Text [Style ["Extinction", FontSize -> 20], {0.2, -0.09 }],
        Text [Style ["Single \n equilibrium", FontSize -> 20], {3.3, 0.6 }]},
    PlotRangePadding -> 0, PlotRangeClipping -> False,
    ImagePadding -> {{110, 20 }, {110, 45 }}, ImageSize -> imageSize / 1.5]
(*GraphicsGrid [{p $\beta$ q, l $\beta$ q}])*

```

Out[469]= {-0.02, 0.02 }

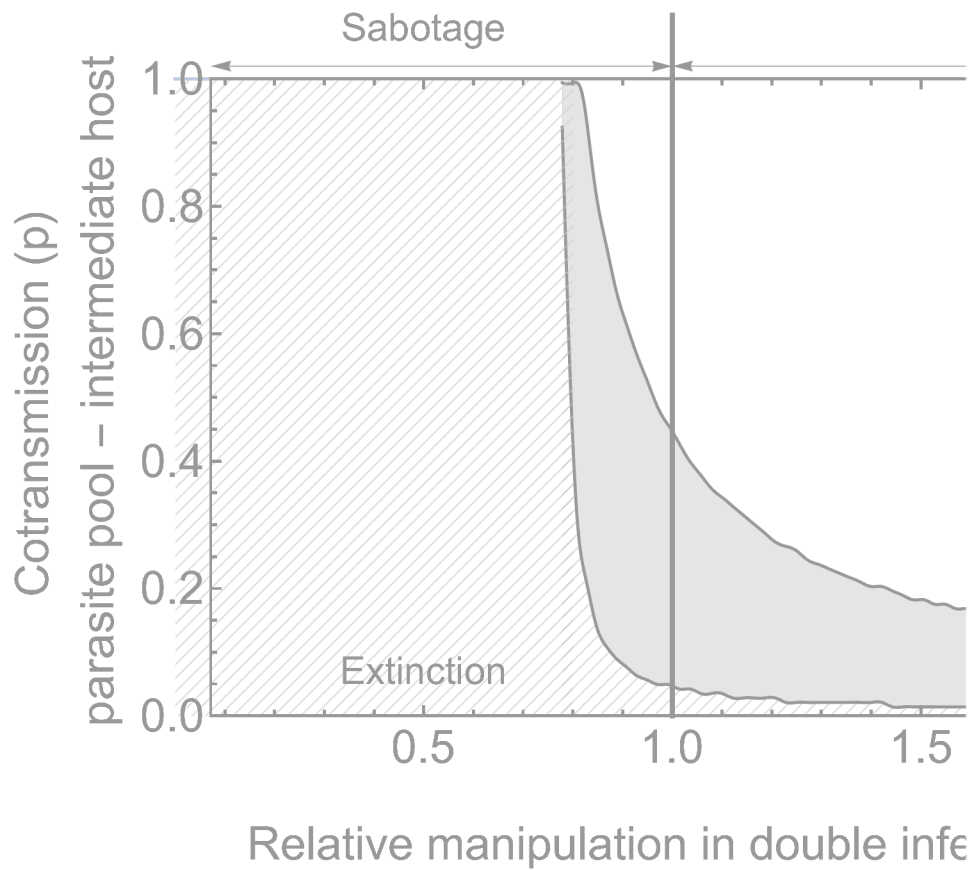
Join: Expression Identity at position 2 is expected to have head List for all expressions at level 2.



## Combined graph

```
GraphicsGrid [{{l $\beta$ p, l $\beta$ q}}, ImageSize -> imageSize * 2.8, Spacings -> -60]
Export ["coinfect_transmission.pdf", %, ImageResolution -> figResolution ]
```

A)



Out[465]=