

Supplementary Information

Regime shifts occur disproportionately faster in larger ecosystems

Cooper et al.

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Supplementary Note 1. Linear model fit of the empirical data

Whilst the main manuscript finds that a significant positive and sublinear relationship exists between system size and regime shift duration, it is reasonable to ask the extent to which a linear fit would better capture the variance in the underlying dataset. Therefore, this section provides a comparative analysis of the empirical sub-linear trend by calculating a linear regression model from the unlogged (i.e. raw values) data, as well as its associated uncertainty and implications for the duration of real-life regime shifts.

We calculate a simple linear regression model between the 42 system areas and regime shifts (Supplementary Table 1) in the statistical software R¹. As per the sub-linear trend, an overarching positive relationship exists between system area and regime shift duration (Supplementary Fig. 1), so that larger systems tend to take longer to shift between two stable regimes. The fit of the linear regression model is superior ($R^2 = 0.947$, $p < 0.001$, $df = 40$) to that of the sub-linear relationship ($R^2 = 0.491$, $p < 0.001$, $df = 40$) when all 42 data points are included. Using the linear model, we are able to derive a second set of projections for the durations of regime shifts across the Amazon rainforest and Caribbean coral reefs. For the Amazon, the linear model (Supplementary Fig. 1) produces a mean duration of 233 years (CI: 210-255 years), which sits in the upper end of our sub-linear uncertainty bounds. Likewise, for the Caribbean coral reefs, the linear model generates a mean duration of 10 years (CI: 5-14 years), which is five years quicker than the sub-linear projection.

However, it is important to note that unlike the sub-linear relationship ($R^2 = 0.423$, $p < 0.001$, $df = 39$), the linear relationship is not robust. The sub-linear relationship is present in all 42 alternative models (see Supplementary Table 2), whilst the linear relationship is only observed in 41 of the alternative models. Essentially, the linear relationship becomes insignificant as soon as the record for the Sahara is removed ($R^2 = 0.050$, $p > 0.1$, $df = 39$), potentially highlighting that the high R^2 value was due to overfitting to this datapoint. In contrast, the sub-linear relationship remains robust to the removal of individual data points (Supplementary Table 2) and random errors between 50-150% of the original duration values (Supplementary Figures 5 and 6). Thus, it is the robust, sub-linear models that we include in the main text.

Supplementary Note 2. Omitted observations from modelling experiments

Supplementary table 9 records the number of model iterations that were omitted from the regression models displayed in Fig. 3. As detailed in the Methods, omitted runs did not reach a stable alternative state over the duration of their simulation. All model runs that demonstrated a regime shift (i.e. reached a stable alternative state) were included in our analyses.

Supplementary Note 3. Amendments to the Game of Life agent-based model code

See Supplementary Table 5 for variable descriptions.

The altered code was as follows:

```
to go
  ask patches
    [if range = 4 [ set live-neighbors count neighbors4 with [living?]]] ;; if range is 4
    then each patch looks at 4 neighbor patches
  ask patches
    [if range = 8 [ set live-neighbors count neighbors with [living?]]] ;; if range is 8
    then each patch looks at 8 neighbor patches
  ;; Starting a new "ask patches" here ensures that all the patches
  ;; finish executing the first ask before any of them start executing
  ;; the second ask. This keeps all the patches in synch with each other,
  ;; so the births and deaths at each generation all happen in lockstep.
  ask patches
    [ ifelse live-neighbors = (range / 2) ;; whatever the range, this is done for the same
    proportion of neighbor patches
      [ cell-birth ]
      [ if live-neighbors != (range / 4) ;; whatever the range, this is done for the
      same proportion of neighbor patches
        [ cell-death ] ] ]
  stop-function
  ifelse stability-score = 100
  [stop] ;; stop the model when the stability score is 100 (i.e. number of alive patches has
  not changed for 100 time-steps
  [tick]
end

to stop-function
  let alive-now (count patches with [pcolor = fgcolor]) ;; how many patches are alive?
  ifelse (alive-now = total-live) ;; if the number of alive patches is the same as the last time-
  step, then add one to the stability score
    [set stability-score (stability-score + 1)]
    [set stability-score 0]
  set total-live (alive-now)
end
```

Supplementary Note 4. Amendments to the Language Change agent-based model code.

See Supplementary Table 6 for variable descriptions.

The altered code was as follows:

```
to create-network
  ;; make the initial network of two nodes and an edge
  let partner nobody
  let first-node one-of nodes
  let second-node one-of nodes with [self != first-node]
  ;; make the first edge
  ask first-node [ create-link-with second-node [ set color white ] ]
  ;; randomly select unattached node to add to network
  let new-node one-of nodes with [not any? link-neighbors]
  ;; and connect it to a partner already in the network
  while [new-node != nobody] [
    set partner find-partner
    ask new-node [ create-link-with partner [ set color white ] ]
    layout
    set new-node one-of nodes with [not any? link-neighbors]
  ]
  ;; The above code is present in the standard NetLogo LC model. It ensures all nodes are in
  the network. The below additions determine how many extra links (determined by num-
  connections) are added on top of this.
  if num-connections < (num-nodes - 1) [set num-connections (num-nodes - 1)]
  ;; stops the number of connections being too small
  while [count links < num-connections] [
    let first-nodeB one-of nodes
    let second-nodeB one-of nodes with [self != first-nodeB]
    ask first-nodeB [create-link-with second-nodeB [ set color white ] ]
    ;; find a node, find a different node and link them if you need more links
    layout
  ]
  ;; while more connections are required, they are continually added
  if count (links) > num-connections [create-network]
  ;; if there are have too many connections the process is restarted
end
```

Supplementary Note 5. Model code to run the SH model.

See Supplementary Table 7 for variable descriptions.

```
to go
  set Zcons ((g * Z * A1) / (A1 + h))
  set A1_increase ((r * A1 * (1 - (A1 / K))) - Zcons + ((d / f) * (A2 - A1)))
  set A2_increase ((r * A2 * (1 - (A2 / K))) - ((d / (1 - f)) * (A1 - A2)))
  set Z_increase ((Zcons * ez) - (m * Z))
  ifelse A1 > 0
    [set A1 (A1 + A1_increase)]
    [set A1 0]
  ifelse A2 > 0
    [set A2 (A2 + A2_increase)]
    [set A2 0]
  ifelse Z > 0
    [set Z (Z + Z_increase)]
    [set Z 0]
  tick
end
```

Supplementary Tables

Supplementary Table 1 | Details of the empirical regime shifts depicted in Fig. 2 (record numbers 1-42) and Supplementary Fig. 2 (record numbers 1-50).

Unless estimated or originally reported as an integer, system surface areas, depths and shift durations are rounded to a maximum of 3 decimal places. The system volumes were calculated by multiplying the surface areas by the mean depths. System depths and volumes shown as a dash (-) could not be reliably estimated.

System	Record number	Case study	System surface area (km ²)	System mean depth (m)	System volume (million m ³)	Regime shift duration (years)	Initial regime	Alternate regime	Reference
Freshwater	1	Lake Erhai, China	250	11	2750	2	Oligotrophic	Eutrophic	^{2,3}
	2	Paul and Peter Lakes, USA	0.020	8.30	0.199	0.077	Undisturbed food web state	Predated food web state	^{4,5}
	3	Lake Veluwe, Netherlands	30	1.30	39	2	Oligotrophic	Eutrophic	^{6,7}
	4	Mwanza gulf, Lake Victoria, Tanzania	500	40	20,000	8	Herbivore dominated	Small predators dominated	^{6,7}
	5	Lake Krankesjön, Sweden	2.90	1.50	6.30	3	Turbid	Oligotrophic	⁸

	6	Lake Stechlin, Germany	4.23	20	84.6	0.030	Oligotrophic	Eutrophic	⁹
	7	Old Danube Lake, Austria	1.60	2.30	3.68	5	Oligotrophic	Eutrophic	¹⁰
	8	Lake Kariba, Zimbabwe/ Zambia	5400	29	156,600	29	Productive fishery	Collapsed fishery	¹¹
	9	Lake of the woods, Canada/USA	4350	3.40	14,790	35	<i>Aulacoseira subarctica</i> diatom dominance	<i>Cyclotella</i> spp. diatom dominance	¹²
	10	Foy Lake, USA	1.95	-	-	5	Low benthic : planktonic diatom ratio	High benthic : planktonic diatom ratio	¹³
	11	Lake Chilika, India	1000	1.50	1500	8	Productive fishery	Collapsed fishery	¹⁴
	12	Lac de Tunis, Tunisia	48.6	1.50	72.9	1	Productive fishery	Collapsed fishery	¹⁵
	13	Victoria Park lake, Australia	0.15	1.10	0.165	0.29	Oligotrophic	Eutrophic	¹⁶
Marine	14	Jamaican coral reef	1000	-		15	Coral reef dominance	Macroalga dominance	^{17,18}

15	Black sea	436,402	1200	5.24×10^8	40	Large predator dominated	Depleted large predators	¹⁹
16	Northern Gulf of Alaska	1250	-	-	17	Shrimp dominated	Depleted shrimp populations	^{20,21}
17	Northern Gulf of Mexico	18,000	1600	2.88×10^7	15	Pre-hypoxic	Hypoxic	^{22,23}
18	Florida Bay, USA	40	1.50	60	5	Sea-grass dominated	Algal eutrophic conditions	^{24,25}
19	Ringkøbing Fjord, Denmark	300	3	900	3	Turbid state	Clear water state	²⁶
20	Frisian Front, Germany	2880	50	144,000	5	Brittle star dominated	Mud shrimp dominated	²⁷
21	Eastern Scotian Shelf	108,000	90	9.72×10^6	5	Productive fishery	Collapsed fishery	²⁸
22	Gulf of Trieste	600	19	11,400	3	Annual red tide events	Repeated mucilage events	^{29,30}
23	Central Baltic sea	100,000	70	2.64×10^7	5	High ecosystem state index	Collapsed ecosystem state index	^{31,32}

24	East Florida Mangroves, USA	20	10	170	4	Salt marshes	Mangrove stands	³³
25	Newfoundland, Canada	400,000	50	2.00 x10 ⁷	28	Productive fishery	Collapsed fishery	³⁴
26	Northern Benguela marine system	24,000	-	-	15	Productive fishery	Collapsed fishery	³⁵
27	Chesapeake Bay, USA	11,600	7	81,200	23	Productive fishery	Collapsed fishery	^{36,37}
28	Aldabra atoll	115	5	575	1	Scleractinian coral dominance	Soft coral dominance	^{38,39}
29	Alamitos Bay, USA	820	-	-	0.96	Stable fishery biomass	Collapsed fishery biomass	⁴⁰
30	Kongsfjorden, Svalbard	230	-	-	2	Low species diversity index	High species diversity index	^{41,42}
31	Izmit Bay, Turkey	310	1.80	540	1	Stable meiofauna community	Collapsed meiofauna community	⁴³

	32	Mariager fjord, Denmark	20	30	600	0.04	Stable macrofauna community	Collapsed macrofauna community	⁴⁴
	33	Gulf of Finland	5700	38	217,000	2	Stable faunal community	Collapsed faunal community	⁴⁵
	34	Georges Bank, USA/Canada	43,000	65	2.80 x10 ⁶	19	Stable haddock fishery	Collapsed haddock fishery	⁴⁶
	35	Nova Scotia kelp beds, Canada	140	-	-	6.00	Kelp dominated ecosystem	Urchin dominated ecosystem	⁴⁷
	36	Peruvian anchovy fishery	14,000	-	-	14.00	Growing anchovy fishery	Collapsed anchovy fishery	⁴⁸
	37	Limfjorden, Denmark	1500	6.5	9750	13.00	Growing oyster fishery	Collapsed oyster fishery	⁴⁹
	38	Osaka Bay ostracods, Japan	1400	28	39,200	50.00	Stable Ostracod population	Collapsed Ostracod population	⁵⁰
Terrestrial	39	The Sahel	9,400,000	1	9.40 x10 ⁶	400	Vegetated Sahel	Desertified Sahel	⁵¹
	40	Maradi agri-system, Niger	35,100	10	351,000	20	Agriculture	Desert	⁵²

	41	Zion National Park, Utah, USA	4500	10	45,000	8	Woody vegetation dominated	Native species collapse	53
	42	Chobe National Park, Botswana	11,700	10	117,000	15	Savanna grassland	Woodland	54
Social	43	Babylon	500,000	-	-	23	Functioning civilisation	Collapsed civilisation	55
	44	Hittite Empire	450,000	-	-	500	Functioning civilisation	Collapsed civilisation	55
	45	Western Chous dynasty	500,000	-	-	163	Functioning civilisation	Collapsed civilisation	57,58
	46	Roman	4,140,000	-	-	432	Functioning civilisation	Collapsed civilisation	56
	47	Assyria	650,000	-	-	217	Functioning civilisation	Collapsed civilisation	55
	48	Harappan civilization	250,000	-	-	300	Functioning civilisation	Collapsed civilisation	55
	49	Mayan	324,000	-	-	100	Functioning civilisation	Collapsed civilisation	57
	50	Almoravid-Almohad	2,300,000	-	-	122	Functioning civilisation	Collapsed civilisation	58

Supplementary Table 2 | Details of the first sensitivity analysis experiment described in Supplementary Note 2. Here, 42 alternative regression models were created by removing one of the empirical records (df = 39 for each alternative model). The ‘original model’ refers to the model detailed in Fig. 2.

Alt. model number	Case study removed from alternative model	Model slope (b-term)	Significance (p-value)
N/A	None – this is the full model displayed in Fig. 2	0.221	1.31 x10 ⁻⁷
1	The Sahel	0.189	2.62 x10 ⁻⁶
2	Central Baltic sea	0.235	3.68 x10 ⁻⁸
3	East Florida Mangroves, USA	0.234	1.85 x10 ⁻⁸
4	Foy Lake, USA	0.233	7.40 x10 ⁻⁸
5	Old Danube Lake, Austria	0.232	8.26 x10 ⁻⁸
6	Eastern Scotian Shelf	0.229	6.59 x10 ⁻⁸
7	Northern Gulf of Alaska	0.229	1.60 x10 ⁻⁷
8	Lake Stechlin, Germany	0.213	4.68 x10 ⁻⁷
9	Mariager fjord, Denmark	0.213	3.14 x10 ⁻⁷
10	Paul and Peter Lakes, USA	0.216	6.19 x10 ⁻⁷
11	Lake Krankesjön, Sweden	0.225	1.79 x10 ⁻⁷
12	Black sea	0.217	4.23 x10 ⁻⁷
13	Lac de Tunis, Tunisia	0.217	2.62 x10 ⁻⁷
14	Aldabra atoll	0.217	2.17 x10 ⁻⁷
15	Victoria Park lake, Australia	0.217	5.29 x10 ⁻⁷
16	Gulf of Finland	0.224	8.28 x10 ⁻⁸
17	Florida Bay, USA	0.223	1.65 x10 ⁻⁷
18	Lake of the woods, Canada/USA	0.218	1.41 x10 ⁻⁷
19	Izmit bay, Turkey	0.218	1.65 x10 ⁻⁷
20	Lake Kariba, Zimbabwe/Zambia	0.218	1.70 x10 ⁻⁷
21	Chesapeake Bay, USA	0.218	2.18 x10 ⁻⁷
22	Kongsfjorden, Svalbard	0.219	2.03 x10 ⁻⁷
23	Lake Erhai, China	0.219	2.00 x10 ⁻⁷
24	Maradi agri-system, Niger	0.219	2.48 x10 ⁻⁷
25	Nova Scotia kelp beds, Canada	0.222	1.73 x10 ⁻⁷
26	Alamitos Bay, USA	0.220	1.16 x10 ⁻⁷
27	Lake Veluwe, Netherlands	0.220	2.51 x10 ⁻⁷
28	Ringkøbing Fjord, Denmark	0.220	2.01 x10 ⁻⁷
29	Chobe National Park, Botswana	0.220	2.15 x10 ⁻⁷
30	Jamaican coral reef	0.221	2.21 x10 ⁻⁷
31	Georges Bank, USA/Canada	0.220	2.41 x10 ⁻⁷
32	Gulf of Trieste	0.220	1.84 x10 ⁻⁷
33	Frisian Front, Germany	0.221	1.69 x10 ⁻⁷
34	Northern Gulf of Mexico	0.220	2.20 x10 ⁻⁷
35	Peruvian anchovy fishery	0.220	2.14 x10 ⁻⁷
36	Newfoundland, Canada	0.220	3.29 x10 ⁻⁷
37	Mwanza gulf, Lake Victoria, Tanzania	0.221	1.86 x10 ⁻⁷
38	Northern Benguela marine system	0.220	2.22 x10 ⁻⁷
39	Osaka Bay ostracods, Japan	0.220	5.68 x10 ⁻⁸
40	Zion National Park, Utah, USA	0.221	1.87 x10 ⁻⁷
41	Limfjorden, Denmark	0.220	1.73 x10 ⁻⁷
42	Lake Chilika, India	0.221	1.85 x10 ⁻⁷

Supplementary Table 3 | The parameters of the Wolf-Sheep Predation agent-based model for the default set-up in NetLogo and the world size experiment.

Experiment	Variable	Description	Default value
Default (simulation outputs not presented here)	World Size	The number of pixels that make up the length and width of the landscape	50 x 50 pixels
	Grass Setting	The setting to select the variant of the WSP model which includes grass	On
	Grass regrowth time	The number of model steps required for grass to regrow after being consumed	30 steps
	Initial number wolves	The initial number of wolves	50
	Wolf gain from food	The energy gained by wolves after consuming a sheep	20
	Sheep gain from food	The energy gained by sheep after consuming grass	4
	Wolf reproduce	The probability each wolf has of reproducing every time-step	5 %
	Sheep reproduce	The probability each sheep has of reproducing every time-step	4 %
	Move	The number of pixels animals can move each time step in a random direction	1
1.1. Effect of system size on shift duration	World Size	The number of pixels that make up the length and width of the landscape	1-100 x 1-100 pixels
	Grass Setting	The setting to select the variant of the WSP model which includes grass	On
	Grass regrowth time	The number of model steps required for grass to regrow after being consumed	1-100
	Initial number wolves	The initial number of wolves	World size / 50
	Initial number sheep	The initial number of sheep	World size / 25
	Wolf gain from food	The energy gained by wolves after consuming a sheep	20
	Sheep gain from food	The energy gained by sheep after consuming grass	4
	Wolf reproduce	The probability each wolf has of reproducing every time-step	5 %
	Sheep reproduce	The probability each sheep has of reproducing every time-step	7 %
Move	The number of pixels animals are able to move each time step in a random direction	1	

Supplementary Table 4 | The parameters of the Wolf-Sheep Predation agent-based model for the modularity and fluidity experiments.

Experiment	Variable	Description	Default value
1.2 Effect of system modularity on shift duration	World Size	The number of pixels that make up the length and width of the landscape	(2, 5 10, 20, 50, or 100) x 100 pixels
	Grass Setting	The setting to select the variant of the WSP model which includes grass	On
	Grass regrowth time	The number of model steps required for grass to regrow after being consumed	1-100
	Initial number wolves	The initial number of wolves	World size / 50
	Initial number sheep	The initial number of sheep	World size / 25
	Wolf gain from food	The energy gained by wolves after consuming a sheep	20
	Sheep gain from food	The energy gained by sheep after consuming grass	4
	Wolf reproduce	The probability each wolf has of reproducing every time-step	5 %
	Sheep reproduce	The probability each sheep has of reproducing every time-step	7 %
	Move	The number of pixels animals are able to move each time step in a random direction	1
1.3 Effect of system fluidity on shift duration	World Size	The number of pixels that make up the length and width of the landscape	100 x 100 pixels
	Grass Setting	The setting to select the variant of the WSP model which includes grass	On
	Grass regrowth time	The number of model steps required for grass to regrow after being consumed	1-100
	Initial number wolves	The initial number of wolves	204
	Initial number sheep	The initial number of sheep	408
	Wolf gain from food	The energy gained by wolves after consuming a sheep	20
	Sheep gain from food	The energy gained by sheep after consuming grass	4
	Wolf reproduce	The probability each wolf has of reproducing every time-step	5 %
	Sheep reproduce	The probability each sheep has of reproducing every time-step	7 %
	Move	The number of pixels animals are able to move each time step in a random direction	1-100

Supplementary Table 5 | The parameters of the Game of Life model for the default set-up and world size, system modularity and fluidity experiments.

Experiment	Variable	Description	Default Value
Default	World size	The number of pixels that make up the length and width of the landscape	100 x 100 pixels
	Initial density	Determines the initial density of cells that are alive. These are randomly placed during setup.	35%
2.1 Effect of system size on regime shift duration	World size	The number of pixels that make up the length and width of the landscape	1-100 x 1-100 pixels
	Initial density	Determines the initial density of cells that are alive. These are randomly placed during setup.	35%
2.2 Effect of system modularity on regime shift duration	World size	The number of pixels that make up the length and width of the landscape	(2, 5, 10, 20, 50 or 100) x 1-100 pixels
	Initial density	Determines the initial density of cells that are alive. These are randomly placed during setup.	35%
2.3 Effect of fluidity on regime shift duration	World size	The number of pixels that make up the length and width of the landscape	1-100 x 1-100 pixels
	Initial density	Determines the initial density of cells that are alive. These are randomly placed during setup.	35%
	Range	This is a new variable to determine how many neighbouring cells are considered by each cell during the decision to live or die.	4 or 8 The latter is equivalent to the default model.

Supplementary Table 6 | The parameters of the Language Change model for the default set-up, and the number of nodes and number of connections experiments. The parameter values for experiment 3.3 (heterogeneity of connections) are also valid for experiment 3.2 (number of connections).

Experiment	Variable	Description	Default value
Default	Number of nodes	The number nodes included in the model	100
	Percent grammar 1	The percentage of nodes with language 1	60 %
	Update algorithm	Select which variant of the model you choose to run	Individual
	Sink state 1	Once an individual adopts language 1, it cannot go back	On
3.1. Effect of the number of nodes on shift duration	Number of nodes	The number nodes included in the model	3-1,000
	Number of connections	The number of connections within the system	(Number of nodes – 1)
	Percent grammar 1	The percentage of nodes with language 1	60 %
	Update algorithm	Select which variant of the model you choose to run	Individual
	Sink state 1	Once an individual adopts language 1, it cannot go back	On
3.3. Effect of connection heterogeneity on shift duration	Number of nodes	The number nodes included in the model	100
	Number of connections	The number of connections within the system	99-4,500
	Percent grammar 1	The percentage of nodes with language 1	60 %
	Update algorithm	Select which variant of the model you choose to run	Individual
	Sink state 1	Once an individual adopts language 1, it cannot go back	On

Supplementary Table 7 | The parameters of the Spatial Heterogeneity predator-prey model for the default set-and the two spatial scale experiments.

Experiment	Variable	Description	Default value
Default	A	Concentration of algae	8
	A1	Proportion of algae inside	0.8
	A2	Proportion of algae outside	1 – A1
	D	Fraction of volume exchanged between parts inside and outside	0.01
	Ez	Efficiency of conversion of food into growth of zooplankton	0.6
	F		0.5
	G	Maximum grazing rate of zooplankton	0.4
	H	Half saturation concentration of algae for Z functional response	0.6
	K	Carrying capacity of algae	10
	Q	Fraction of total lake volume occupied by zooplankton (i.e. inside)	A1
	M	Mortality rate of zooplankton	0.15
	R	Maximum growth rate of algae	0.5
	Z	Concentration of large herbivorous zooplankton	A / 4
	Zcons		0
5.1 Effect of system size (K) on regime shift duration	A	Concentration of algae	8
	A1	Proportion of algae inside	0.8
	A2	Proportion of algae outside	1 – A1
	D	Fraction of volume exchanged between parts inside and outside	0.01
	Ez	Efficiency of conversion of food into growth of zooplankton	0.6
	F		0.5
	G	Maximum grazing rate of zooplankton	0.4
	H	Half saturation concentration of algae for Z functional response	0.6
	K	Carrying capacity of algae	1-100
	Q	Fraction of total lake volume occupied by zooplankton (i.e. inside)	A1
	M	Mortality rate of zooplankton	0.15
	R	Maximum growth rate of algae	0.5
	Z	Concentration of large herbivorous zooplankton	A / 4
	Zcons		0
5.2 Effect of fluidity (d) on regime shift duration	A	Concentration of algae	8
	A1	Proportion of algae inside	0.8
	A2	Proportion of algae outside	1 – A1
	D	Fraction of volume exchanged between parts inside and outside	0-1
	Ez	Efficiency of conversion of food into growth of zooplankton	0.6

F		0.5
G	Maximum grazing rate of zooplankton	0.4
H	Half saturation concentration of algae for Z functional response	0.6
K	Carrying capacity of algae	10
Q	Fraction of total lake volume occupied by zooplankton (i.e. inside)	A1
M	Mortality rate of zooplankton	0.15
R	Maximum growth rate of algae	0.5
Z	Concentration of large herbivorous zooplankton	A / 4
Zcons		0

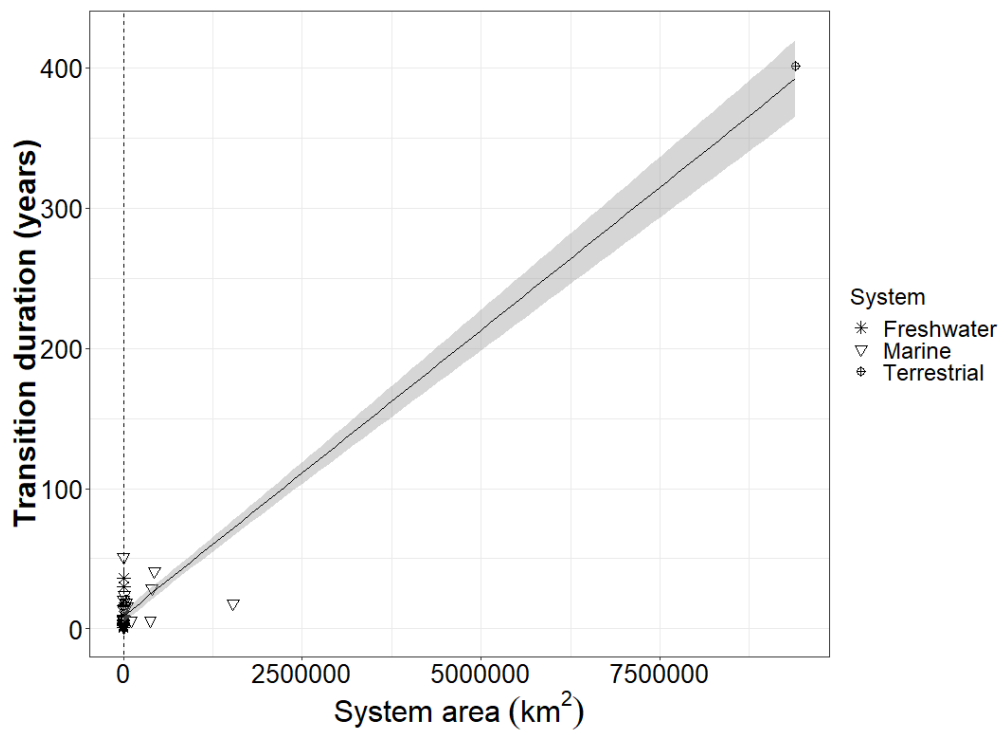
Supplementary Table 8 | The regression coefficients of the linear models describing the association between the different spatial properties and the duration of system shift. These models are created from the raw (unlogged) data, with the final column of the table comparing the strength of these linear models to the strength of the nonlinear models presented in the main manuscript (see Fig. 2). Significance levels: ‘-’ $p > 0.05$; $0.05 > *$ ≥ 0.01 ; $0.01 > **$ ≥ 0.001 ; $0.001 > p$ ‘****’.

Model name	Model type	Parameter varied	Model slope (b-term)	Coefficient of determination (R ²)	Significance (p-value)	Degrees of freedom (df)
Wolf- Sheep -Predation (WSP)	Agent-based	(1.1) Model total area	0.53	0.04	***	179,096
		(1.2) Module size (divide constant 100 x 100 area into sub-worlds)	49.0	0.23	***	19,121
		(1.3) Maximum distance wolves and sheep can move per timestep	0.43	0.00	-	7177
Game of Life (GoL)	Cellular automata	(2.1) Model total area	0.25	0.21	***	246,235
		(2.2) Module size (divide constant 100 x 100 area into discrete sub-worlds)	127	0.51	***	598
		(2.3) Number of neighbouring cells any one cell can interact with	-5635	0.63	***	102
Language Change (LC)	Network-structured	(3.1) Number of network nodes	0.00	0.00	-	99,838
		(3.2) Number of inter-nodal connections	0.00	0.00	-	440,198
		(3.3) Standard deviation of connections measured from experiment 3.2.	0.97	0.27	***	440,198
Lake Chilika fishery (LCH)	System dynamics model	(4) Model total area	0.01	0.89	***	2751
Spatial Heterogeneity (SH)	Ordinary differential equation	(5.1) Carrying capacity for algae (i.e. model size)	0.18	0.97	***	92
		(5.2) Fraction of volume exchanged between model parts (i.e. diffusion of stress)	-14.9	0.22	***	84

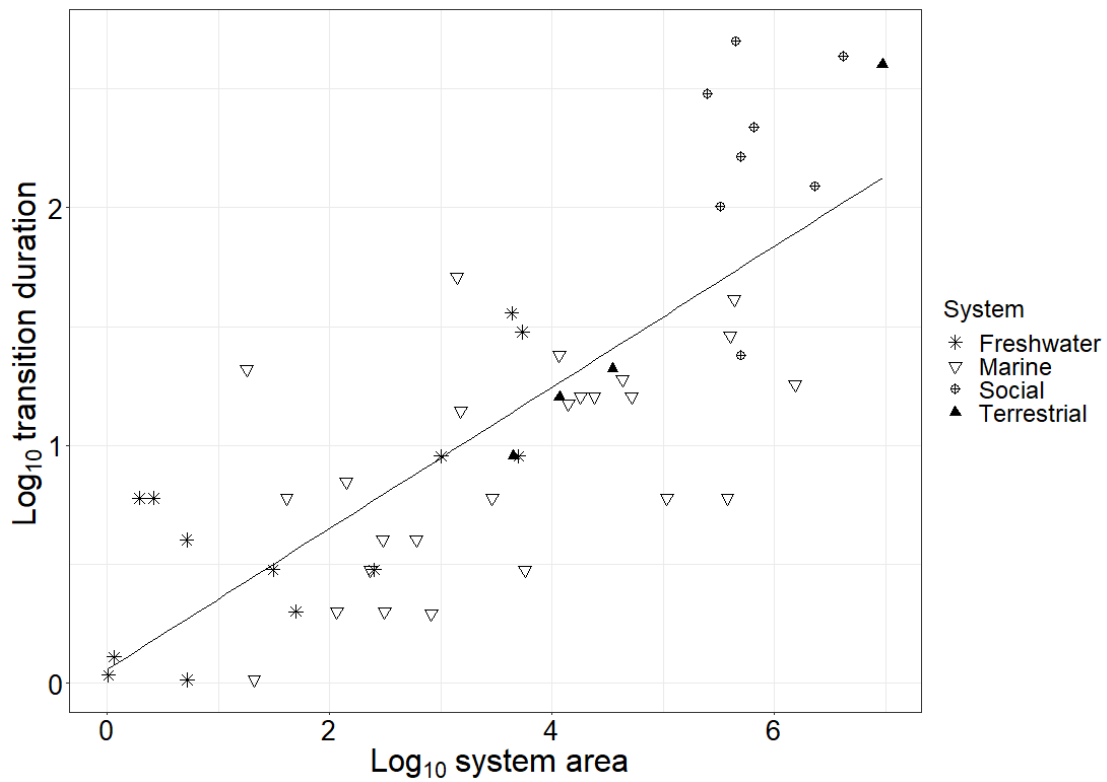
Supplementary Table 9 | The omitted model runs expressed as an absolute number and as the percentage of total model runs per experiment. †In GoL-2.2, the two simulations with the highest degrees of modularity (e.g. module sizes '2' and '5') did not record any shifts. Therefore, any model runs in GoL-2.2 that did not record regime shifts over the 5000 timesteps were given a regime shift duration of 5001 time-steps. This conservative estimate ensured that results from all six module sizes could be included in the analysis.

Experiment number	Parameter varied	Number of omitted model runs	Proportion of total model runs omitted (%)
WSP-1.1	Model total area	80,912	31.1
WSP-1.2	Module size	41,479	68.4
WSP-1.3	Maximum distance wolves and sheep can move per timestep	2821	28.1
GoL-2.1	Model total area	13,765	5.3
GoL-2.2	Module size	0 [†]	0
GoL-2.3	Number of neighbouring cells any one cell can interact with	96	48.0
LC-3.1	Number of network nodes	0	0
LC-3.2	Number of inter-nodal connections	0	0
LC-3.3	Standard deviation of connections measured from experiment 3.2.	0	0
LCH-4	Model total area	2249	45.0
SH-5.1	Carrying capacity for phytoplankton (i.e. model size)	7	6.9
SH-5.2	Fraction of volume exchanged between model parts (i.e. diffusion of stress)	15	14.9

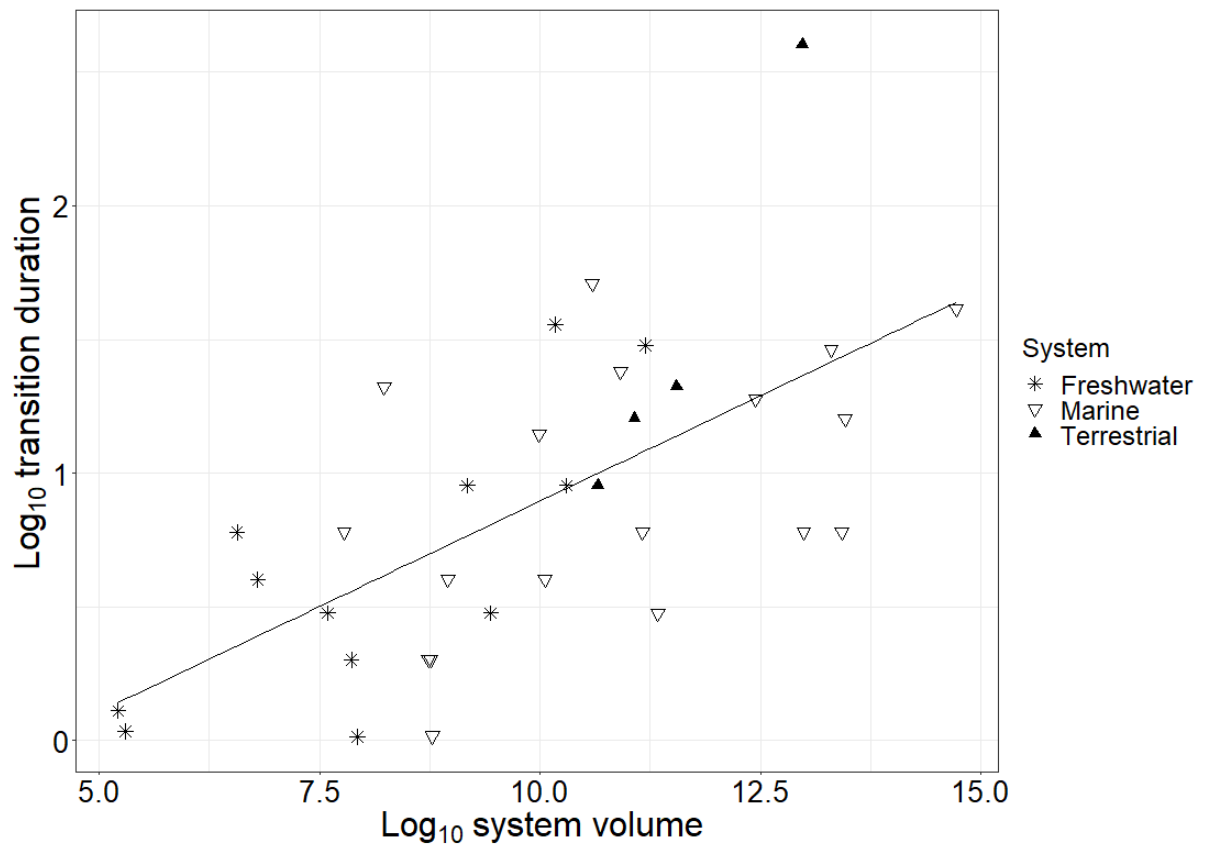
Supplementary Figures



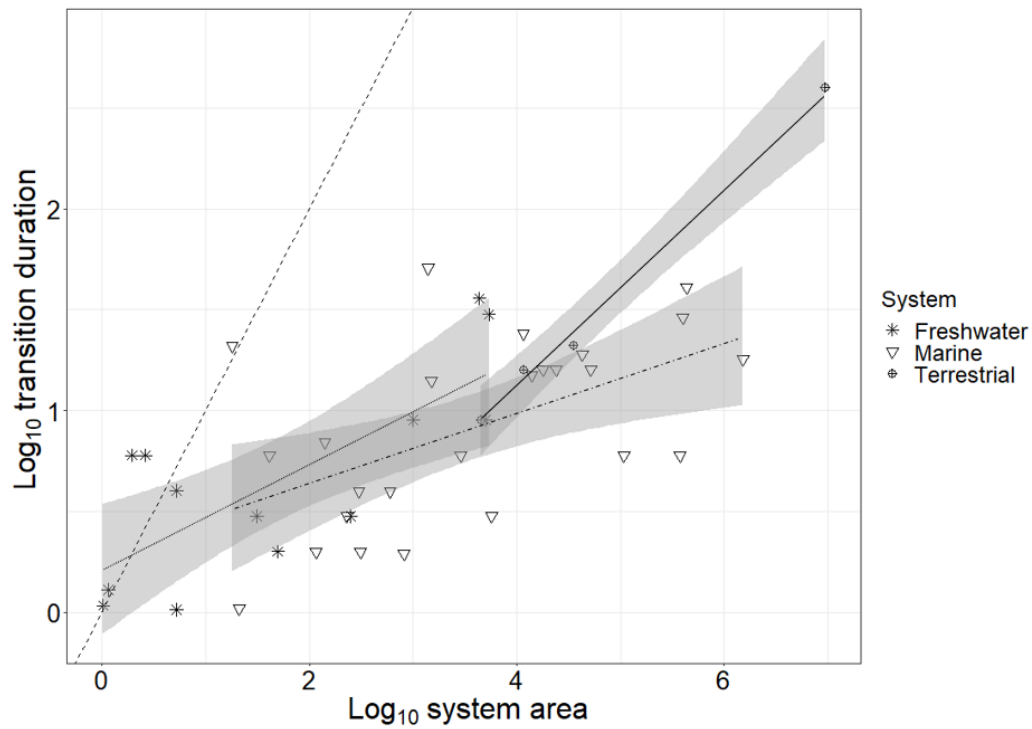
Supplementary Figure 1 | The unlogged, linear relationship between system area and transition duration. The shaded region equals the standard error around the regression line for the 42 empirical data points ($R^2 = 0.95$, $p < 0.001$, $df = 40$).



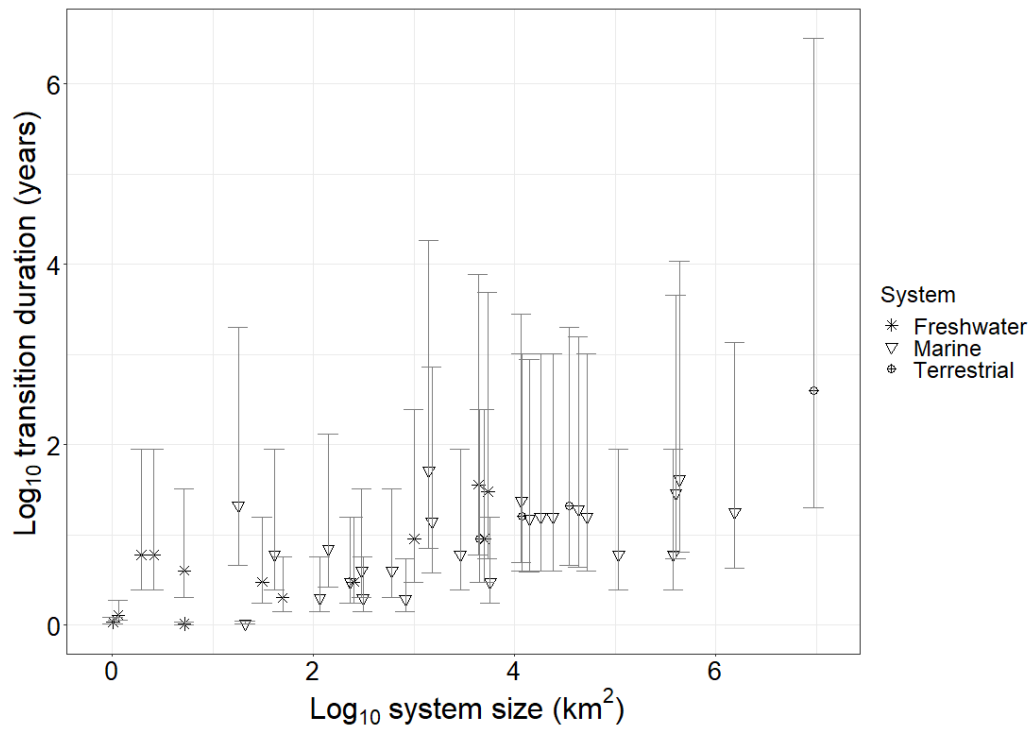
Supplementary Figure 2 | The log-log linear relationship between the spatial extent and temporal duration of regime shifts observed in nature and 8 historical societal collapses. The relationship is significantly positive ($R^2 = 0.607$, $p < 0.05$, $df = 48$) and sub-linear (slope = 0.297), providing tentative evidence that relative to shifts in nature, the collapse of social systems become disproportionately slower with increasing system area. The untransformed unit of the x-axis is *kilometres-squared*, whilst the y-axis is *years*.



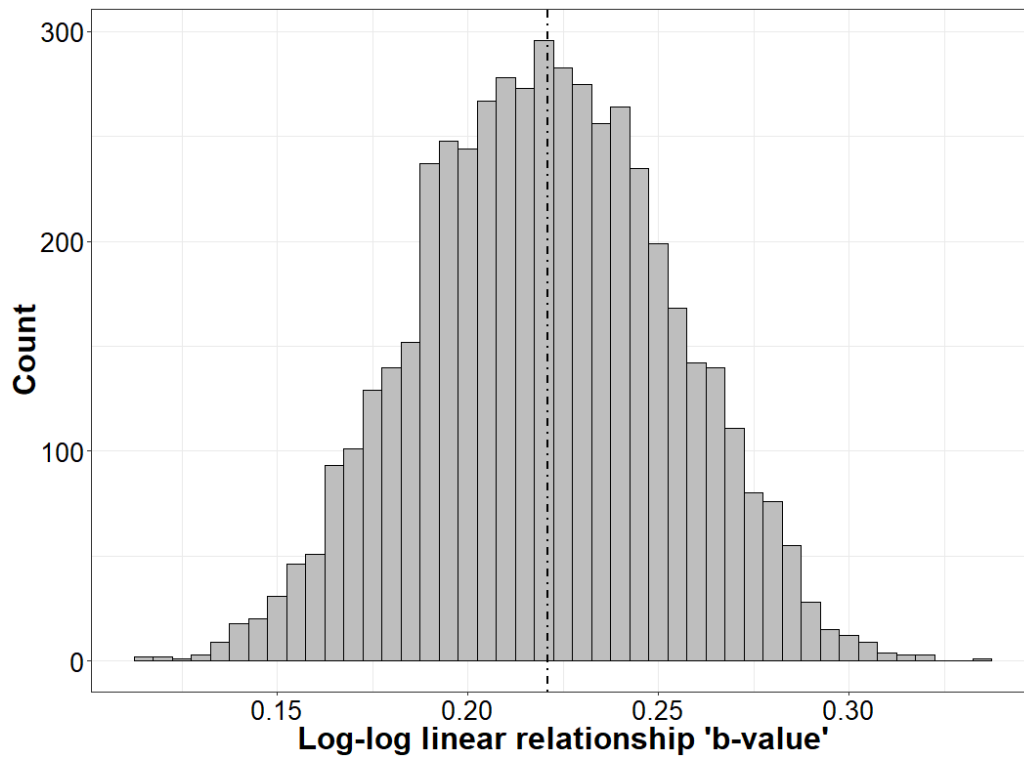
Supplementary Figure 3 | The log-log linear relationship between the system volume and temporal duration of regime shifts observed in nature. System volume exhibits a significant positive ($R^2 = 0.404$, $p < 0.05$, $df = 32$) and sub-linear power law relationships (slope = 0.158) with shift duration. Only the ecosystems with volumes that could be reliably estimated are shown. The original x-axis unit was *million cubic-metres*, whilst the y-axis unit was *years*.



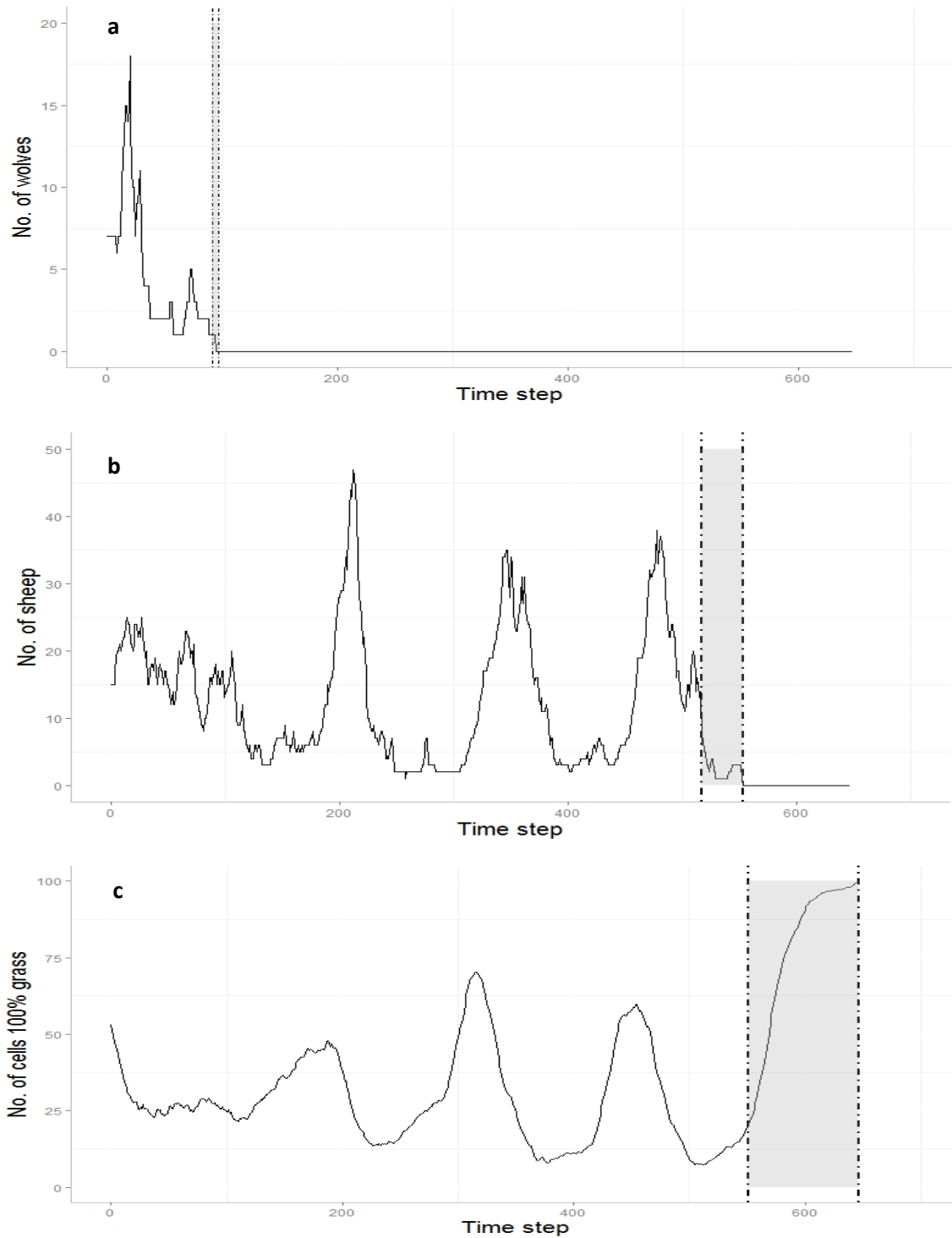
Supplementary Figure 4 | The log-log linear relationship between the spatial extent and temporal duration of regime shifts observed in nature by system type. The empirical dataset presented in Fig. 1 with the individual regression models grouped by system type. Consistent with the overall trend presented in Fig. 1, all three of the regression models exhibit significant positive sublinear trends ($p < 0.05$). Freshwater (dotted line) – $R^2 = 0.543$, slope = 0.262, $p < 0.05$, $df = 11$; Marine (dot dash line) – $R^2 = 0.267$, slope = 0.174, $p < 0.05$, $df = 23$; Terrestrial (solid line) – $R^2 = 0.996$, slope = 0.495, $p < 0.05$, $df = 2$. The dashed line represents the 1:1 reference line.



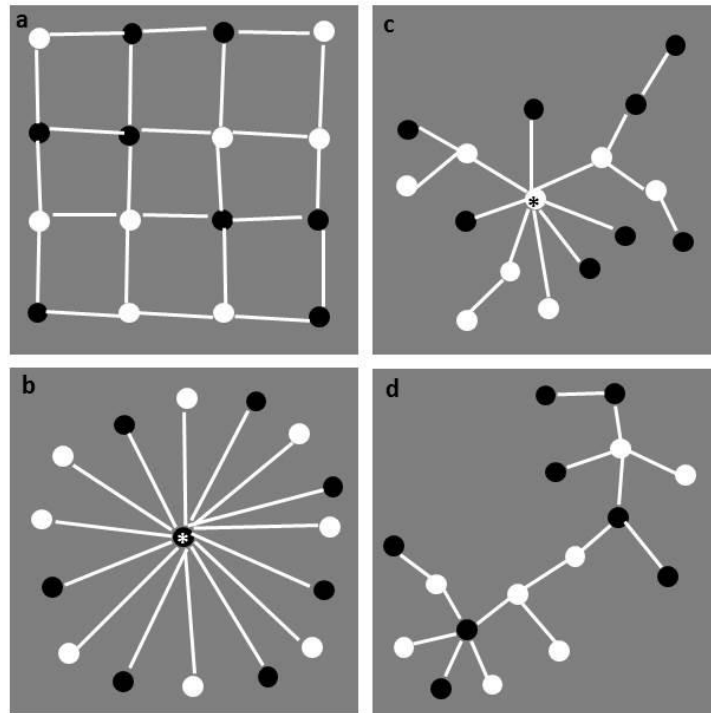
Supplementary Figure 5 | The empirical dataset depicted with the 50-150% ranges resulting from the Monte Carlo sensitivity analysis described in the Methods section ($n = 210,000$; equal to 42 empirical records multiplied by 5000 error ranges). The lower ranges of the error bars were created by multiplying the empirical values by 0.5, whilst the upper ranges were created by multiplying the empirical ranges by 1.5 (see Methods for more details).



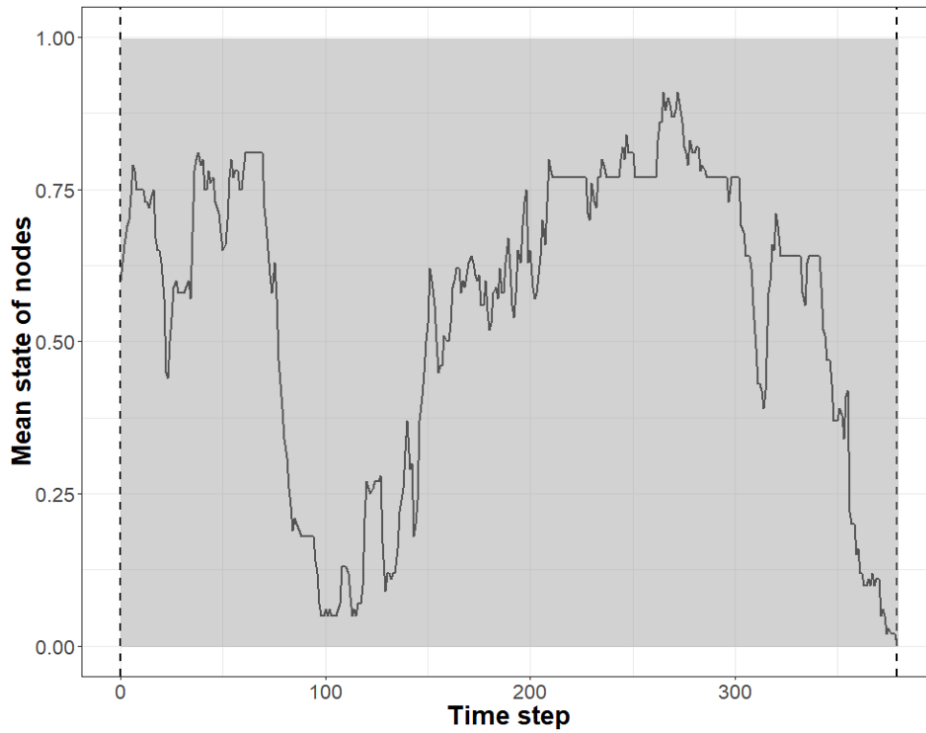
Supplementary Figure 6 | Histogram of the b-terms (regression slope values) resulting from the production of 5,000 Monte Carlo simulations where the shift durations depicted in the original empirical model were multiplied by random error ranges between 50-150% (see Supplementary Fig. 4). The dashed line represents the median b-value of the 5,000 simulations.



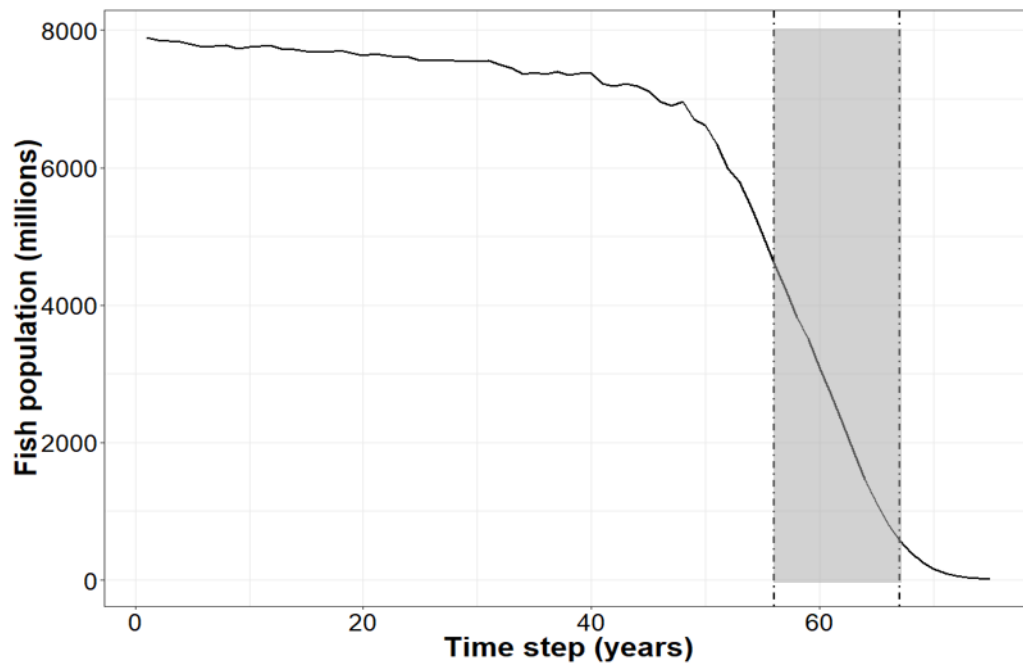
Supplementary Figure 7 | Visualisation of the trends and regime shifts (shaded regions) of (a) wolves, (b) sheep and (c) grass, for a single run of the WSP model within the world size experiments. The dashed lines represent the starts and finishes of the respective shifts. The starts of all shifts are identified by the breakpoint function described in Methods. The end of the wolves and sheep shifts are when their respective abundances equal 0; the end of the grass shift is once the entire world area is covered by grass. Here, the corresponding shift lengths are: 3, 30, 57 (measured in time-steps).



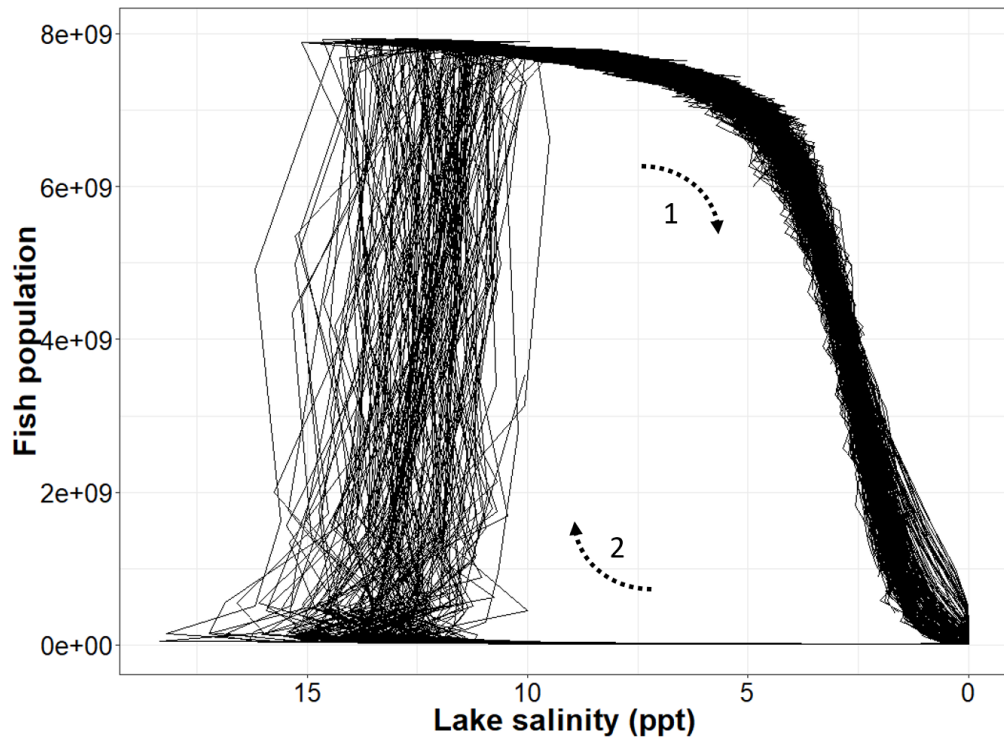
Supplementary Figure 8 | Four LC model structures comprised of 16 nodes but with different connection distributions. The most heterogeneously wired system (*b*) has the greatest variation in the number of connections between nodes (standard deviation = 3.389, as opposed to 0.707, 1.759 and 1.183 for networks *a*, *c* and *d*, respectively). The asterisks mark the keystone nodes of the two most heterogeneously wired networks.



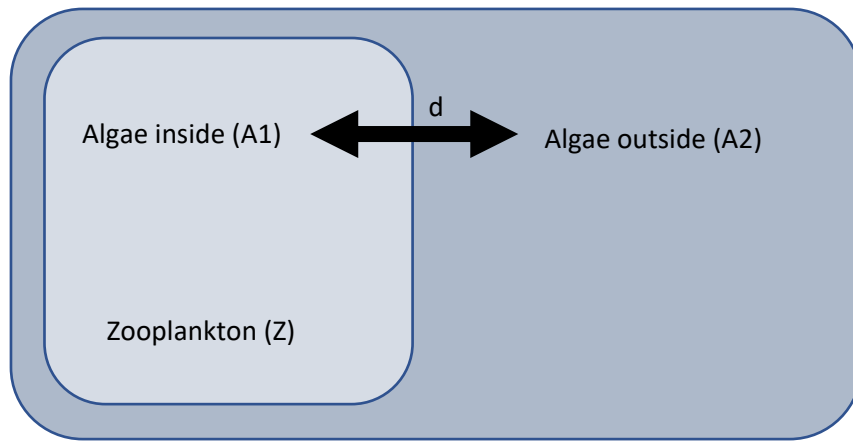
Supplementary Figure 9 | Visualisation of the timeseries trend and shift (grey shaded) from a representative run of the LC model (Methods). As the model starts unstable, the shift is measured from the first timestep until the timestep when all nodes have the same language. The same strategy to measure the duration of shift is used for the GoL model (Methods).



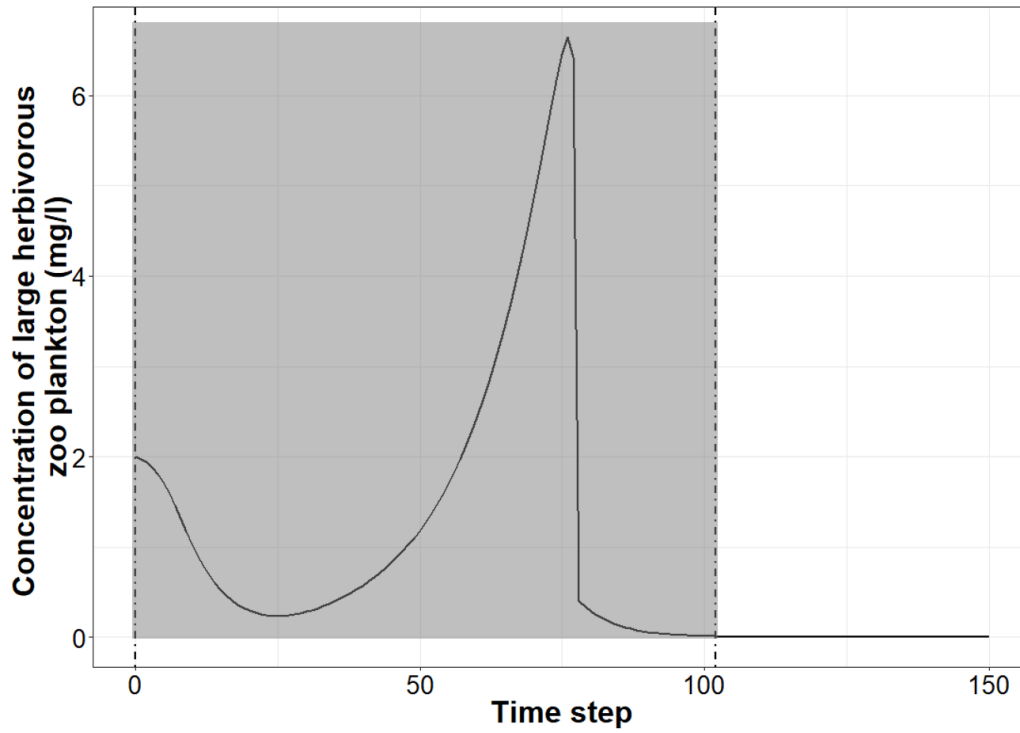
Supplementary Figure 10 | Visualisation of the fish population trend and shift (grey shaded) from a representative run of the LCH model (Methods). The first dashed line represents the onset of the shift, as detected by the breakpoint function (Methods), and the second dashed line represents the end of the shift (i.e. fish population is less than 1% of the original fish population).



Supplementary Figure 11 | Visualisation of the hysteresis displayed by the LCH model (note the x-axis has been reversed). The fish population collapses from left to right (dashed arrow number 1) as lake water salinity declines below ~5 parts per trillion (ppt). The recovery is stimulated by opening the tidal outlet (dashed arrow number 2); however, the recovery does not occur until lake salinity has been increased to >5 ppt past the point associated with the initial collapse. The figure shows 100 random model runs that displayed hysteresis.



Supplementary Figure 12 | Simplification of spatial complexity in the SH model (see Methods)



Supplementary Figure 13 | Visualisation of the timeseries trend and shift (grey shaded) from a representative run of the SH model (Methods). As the model starts unstable, the shift is measured from the first timestep until the timestep when the concentration of zooplankton equals zero.

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