Anaylsis of the Subfunctionalization Model for the Fate of Gene Duplicates

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• in any of the z regulatory regions of either copy. We assume this occurs at equal Poisson rate u_r for all 2z regions.



• in the coding region of either gene. We assume this occurs at Poisson rate u_c for each gene.





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- After first mutation, drops to *u_c*
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- Rate of Subfunctionalization equals rate of transition to i + 1 equals $(z i)u_r$.



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$$q_{ij} = \begin{cases} 2u_c & \text{if } i = 0, j = P \\ 2zu_r & \text{if } i = 0, j = 1 \\ u_c & \text{if } 1 \le i \le z - 2, j = P \\ (z - i)u_r & \text{if } 1 \le i \le z - 2, j = i + 1 \text{ or } j = S \\ u_r + u_c & \text{if } i = z - 1, j = P \\ u_r & \text{if } i = z - 1, j = S. \end{cases}$$
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The structure of this CTMC is much like those that give rise to the phase-type distribution.



For CTMCs of this structure, it is convenient to write

$$\mathbf{Q} = \begin{bmatrix} \mathbf{Q}^* & \mathbf{V} \\ \hline \mathbf{O} & \mathbf{O} \end{bmatrix},\tag{2}$$

where ${\bm Q}^*$ contains the entries corresponding to transitions between transient states, and ${\bm V}$ transitions to absorbing states.



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Cumulative distribution function

$$\begin{aligned} \overline{F}_i(t) &= \int_{u=0}^t f_i(u) du \\ &= \int_0^t \underline{\mathbf{e}}_i e^{\mathbf{Q}^* u} \mathbf{V} \underline{\mathbf{1}} du \end{aligned}$$

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$$F_i(t) = 1 - \underline{\mathbf{e}_i} e^{\mathbf{Q}^* t} \underline{\mathbf{1}}.$$

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Fate of Gene Duplicates

(4)



With the density and cumulative distribution functions, we're able to derive results for various measures

• Probability of absorption into $j \in \{S, P\}$

$$p_{i,j} = \int_{t=0}^{\infty} \underline{\mathbf{e}_{i}} e^{\mathbf{Q}^{*}t} \mathbf{V}_{j} dt$$
$$= -\underline{\mathbf{e}_{i}} (\mathbf{Q}^{*})^{(-1)} \mathbf{V}_{j}$$
(5)

• The k^{th} moment of time until absorption

$$m_{i}^{(k)} = \int_{t=0}^{\infty} t^{k} \underline{\mathbf{e}}_{i} e^{\mathbf{Q}^{*t}} \mathbf{V} \underline{\mathbf{1}} dt$$
$$= (-1)^{k} k! \underline{\mathbf{e}}_{i} (\mathbf{Q}^{*})^{(-k)} \underline{\mathbf{1}}, \qquad (6)$$

• Variance of time until absorption

$$var_i = m_i^{(2)} - (m_i)^2.$$
 (7)

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$$\lambda_{ij}(t) = \lim_{h \to 0^+} \frac{P(t < T_{\{S,P\}} < t + h, X(T_{\{S,P\}}) = j | T_{\{S,P\}} > t, X(0) = i)}{h}$$
$$= \frac{f_{ij}(t)}{1 - F_i(t)} = \frac{\underline{\mathbf{e}}_i e^{\mathbf{Q}^* t} \mathbf{V}_j}{\underline{\mathbf{e}}_0 e^{\mathbf{Q}^* t} \underline{\mathbf{1}}}$$
(8)

$$f_i(t) = \sum_{j \in \{S,P\}} f_{ij}(t),$$

$$\lambda_i(t) = \sum_{j \in \{S,P\}} \lambda_{ij}(t).$$
(9)





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So
$$\lim_{t\to\infty} \lambda_{ij}$$
 is surely $u_c + u_r$ for $j = P$ u_r for $j = S$.



Some events

• The event that processes has not been absorbed by time t, but is absorbed by later time t + h $A^h - \{t \in T_{table}, t \neq h\}$

$$A_t^n = \{t < T_{\{S,P\}} < t+h\}.$$



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- The event that the process has not entered state z 1 by time t $\overline{C}_t = \{T_{z-1} > t\}$


First, note that

$$\lim_{t\to\infty} P(C_t|B_t, X(0)=i) = 1 \quad \text{and} \quad \lim_{t\to\infty} P(\overline{C}_t|B_t, X(0)=i) = 0,$$



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Now,

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By Markov Property we can drop the t's, and we're left with $q_{z-1,j}$

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(10)

Cause-specific hazard rates





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Something slightly different



We might be interested in the rate of absorption into state P at time t conditional only on not having been absorbed into P.

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We might be interested in the rate of absorption into state P at time t conditional only on not having been absorbed into P. We define the following rate

Pseudogenization rate

$$h_{P}^{z}(t) = \lim_{h \to 0^{+}} \frac{P(t < T_{P} < t + h | T_{P} > t, X(0) = 0)}{h}$$

$$= \frac{f(t)}{1 - F(t)}$$

$$= \frac{\underline{\mathbf{e}}_{0} e^{\mathbf{Q}^{*} t} \mathbf{V}_{P}}{1 - \int_{u=0}^{t} \underline{\mathbf{e}}_{0} e^{\mathbf{Q}^{*} u} \mathbf{V}_{P} du}$$

$$= \frac{\underline{\mathbf{e}}_{0} e^{\mathbf{Q}^{*} t} \underline{\mathbf{V}}_{P}}{1 - \underline{\mathbf{e}}_{0} (e^{\mathbf{Q}^{*} t} - \mathbf{I}) (\mathbf{Q}^{*})^{(-1)} \mathbf{V}_{P}}.$$
 (11)

Here T_P is RV tracking time to absorption into P, and could be infinity.

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Wake up! A picture.





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This motivates us to look into the behaviour of the model in negative time!

Sigmoid Function





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Using l'Hôpital's rule we get

$$\lim_{t \to -\infty} h_P(t; \underline{\alpha}) = \lim_{t \to -\infty} \frac{\underline{\alpha} e^{\mathbf{Q}^*} \mathbf{Q}^* \underline{\mathbf{V}}_P}{-\underline{\alpha} e^{\mathbf{Q}^* t} \mathbf{v}_P}.$$
 (15)

Limit as $\overline{t \to -\infty}$



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Numerator

$$\sum_{k} [\alpha \mathbf{A}^{-1}]_{k} e^{(\lambda_{k} - \lambda_{m})t} \lambda_{k} (\mathbf{A} \mathbf{v}_{p})$$

Which in the limit is just λ_m

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Denomenator

$$-\sum_{l} [\alpha \mathbf{A}^{-1}]_{l} e^{(\lambda_{l} - \lambda_{m})t} \lambda_{l} (\mathbf{A} \mathbf{V}_{p})$$



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Result

$$\lim_{t \to -\infty} h_{\mathcal{P}}(t) = -\lambda_m = \mathcal{S}_{\mathcal{P}}(\mathbf{Q})$$
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We can use this to fit the phenomenological approximation of Tuefel et al (2014) to our exact mechanistically function.

Phenom. vs Exact





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Summary



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- Expanding the model to allow for a mixture of sub- and neofunctionalization.