# Component response rate variation drives stability in

# large complex systems

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- 5 The stability of a complex system generally decreases with increasing system size and inter-
- 6 connectivity, a counter-intuitive result of widespread importance across the physical, life, and
- <sub>7</sub> social sciences. Despite recent interest in the relationship between system properties and sta-
- bility, the effect of variation in the response rate of individual system components remains
- $_{9}$  unconsidered. Here I vary the component response rates  $(\gamma)$  of randomly generated complex
- systems. I show that when component response rates vary, the potential for system stability
- is markedly increased. Variation in  $\gamma$  becomes increasingly important as system size increases,
- such that the largest stable complex systems would be unstable if not for  $Var(\gamma)$ . My results
- 13 reveal a previously unconsidered driver of system stability that is likely to be pervasive across
- 14 all complex systems.

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- In 1972, May<sup>1</sup> first demonstrated that randomly assembled systems of sufficient complexity are almost
- inevitably unstable given infinitesimally small perturbations. Complexity in this case is defined by the
- 17 size of the system (i.e., the number of potentially interacting components; S), its connectance (i.e., the
- probability that one component will interact with another; C), and the variance of interaction strengths
- $(\sigma^2)^2$ . May's finding that the probability of local stability falls to near zero given a sufficiently high threshold
- of  $\sigma\sqrt{SC}$  is broadly relevant for understanding the dynamics and persistence of systems such as ecological 1-5,
- neurological<sup>6,7</sup>, biochemical<sup>8,9</sup>, and socio-economic<sup>10–13</sup> networks. As such, identifying general principles that

drive stability in complex systems is of wide-ranging importance.

Randomly assembled complex systems can be represented as large square matrices (**M**) with S components (e.g., networks of species<sup>2</sup> or banks<sup>11</sup>). One element of such a matrix,  $M_{ij}$ , defines how component j affects component i in the system at a point of equilibrium<sup>2</sup>. Off-diagonal elements ( $i \neq j$ ) therefore define interactions between components, while diagonal elements (i = j) define component self-regulation (e.g., carrying capacity in ecological communities). Traditionally, off-diagonal elements are assigned non-zero values with a probability C, which are sampled from a distribution with variance  $\sigma^2$ ; diagonal elements are set to  $-1^{1,2,4}$ . Local system stability is assessed using eigenanalysis, with the system being stable if the real parts of all eigenvalues ( $\lambda$ ) of **M** are negative (max ( $\Re(\lambda)$ ) < 0)<sup>1,2</sup>. In a large system (high S), eigenvalues are distributed uniformly<sup>14</sup> within a circle centred at  $\Re = -1$  (the mean value of diagonal elements) and  $\Im = 0$ , with a radius of  $\sigma\sqrt{SC}$ . (Figs 1a and 2a). Local stability of randomly assembled systems therefore becomes increasingly unlikely as S, C, and  $\sigma^2$  increase.

May's<sup>1,2</sup> stability criterion  $\sigma\sqrt{SC} < 1$  assumes that individual components respond to perturbations of the system at the same rate ( $\gamma$ ), but this is highly unlikely in any complex system. In ecological communities, for

May  $s^{\alpha\beta}$  stability criterion  $\sigma_V SC < 1$  assumes that individual components respond to perturbations of the system at the same rate  $(\gamma)$ , but this is highly unlikely in any complex system. In ecological communities, for example, the rate at which population density changes following perturbation will depend on the generation time of individuals, which might vary by orders of magnitude among species. Species with short generation times will respond quickly (high  $\gamma$ ) to perturbations relative to species with long generation times (low  $\gamma$ ). Similarly, the speed at which individual banks respond to perturbations in financial networks, or individuals or institutions respond to perturbations in complex social networks, is likely to vary. The effect of such variance on stability has not been investigated in complex systems theory. Intuitively, variation in  $\gamma$  might be expected to decrease system stability by introducing a new source of variation into the system and thereby increasing  $\sigma$ . Here I show why, despite higher  $\sigma$ , complex systems in which  $\gamma$  varies are actually more likely to be stable, especially when S is high.

### 45 Results

Component response rates of random complex systems. Rows in M define how a given component i is affected by other components of the system, meaning that the rate of component response time can be modelled by multiplying all row elements by a real scalar value  $\gamma_i^{15}$ . The distribution of  $\gamma$  over S components thereby models the distribution of component response rates. An instructive example compares one M where  $\gamma_i = 1$  for all i in S to the same **M** when half of  $\gamma_i = 1.95$  and half of  $\gamma_i = 0.05$ . This models one system in which  $\gamma$  is invariant and one in which  $\gamma$  varies, but systems are otherwise identical (note  $E[\gamma_i] = 1$  in both cases). I assume S = 200, C = 0.05, and  $\sigma = 0.4$ ; diagonal elements are set to -1 and non-zero off-diagonal 52 elements are drawn randomly from  $\mathcal{N}(0, \sigma^2)$ . Rows are then multiplied by  $\gamma_i$  to generate **M**. When  $\gamma_i = 1$ , eigenvalues of  $\mathbf{M}$  are distributed uniformly within a circle centred at (-1,0) with a radius of 1.265 (Fig. 1a). Hence, the real components of eigenvalues are highly unlikely to all be negative when all  $\gamma_i = 1$ . But when  $\gamma_i$  values are separated into two groups, eigenvalues are no longer uniformly distributed (Fig. 1b). Instead, two distinct clusters of eigenvalues appear (red circles in Fig. 1b), one centred at (-1.95,0) and the other 57 centred at (-0.05, 0). The former has a large radius, but the real components have shifted to the left (in comparison to when  $\gamma = 1$ ) and all  $\Re(\lambda) < 0$ . The latter cluster has real components that have shifted to the right, but has a smaller radius. Overall, for 1 million randomly assembled M, this division between slow and fast component response rates results in more stable systems: 1 stable given  $\gamma = 1$  versus 32 stable given 61  $\gamma = \{1.95, 0.5\}.$ Higher stability in systems with variation in  $\gamma$  can be observed by sampling  $\gamma_i$  values from various distributions. I focus on a uniform distribution where  $\gamma \sim \mathcal{U}(0,2)$  (see Supplementary Information for other distributions, which give similar results). As with the case of  $\gamma = \{1.95, 0.5\}$  (Fig. 1b),  $E[\gamma] = 1$  when  $\gamma \sim \mathcal{U}(0, 2)$ , allowing comparison of M before and after the addition of variation in component response rate. Figure 2 shows a comparison of eigenvalue distributions given S = 1000, C = 0.05, and  $\sigma = 0.4$ . As expected<sup>14</sup>, when  $\gamma = 1$ , 67 eigenvalues are distributed uniformly in a circle centred at (-1,0) with a radius of  $\sigma\sqrt{SC}=2.828$ . Uniform 68 variation in  $\gamma$  leads to a non-uniform distribution of eigenvalues  $^{16-18}$ , some of which are clustered locally near the centre of the distribution, but others of which are spread outside the former radius of 2.828 (Fig 2b). The 70

- clustering and spreading of eigenvalues introduced by  $Var(\gamma)$  can destabilise previously stable systems or stabilise systems that are otherwise unstable. But where systems are otherwise too complex to be stable given  $\gamma = 1$ , the effect of  $Var(\gamma)$  can often lead to stability above May's<sup>1,2</sup> threshold  $\sigma\sqrt{SC} < 1$ .
- Simulation of random M across S. To investigate the effect of  $Var(\gamma)$  on system stability, I simulated random M matrices at  $\sigma=0.4$  and C=1 across S ranging from 2-50. One million M were simulated for each S, and the stability of M was assessed given  $\gamma=1$  versus  $\gamma\sim\mathcal{U}(0,2)$ . For all S>10, I found that the number of stable random systems was higher given  $Var(\gamma)$  than when  $\gamma=1$  (Fig. 3; see Supplementary Information for full table of results), and that the difference between the probabilities of observing a stable system increased with an increase in S; i.e., the potential for  $Var(\gamma)$  to drive stability increased with system complexity. For the highest values of S, nearly all systems that were stable given  $Var(\gamma)$  would not have been stable given  $\gamma=1$ .
- Targeted manipulation of  $\gamma$ . To further investigate the potential of  $Var(\gamma)$  to be stabilising, I used a genetic algorithm. Genetic algorithms are heuristic tools that mimic evolution by natural selection, and are useful when the space of potential solutions (in this case, possible combinations of  $\gamma$  values leading to stability in a large complex system) is too large to search exhaustively<sup>19</sup>. Generations of selection on  $\gamma$  value combinations to minimise max  $(\Re(\lambda))$  demonstrated the potential for  $Var(\gamma)$  to increase system stability. Across  $S = \{2, 3, ..., 39, 40\}$ , sets of  $\gamma$  values were found that resulted in stable systems with probabilities that were up to four orders of magnitude higher than when  $\gamma = 1$  (Fig. 4), meaning that stability could often be achieved by manipulating S  $\gamma$  values rather than  $S \times S$   $\mathbf{M}$  elements.
- System feasibility given  $Var(\gamma)$  For complex systems in which individual system components represent the density of some tangible quantity, it is relevant to consider the feasibility of the system. Feasibile equilibria assume that the values of all system components are positive at equilibrium<sup>5,20,21</sup>. This is of particular interest for ecological communities because population density cannot take negative values, meaning that ecological systems need to be feasible for stability to be biologically realistic<sup>20</sup>. While my results are intended to be general to all complex systems, and not restricted to species networks, I have also performed a feasibility analysis on all matrices M tested for stability, and additionally for specific types of ecological communities<sup>2</sup>

- 97 (e.g., competitive, mutualist, predator-prey; see Supplementary Information). Feasibility was unaffected
- by  $Var(\gamma)$ , meaning that for pure interacting species networks, variation in component response time (i.e.,
- 99 species generation time) does not affect stability at biologically realistic species densities.

# Discussion

- Here I have shown that the stability of large systems might be often contigent upon variation in the response
- $_{102}$  rates of their individual components, meaning that factors such as generation time and rate of trait evolution
- (in biological networks), transaction speed (in economic networks), or communication speed (in social networks)
- need to be considered when investigating the stability of complex systems.
- It is important to point out that  $Var(\gamma)$  is not stabilising per se; that is, adding variation in  $\gamma$  to a particular
  - $_{16}$  system M does not necessarily increase the probability that the system will be stable. Rather, systems that
- are observed to be stable are more likely to vary in  $\gamma$ , and for this  $Var(\gamma)$  to be critical to their stability.
- This is caused by the shift in the distribution of eigenvalues that occurs by introducing  $Var(\gamma)$  (Fig. 1b,
- 2b), which can sometimes result in all  $\Re(\lambda) < 0$  but might also increase  $\Re(\lambda)$  values. The mathematics
- underlying this shift in eigenvalue distribution have been investigated <sup>16</sup> and recently applied to questions
- concerning species density and feasibility<sup>17,18</sup>, but have not been interpreted as rates of response of individual
- componetents to system perturbation.
- Nevertheless, ecological interactions do not exist in isolation in empirical systems, but instead interact with
- evolutionary  $^{15}$ , abiotic, or social-economic systems. The relevance of  $\gamma$  for complex system stability presented
- in the main text should therefore not be ignored in the broader context of ecological communities.
- 116 I have focused broadly on random complex systems, but it is also worthwhile to consider more restricted
- interactions such as those of specific ecological networks<sup>2</sup>. These include systems in which all interactions
- are negative (competitive networks), positive (mutualist networks), or i and j pairs have opposing signs
- 119 (predator-prey networks). In general, competitive and mutualist networks tend to be destabilising, and
- predator-prey network tend to be stabilising<sup>22</sup>. When  $Var(\gamma)$  is applied to each, the proportion of stable

competitive and predator-prey networks increases, but the proportion of stable mutualist networks does not (see Supplementary Information). Additionally, when each component of  $\mathbf{M}$  is interpreted as a unique species and given a random intrinsic growth rate<sup>20</sup>, feasibility is not increased by  $Var(\gamma)$ , suggesting that variation in species generation time might be unlikely to drive stability in purely multi-species networks (see Supplementary Information).

Hence, managing the response rates of system components in a targeted way can potentially facilitate the stabilisation of complex systems through a reduction in dimensionality.

My results show that complex systems are more likely to be stable when the response rates of system components vary. These results are broadly applicable to understanding stability of complex networks in the physical, life, and social sciences.

## 31 Methods

Component response rate variation ( $\gamma$ ). In a synthesis of eco-evolutionary feedbacks on community stability, Patel et al. model a system that includes a vector of potentially changing species densities ( $\mathbf{N}$ ) and a vector of potentially evolving traits ( $\mathbf{x}$ )<sup>15</sup>. For any species i or trait j, change in species density ( $N_i$ ) or trait value ( $x_j$ ) with time (t) is a function of the vectors  $\mathbf{N}$  and  $\mathbf{x}$ ,

$$\frac{dN_i}{dt} = N_i f_i(\mathbf{N}, \mathbf{x}),$$

$$\frac{dx_j}{dt} = \epsilon g_j(\mathbf{N}, \mathbf{x}).$$

In the above,  $f_i$  and  $g_j$  are functions that define the effects of all species densities and trait values on the density of a species i and the value of trait j, respectively. Patel et al. were interested in stability when the evolution of traits was relatively slow or fast in comparison with the change in species densities  $^{15}$ , and

this is modulated in the above by the scalar  $\epsilon$ . The value of  $\epsilon$  thereby determines the timescale separation between ecology and evolution, with high  $\epsilon$  modelling relatively fast evolution and low  $\epsilon$  modelling relative slow evolution<sup>15</sup>.

I use the same principle that Patel et al. use to modulate the relative rate of evolution to modulate rates of
component responses for S components. Following May<sup>1,23</sup>, the value of a component i at time t ( $v_i(t)$ ) is
affected by the value of j ( $v_j(t)$ ) and j's marginal effect on i ( $a_{ij}$ ), and by i's response rate ( $\gamma_i$ ),

$$\frac{dv_i(t)}{dt} = \gamma_i \sum_{j=1}^{S} a_{ij} v_j(t).$$

In matrix notation<sup>23</sup>,

$$\frac{d\mathbf{v}(t)}{dt} = \gamma \mathbf{A}\mathbf{v}(t).$$

In the above,  $\gamma$  is a diagonal matrix in which elements correspond to individual component response rates.

Therefore,  $\mathbf{M} = \gamma \mathbf{A}$  modulates the values of components and can be analysed using the techniques of May<sup>1,16,23</sup>.

Genetic algorithm. Ideally, to investigate the potential of  $Var(\gamma)$  for increasing the proportion of stable complex systems, the search space of all possible  $\gamma$  vectors would be evaluated for each unique  $\mathbf{M} = \gamma \mathbf{A}$ .

This is technically impossible because  $\gamma_i$  can take any real value between 0-2, but even rounding  $\gamma_i$  to reasonable values would result in a search space too large to practically explore. Under these conditions, genetic algorithms are highly useful tools for finding practical solutions by mimicking the process of biological evolution<sup>19</sup>. In this case, the practical solution is finding vectors of  $\gamma$  that decrease the most positive real eigenvalue of  $\mathbf{M}$ . The genetic algorithm used achieves this by initialising a large population of 1000 different potential  $\gamma$  vectors and allowing this population to evolve through a process of mutation, crossover (swaping  $\gamma_i$  values between vectors), selection, and reproduction until either a  $\gamma$  vector is found where all  $\Re(\lambda) < 0$  or

some "giving up" critiera is met.

For each  $S = \{2, 3, ..., 39, 40\}$ , the genetic algorithm was run for 100000 random  $\mathbf{M}$  ( $\sigma = 0.4, C = 1$ ), where  $\mathbf{M} = \gamma \mathbf{A}$ . The genetic algorithm was initialised with a population of 1000 different  $\gamma$  vectors with elements 160 sampled i.i.d from  $\gamma_i \sim \mathcal{U}(0,2)$ . Eigenanalysis was performed on the M resulting from each  $\gamma$  vector, and the 161 20  $\gamma$  vectors resulting in M with the lowest max  $(\Re(\lambda))$  each produced 50 clonal offspring with subsequent 162 random mutation and crossover between the resulting new generation of 1000  $\gamma$  vectors. Mutation of each  $\gamma_i$ 163 in a  $\gamma$  vector occurred with a probability of 0.2, resulting in a mutation effect of size  $\mathcal{N}(0,0.02)$  being added 164 to generate the newly mutated  $\gamma_i$  (any  $\gamma_i$  values that mutated below zero were multiplied by -1, and any 165 values that mutated above 2 were set to 2). Crossover occurred between two sets of 100  $\gamma$  vectors paired in 166 each generation; vectors were randomly sampled with replacement among but not within sets. Vector pairs 167 selected for crossover swapped all elements between and including two  $\gamma_i$  randomly selected with replacement 168 (this allowed for reversal of vector element positions during crossover; e.g.,  $\{\gamma_4, \gamma_5, \gamma_6, \gamma_7\} \rightarrow \{\gamma_7, \gamma_6, \gamma_5, \gamma_4\}$ ). 169 The genetic algorithm terminated if a stable M was found, 20 generations occurred, or if the mean  $\gamma$  fitness 170 increase between generations was less than 0.01 (where fitness was defined as  $W_{\gamma} = -\max(\Re(\lambda))$  for M). 171 System feasibility. Dougoud et al.<sup>20</sup> define the following feasibility criteria for ecological systems charac-172 terised by S interacting species with varying densities in a classical Lotka-Volterra model, 173

$$\mathbf{x}^* = -\left(\theta \mathbf{I} + (CS)^{-\delta} \mathbf{J}\right)^{-1} \mathbf{r}.$$

In the above,  $\mathbf{x}^*$  is the vector of species abundances at equilibrium. Feasibility is satisfied if all elements in  $\mathbf{x}^*$  are positive. The matrix  $\mathbf{I}$  is the identity matrix, and the value  $\theta$  is strength of intraspecific competition (diagonal elements). Diagonal values are set to -1, so  $\theta = -1$ . The variable  $\delta$  is a normalisation parameter that modulates the strength of interactions ( $\sigma$ ) for  $\mathbf{J}$ . Implicitly, here  $\delta = 0$  underlying strong interactions. Hence,  $(CS)^{-\delta} = 1$ , so in the above, a diagonal matrix of -1s ( $\theta \mathbf{I}$ ) is added to  $\mathbf{J}$ , which has a diagonal of all zeros and an off-diagonal affecting species interactions (i.e., the expression  $(CS)^{-\delta}$  relates to May's<sup>1</sup> stability criterion<sup>20</sup> by  $\frac{\sigma}{(CS)^{-\delta}}\sqrt{SC} < 1$ , and hence for my purposes  $(CS)^{-\delta} = 1$ ). Given  $\mathbf{A} = \theta \mathbf{I} + \mathbf{J}$ , the above

criteria is therefore reduced to the below,

$$\mathbf{x}^* = -\mathbf{A}^{-1}\mathbf{r}.$$

To check the feasibility criteria for  $\mathbf{M} = \gamma \mathbf{A}$ , I therefore evaluated  $-\mathbf{M}^{-1}\mathbf{r}$  ( $\mathbf{r}$  elements were sampled i.i.d.

from  $r_i \sim \mathcal{N}(0, 0.4^2)$ ). Feasibility is satisfied if all of the elements of the resulting vector are positive.

Acknowledgements: I am supported by a Leverhulme Trust Early Career Fellowship (ECF-2016-376).

Conversations with L. Bussière and N. Bunnefeld, and comments from J. J. Cusack and I. L. Jones, improved

the quality of this work.

Supplementary Information: Full tables of stability results for simulations across different system size (S)

values, ecological community types, connectance (C) values, interaction strengths ( $\sigma$ ), and  $\gamma$  distributions are

provided as supplementary material. An additional table also shows results for how feasibility changes across

<sup>190</sup> S. All code and simulation outputs are publicly available as part of the RandomMatrixStability package on

GitHub (https://github.com/bradduthie/RandomMatrixStability).

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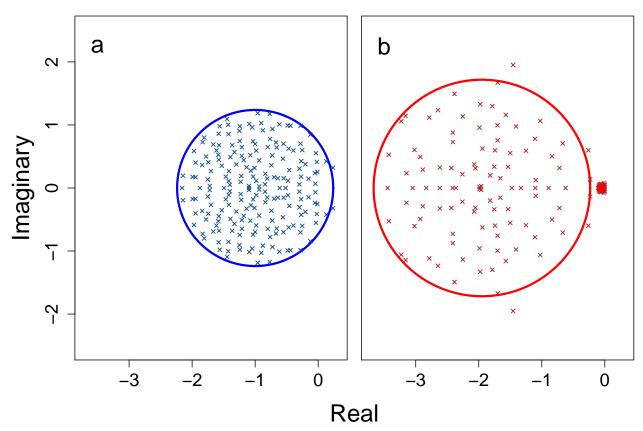
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Figure 1: Example distribution of eigenvalues before (a) and after (b) separating a randomly 235 generated complex system into fast ( $\gamma = 1.95$ ) and slow ( $\gamma = 0.05$ ) component response rates. 236 Each panel shows the same system where S = 200, C = 0.05, and  $\sigma = 0.4$ , and in each case  $E[\gamma] = 1$  (i.e., 237 only the distribution of  $\gamma$  differs between panels). a. Eigenvalues plotted when all  $\gamma = 1$ ; distributions of 238 points are uniformly distributed within the blue circle with a radius of  $\sigma\sqrt{SC}=1.238$  centred at -1 on 239 the real axis. b. Eigenvalues plotted when half  $\gamma = 1.95$  and half  $\gamma = 0.05$ ; distributions of points can be 240 partitioned into one large circle centred at  $\gamma = -1.95$  and one small circle centred at  $\gamma = -0.05$ . In a, the 241 maximum real eigenvalue max  $(\Re(\lambda)) = 0.2344871$ , while in b max  $(\Re(\lambda)) = -0.0002273135$ , meaning that the complex system in b but not a is stable because in b max  $(\Re(\lambda)) < 0$ . In 1 million randomly generated complex systems under the same parameter values, 1 was stable when  $\gamma = 1$  while 32 were stable when  $\gamma = \{1.95, 0.05\}$ . Overall, complex systems that are separated into fast versus slow components tend to be more stable than otherwise identical systems with identical component response rates.



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Figure 2: Distributions of eigenvalues before (a) and after (b) introducing variation in component response rate ( $\gamma$ ) in complex systems. Each panel show the same system where S=1000, C=0.05, and  $\sigma=0.4$ . a. Eigenvalues plotted in the absence of  $Var(\gamma)$  where  $E[\gamma]=1$ , versus b. eigenvalues
plotted given  $\gamma \sim \mathcal{U}(0,2)$ , which increases the variance of interaction strengths ( $\sigma^2$ ) but also creates a cluster
of eigenvalues toward the distribution's centre (-1, 0). Blue elipses in both panels show the circle centred on
the distribution in panel a. Proportions of  $\Re(\lambda) < 0$  are 0.718 and 0.744 for a and b, respectively.

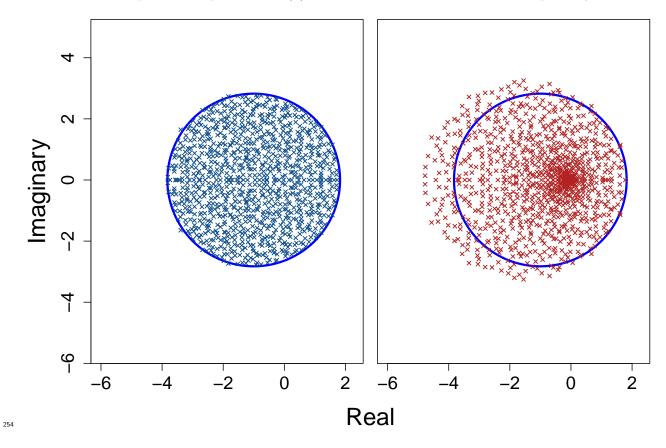


Figure 3: Stability of large complex systems with and without variation in component response rate  $(\gamma)$ . The ln number of systems that are stable across different system sizes  $(S, \max S = 50)$  given C = 1, and the proportion of systems in which variation in  $\gamma$  is critical for system stability. For each S, 1 million complex systems are randomly generated. Stability of each complex system is tested given variation in  $\gamma$  by randomly sampling  $\gamma \sim \mathcal{U}(0,2)$ . Stability given  $Var(\gamma)$  is then compared to stability in an otherwise identical system in which  $\gamma = E[\mathcal{U}(0,2)]$  for all components. Blue and red bars show the number of stable systems in the absence and presence of  $Var(\gamma)$ , respectively. The black line shows the proportion of systems that are stable when  $Var(\gamma) > 0$ , but would be unstable if  $Var(\gamma) = 0$ .

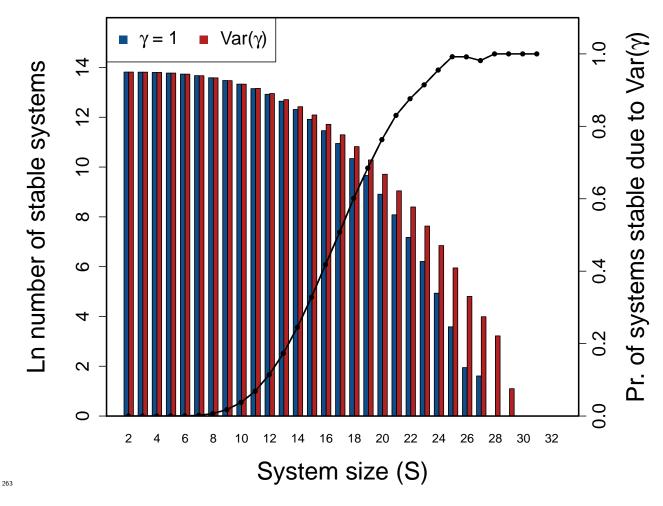


Figure 4: Stability of large complex systems given  $\gamma = 1$  versus targeted  $Var(\gamma)$ . The ln number of systems that are stable across different system sizes  $(S, \max S = 40)$  for C = 1, and the proportion of systems wherein a targeted search of  $\gamma$  values successfully resulted in system stability. For each S, 100000 complex systems are randomly generated. Stability of each complex system is tested given variation in  $\gamma$  using a genetic algorithm to maximise the effect of  $\gamma$  values on increasing stability, as compared to stability in an otherwise identical system in which  $\gamma$  is the same for all components. Blue bars show the number of stable systems in the absence of component response rate variation, while red bars show the number of stable systems that can be generated if component response rate is varied to maximise system stability. The black line shows the proportion of systems that are stable when component response rate is targeted to increase stability, but would not be stable if  $Var(\gamma) = 0$ .

