



Article Modelling Functional Shifts in Two-Species Hypercycles

Bernat Bassols ^{1,*}^(D), Ernest Fontich ²^(D), Daniel Oro ³^(D), David Alonso ³^(D) and Josep Sardanyés ^{4,*}^(D)

- ¹ Department of Mathematics, Imperial College London, London SW7 2AZ, UK ² Department de Matemàtiques i Informàtica, Universitat de Barcolone (UB). Co
 - Departament de Matemàtiques i Informàtica, Universitat de Barcelona (UB), Gran Via de les Corts Catalanes 585, 08007 Barcelona, Spain; fontich@ub.edu
- ³ Theoretical and Computational Ecology Laboratory, Center for Advanced Studies of Blanes (CEAB-CSIC),
- Accés Cala Sant Francesc 14, 17300 Blanes, Spain; d.oro@csic.es (D.O.); dalonso@ceab.csic.es (D.A.)
- ⁴ Centre de Recerca Matemàtica, Edicici C, Campus de Bellatera, 08193 Cerdanyola del Vallès, Spain
- Correspondence: bb420@ic.ac.uk (B.B.); jsardanyes@crm.cat (J.S.)

Abstract: Research on hypercycles focuses on cooperative interactions among replicating species, including the emergence of catalytic parasites and catalytic shortcircuits. Further interactions may be expected to arise in cooperative systems. For instance, molecular replicators are subject to mutational processes and ecological species to behavioural shifts due to environmental and ecological changes. Such changes could involve switches from cooperative to antagonistic interactions, in what we call a functional shift. In this article, we investigate a model for a two-member hypercycle model, considering that one species performs a functional shift. First, we introduce the model dynamics without functional shifts to illustrate the dynamics only considering obligate and facultative cooperation. Then, two more cases maintaining cross-catalysis are considered: (i) a model describing the dynamics of ribozymes where a fraction of the population of one replicator degrades the other molecular species while the other fraction still receives catalytic aid; and (ii) a system in which a given fraction of the population predates on the cooperating species while the rest of the population still receives aid. We have characterised the key bifurcation parameters determining extinction, survival, and coexistence of species. We show that predation, regardless of the fraction that benefits from it, does not significantly change dynamics with respect to the degradative case (i), thus conserving dynamics and bifurcations. Their biological significance is interpreted, and their potential implications for the dynamics of early replicators and ecological species are outlined.

Keywords: cooperation; dynamical systems; functional shifts; ribozymes; origins of life; behavioural ecology

1. Introduction

Hypercycles are catalytic networks of self-replicating species able to catalyse the replication of another single species as they all form a closed loop (see Figure 1a,b) [1,2]. Originally, the hypercycle was suggested to allow for the cooperative selection of competing replicators in the origins of life, ensuring the stability of broad contents of information, contrarily to non-catalytically joint replicators, i.e., quasispecies [3,4]. Hypercycles have been suggested as a possible molecular network of prebiotic replicators involving a crucial step from the transition from inanimate to living chemistry [3,4] due to their potential capacity of maintaining larger stable genetic contents as compared to quasispecies populations under large mutations [1,3]. The interest in hypercycles goes beyond the origin of life problem, being a canonical model to investigate different dynamical systems, including cooperation. The generality of the hypercycle replicator equations has allowed using this model to explore virus replication dynamics [5–8], neural networks [9,10], the immune system [11], or the emergence of language [12]. Moreover, several hypercycle-like systems have been implemented experimentally. These include coiled-coil peptides [13], yeast cell populations [14], and engineered bacteria growing with catalytic parasites [15].



Citation: Bassols, B.; Fontich, E.; Oro, D.; Alonso, D.; Sardanyés, J. Modelling Functional Shifts in Two-Species Hypercycles. *Mathematics* **2021**, *9*, 1809. https:// doi.org/10.3390/math9151809

Academic Editors: Jan Awrejcewicz and José A. Tenreiro Machado

Received: 2 July 2021 Accepted: 23 July 2021 Published: 30 July 2021

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It has been argued that hypercycle units may need two minimal conditions to be evolutionary stable [4,16], namely, catalytic activity and capacity to store information. Good candidates fulfilling these two conditions are catalytic RNA molecules (ribozymes). RNAs with stem and loop structures, similar to transference RNAs [17], are known to be resistant to hydrolysis [18] and also have replicability potential [19,20]. Ribozymes are short RNA molecules with catalytic activity similar to protein enzymes [21,22] and have been considered as potential candidates for forming the first autonomous, self-replicating molecular systems involved in the origins of life [4,16,19,23–26]. Different activities have been described for natural and in vitro (e.g., peptide-bond formation [27]) evolved ribozymes. Certain introns can catalyse their own excision (self-cleavage) from single-stranded RNA (ssRNA) [20] and ligase reactions by RNA catalysts can occur even with short RNA sequences [28]. Moreover, the same RNA sequences can catalyse trans-esterification reactions for the elongation of one monomer [20], ligation of two independent ssRNAs [29,30], and cleavage of RNA into smaller sequences [20,22,31,32] (see also [26,33] for reviews). Recent experiments have revealed that the combination of RNAs with cold-adaptative mutations enabled catalysing the synthesis of an RNA sequence longer than itself (adding about 200 nucleotides) [34]. Other experiments have shown the spontaneous formation of catalytic cycles and networks from mixtures of RNA fragments able to self-assemble into self-replicating ribozymes [35], providing evidence for selective advantage of cooperative systems composed of ribozymes.

Hypercycle equations have also been suggested as a suitable modelling framework to investigate the dynamics of cooperation in complex ecosystems [36]. More recently, the hypercycle theory and the spatio-temporal dynamics of cooperative ecological interactions have been reviewed in the framework of landscape ecology [37]. Cooperation arises in a multitude of ecological systems [38–42]. These include alarm calls, coalition formation, predator inspection, cooperative breeding, protection against attacks by predators or conspecifics, or cooperative hunting, among others. Examples of intra-specific cooperation (individuals of the same species cooperate) are found, for instance, in females of the vampire species Desmodus rotundus [43-45], which perform cooperative breeding (i.e., reciprocal altruism) by sharing blood between themselves. Cooperative hunting, in which the individuals of a given species cooperate to catch the prey, has been described in lions (Panthera leo) [46], brown hyenas (Hyaena brunnea) [47], wild chimpanzees (Pan troglodites) [48], or the fish species Caranx ignobilis [49]. Invertebrate species, such as arthropods, can also hunt cooperatively: Golden-web spiders (Nephila clavipes) [50], the Stegodyphid spider (Stegodyphus mimosarum) [51], or the heteroptera Microvelia douglasi atrolineata [52]. Examples of inter-specific cooperation (cooperation between individuals of different species) have been described in the deep-sea tube worm Lamellibranchia luymesi and microbial symbionts [40], as well as in arbuscular-mycorrhizal fungi, which form an obligate symbiosis with more than 80% of plant species in all terrestrial habitats [42].

Up to now, most of the research on hypercycles has focused on the dynamics arising from the processes of cooperation between replicators, and some works have also included competition for resources (e.g., available mononucleotides or available space) [3,53] and mutation processes [54–56]. In this sense, different architectural changes in hypercycles jeopardising their stability and persistence have been thoroughly investigated to date, mainly considering the so-called catalytic parasites [15,57–60] and, to a lesser extent, catalytic short-circuits [61,62]. Both catalytic parasites and short-circuits involve modifications in the cyclic catalytic patterns. The former involves the emergence (e.g., via mutation) of a replicator that receives catalysis but does not reciprocate the catalytic aid. The latter involves an internal catalytic connection generating a smaller catalytic cycle within the full system.

Some authors have recently been interested in possible shifts in cooperation among species and have used the hypercycle equations to evaluate the dynamics arising from such shifts [63]. For instance, molecular replicators are subject to mutational processes and ecological species to behavioural shifts due to environmental or ecological changes, and one

might expect such species to switch from cooperative to antagonistic interactions in what we call a *functional shift*. The dynamics arising due to functional shifts in hypercycles are still unexplored. The impact of a shift from catalytic cooperation to directed degradation has recently been studied in small discrete-time hypercycles [63]. For this particular case, it was assumed that one of the hypercycle replicators (ribozyme) became a degrading species instead of being cooperative. To the extent of our knowledge, no works have investigated how these shifts may affect the stability of small hypercycles in time-continuous systems.



Figure 1. The different hypercycle architectures studied in this article: (**a**) obligate two-species hypercycle with heterocatalysis; (**b**) facultative system with heterocatalysis and autonomous self-replication; (**c**) hypercycle including directed degradation of S_1 by S_2 while still keeping some fractions of the population receiving catalytic aid; (**d**) two-species ecological hypercycle (e.g., wader and seagull) including opportunistic predation (especially on eggs and chicks). Here, S_2 consumes S_1 , while some individuals still receive cooperation from S_1 .

Examples of functional shifts between cooperating species are found in different ecosystems. For instance, large fish and mammal pelagic predators (tuna, sharks, dolphins) likely cooperate to locate and handle small pelagic shoals [64], and the predation of sharks on dolphins and tuna has been recorded [65,66]. Seabirds form inter-specific flocks that cooperate to locate food at sea, and the predation that occurs at land while breeding in colonies. Body size drives the predatory matrix, which increases its values when environmental conditions worsen [67,68]. Waterbirds form inter-specific mixed colonies to protect against predators (they can contain several different species of waders, terns, and gulls), but some species may predate opportunistically (especially on eggs and chicks) on heterospecifics or may exert kleptoparasitism for food, especially when the environment is harsher [69–75]. Although this may be anecdotal, and the cooperation may be subtle, marine mammals may "cooperate" to locate and capture prey, and at the same time, larger species may exert some predation, e.g., killer whales on elephant seals [76].

In this article, we analyse two-species (labeled as S_1 and S_2) hypercycles considering, together with cooperation, directed degradation and predation (see Figure 1). The paper is organised as follows. In Section 2, we introduce the general model that considers crossed cooperation, competition, and decay of the species and includes the interactions due to the functional shifts. Section 3 contains the results of four different systems that can be studied from the general model by removing some of the interactions. This allows a better comprehension of the dynamics only considering cooperation (plus competition and decay) between species and the dynamical changes arising from directed degradation or predation. First, an obligate two-member hypercycle (Figure 1a), which has been thoroughly investigated in Ref. [53] (see also [77]), is briefly commented. The same system including autonomous self-replication for the two species (see Figure 1b) is explored. Then we explore a hypercycle with a functional shift in a given fraction of S_2 population, which is able to degrade S_1 while still providing catalytic aid to the first species (Figure 1c). This system is inspired in two populations of ribozymes in which some molecules of one of the species (the rest still receive catalysis) acquires the capacity to degrade the other one, still maintaining the catalytic aid. Finally, we explore a similar system in an ecological context considering that a given fraction of S_2 population predates on S_1 (the other fraction receiving aid from S_1) while maintaining the crossed cooperation. Section 4 is devoted to some conclusions.

2. General Mathematical Model

Let us consider a two-member hypercycle with species S_1 and S_2 , and let x_1 and x_2 represent their respective concentrations (population numbers). The mathematical model we investigate adds further interactions to a standard two-member hypercycle [53] considering processes of directed degradation or predation between the two species while keeping some degree of cooperation between the hypercycle members. This model is given by the following dynamical system:

$$\begin{cases} \dot{x}_1 = (\alpha_1 x_1 + k x_1 x_2) \left(1 - \frac{x_1 + x_2}{c_0} \right) - \varepsilon x_1 - (1 - \beta) \varepsilon_{12} x_1 x_2, \\ \dot{x}_2 = (\alpha_2 x_2 + \beta \eta_k k x_1 x_2) \left(1 - \frac{x_1 + x_2}{c_0} \right) - \eta_\varepsilon \varepsilon x_2 + \gamma (1 - \beta) \varepsilon_{12} x_1 x_2, \end{cases}$$
(1)

where all parameters except η_k and η_{ε} are assumed to be bound in I = [0, 1] (see below), and $\eta_k, \eta_{\varepsilon} \ge 0$. That is, parameters have non-negativity constraints: $\alpha_i, k, \varepsilon, \beta, \varepsilon_{ij}, \eta_{k,\varepsilon}, \gamma \ge 0$, and $c_0 > 0$, with i, j = 1, 2 and $i \neq j$. Each of the model parameters corresponds to the following biological processes (see also Figure 1):

- α_i : autonomous self-replication rate (Malthusian growth) of species i = 1, 2. $\alpha_i \neq 0$ implies that species can replicate themselves without the catalytic aid of the other replicator.
- *k*: the cross-catalytic replication parameter between S_1 and S_2 . In the case of S_2 , the term η_k allows considering a non-symmetric case in which both species are kinetically different. The case $\eta_k = 1$ involves symmetric catalytic replication.
- c_0 : known as the carrying capacity, which limits the population of replicators due to finite resources or (implicit) space. Due to the large number of parameters, we will set $c_0 = 1$.
- ε : density-independent spontaneous degradation or death of the species. Analogously to the characteristics of η_k , $\eta_{\varepsilon} \neq 1$ allows considering a non-symmetric decay scenario for S_1 and S_2 .
- ε_{12} : the density-dependent degradation rate due to the cleavage (or predation, see below) of species S_1 by S_2 . We notice that the degradative and predatory dynamics are exclusive. That is, we do not consider a case with directed degradation and predation taking place at the same time (these two different cases will be considered with $\gamma = 0$ or $\gamma \neq 0$, respectively).

- β: the fraction of the S₂ population still receiving cooperation from S₁ replication. The term 1 β corresponds to the rest of the S₂ population that exerts the degradation or the predation of species S₁.
- γ : the energetic efficiency coefficient from predation. It can be understood as the amount of energy species S_2 can gain from predating S_1 and investing it for reproduction (due to energetic constraints $\gamma < 1$). As mentioned, the investigation of the degradative model will be performed by setting $\gamma = 0$, while the predatory system will consider $0 < \gamma < 1$.

As previously mentioned, we focus on parameter values restricted to the interval I = [0, 1], except for η_k , η_ε since we are interested in asymmetric cases. Since the qualitative dynamics mainly depends upon the relative relations between replication/growth and decay/predation parameters, with such an interval we expect to gather all possible dynamics. As mentioned, parameters β and γ are naturally found within this interval in order to be biologically meaningful. Moreover, the carrying capacity has been set to 1 for simplicity. The choice of a different value for c_0 does not change dynamics qualitatively.

It is clear that for the system to hold biological meaning, the phase space is bounded only to the first quadrant of \mathbb{R}^2 . In general terms, the system Equation (1) is able to model the directed degradation of S_1 by S_2 for values of $0 \le \beta < 1$ and $\gamma = 0$ that becomes fully predatory when $\gamma > 0$. The term $1 - (x_1 + x_2)/c_0$ in Equation (1) is a logistic-like function that introduces competition between the two hypercycle species, also bounding the dynamics.

To provide a general modeling framework and investigate the interplay between cooperation (plus competition and decay) and the functional shifts, i.e., directed degradation or predation, we have kept the cross-catalytic terms. A simpler system considering that species S_2 receives aid from S_1 and a given fraction $(1 - \beta)$, with $0 \le \beta < 1$, of the S_2 population only exerts directed degradation (or predation) on S_1 and the rest of the population catalyses S_1 can be also investigated by changing kx_1x_2 to βkx_1x_2 in \dot{x}_1 ; and $\beta \eta_k kx_1x_2$ to $\mu \eta_k x_1x_2$ in \dot{x}_2 , μ being the rate of replication due to the catalysis performed by S_1 on S_2 .

3. Results and Discussion

In this Section, we will investigate the dynamics of Equation (1) considering four different systems. Section 3.1 summarises the results for the obligate two-member hypercycle in which the two species can only replicate catalytically (Figure 1a, see [53,77] for a detailed investigation of this system). Section 3.2 provides further results on this system, including the exponential (non-catalytic) replication of each species. In Section 3.3, we investigate the model considering directed degradation, while Section 3.4 provides further results on the model with predation.

3.1. Obligate Two-Member Hypercycle

Before analysing the models of interest given by directed degradation and predation, it is interesting to start the analysis at a much simpler level. Here, we briefly summarise the dynamics of an obligate two-member hypercycle (Figure 1a). This system, which considers that a species can only replicate due to catalytic (cooperative) processes, can be obtained from Equation (1), setting $\alpha_1 = \alpha_2 = \gamma = 0$ and $\beta = 1$. This model is given by

$$\begin{cases} \dot{x}_1 = \Phi_1(x_1, x_2) = k x_1 x_2 (1 - x_1 - x_2) - \varepsilon x_1, \\ \dot{x}_2 = \Phi_2(x_1, x_2) = \eta_k k x_1 x_2 (1 - x_1 - x_2) - \eta_\varepsilon \varepsilon x_2. \end{cases}$$
(2)

This dynamical system was thoroughly investigated in Ref. [53] (see also [77]). The system has three fixed points: the origin (0,0) and the pair $((\eta_{\varepsilon}/\eta_k)\Gamma_{\pm},\Gamma_{\pm})$, where

$$\Gamma_{\pm} = \frac{1}{2\xi} \left(1 \pm \sqrt{1 - \frac{4\varepsilon\xi}{k}} \right),\tag{3}$$

and $\xi := 1 + \eta_{\varepsilon}/\eta_k$. Note, however, that the expression Γ_{\pm} is well defined only for $\varepsilon \le k/4\xi =: \varepsilon_c$. Otherwise, $\Gamma_{\pm} \in \mathbb{C}$. The origin is a locally asymptotically stable equilibrium, while Γ_{-} is a saddle point and Γ_{+} can be either a stable node [53] or a weak stable focus (see Appendix A). The pair Γ_{\pm} suffers a saddle-node bifurcation at the critical value ε_c . Above this threshold, the origin becomes globally asymptotically stable (see Ref. [53] for further details and the study of this system using spatially-explicit models).

3.2. Facultative Two-Member Hypercycle

The obligate two-member hypercycle is a simple case in which species depend explicitly on the presence of the other to survive. For S_1 to replicate, S_2 must be present in the system, and vice versa [53]. We shall extend the previous model to a system that considers both exponential replication and cross-catalysis, i.e., facultative hypercycle, proposing a setting that has not been analysed in detail so far (see Refs. [78–80] for two-species facultative hypercycles with mutation and error tail). The coupled system of ODEs now involves a parameter of autonomous replication (Malthusian growth), which can be introduced in (1) setting $\alpha_{1,2} > 0$, while maintaining with $\beta = 1$ and $\gamma = 0$. We thus consider:

$$\begin{cases} \dot{x}_1 = \Phi_1(x_1, x_2) = (\alpha_1 x_1 + k x_1 x_2)(1 - x_1 - x_2) - \varepsilon x_1, \\ \dot{x}_2 = \Phi_2(x_1, x_2) = (\alpha_2 x_2 + \eta_k k x_1 x_2)(1 - x_1 - x_2) - \eta_\varepsilon \varepsilon x_2. \end{cases}$$
(4)

To understand this system, we may differentiate three cases depending on the relative value of α_i and ε . We will consider $\alpha_i > \varepsilon$ for i = 1, 2, $\alpha_i < \varepsilon < \alpha_j$ for $i \neq j$, and $\alpha_i < \varepsilon$ for i = 1, 2.

Solving Equation (4) for $\dot{x}_1 = \dot{x}_2 = 0$ gives five critical, i.e., equilibrium, points:

$$(0,0), \qquad \left(0,\frac{\alpha_2-\eta_{\varepsilon}\varepsilon}{\alpha_2}\right), \qquad \left(\frac{\alpha_1-\varepsilon}{\alpha_1},0\right), \qquad (\Gamma_{\pm},\Omega_{\pm}),$$

where

$$\Gamma_{\pm} = \frac{\eta_{\varepsilon}\eta_{k}k + \eta_{\varepsilon}\eta_{k}\alpha_{1} - \eta_{\varepsilon}\alpha_{2} - 2\eta_{k}\alpha_{2} \pm \eta_{\varepsilon}\sqrt{\Delta}}{2k\eta_{k}(\eta_{\varepsilon} + \eta_{k})},$$
(5)

$$\Omega_{\pm} = \frac{-2\eta_{\varepsilon}\alpha_1 + \eta_k k - \eta_k \alpha_1 + a_2 \pm \sqrt{\Delta}}{2k(\eta_{\varepsilon} + \eta_k)},\tag{6}$$

and

$$\Delta = (k^2 + (-4\varepsilon + 2\alpha_1)k + \alpha_1^2)\eta_k^2 + ((-4\eta_\varepsilon \varepsilon + 2\alpha_2)k + s\alpha_1\alpha_2)\eta_k + \alpha_2^2.$$
(7)

The nullclines of the system are the set of solutions of $\Phi_1(x_1, x_2) = 0$ and $\Phi_2(x_1, x_2) = 0$ and determine the direction of the flow in the phase space. Most of the results presented below will be obtained from the nullclines, which are given by:

$$x_1 = 0, \qquad x_1 = -\frac{kx_2^2 - kx_2 + x_2\alpha_1 + \varepsilon - \alpha_1}{kx_2 + \alpha_1} =: g(x_2)$$
 (8)

for Φ_1 , and

$$x_{2} = 0, \qquad x_{2} = -\frac{\eta_{k}kx_{1}^{2} - \eta_{k}kx_{1} + x_{1}\alpha_{2} + \eta_{\varepsilon}\varepsilon - \alpha_{2}}{\eta_{k}kx_{1} + \alpha_{2}} =: h(x_{1})$$
(9)

for Φ_2 . The intersection point of a curve in Equation (8) with another in Equation (9) is an equilibrium point, here denoted by (x_1^*, x_2^*) , where the flow vanishes. Hence, we are interested in the existence of such equilibria in the phase space $(x_1, x_2 > 0)$, as well as in their stability.

Let us consider $\alpha_1, \alpha_2/\eta_{\varepsilon} > \varepsilon$. The nullclines from Equations (8) and (9) allow for three possible configurations, which are shown in Figure 2 and will be shown in the following construction. These can be analytically separated by studying the relative position of each

curve's intersection with the axes. On one hand, $g(x_2)$ crosses the x_1 axis at the critical point $((\alpha_1 - \varepsilon)/\alpha_1, 0)$ and intersects with the x_2 axis at $(0, \Theta_{\pm})$, where

$$\Theta_{\pm} = \frac{k - \alpha_1 \pm \sqrt{k^2 + (-4\varepsilon + 2\alpha_1)k + \alpha_1^2}}{2k}.$$
(10)

Note that Θ_{\pm} is well defined since for its square root to be real, it is needed that $\varepsilon < (k + \alpha_1)^2/4k$, and this holds since $\varepsilon < \alpha_1 \le (k + \alpha_1)^2/4k$, indeed

$$(k+\alpha_1)^2 - 4k\alpha_1 = k^2 + \alpha_1^2 - 2k\alpha_1 = (k-\alpha_1)^2 \ge 0$$

$$\Rightarrow \alpha_1 \le \frac{(k+\alpha_1)^2}{4k}.$$
 (11)

Similarly, $h(x_1)$ intersects with the x_1 axis at $(\Lambda_{\pm}, 0)$ with

$$\Lambda_{\pm} = \frac{k\eta_k - \alpha_2 \pm \sqrt{\eta_k^2 k^2 + (-4\eta_\varepsilon \varepsilon + 2\alpha_2)k\eta_k + \alpha_2^2}}{2\eta_k k}$$
(12)

and at $(0, (\alpha_2 - \eta_{\varepsilon}\varepsilon)/\alpha_2)$ with the x_2 axis. Again, Λ_{\pm} is well defined since for its square root to be real we need

$$\varepsilon \le \frac{(k\eta_k + \alpha_2)^2}{4\eta_k k\eta_\varepsilon} \tag{13}$$

and we have

$$(k\eta_k + \alpha_2)^2 - 4\eta_k k\eta_\varepsilon \frac{\alpha_2}{\eta_\varepsilon} = k^2 \eta_k^2 + \alpha_2^2 - 2k\eta_k \alpha_2 = (k\eta_k - \alpha_2)^2 \ge 0$$

$$\Rightarrow \frac{\alpha_2}{\eta_\varepsilon} \le \frac{(h\eta_k + \alpha_2)^2}{4\eta_k k\eta_\varepsilon}.$$
 (14)

Thus, as expected

$$\varepsilon < \frac{\alpha_2}{\eta_\varepsilon} \le \frac{(k\eta_k + \alpha_2)^2}{4\eta_k k\eta_\varepsilon}.$$
 (15)

In order for the case in Figure 2a to take place, the following conditions must be fulfilled:

$$\frac{\alpha_1 - \varepsilon}{\alpha_1} < \Lambda_+ \Rightarrow \varepsilon < \varepsilon_1 := \frac{-\alpha_1(\eta_\varepsilon \alpha_1 - \eta_k k - \alpha_2)}{\eta_k k},\tag{16}$$

and

$$\frac{\alpha_2 - \eta_{\varepsilon}\varepsilon}{\alpha_2} < \Theta_+ \Rightarrow \varepsilon < \varepsilon_2 := \frac{\alpha_2(k\eta_{\varepsilon} + \eta_{\varepsilon}\alpha_1 - \alpha_2)}{\eta_{\varepsilon}^2 k}.$$
(17)

Therefore, we will encounter Figure 2b if $\varepsilon_1 < \varepsilon < \varepsilon_2$ and Figure 2c when $\varepsilon_2 < \varepsilon < \varepsilon_1$. Given the nature of the considered parameters and the curves $g(x_2)$ and $h(x_1)$, ε can not be greater than both ε_1 and ε_2 simultaneously: Let us assume $\varepsilon > \varepsilon_1$. We can express ε_1 and ε_2 as

$$\varepsilon_{1} = \alpha_{1} \left(1 + \frac{\alpha_{2} - \eta_{\varepsilon} \alpha_{1}}{\eta_{k} k} \right),$$

$$\varepsilon_{2} = \frac{\alpha_{2}}{\eta_{\varepsilon}} \left(1 - \frac{\alpha_{2} - \eta_{\varepsilon} \alpha_{1}}{\eta_{\varepsilon} k} \right),$$
(18)

and since $\varepsilon < \alpha_1$, we have $\alpha_2 - \eta_{\varepsilon} \alpha_1 < 0$. This implies $\varepsilon_2 > \alpha_2 / \eta_{\varepsilon}$, and since $\varepsilon < \alpha_2 / \eta_{\varepsilon}$, the proof is finished. The negation of this statement allows us to justify the impossibility of $\varepsilon < \varepsilon_1, \varepsilon_2$. Therefore, only the three scenarios presented in Figure 2 will be considered.



Figure 2. Possible dynamical scenarios according to the shape of the nullclines. Solutions of $\dot{x}_1 = 0$ are plotted in black and in red for $\dot{x}_2 = 0$. The symmetry coefficients η_k and η_{ε} have been set to 1 for the three plots.

The corresponding phase portraits of the three presented cases can be obtained by further extending the study on the nullclines and the sign of Φ_1 and Φ_2 in the phase space, as we show in Figure 3. It is interesting to note that the origin is a repulsive node for all three scenarios, as its associated Jacobian matrix has positive eigenvalues $\alpha_1 - \varepsilon$ and $\alpha_2 - \eta_{\varepsilon}\varepsilon$. This implies that critical points located on each axis are attractive, i.e., there will never be a complete extinction of both species simultaneously as they can survive without the need of the other.



Figure 3. The direction of the dynamical system vector field for all possible situations regarding $\varepsilon < \alpha_1, \alpha_2/\eta_\varepsilon$ together with nullclines $g(x_2)$ and $h(x_1)$ plotted in black and red, respectively. Parameters used: (**a**) $\varepsilon = 0.16, k = 0.74, \alpha_1 = 0.28, \alpha_2 = 0.4$; (**b**) $\varepsilon = 0.17, k = 0.59, \alpha_1 = 0.96, \alpha_2 = 0.27$; (**c**) $\varepsilon = 0.17, k = 0.40, \alpha_1 = 0.28, \alpha_2 = 0.78$. Below, we display all possible phase portraits for the two-member hypercycle with exponential replication parameter higher than its decay rate in both species. In particular, the plots have been obtained for the values: (**a**) $\alpha_1 = 0.39, \alpha_2 = 0.94, \varepsilon = 0.14, k = 1, \eta_k = 3.6, \eta_{\varepsilon} = 2.95$; (**b**) $\alpha_1 = 0.77, \alpha_2 = 0.51, \varepsilon = 0.14, k = 0.28, \eta_k = 3.6, \eta_{\varepsilon} = 2.95$; (**c**) $\alpha_1 = 0.27, \alpha_2 = 0.93, \varepsilon = 0.16, k = 0.3, \eta_k = 0.35, \eta_{\varepsilon} = 1.45$. Stable equilibria are denoted with red solid circles.

In conclusion, parameters ε_1 and ε_2 allow us to have certain control on the ratio of births versus deaths of each species. As long as they are both higher than ε , an attractive

coexistence will take place without them being able to become extinct. However, as soon as $\varepsilon_i < \varepsilon$ (i = 1, 2), species S_i , $j \neq i$, will become extinct.

Let us now consider the case $\alpha_1 > \varepsilon$, $\alpha_2 < \varepsilon/\eta_{\varepsilon}$. The nullclines allow the five qualitative distributions represented in Figure 4. Since they cannot be all obtained only by variations of ε , it is more adequate to use a different control parameter, specifically α_1 . Since $\alpha_1 > \varepsilon$, there will always be at least another critical point apart from the origin located at $((\alpha_1 - \varepsilon)/\alpha_1, 0)$. This is a result of the exponential self-replication rate for species S_1 being higher than its decay and not needing species S_2 for replication.



Figure 4. The direction of the dynamical system vector field for all possible situations regarding $\alpha_1 > \varepsilon$ and $\alpha_2 < \eta_{\varepsilon}\varepsilon$. Below, we display a phase portrait of the system in the five possible situations studied. Specifically, the plots have been obtained using the following parameters: (**a**) $\alpha_1 = 0.3$, $\alpha_2 = 0.18$, $\varepsilon = 0.23$, k = 0.29, $\eta_k = 5.5$, $\eta_{\varepsilon} = 2.3$; (**b**) $\alpha_1 = 0.24$, $\alpha_2 = 0.05$, $\varepsilon = 0.21$, k = 0.37, $\eta_k = 4.3$, $\eta_{\varepsilon} = 1.2$; (**c**) $\alpha_1 = 0.34$, $\alpha_2 = 0.13$, $\varepsilon = 0.32$, k = 1, $\eta_k = 7.5$, $\eta_{\varepsilon} = 2.75$; (**d**) $\alpha_1 = 0.56$, $\alpha_2 = 0.13$, $\varepsilon = 0.32$, k = 1, $\eta_k = 7.5$, $\eta_{\varepsilon} = 2.75$.

It has already been seen in Equation (13) that $h(x_1)$ (red nullcline) intersects with axis x_1 as long as $\varepsilon < (k\eta_k + \alpha_2)^2/(4\eta_k\eta_{\varepsilon}k)$. If ε is greater than or equal to this value then, as long as $\alpha_1 > \varepsilon$, there will only be two equilibrium points: the origin and $((\alpha_1 - \varepsilon)/\alpha_1, 0)$, as shown in Figure 4a. In this case, the origin is a saddle since its eigenvalues are $\alpha_1 - \varepsilon > 0$ in the x_1 direction, and $\alpha_2 - \eta_{\varepsilon}\varepsilon < 0$ in the x_2 direction. The other equilibrium corresponds to an attractive node since its eigenvalues are $\lambda_1 = \varepsilon - \alpha_1 < 0$ in the x_1 direction and

$$\lambda_2 = -\frac{\varepsilon(\varepsilon\eta_k k + \eta_\varepsilon \alpha_1^2 - \eta_k k\alpha_1 - \alpha_1 \alpha_2)}{\alpha_1^2} < 0, \tag{19}$$

given the restriction on ε from Equation (13), in another direction, we can leave unspecified.

If ε is lower than the value given in (13), there is a possibility that more critical points appear. As seen below, for the scenario in Figure 4b to take place, potential critical points ($\Gamma_{\pm}, \Omega_{\pm}$) must be undefined, i.e., $\varepsilon > \varepsilon_c$, which is equivalent to

$$\alpha_1 < \frac{-k\eta_k - \alpha_2 + 2\sqrt{\varepsilon\eta_k k(\eta_\varepsilon + \eta_\varepsilon)}}{\eta_k} =: \alpha_1^c.$$
(20)

In order to characterise cases in Figure 4c–e, we need to find the value of α_1 for which the nullclines intersect right on top of the x_1 axis. After solving the corresponding equation, the intersection at $(\Lambda_-, 0)$ happens if

$$\alpha_1 = \frac{2\varepsilon\eta_k k}{k\eta_k + \alpha_2 + \sqrt{\eta_k^2 k^2 - (4\eta_\varepsilon \varepsilon - 2\alpha_2)k\eta_k + \alpha_2^2}} =: \alpha_1^-.$$
(21)

and at $(\Lambda_+, 0)$ if

$$\alpha_1 = \frac{2\varepsilon\eta_k k}{k\eta_k + \alpha_2 - \sqrt{\eta_k^2 k^2 - (4\eta_\varepsilon \varepsilon - 2\alpha_2)k\eta_k + \alpha_2^2}} =: \alpha_1^+.$$
(22)

Note α_1^{\pm} are well defined since $\varepsilon < (k\eta_k + \alpha_2)^2/(4\eta_k\eta_{\varepsilon}k)$. Therefore, we will see a nullcline distribution similar to the one presented in Figure 4c if $a_1^c < a_1 < a_1^-$, in Figure 4d if $a_1^- < a_1 < a_1^+$, and in Figure 4e if $\alpha_1^+ < \alpha_1$.

As already stated, the origin is a saddle with positive eigenvalue in the x_1 direction and negative in the x_2 . As for the critical point on the x_1 axis, $((\alpha_1 - \varepsilon)/\alpha_1, 0)$, its eigenvalues are $\lambda_1 = \varepsilon - \alpha_1 < 0$ in the x_1 direction and λ_2 from Equation (19) in another unspecified direction. This last expression is positive for $\alpha_1 \in (\alpha_1^-, \alpha_1^+)$ and negative elsewhere. By Hartman's theorem, we can conclude that it is a saddle in the case Figure 4d and an attractive node in the rest. Note the degenerate situation for $\alpha_1 = \alpha_1^-, \alpha_1^+$ with a null eigenvalue, while the other is still negative, thus being two stable equilibria. Representing the vector field together with the nullclines allows us to see, as expected, that the critical point (Γ_{-}, Ω_{-}) present when $\alpha_1^c < \alpha_1 < \alpha_1^-$ is a saddle and (Γ_{+}, Ω_{+}) a stable node. Figure 4 shows the corresponding phase portraits for all five possible scenarios commented. In Figure 5, we present the bifurcation diagram in which all cases are probed using α_1 as the control parameter. First of all, the only stable critical point is the one located at the x_1 axis, as both nullclines do not intersect at $(\Gamma_{\pm}, \Omega_{\pm})$. After this region, there is a range, $(\alpha_1^c, \alpha_1^-) = (0.3242, 0.3345)$, in which two equilibria are found. Indeed, one of them has $x_2^* = 0$ corresponding to $((\alpha_1 - \varepsilon)/\alpha_1, 0)$, while the other is (Γ_+, Ω_+) . Finally, the higher equilibrium is lowered until it reaches the axis again and x_2^* becomes zero.



Figure 5. (Left) A bifurcation diagram when $\varepsilon > (k\eta_k + \alpha_2)^2 / (4\eta_k\eta_{\varepsilon}k)$ using $\alpha_1 \in (\varepsilon, 1)$ as control parameter for a set of parameters such that the four cases in Figure 4b–e are present. Specifically, $\alpha_2 = 0.14$, $\varepsilon = 0.26$, k = 1, $\eta_k = 2.75$, $\eta_{\varepsilon} = 2.25$. Stable equilibria are represented in blue for x_1^* and in red for x_2^* coordinates. (**Right**) Close up of the diagram for $\alpha_1 \in (\alpha_1^c, \alpha_1^-)$ showing the coexistence of two stable equilibria; one of which implies the extinction of x_2 .

A study regarding $\alpha_1 < \varepsilon$ and $\alpha_2 > \eta_{\varepsilon}\varepsilon$ would bring us to a set of results analogous to the ones exposed in the previous paragraphs because the symmetry of the problem allows us to use the arguments already presented.

The last case we need to look into is when $\alpha_1 < \varepsilon$ and $\alpha_2 < \eta_{\varepsilon}\varepsilon$. Apart from the origin, the other critical points there can be are $(\Gamma_{\pm}, \Omega_{\pm})$. These will be present as long as $\varepsilon \leq \varepsilon_c$ where the higher is an attractive node and the lower a saddle. At $\varepsilon = \varepsilon_c$, the bifurcation point, the critical point becomes a saddle-node, just as studied in Section 3.1.

3.3. Directed Degradation and Cooperation in Ribozymes

The next step towards the generalisation of the dynamical system to describe all considered interactions is the addition of density-dependent degradation, describing the

process by which some replicators (ribozymes) can degrade the other hypercycle member due to trans-cleaving activity. In order to do so, a new parameter must be introduced, β , which may be regarded as the fraction of species S_2 that receives catalysis from species S_1 , while $(1 - \beta)$ does exactly the opposite with S_2 degrading S_1 at a rate ε_{12} . In this case, the cross-catalytic replication for species S_2 is affected by the presence of this parameter as only a fraction β of this species will receive catalysis.

For the sake of length and simplicity, this section is studied under the assumption that $\alpha_1 = \alpha_2 = \eta_{\varepsilon} = k = c_0 = 1$, while ε , ε_{12} , η_k and β are allowed to move freely. This lets us focus strictly on the effect of the degradation behaviour led by S_2 without over-complicating the whole model. Furthermore, such restrictions have not been chosen arbitrarily but, in fact, hold specific biological meaning and build the cases we are most interested in when considering ribozyme interactions. Hence, the system we will focus on along this section is:

$$\begin{cases} \dot{x}_1 = (x_1 + x_1 x_2)(1 - (x_1 + x_2)) - \varepsilon x_1 - (1 - \beta)\varepsilon_{12} x_1 x_2, \\ \dot{x}_2 = (x_2 + \beta \eta_k x_1 x_2)(1 - (x_1 + x_2)) - \varepsilon x_2. \end{cases}$$
(23)

An analytical study, as has been carried out in the previous sections, shows that up to three critical points in the first quadrant apart from the origin exist. As previously stated, given $\alpha_1 = \alpha_2 = 1 > \varepsilon = \eta_{\varepsilon}\varepsilon$, global extinction is impossible since the origin is a repulsive node. As for the other equilibria, one is located on each axis representing the potential survival of a single species without the need of the other. The third equilibrium is only present for a set of values such that $\eta_k > 0$ and

$$\beta > \beta_{\rm bif} = 1 - \frac{\varepsilon}{\varepsilon_{12}} > 0, \tag{24}$$

stating that a certain amount of S_2 higher than this specific threshold must collaborate with S_1 in order to achieve coexistence. Interestingly, for $\varepsilon > \varepsilon_{12}$, both species evolve towards their coexistence regardless of the initial conditions, as long as they are non-zero.

To prove the attractive behaviour of the critical point located outside the axes, we can once again focus on the nullcline distribution. It is quite straightforward to see that bounded to the *x*-axis, $g(x) > \max\{h(0)\}$, where $g(x_1)$ and $h(x_2)$ have been defined analogously to the previous sections. This describes the same scenario presented in Figure 3a, which allows us to conclude this equilibrium is stable. The fact that $g(x) > \max\{h(0)\}$ on the abscissa tells us that shifting β to lower values results in the middle equilibrium to move towards the critical point on the *y*-axis as long as $\varepsilon \leq \varepsilon_{12}$, i.e., $\beta_{\text{bif}} \geq 0$, until the critical point leaves the phase space and a transcritical bifurcation takes place ($\beta \leq \beta_{\text{bif}}$). See Figure 6a,b. On the other hand, if $\varepsilon > \varepsilon_{12}$, lowering β results in the approach of the critical point present on the *x*-axis until $\beta = 0$, where they collide and a transcritical bifurcation takes place as well. See Figure 6c,d.

Finally, a different behaviour is seen for $\eta_k = 0$, i.e., when S_2 is unable to catalytically replicate with the help of S_1 . In this case, there are only three critical points in the region of interest: the origin, which remains a repulsive node, and each individual survival. If $\beta > \beta_{\text{bif}}$, then the previous critical point approaches the equilibrium on the *x*-axis as η_k is reduced until it reaches zero and is absorbed by it. However, for $\beta < \beta_{\text{bif}}$, it is the other equilibrium that becomes the ω -limit of the system. See Figure 6e and Figure 6f, respectively.



Figure 6. Phase portraits of the system in the qualitatively different identified situations. Specifically, we have used: (a) $\varepsilon = 0.13$, $\varepsilon_{12} = 0.72$, $\beta = 0.9$, $\eta_k = 1.7$; (b) $\varepsilon = 0.13$, $\varepsilon_{12} = 0.72$, $\beta = 0.25$, $\eta_k = 1.7$; (c) $\varepsilon = 0.52$, $\varepsilon_{12} = 0.22$, $\beta = 0.55$, $\eta_k = 0.8$; (d) $\varepsilon = 0.52$, $\varepsilon_{12} = 0.22$, $\beta = 0$, $\eta_k = 0.8$; (e) $\varepsilon = 0.25$, $\varepsilon_{12} = 0.49$, $\beta = 0.8$, $\eta_k = 0$; (f) $\varepsilon = 0.25$, $\varepsilon_{12} = 0.49$, $\beta = 0.14$, $\eta_k = 0$. Again, a red dot indicates the ω -limit of each orbit and a blue dot its initial condition.

The behaviour described above can be very well visualised by plotting three-dimensional and heatmap bifurcation diagrams where the x_1^* or x_2^* equilibrium component is computed for given parameters. This is possible given the fact that all orbits outside the axes will evolve to the same equilibrium point and because the origin acts as a repulsive node at all times. Thus, if we choose to plot x_1^* , i.e., the abscissa component of the equilibrium, and it falls on 0, we can read that S_1 has been extinguished, while S_2 has survived. The annihilation of S_2 when $\eta_k > 0$ is only possible for $\varepsilon > \varepsilon_{12}$ and $\beta = 0$; therefore, it is more adequate to represent x_1^* instead of x_2^* as the coexistence will be present as long as $x_1^* > 0$ and $\beta \neq 0$.

Keeping this argument in mind, Figure 7 collects all information presented in Figure 6a–c. Indeed, we can see how the coexistence is present for $\beta > \beta_{\text{bif}}$, while S_1 ceases to exist if $\beta \leq \beta_{\text{bif}}$. For $\eta_k = 0$, Figure 8 shows how x_1^* becomes zero if $\beta < \beta_{\text{bif}}$, indicating that S_1 vanishes while S_2 remains, and vice versa for $\beta > \beta_{\text{bif}}$. In order to demonstrate that coexistence is not possible in these scenarios, both x_1^* and x_2^* have been represented in separate plots (Figure 8a,c and Figure 8b,d, respectively). Note that the coexistence that can be observed on the curve $\beta = \beta_{\text{bif}}$ is not a mere casualty and is thoroughly explained below.

As presented in Figure 8, the case $\beta = \beta_{\text{bif}}$ under $\eta_k = 0$ is quite interesting as well. Here, both equilibria located on the axes are present; hence, the nullclines become the exact same curves in the phase space. This translates to a continuum of fixed points filling the attractive linear manifold $x_2 = 1 - \varepsilon - x_1$. The dynamical system under these conditions and the adequate simplification becomes:

$$\begin{cases} \dot{x}_1 = -x_1(x_2+1)(x_1+x_2+\varepsilon-1), \\ \dot{x}_2 = -x_2(x_1+x_2+\varepsilon-1). \end{cases}$$
(25)

0.8

0.6 & 0.4

0.2



0.2

0.4 ß

Figure 7. Two-parameter bifurcation diagram representing x_1^* as a function of ε_{12} and β plotted as a surface (**a**) and as a heat map (**b**) for a more intuitive understanding. Parameters used: $\varepsilon = 0.2$, $\eta_k = 2$.

0.0

Assuming $(x_1 + x_2 + \varepsilon - 1) \neq 0$, we can compute the quotient \dot{x}_1/\dot{x}_2 , allowing us to find a first integral of the problem:

$$\frac{\dot{x}_1}{\dot{x}_2} = \frac{x_1(x_2+1)}{x_2} \Rightarrow \frac{\dot{x}_1}{x_1} = \left(1 + \frac{1}{x_2}\right)\dot{x}_2,$$
(26)

0.6

(b)

0.8

integrating on both sides, we obtain

02

0.4 0.6 ^{εl2}

0.8

0.0 1.0 (a)

$$\ln(x_1) = x_2 + \ln(x_2) + C, \tag{27}$$

such that if initial conditions $x_1(0)$, $x_2(0)$ are the species initial conditions, Equation (27) can be rewritten as

$$x_1 = \frac{x_1(0)}{x_2(0)} x_2 e^{x_2 - x_2(0)}.$$
(28)

Taking $H = \ln(x_1) - x_2 - \ln(x_2) = 0$ and *X* the system's field, we have

$$H' = \langle \nabla H, X \rangle = \frac{\dot{x}_1}{x_1} - \left(1 + \frac{1}{x_2}\right) \dot{x}_2 = 0$$
(29)

which confirms that the expression is a first integral. This result tells us that all orbits outside the axes follow Equation (27) until they reach the stable line $x_2 = 1 - \varepsilon - x_1$. This is of course attractive since $\dot{x}_1, \dot{x}_2 > 0$ for $x_1 + x_2 < 1 - \varepsilon$ and $\dot{x}_1, \dot{x}_2 < 0$ for $x_1 + x_2 > 1 - \varepsilon$. The corresponding phase portrait for this setting is shown in Figure 9, where the solutions can be seen to follow exactly the level curves of the first integral.

Interestingly enough, we are in front of a case of extreme equilibrium in which the degrading species, S_2 , is unable to cross-catalytically replicate with S_1 , while S_1 benefits from this interaction with a fraction β of S_2 but is degraded by a fraction $1 - \beta$.

0.1

0.0

1.0



Figure 8. Two-parameter bifurcation diagram representing x_1^* (**a**,**c**) and x_2^* (**b**,**d**) as a function of ε_{12} and β plotted as a surface (**a**,**b**) and a heat map (**c**,**d**) for a more intuitive understanding. Parameters used: $\varepsilon = 0.13$, $\eta_k = 0$.



Figure 9. Phase portraits showing a continuum of equilibrium points following $x_1 + x_2 = 1 - \varepsilon$ (line of quasi-neutral equilibria). Initial conditions are represented with a blue dot and their ω -limit with a red dot. Specific parameters used are: (**a**) $\varepsilon = 0.25$, $\varepsilon_{12} = 0.49$, $\eta_k = 0$, and $\beta = \beta_{\text{bif}} = 0.8194$; and (**b**) $\varepsilon = 0$, $\varepsilon_{12} = 0$, $\eta_k = 0.8$, and $\beta = 0.5$.

Another case worth considering is that in which $\varepsilon = 0$ and $\varepsilon_{12} \ll 1$, meaning that natural degradation is no longer present. This leaves us with a system in which both

species will only decrease as a consequence of the coupling term as long as $x_1 + x_2 > 1$. For $\varepsilon_{12} = 0$, the ODE system (1) becomes:

$$\begin{cases} \dot{x}_1 = (x_1 + x_1 x_2)(1 - (x_1 + x_2)), \\ \dot{x}_2 = (x_2 + \beta \eta_k x_1 x_2)(1 - (x_1 + x_2)). \end{cases}$$
(30)

These equations can be solved assuming $1 - (x_1 + x_2) \neq 0$, dividing \dot{x}_1 over \dot{x}_2 , and organising both sides of the expression to find:

$$\left(\beta\eta_k + \frac{1}{x_1}\right)\dot{x}_1 = \left(1 + \frac{1}{x_2}\right)\dot{x}_2,\tag{31}$$

which can be integrated resulting in

$$\beta \eta_k x_1 + \ln(x_1) = x_2 + \ln(x_2) + C, \tag{32}$$

where $C = \beta \eta_k x_1(0) + \ln(x_1(0)) - x_2(0) - \ln(x_2(0))$. This curve describes the orbits followed by an initial condition $(x_1(0), x_2(0))$ in the phase space such that $1 - (x_1(0) + x_2(0)) \neq 0$. Note that $1 - (x_1 + x_2) = 0$ is a continuum of attractive equilibria and that the origin acts as a repulsive node. Figure 9b explicitly shows this behaviour. Finally, as long as $\beta > 0$, a small variation in ε_{12} brings us straight to the case studied in Figure 6b, implying the extinction of S_1 , and a minor variation in ε recovers Figure 6c, showing coexistence.

3.4. Interplay between Predation and Cooperation in Ecology

All systems studied above have been analysed in order to build up a general hypercycle model that allows us to consider all interactions presented in Section 3.3 while S_2 also behaves as a predator. Both species are now under the influence of self-replication ($\alpha_i > 0$ for i = 1, 2) and cross-catalytic replication ($k, \eta_k > 0$), they are also degraded by natural causes ($\varepsilon, \eta_{\varepsilon} > 0$), a fraction β of S_2 receives cooperation from S_1 , the rest of S_2 ($1 - \beta$) consumes S_1 at a rate ε_{12} , and for the first time, S_2 benefits from this consumption with an efficiency of $\gamma \in (0, 1)$, acting as a predator. We now reached the system presented in Equation (1), Section 2, with all parameters playing their corresponding biological roles:

$$\begin{cases} \dot{x}_1 = (\alpha_1 x_1 + k x_1 x_2)(1 - (x_1 + x_2)) - \varepsilon x_1 - (1 - \beta)\varepsilon_{12} x_1 x_2, \\ \dot{x}_2 = (\alpha_2 x_2 + \beta \eta_k k x_1 x_2)(1 - (x_1 + x_2)) - \eta_\varepsilon \varepsilon x_2 + \gamma (1 - \beta)\varepsilon_{12} x_1 x_2. \end{cases}$$
(33)

As we did above, we shall consider a simple scenario setting $\alpha_1 = \alpha_2 = k = \eta_{\varepsilon} = c_0 = 1$ to be able to analyse it mathematically and to obtain clearer results arising from predation. Even though a new process is being considered in this case, surprisingly, the dynamics of the predatory scenario behave exactly as those described in Section 3.3. Curiously enough, although species S_2 now grows at the expense of S_1 , studying the relative positions of the nullclines shows that for $\beta > \beta_{\text{bif}} = 1 - \varepsilon/\varepsilon_{12}$, there is coexistence while S_1 vanishes for $\beta \leq \beta_{\text{bif}}$ (see Figures 10 and 11). As previously explained, the existence of the coexistence equilibrium is given by the position of all four positive axis crossings of the nullclines. We shall refer to the intersection of the x_j nullcline with the x_i -axis as $x_{i,j}^*$ and claim that $x_{1,1}^* < x_{1,2}^*$ and $x_{2,1}^* > x_{2,2}^*$ for $\beta > \beta_{\text{bif}}$, which implies the desired coexistence. Let us compute these values explicitly. On the x_1 axis, the intersections of interest happen at

$$x_{1,1}^* = 1 - \varepsilon,$$

$$x_{1,2}^* = \frac{1}{2\beta\eta_k} \bigg(-\beta\gamma\varepsilon_{12} + \beta\eta_k + \varepsilon_{12}\gamma - 1 + \sqrt{(-\beta\gamma\varepsilon_{12} + \beta\eta_k + \varepsilon_{12}\gamma - 1)^2 + 4\beta\eta_k(1 - \varepsilon)} \bigg),$$
(34)

for $\dot{x}_1 = 0$ and \dot{x}_2 , respectively. These values will coincide if

$$\beta = \frac{\varepsilon_{12}\gamma}{\varepsilon_{12}\gamma - \varepsilon\eta_k},\tag{35}$$

which is greater than 1, positive for $\varepsilon_{12}\gamma > \varepsilon\eta_k$, and negative for $\varepsilon_{12}\gamma < \varepsilon\eta_k$. Notice that the change of sign implies a change in the inequality such that for the considered range of β , it can only be that $x_{1,1}^* < x_{1,2}^*$. This is precisely the reason behind the non-alteration of the dynamics under predatory behaviour. On the vertical axis, the crossings are given by

$$x_{2,1}^* = \frac{1}{2} \left(-\varepsilon_{12}(1-\beta) + \sqrt{\varepsilon_{12}^2(1-\beta)^2 + 4(1-\varepsilon)} \right),$$

$$x_{2,2}^* = 1 - \varepsilon.$$
(36)

In this case, $x_{2,1}^* = x_{2,2}^*$ occurs at the critical value

$$\beta = 1 - \frac{\varepsilon}{\varepsilon_{12}} = \beta_{\text{bif}},\tag{37}$$

and $x_{2,1}^* > x_{2,2}^*$ for $\beta > \beta_{\text{bif}}$, a value that can very well be assumed by β . Thus, the claim follows, and we conclude that a dual behaviour within S_2 concerning its interaction with S_1 does not modify the overall dynamics of the system. As expected, all equilibria are found at values of S_2 just above those studied in the preceding section since S_2 is now replicating from the extinction of S_1 . Moreover, the results found for $\beta = \beta_{\text{bif}}$ are now independent of η_k as the influence of γ on the system makes coexistence impossible under such assumptions (see Figure 11a). Of course, since γ multiplies ε_{12} in \dot{x}_2 , considering $\varepsilon = \varepsilon_{12} = 0$ brings us to the same results shown in Equation (32) and small perturbation of ε_{12} will also result in the S_1 to extinction given $\gamma \varepsilon_{12} > 0$ (see Figure 11b).



Figure 10. Two-parameter bifurcation diagram representing x_1^* as a function of ε_{12} and β plotted as a surface (**a**) and a heat map (**b**) for a more intuitive understanding. Parameters used: $\varepsilon = 0.13$, $\eta_k = 1.4$, $\gamma = 0.5$.



Figure 11. Phase portraits for different cases: (a) $\eta_k = 0$, $\beta = \beta_{\text{bif}} = 0.375$, $\varepsilon = 0.3$, $\varepsilon_{12} = 0.48$, $\gamma = 0.7$; and (b) $\eta_k = 3$, $\beta = 0.6$, $\varepsilon = 0$, $\varepsilon_{12} = 0.48$, $\gamma = 0.2$. As done in all previous plots, the red solid circles denote the ω -limits while the small blue dots mark the initial condition for each orbit.

4. Conclusions

Research on hypercycles has focused on the dynamics arising from the processes of cooperation (e.g., catalysis) between replicators [3,53–56]. The cyclic architecture of cooperative interactions is known to provide stability and survival of all the species. However, other different architectures able to impair such stability have been thoroughly investigated. These include the so-called catalytic parasites [15,57–60] and the catalytic short-circuits [61,62]. Recent research has combined cooperative interactions with other antagonistic processes between species [63]. This is based on the rationale that molecular replicators are subject to mutational processes and ecological species to behavioural shifts due to environmental or ecological changes. These changes could make species switch from cooperative to antagonistic interactions, in, what we have called here, a functional shift. These functional shifts could arise in ribozymes due to mutational processes and could involve a shift from cooperation (hetero-catalysis) to directed degradation (trans-cleavage) [63]. In the context of complex ecosystems, several species undergoing inter-specific cooperation are known to switch to predation (see for example Refs. [65–68]). Even though the drivers of such functional shifts may not be trivial and respond to complex behavioural patterns among species or both biotic and abiotic changes, the dynamics arising due to such functional shifts are still an unexplored subject. As mentioned, some authors have used the hypercycle theory to investigate the impact of these functional shifts in ecosystems dynamics [63]. Specifically, the impact of a shift from catalytic cooperation to directed degradation was studied in small (one to four species) discrete-time hypercycles [63]. As far as we know, no works have investigated how these shifts may affect the stability of small hypercycles in time-continuous systems.

In this article, we have analysed two-species hypercycles considering, together with cooperation, directed degradation or predation (see Figure 1). The changes due to functional shifts have been studied using a general hypercycle model that allows for different architectures by changing different parameter conditions. To understand the changes due to these functional shifts, we have summarised and extended previous results on obligate two-member hypercycles [53] while also analysing a facultative case.

The model considering directed degradation shows no full extinction when both autocatalytic replication rates are higher than their natural decays. We have found a critical value for β , β_{bif} , below which only the dominant species S_2 survives, while coexistence is present for $\beta > \beta_{\text{bif}}$. We have also identified parametric scenarios (i.e., considering no catalytic support of the degrader species by the other one ($\eta_k = 0$) or no decay rate of any of the two species ($\epsilon = 0$)), giving place to a continuum of fixed critical points following a straight line at $\beta = \beta_{\text{bif}}$. For these two scenarios, the complete analytical expressions describing all orbits have been found. Finally, we considered a cooperative system with

predation. Interestingly, bifurcation parameters found for the directed degradation system re-appeared in the same form, stating that no natural benefit towards S_2 is enough to change the dynamics of the system as long as $\eta_k \varepsilon > 0$. However, these extreme cases with $\eta_k = 0$ or $\varepsilon = 0$ stopped presenting a continuum of critical points and turned into the extinction of S_1 as a consequence of the predatory term.

Future research directions may consider functional shifts in non-autonomous systems, considering that changes among cooperative and antagonistic interactions may vary in time due to seasonality or external perturbations.

Author Contributions: Conceptualization, B.B., E.F., D.O., D.A. and J.S.; methodology, B.B., E.F. and J.S.; software, B.B.; validation, B.B., E.F. and J.S; formal analysis, B.B., E.F. and J.S.; investigation, B.B., E.F. and J.S.; resources, J.S.; data curation, B.B.; writing–original draft preparation, B.B., E.F., D.O., D.A. and J.S.; writing—review and editing, B.B., E.F., D.O., D.A. and J.S.; supervision, B.B., E.F., D.O., D.A. and J.S.; project administration, E.F. and J.S; funding acquisition, J.S. All authors have read and agreed to the published version of the manuscript.

Funding: E.F. has been funded by the Spanish Government grant PID2019-104851GB-I00 (MICINN/FEDER, UE) and the Catalan Government grant 2017-SGR-1374. D.O. was funded by the Spanish Government grant CGL2017-85210-P (MICIN/FEDER, UE). D.A. has been partially funded by the Spanish "Ministerio de Ciencia, Innovación y Universidades" under the project CRISIS (PGC2018-096577-B-I00) and the European Regional Development Fund. J.S. has been partially funded by the CERCA Programme of the "Generalitat de Catalunya", by "Agencia Estatal de Investigacíon" grant RTI2018-098322-B-I00, and the "Ramón y Cajal" contract RYC-2017-22243.

Institutional Review Board Statement: Not applicable.

Informed Consent Statement: Not Applicable.

Data Availability Statement: Not Applicable.

Conflicts of Interest: The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript, or in the decision to publish the results.

Appendix A. Obligate Two-Member Hypercycle: Coexistence Equilibrium

Here, we extend previous research for the obligate two-member hypercycle reported in Ref. [53], where numerical results suggested species coexistence via an internal stable node. Here, we show that such a coexistence is also possible through a weak stable focus. Specifically, we compute the stability of the coexistence equilibrium for the asymmetric case of system (2), given by $((\eta_{\varepsilon}/\eta_k)\Gamma_+, \Gamma_+)$. We pay attention to the system's Jacobian Matrix eigenvalues evaluated at such equilibria. Assuming that these two equilibria exist, i.e., $\varepsilon \leq \varepsilon_c = k/(4(1 + \eta_{\varepsilon}/\eta_k))$ (see Section 3.1), the eigenvalues are given by:

$$\lambda_{+,\pm} = \frac{1}{4(\eta_{\varepsilon} + \eta_k)^2} (A \pm B) \tag{A1}$$

for $((\eta_{\varepsilon}/\eta_k)\Gamma_+,\Gamma_+)$ and

$$\lambda_{-,\pm} = \frac{1}{4(\eta_{\varepsilon} + \eta_k)^2} (C \pm D), \tag{A2}$$

for $((\eta_{\varepsilon}/\eta_k)\Gamma_-,\Gamma_-)$, where

$$\begin{split} A &= -\eta_{\varepsilon} \left((k-2\varepsilon)\eta_{k} - 2\eta_{\varepsilon}\varepsilon + \sqrt{((k-4\varepsilon)\eta_{k} - 4\eta_{\varepsilon}\varepsilon)\eta_{k}k} \right) (\eta_{k} + 1), \\ B &= \sqrt{2} \left(\eta_{\varepsilon} \left(\left(-4 \left((\eta_{\varepsilon}/2 + 1)\eta_{k}^{2} + 3\eta_{\varepsilon}\eta_{k} + \eta_{\varepsilon}^{2} + \eta_{\varepsilon}/2 \right) (\eta_{\varepsilon} + \eta_{k})\varepsilon + \eta_{k}k\eta_{\varepsilon}(\eta_{k} + 1)^{2} \right) \right) \\ &\times \sqrt{((k-4\varepsilon)\eta_{k} - 4\eta_{\varepsilon}\varepsilon)\eta_{k}k} + 16 \left((\eta_{\varepsilon}/8 + 1)\eta_{k}^{2} + 9\eta_{\varepsilon}\eta_{k}/4 + \eta_{\varepsilon}^{2} + \eta_{\varepsilon}/8 \right) (\eta_{\varepsilon} + \eta_{k})^{2}\varepsilon^{2}} \\ &- 4\eta_{k}(\eta_{\varepsilon} + \eta_{k})((\eta_{\varepsilon} + 1)\eta_{k}^{2} + 4\eta_{\varepsilon}\eta_{k} + \eta_{\varepsilon}^{2} + \eta_{\varepsilon})k\varepsilon + \eta_{\varepsilon}\eta_{k}^{2}k^{2}(\eta_{k} + 1)^{2} \right) \right)^{1/2}, \\ C &= - \left((k-2\varepsilon)\eta_{k} - 2\eta_{\varepsilon}\varepsilon - \sqrt{((k-4\varepsilon)\eta_{k} - 4\eta_{\varepsilon}\varepsilon)\eta_{k}k} \right) \eta_{\varepsilon}(\eta_{k} + 1), \\ D &= \sqrt{2} \left(\eta_{\varepsilon} \left(\left(4 \left((\eta_{\varepsilon}/2 + 1)\eta_{k}^{2} + 3\eta_{\varepsilon}\eta_{k} + \eta_{\varepsilon}^{2} + \eta_{\varepsilon}/2 \right) (\eta_{\varepsilon} + \eta_{k})\varepsilon - \eta_{k}k\eta_{\varepsilon}(\eta_{k} + 1)^{2} \right) \right) \\ &\times \sqrt{((k-4\varepsilon)\eta_{k} - 4\eta_{\varepsilon}\varepsilon)\eta_{k}k} + 16 \left((\eta_{\varepsilon}/8 + 1)\eta_{k}^{2} + 9\eta_{\varepsilon}\eta_{k}/4 + \eta_{\varepsilon}^{2} + \eta_{\varepsilon}/8 \right) (\eta_{\varepsilon} + \eta_{k})^{2}\varepsilon^{2}} \\ &- 4\eta_{k}(\eta_{\varepsilon} + \eta_{k})((\eta_{\varepsilon} + 1)\eta_{k}^{2} + 4\eta_{\varepsilon}\eta_{k} + \eta_{\varepsilon}^{2} + \eta_{\varepsilon})k\varepsilon + \eta_{\varepsilon}\eta_{k}^{2}k^{2}(\eta_{k} + 1)^{2} \right) \right)^{1/2}. \end{split}$$

Note that, although *A* and *B* may look very similar to *C* and *D*, respectively, these are not the same expression. To establish in which cases these eigenvalues become imaginary, we shall use the fact that the Jacobian Matrix is 2×2 and real, implying that, for example, if $\lambda_{-,+} = a + ib$, then $\lambda_{-,-} = a - ib$. Therefore, we can consider the potential real, *a*, and imaginary times, *i*, *ib*, of these eigenvalues such that:

$$a_{\pm} = \frac{\lambda_{\pm,-} + \lambda_{\pm,+}}{2} \tag{A4}$$

$$ib_{\pm} = \frac{\lambda_{\pm,-} - \lambda_{\pm,+}}{2}.\tag{A5}$$

where $\lambda_{\pm,\pm} = a_{\pm} \pm ib_{\pm}$. We are most interested in the values that the imaginary part of $\lambda_{\pm,\pm}$ can take. To simplify the problem, note that the term $4(\eta_{\varepsilon} + \eta_k)^2 / \sqrt{2\eta_{\varepsilon}}$ can be factored out, leaving a potential imaginary part squared of the form:

$$\left(\frac{4(\eta_{\varepsilon}+\eta_{k})^{2}}{\sqrt{2\eta_{\varepsilon}}}ib_{+}\right)^{2} = 16((\eta_{\varepsilon}/8+1)\eta_{k}^{2}+(9\eta_{\varepsilon}\eta_{k})/4+\eta_{\varepsilon}^{2}+\eta_{\varepsilon}/8)(\eta_{\varepsilon}+\eta_{k})^{2}\varepsilon^{2} -2(\eta_{\varepsilon}+\eta_{k})(2k(\eta_{\varepsilon}+1)\eta_{k}^{3}) +(8k\eta_{\varepsilon}+\sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k})(\eta_{\varepsilon}+2))\eta_{k}^{2} +2\eta_{\varepsilon}(k(\eta_{\varepsilon}+1)+3\sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k})\eta_{k} +\sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k}\eta_{\varepsilon}(2\eta_{\varepsilon}+1))\varepsilon +k\eta_{\varepsilon}\eta_{k}(\eta_{k}+1)^{2}(\eta_{k}k+\sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k}),$$
(A6)

and

$$\left(\frac{4(\eta_{\varepsilon}+\eta_{k})^{2}}{\sqrt{2\eta_{\varepsilon}}}ib_{-}\right)^{2} = 16((\eta_{\varepsilon}/8+1)\eta_{k}^{2}+(9\eta_{\varepsilon}\eta_{k})/4+\eta_{\varepsilon}^{2}+\eta_{\varepsilon}/8)(\eta_{\varepsilon}+\eta_{k})^{2}\varepsilon^{2} + 2(\eta_{\varepsilon}+\eta_{k})(-2k(\eta_{\varepsilon}+1)\eta_{k}^{3} + (-8k\eta_{\varepsilon}+\sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k})(\eta_{\varepsilon}+2))\eta_{k}^{2} + 2\eta_{\varepsilon}(-k(\eta_{\varepsilon}+1)+3\sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k})\eta_{k} + \sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k}\eta_{k} + \sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k}\eta_{\varepsilon} - k\eta_{\varepsilon}\eta_{k}(\eta_{k}+1)^{2}(-\eta_{k}k+\sqrt{((k-4\varepsilon)\eta_{k}-4\eta_{\varepsilon}\varepsilon)\eta_{k}k}).$$

Equalling these two expressions to zero shows that there exist two solutions for $ib_{+} = 0$ and none for $ib_{-} = 0$, stating that for ib_{+} , there is a change in its sign, while ib_{-} stays always with the same one. Notice that omputations to show such zeros are non-double are not included for the sake of length. After plugging any set of acceptable parameters in the second expression, we can see that ib_{-} is, in fact, not an imaginary value. Therefore, it is only necessary to study the eigenvalue sign, which easily shows that the critical point under inspection is a saddle. As for the other equilibrium, we have found two solutions for ε such that the imaginary part changes sign. These are $\xi_{\pm} = (R \pm S)/T$, where

$$R = 8\eta_k k(\eta_\varepsilon + \eta_k) \left(\eta_\varepsilon^2 + \left(\frac{5}{4}\eta_k^2 + \frac{9}{2}\eta_k + \frac{5}{4}\right)\eta_\varepsilon + \eta_k^2 \right), \tag{A8}$$

$$S = 8\eta_k k \sqrt{(\eta_{\varepsilon} - 1)(-\eta_k^2 + \eta_{\varepsilon})(4\eta_{\varepsilon}^2 + (-\eta_k^2 + 6\eta_k - 1)\eta_{\varepsilon} + 4\eta_k^2)},$$
(A9)

$$T = (\eta_k^2 \eta_\varepsilon + 8\eta_\varepsilon^2 + 18\eta_\varepsilon \eta_k + 8\eta_k^2 + \eta_\varepsilon)^2.$$
(A10)

For $\varepsilon \in (\xi_{-}, \xi_{+})$, $(ib_{+})^2$ is a negative expression, indicating that the eigenvalue is indeed imaginary and that, in this region, the analysed equilibrium behaves as a focus. Note that $\xi_{+} \leq \varepsilon_{c}$, thus always defining a region below the critical value of ε in which the stable point is a focus. This result holds for all values of η_{ε} and η_{k} as long as *S* is real, i.e., $(\eta_{\varepsilon} - 1)(-\eta_{k}^{2} + \eta_{\varepsilon}) \geq 0$.

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