

B4. Constrained CR model:

Recall for single species (i), two nutrients (A, B)

$$\begin{array}{l|l|l} \omega_{iA} M_c^{i,A} = -\dot{n}_A & M_c^{i,A} = \gamma_{i,A} M_c^i & \frac{M_c^i}{M_i^i} = \phi_{\max} \left(1 - \frac{\dot{n}_i}{r_c^i}\right) \\ \omega_{iB} M_c^{i,B} = -\dot{n}_B & M_c^{i,B} = \gamma_{i,B} M_c^i & \\ \end{array} \quad \hookrightarrow \sum_{\alpha} \gamma_{i,\alpha} = 1.$$

flux matching: $\omega_{iA} M_c^{i,A} Y_A + \omega_{iB} M_c^{i,B} Y_B = r_i M_i^i$

$$f_c^{i,\alpha} = \omega_{\alpha} Y_{\alpha} \quad f_c^{i,A} \cdot \frac{M_c^{i,A}}{M_i^i} + f_c^{i,B} \frac{M_c^{i,B}}{M_i^i} = r_i$$

$$(f_c^{i,A} \cdot \gamma_{iA} + f_c^{i,B} \gamma_{iB}) \frac{M_c^i}{M_i^i} = r_i$$

$$(f_c^{i,A} \cdot \gamma_{iA} + f_c^{i,B} \gamma_{iB}) \phi_{\max}^i = \frac{r_i}{1 - \frac{r_i}{r_c^i}} \approx r_i$$

MM kinetics of uptake proteins:

$$f_c^{i,\alpha} = f_{c,\max}^{i,\alpha} \frac{n_{\alpha}}{n_{\alpha} + K_{i\alpha}}$$

for small n_{α} , $r_i \approx \underbrace{\frac{f_{c,\max}^{i,A}}{K_{iA}} \phi_{\max}^i \gamma_{iA} n_A}_{V_{iA}^0} + \underbrace{\frac{f_{c,\max}^{i,B}}{K_{iB}} \phi_{\max}^i \gamma_{iB} n_B}_{V_{iB}^0}$

Hypothesis 1: Similar properties for catabolic enzymes for different species in community

(due to, e.g. Chemical limit, HGT)

Hypothesis 2: Similar constraint for different species in community
(i.e., same ϕ_{max} for diff. species)

$$\Rightarrow \nu_{i,d}^o = \nu_d^o \quad (\text{independent of sp } i)$$

$$\Rightarrow \nu_{i,d} = \nu_d^o \cdot \gamma_{i,d}$$

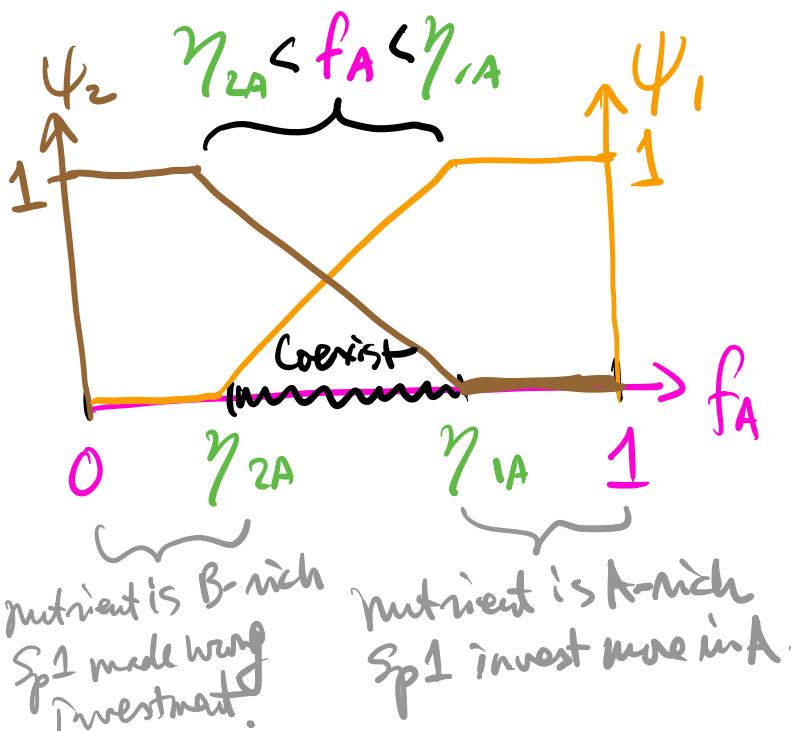
$$\Psi_1 = \frac{f_A}{1 - \frac{\nu_{1B}}{\nu_{2B}}} - \frac{f_B}{\frac{\nu_{1A}}{\nu_{2A}} - 1} = \frac{f_A - \gamma_{2A}}{\gamma_{1A} - \gamma_{2A}}$$

$$\begin{aligned}\gamma_{1B} &= 1 - \gamma_{1A} \\ \gamma_{2B} &= 1 - \gamma_{2A}\end{aligned}$$

Ecological phase diagram

$$\gamma_{1A} > \gamma_{2A}$$

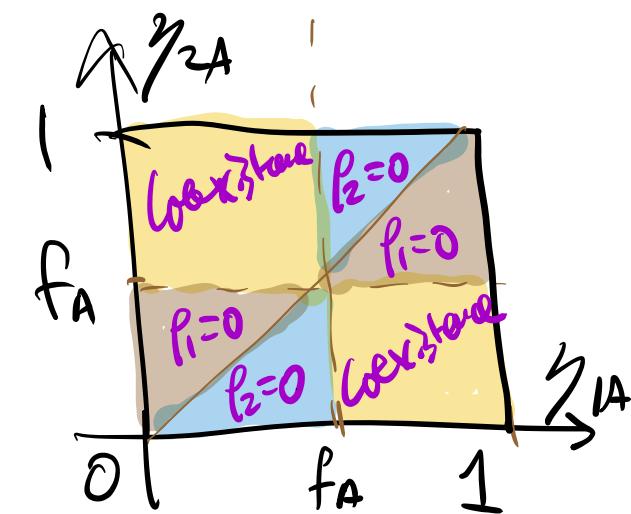
(Sp. 1 invests more in nutrient A than Sp 2)



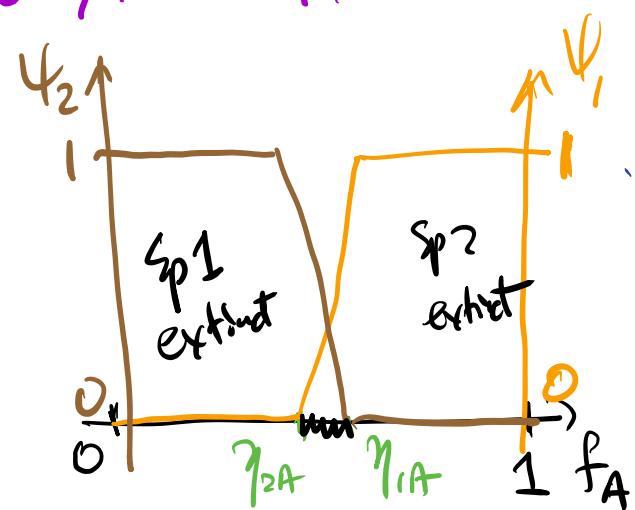
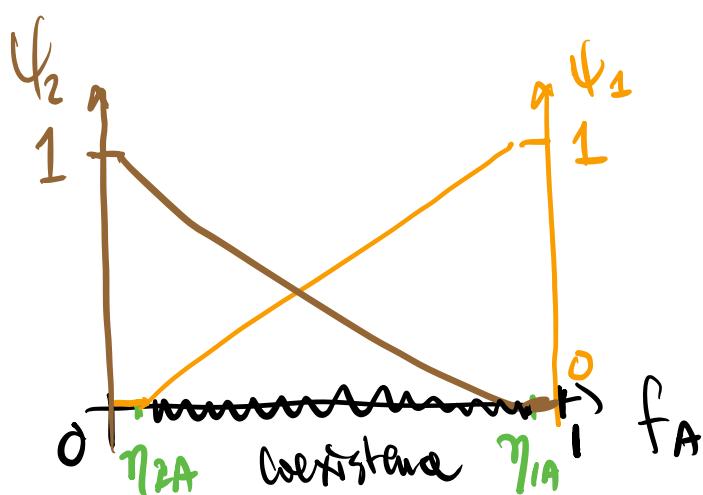
feasibility space (fixed f_A)

$$\gamma_{2A} < \gamma_{1A} : \quad \gamma_{2A} < f_A < \gamma_{1A}$$

$$\gamma_{1A} < \gamma_{2A} : \quad \gamma_{1A} < f_A < \gamma_{2A}$$



- Coexistence favored if $\gamma_{2A} \rightarrow 0, \gamma_{1A} \rightarrow 1$
or $\gamma_{2A} \rightarrow 1, \gamma_{1A} \rightarrow 0$



- Starting from $\gamma_{2A} \approx 0, \gamma_{1A} \approx 1$
if γ_{2A} increases toward f_A , Sp 2 removes Sp 1
or if γ_{1A} decreases toward f_A , Sp 1 removes Sp 2.
 - However, if $\gamma_{1A} \rightarrow f_A + \gamma_{2A} \rightarrow f_A$, then each sp.
risks extinction if f_A fluctuates
- ⇒ Given a distribution of f_A
What is the evol. stable dist of γ_1, γ_2 ?

Extend ecological phase diagram to 3 nutrients (A, B, C)

$$\begin{cases} \dot{S}_i = (\sum_{\alpha} \gamma_{i\alpha} n_{\alpha} - \mu) S_i = (\sum_{\alpha} \gamma_{i\alpha}^0 \eta_{i\alpha} n_{\alpha} - \mu) S_i \\ \dot{n}_{\alpha} = \mu (n_{\alpha}^0 - n_{\alpha}) - \sum_i \gamma_{i\alpha}^0 \eta_{i\alpha} n_{\alpha} S_i / Y_{\alpha} \end{cases}$$

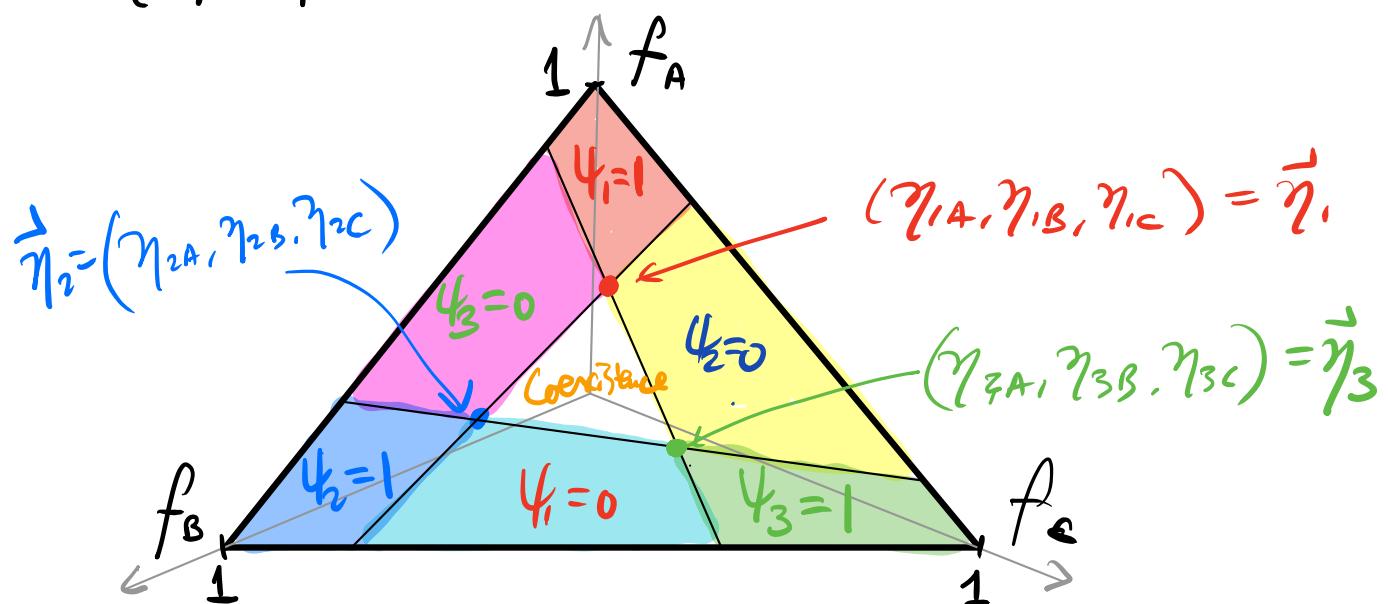
* Consider 3 species P_1, P_2, P_3 with nutrient uptake rates

$$\gamma_{1\alpha} = \gamma_{\alpha}^0 \eta_{1\alpha}, \gamma_{2\alpha} = \gamma_{\alpha}^0 \eta_{2\alpha}, \gamma_{3\alpha} = \gamma_{\alpha}^0 \eta_{3\alpha} \quad (\sum_{\alpha} \eta_{i\alpha} = 1; 6 \text{ independent parameters})$$

Phase space: $f_{\alpha} = n_{\alpha}^0 Y_{\alpha} / \sum_{\alpha} n_{\alpha}^0 Y_{\alpha}$, with $f_A + f_B + f_C = 1$

→ Can represent results succinctly in simplex

(Posfai et al, 2017; working in HW)



- each position in this space represents the value of $\{f_{\alpha}\}$
 - Strain property shown as colored dots ($f_{\alpha} = \eta_{i\alpha}$)
 - Colored regions: phases of partial coexistence
- ⇒ phase boundary obtained simply by
Connecting $\overline{\eta_1 \eta_2}, \overline{\eta_2 \eta_3}, \overline{\eta_3 \eta_1}$

* Important observation by Posfai et al
for the class of constrained CR model $V_{i\alpha} = V_\alpha^0 \gamma_{i\alpha}$

fixed point condition :

$$\dot{p}_i=0 = p_i^* \cdot \left(\sum_\alpha V_\alpha^0 \gamma_{i\alpha} n_\alpha^* - \mu \right)$$

$$\dot{n}_\alpha=0 = \mu(n_\alpha^0 - n_\alpha^*) - \sum_i V_\alpha^0 \gamma_{i\alpha} n_\alpha^* p_i^*/Y_\alpha$$

if $n_\alpha^* = \mu/V_\alpha^0$,

then $\sum_\alpha \gamma_{i\alpha} = 1$ guarantees $\dot{p}_i = 0$ if $p_i^* \neq 0$
for arbitrary # species (even if $N_c > N_R$)

$$\dot{n}_\alpha=0 \rightarrow n_\alpha^0 Y_\alpha = \sum_i \gamma_{i\alpha} p_i^*$$

$$\text{or } \sum_i \gamma_{i\alpha} p_i^* = f_\alpha \quad (\text{feasibility condition})$$

→ can show for each species j whose $\gamma_{j\alpha}$
lies in the white region defined by $\gamma_{1\alpha}, \gamma_{2\alpha}, \gamma_{3\alpha}$

$f_j > 0$, and $\dot{p}_j = 0$ satisfied by $\sum_\alpha \gamma_{j\alpha} = 1$

→ with many species, the coexistence region
is specified by the "convex hull" of all $\gamma_{i\alpha}$

→ 3 "keystone" sp. with $\gamma_{i\alpha} = (1,0,0), (0,1,0), (0,0,1)$

would suffice to support the coexistence
of infinite # of intermediate species.