Substitution Models

Tomáš Vinař November 4, 2021

# Substitution models, notation

 $P(b|a,t)$ : probability that if we start with symbol  $a$ , after time  $t$  we will see symbol  $b$ 

Transition probability matrix:

$$
S(t) = \begin{pmatrix} P(A|A,t) & P(C|A,t) & P(G|A,t) & P(T|A,t) \\ P(A|C,t) & P(C|C,t) & P(G|C,t) & P(T|C,t) \\ P(A|G,t) & P(C|G,t) & P(G|G,t) & P(T|G,t) \\ P(A|T,t) & P(C|T,t) & P(G|T,t) & P(T|T,t) \end{pmatrix}
$$

## Substitution models, basic properties

 $\bullet S(0) = I$ 

$$
\bullet \lim_{t \to \infty} S(t) = \begin{pmatrix} \pi_A & \pi_C & \pi_G & \pi_T \\ \pi_A & \pi_C & \pi_G & \pi_T \\ \pi_A & \pi_C & \pi_G & \pi_T \\ \pi_A & \pi_C & \pi_G & \pi_T \end{pmatrix}
$$

Distribution  $\pi$  is called stationary (equilibrium)

 $\bullet \ \ S(t_1+t_2)=S(t_1)S(t_2) \ \text{(multiplicativity)}$ 

• Jukes-Cantor model should also satisfy

$$
S(t) = \begin{pmatrix} 1 - 3s(t) & s(t) & s(t) & s(t) & s(t) \\ s(t) & 1 - 3s(t) & s(t) & s(t) & s(t) \\ s(t) & s(t) & 1 - 3s(t) & s(t) & s(t) \\ s(t) & s(t) & s(t) & 1 - 3s(t) \end{pmatrix}
$$

$$
S(t) = \begin{pmatrix} 1 - 3s(t) & s(t) & s(t) & s(t) \\ s(t) & 1 - 3s(t) & s(t) & s(t) \\ s(t) & s(t) & 1 - 3s(t) & s(t) \\ s(t) & s(t) & s(t) & 1 - 3s(t) \end{pmatrix}
$$

$$
S(2t) = S(t)^2 =
$$
\n
$$
= \begin{pmatrix}\n1 - 6s(t) + 12s(t)^2 & 2s(t) - 4s(t)^2 & 2s(t) - 4s(t)^2 & 2s(t) - 4s(t)^2 \\
2s(t) - 4s(t)^2 & 1 - 6s(t) + 12s(t)^2 & 2s(t) - 4s(t)^2 & 2s(t) - 4s(t)^2 \\
2s(t) - 4s(t)^2 & 2s(t) - 4s(t)^2 & 1 - 6s(t) + 12s(t)^2 & 2s(t) - 4s(t)^2 \\
2s(t) - 4s(t)^2 & 2s(t) - 4s(t)^2 & 2s(t) - 4s(t)^2 & 1 - 6s(t) + 12s(t)^2\n\end{pmatrix}
$$
\n
$$
\approx \begin{pmatrix}\n1 - 6s(t) & 2s(t) & 2s(t) & 2s(t) \\
2s(t) & 1 - 6s(t) & 2s(t) & 2s(t) \\
2s(t) & 2s(t) & 1 - 6s(t) & 2s(t) \\
2s(t) & 2s(t) & 1 - 6s(t) & 1 - 6s(t)\n\end{pmatrix}
$$

for  $t \to 0$ 

### Substitution rate matrix (matica rýchlostí, matica intenzít)

\n- Substitution rate matrix for Jukes-Cantor model:
\n- \n
$$
R = \begin{pmatrix}\n -3\alpha & \alpha & \alpha & \alpha \\
 \alpha & -3\alpha & \alpha & \alpha \\
 \alpha & \alpha & -3\alpha & \alpha \\
 \alpha & \alpha & \alpha & -3\alpha\n \end{pmatrix}
$$
\n
\n

- $\bullet\,$  For very small  $t$  we have  $S(t)\approx I+Rt$
- $\bullet\,$  Rate  $\alpha$  is the probablity of a change per unit of time for very small  $t$ , or derivative of  $s(t)$  with respect to  $t$  at  $t=0$
- Solving the differential equation for the Jukes-Cantor model <sup>w</sup> e  $\textsf{get}\,\, s(t) = (1-e)$  $\frac{-4\alpha t}{4}$

#### Jukes-Cantor model

$$
S(t) = \begin{pmatrix} (1+3e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 \\ (1-e^{-4\alpha t})/4 & (1+3e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 \\ (1-e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 & (1+3e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 \\ (1-e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 & (1-e^{-4\alpha t})/4 & (1+3e^{-4\alpha t})/4 \end{pmatrix}
$$

The rate matrix is typically normalized so that there is on average one substitution per unit of time, here  $\alpha = 1/3$ 

#### Jukes-Cantor model, summary

- $\bullet\; \; S(t) \colon \mathsf{matrix}\;4 \times 4,$  where  $S(t)_{a,b} = P(b|a,t)$  is the probability that if we start with base  $a$ , after time  $t$  we have base  $b$ .
- $\bullet\,$  Jukes-Cantor model assumes that  $P(b|a,t)$  is the same for all  $a\neq b$
- $\bullet\,$  For a given time  $t$ , off-diagonal elements are  $s(t)$ , diagonal  $1-3s(t)$
- $\bullet\,$  Rate matrix  $R$ : for J-C off-diagonal  $\alpha$ , diagonal  $-3\alpha$
- $\bullet\,$  For very small  $t$  we have  $S(t)\approx I-Rt$
- $\bullet~$  Rate  $\alpha$  is the probablity of a change per unit of time for very small  $t,$ or derivative of  $s(t)$  with respect to  $t$  for  $t=0$
- Solving the differential equation for the Jukes-Cantor model, we get  $s(t)=(1-e)$  $\frac{-4\alpha t}{4}$
- The rate matrix is typically normalized so that there is on average one substitution per unit of time, that is,  $\alpha=1/3$

#### Correction of evolutionary distances

$$
\Pr(X_{t_0+t} = C \,|\, X_{t_0} = A) = \frac{1}{4}(1 - e^{-\frac{4}{3}t})
$$

The expected number of observed changes per base in time  $t$ :  $D(t) = \Pr(X_{t_0+t} \neq X_{t_0}) = \frac{3}{4}(1-e)$ − 4  $\frac{4}{3}t\big)$ 



Branch length (time)

Correction of observed distances

$$
D = \frac{3}{4} \left( 1 - e^{-\frac{4}{3}t} \right) \qquad \Rightarrow \qquad t = -\frac{3}{4} \ln \left( 1 - \frac{4}{3}D \right)
$$

#### More complex models

• General rate matrix  $R$ 

$$
R = \left(\begin{array}{cccc} \cdot & \mu_{AC} & \mu_{AG} & \mu_{AT} \\ \mu_{CA} & \cdot & \mu_{CG} & \mu_{CT} \\ \mu_{GA} & \mu_{GC} & \cdot & \mu_{GT} \\ \mu_{TA} & \mu_{TC} & \mu_{TG} & \cdot \end{array}\right)
$$

•  $\mu_{xy}$  is the rate at which base x changes to a different base y

• Namely, 
$$
\mu_{xy} = \lim_{t \to 0} \frac{\Pr(y \mid x, t)}{t}
$$

- The diagonal is added so that the sum of each row is 0
- There are models with <sup>a</sup> smaller number of parameters (compromise between J-C and an arbitrary matrix)

## Kimura model

- A and G are purines, C and T pyrimidines
- Purines more often change to other purines and pyrimidines to pyrimidines
- Transition: change within group  $A \Leftrightarrow G, C \Leftrightarrow T$ , Transversion: change to a different group  $\{A, G\} \Leftrightarrow \{C, T\}$
- Two parameters: rate of transitions  $\alpha$ , rate of transversions  $\beta$

$$
\bullet \ \ R = \left(\begin{array}{cccc} -2\beta-\alpha & \beta & \alpha & \beta \\ \beta & -2\beta-\alpha & \beta & \alpha \\ \alpha & \beta & -2\beta-\alpha & \beta \\ \beta & \alpha & \beta & -2\beta-\alpha \end{array}\right)
$$

### HKY model (Hasegawa, Kishino, Yano)

- Extension of Kimura model, which allows different probabilities of A, C, G, <sup>T</sup> in the equilibrium
- If we set time to infinity, original base is not important, base frequencies stabilize in an equilibrium.
- Jukes-Cantor has probability of each base in the equilibrium 1/4.
- $\bullet\,$  In HKY the equilibrium frequencies  $\pi_A,\pi_C,\pi_G,\pi_T$  are parameters (summing to 1)
- $\bullet\,$  Parameter  $\kappa\colon$  transition  $/$  transversion ratio  $(\alpha/\beta)$
- Rate matrix: at<br>1

 $\mu_{x,y} =$  $\left\{\right.$  $\begin{matrix} \end{matrix}$  $\kappa\pi_y$  ) if mutation from  $x$  to  $y$  is transition  $\pi_y$  if mutation from  $x$  to  $y$  is transversion

# From rate matrix R to transition probabilities  $S(t)$

- J-C and some other models have explicit formulas for  $S(t)$
- For more complex models, such formulas are not available
- In general,  $S(t) = e^{Rt}$
- Exponential of a matrix A is defined as  $e^A = \sum_{k=0}^{\infty} \frac{1}{k!} A^k$
- If R is diagonalized  $R = UDU^{-1}$ , where D is a diagonal matrix, then  $e^{Rt} = Ue^{Dt}U^{-1}$  and the exponential function is applied to the diagnal elements of  $D$
- Diagonalization always exists for symmetric matrices  $R$ (the diagonal contains eigenvalues)