Algorithms, Games, and Evolution

Theoretical Computer Science Meets other Disciplines

"Nothing in biology makes sense except in the light of evolution."

Theodosius Dobzhansky

Contribution of this paper

Evolution

=

Multiplicative Weight Update Algorithm

applied to

Finding Nash equilibrium of a coordination game

Theoretical Biology

Modelling Evolution

- Population of an infinite number of members
- Each member has a genotype consisting of two genes



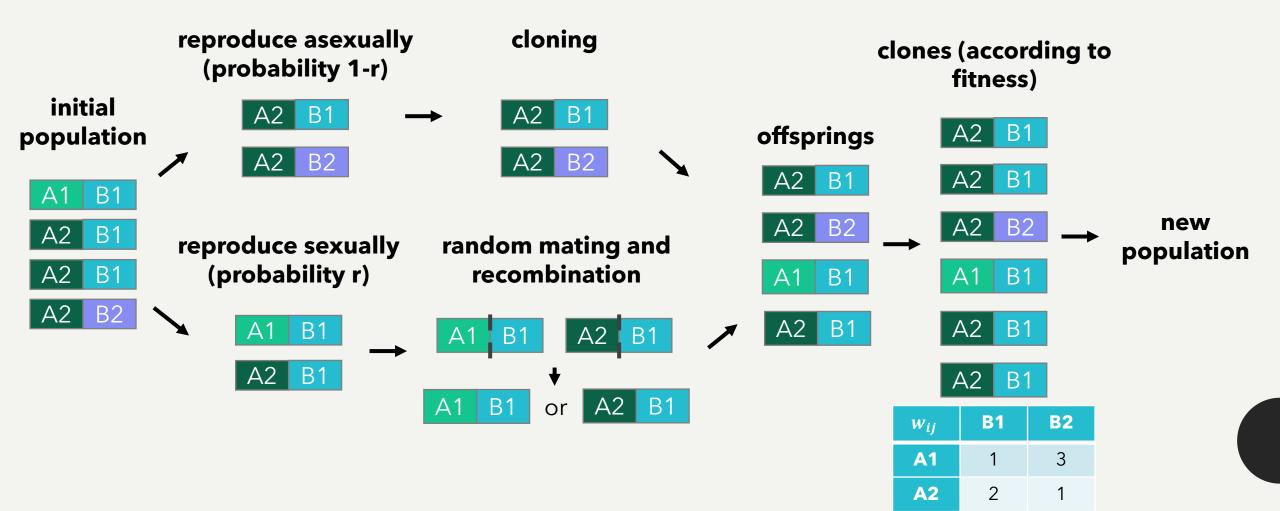
• Two possible alleles per gene



• (Generalisation to an arbitrary number of genes with arbitrarily many alleles is not difficult)

• fitness value of genotypes

w _{ij}	B1	B2
A1	1	3
A2	2	1

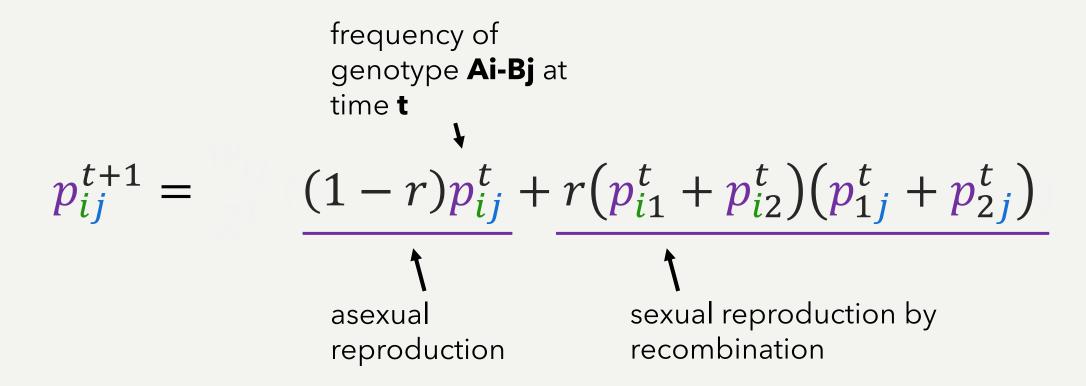


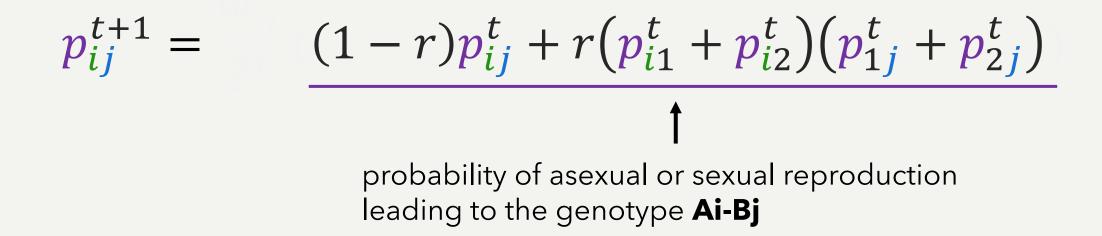
frequency of genotype **Ai-Bj** at time **t+1**

 $p_{ij}^{t+1} =$

frequency of genotype **Ai-Bj** at time **t** $(1 - r)p_{ij}^t$ asexual reproduction

 $p_{ij}^{t+1} =$





fitness of genotype Ai-Bj

(«number of clones of each offspring with this genotype»)

$$p_{ij}^{t+1} = \frac{W_{ij}}{((1-r)p_{ij}^t + r(p_{i1}^t + p_{i2}^t)(p_{1j}^t + p_{2j}^t))}$$

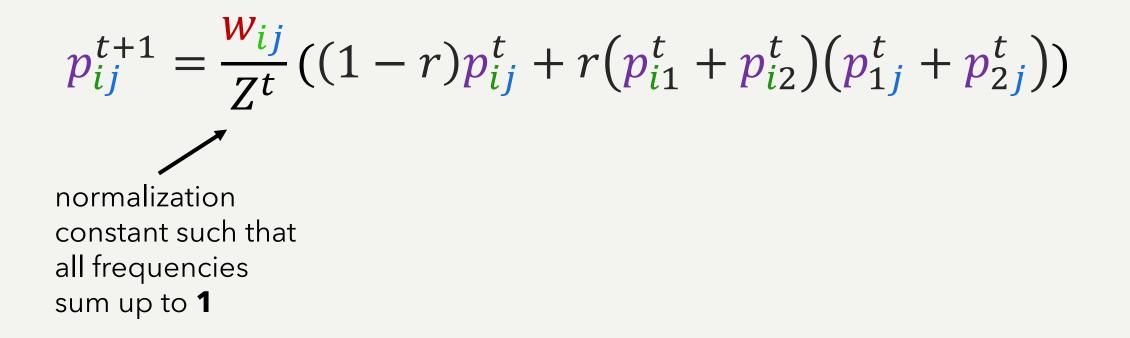
$$\uparrow$$
probability of asexual or sexual reproduction
leading to the genotype **Ai-Bi**

fitness of genotype Ai-Bj

(«number of clones of each offspring with this genotype»)

$$p_{ij}^{t+1} = \frac{W_{ij}}{((1-r)p_{ij}^t + r(p_{i1}^t + p_{i2}^t)(p_{1j}^t + p_{2j}^t))}$$

$$\uparrow$$
Expected number of offsprings with the genotype **Ai-Bj**



Evolution implements MWUA

Evolution

=

Multiplicative Weight Update Algorithm

applied to

Finding Nash equilibrium of a coordination game

Computer Science

Multiplicative Weight Update Algorithm

MWUA

- Simple and general algorithm used in many different areas:
- Machine Learning (e.g. AdaBoost)
- Approximations for NP-hard problems
- Solving linear programs
- Finding Nash equilibria for some types of games

Game Theory

MWUA and Coordination Games

Coordination Games

- Two players A and B play a game
- Reward for player A = reward for player B
- \rightarrow They strive to reach the same goal!

Two friends decided to go to a concert together, but they both forgot if they had agreed to see **The Chainsmokers** or **Adele**. They can't communicate until they reach the location.

	The Chainsmokers	Adele
The Chainsmokers	2/2	0 / 0
Adele	0 / 0	1 / 1

Coordination Games

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ocation.		The Chainsmokers	Adele
	The Chainsmokers	2/2	0 / 0
	Adele	0 / 0	1 / 1

- How to find the perfect action? (knowing all rewards)
- (Besides the obvious way...)

0

• They can use the **multiplicative weight update algorithm (MWUA)**

- We start with any strategies $\boldsymbol{p_A^0}$ and $\boldsymbol{p_B^0}$
- Apply iteratively:

$$p_A^{t+1}(i) =$$

probability of player **A** playing
action **i** following the strategy p_A^{t+1}

- We start with any strategies $\boldsymbol{p_A}^0$ and $\boldsymbol{p_B}^0$
- Apply iteratively:

$$p_{A}^{t+1}(i) = p_{A}^{t}(i)$$

$$\uparrow$$
probability of player **A** playing
action **i** following the strategy **P**_A^t

- We start with any strategies $\boldsymbol{p_A}^0$ and $\boldsymbol{p_B}^0$
- Apply iteratively:

$$p_{A}^{t+1}(i) = p_{A}^{t}(i) (\uparrow$$

- We start with any strategies $\boldsymbol{p_A^0}$ and $\boldsymbol{p_B^0}$
- Apply iteratively:

$$p_{A}^{t+1}(i) = p_{A}^{t}(i)(1 + \mathbb{E}_{j \sim p_{B}^{t}}[r(i,j)|i])$$

$$(t) = p_{A}^{t}(i)(1 + \mathbb{E}_{j \sim p_{B}$$

- We start with any strategies $\boldsymbol{p_A^0}$ and $\boldsymbol{p_B^0}$
- Apply iteratively:

$$p_{A}^{t+1}(\mathbf{i}) = p_{A}^{t}(\mathbf{i})(1 + \epsilon \cdot \mathbb{E}_{j \sim p_{B}^{t}}[r(\mathbf{i}, j)|\mathbf{i}])$$

$$\mathbf{1}$$
small positive **«learning rate»**

Given reward function $\mathbf{r} \rightarrow \text{find the optimal strategies } \mathbf{p}_{\mathbf{A}} \text{ and } \mathbf{p}_{\mathbf{B}}$

- We start with any strategies $\boldsymbol{p_A}^0$ and $\boldsymbol{p_B}^0$
- Apply iteratively:

r

$$p_A^{t+1}(\mathbf{i}) = \frac{1}{Z^t} p_A^t(\mathbf{i}) (1 + \epsilon \cdot \mathbb{E}_{j \sim p_B^t}[r(\mathbf{i}, j) | \mathbf{i}])$$

$$\mathbf{k}$$
normalization constant such that \mathbf{p}_A^{t+1} is a

probability distribution

$$MWUA for games$$
$$p_A^{t+1}(i) = \frac{1}{Z^t} p_A^t(i) (1 + \epsilon \cdot \mathbb{E}_{j \sim p_B^t}[r(i, j)|i])$$

Theorem

 p_A^t and p_B^t converge to a nash equilibrium.

But why is this relevant for evolution?

Evolution

$$p_{ij}^{t+1} = \frac{w_{ij}}{Z^t} \left((1-r)p_{ij}^t + r(p_{i1}^t + p_{i2}^t)(p_{1j}^t + p_{2j}^t) \right)$$

MWUA for coordination games

$$p_A^{t+1}(\mathbf{i}) = \frac{1}{Z^t} p_A^t(\mathbf{i}) (1 + \epsilon \cdot \mathbb{E}_{j \sim p_B^t}[r(\mathbf{i}, j) | \mathbf{i}])$$

Not quite there!

Theoretical Biology

Modelling Evolution (again)

Weak Selection

 $\forall i, j. \ w_{ij} \in [1 - s, 1 + s]$

for a small **selection strength** *s* > 0

$$\Rightarrow w_{ij} = 1 + s \cdot \Delta_{ij}, \ \Delta_{ij} \in [-1, 1]$$

Weak Selection

assumption of weak selection

 $w_{ij} = 1 + s \cdot \Delta_{ij}, \qquad \Delta_{ij} \in [-1, 1]$

allele frequencies

$$p_A^{t+1}(i) \approx \frac{1}{Z^t} p_A^t(i) (1 + s \cdot \mathbb{E}_{j \sim p_B^t}[\Delta_{ij}|i])$$

probability of **allele i**

at **gene A** at **time t+1**

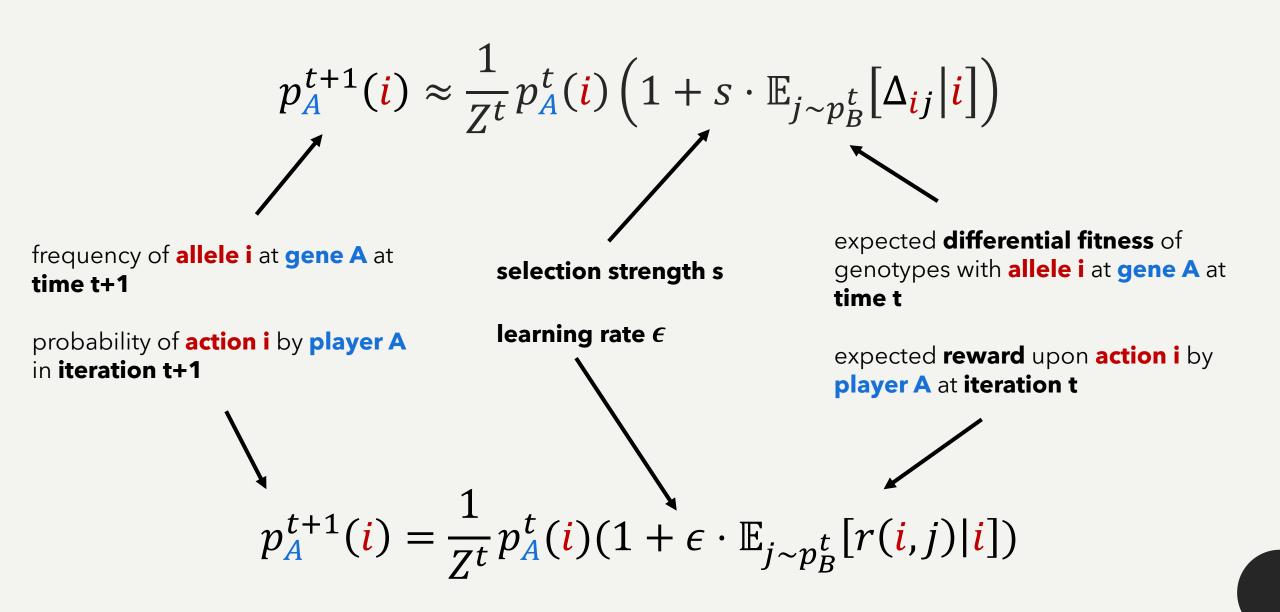
Evolution

$$p_A^{t+1}(i) \approx \frac{1}{Z^t} p_A^t(i) (1 + s \cdot \mathbb{E}_{j \sim p_B^t}[\Delta_{ij}|i])$$

MWUA for coordination games

$$p_A^{t+1}(i) = \frac{1}{Z^t} p_A^t(i) (1 + \epsilon \cdot \mathbb{E}_{j \sim p_B^t}[r(i,j)|i])$$

Looks similar?



Evolution implements MWUA

Evolution «implements» the MWU-algorithm

- Genes are the players
- The alleles are the actions
- The allele frequencies are the strategies
- The reward is the differential fitness

What we learned

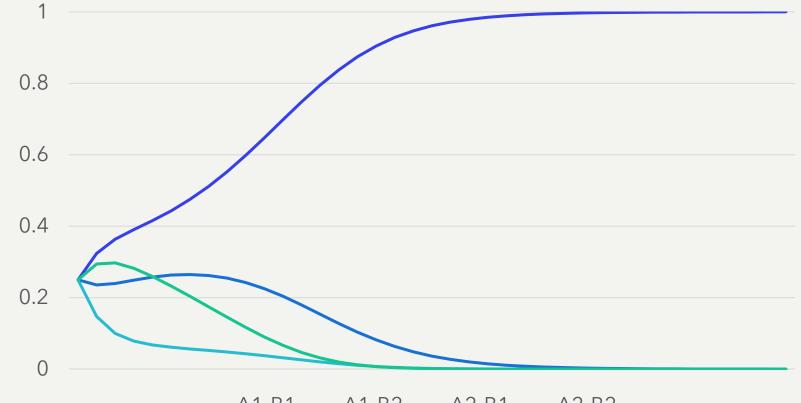
- Evolution under weak selection has the same behavior as the MWUA for coordination games
- Until convergence, there is a trade-off between fitness and diversity
- Link between Theoretical Biology, Game Theory, Theoretical Computer Science, and Information Theory!

Thank you for your attention!

Information Theory

Source of Diversity

w _{ij}	B1	B2
A1	1.1	0.8
A2	0.5	1.0



$$p_{ij}^{t+1} = \frac{w_{ij}}{Z^t} \left((1-r)p_{ij}^t + r(p_{i1}^t + p_{i2}^t)(p_{1j}^t + p_{2j}^t) \right)$$

There's more!

• Weak selection maximizes the following quantity:

$$x_{A}^{t}(1)\sum_{\tau=0}^{t} \mathbb{E}_{j\sim x_{B}^{\tau}}[\Delta_{1j}] + x_{A}^{t}(2)\sum_{\tau=0}^{t} \mathbb{E}_{j\sim x_{B}^{\tau}}[\Delta_{2j}] + \frac{1}{s}H(x_{A}^{t})$$

cumulative expected differential fitness

entropy of the allele distribution (high entropy equals high uncertainty)

There's more!

- Allele frequencies will eventually collapse
- Until collapse, allele distributions with high uncertainty are preferred
- This possibly explains the huge diversity in nature:
- Trade-off between fitness and diversity until convergence!

Weak Selection

$$\forall i, j. \ w_{ij} \in [1 - s, 1 + s]$$

for a small **selection strength** *s* > 0

 $\Rightarrow w_{ij} = 1 + s \cdot \Delta_{ij}, \ \Delta_{ij} \in [-1, 1]$

Theorem (Nagylaki)

Under weak selection, the frequency p_{ij}^t of a genotype **Ai-Bj** can be approximated as a product of the marginal frequencies of the two alleles:

$$p_{ij}^{t} \approx \left(p_{i1}^{t} + p_{i2}^{t}\right) \cdot \left(p_{1j}^{t} + p_{2j}^{t}\right) = p_{A}^{t}(i) \cdot p_{B}^{t}(j)$$

The error of this approximation is in O(s).

probability of **allele i** at **gene A** at **time t**

$$w_{ij} \in [1 - s, 1 + s]$$

$$p_{ij}^t \approx \left(p_{i1}^t + p_{i2}^t\right) \cdot \left(p_{1j}^t + p_{2j}^t\right) = p_A^t(i) \cdot p_B^t(j)$$

Weak Selection

$$\begin{split} p_{ij}^{t+1} &= \frac{w_{ij}}{Z^t} \Big((1-r) p_{ij}^t + r \big(p_{i1}^t + p_{i2}^t \big) \big(p_{1j}^t + p_{2j}^t \big) \Big) \\ &\approx \frac{w_{ij}}{Z^t} \Big((1-r) \cdot p_A^t(i) \cdot p_B^t(j) + r \cdot p_A^t(i) \cdot p_B^t(j) \Big) \\ &= \frac{w_{ij}}{Z^t} \cdot p_A^t(i) \cdot p_B^t(j) \\ &= \frac{1+s \cdot \Delta_{ij}}{Z^t} \cdot p_A^t(i) \cdot p_B^t(j) & \stackrel{(\Delta_{ij} \in [-1,1] = \text{differential fitness}}{\text{of genotype Ai-Bj}} \end{split}$$

$$p_{ij}^{t+1} \approx \frac{1 + s \cdot \Delta_{ij}}{Z^t} \cdot x_A^t(i) \cdot x_B^t(j)$$

$$\begin{aligned} Allele frequencies \\ p_A^{t+1}(i) &= p_{i1}^{t+1} + p_{i2}^{t+1} \\ &\approx \frac{1+s \cdot \Delta_{i1}}{Z^t} \cdot p_A^t(i) \cdot p_B^t(1) + \frac{1+s \cdot \Delta_{i2}}{Z^t} \cdot p_A^t(i) \cdot p_B^t(2) \\ &= \frac{1}{Z^t} \cdot p_A^t(i) \cdot (p_B^t(1) + p_B^t(2) + s \cdot (\Delta_{i1} p_B^t(1) + \Delta_{i2} p_B^t(2))) \\ &= \frac{1}{Z^t} p_A^t(i)(1+s \cdot \mathbb{E}_{j \sim p_B^t}[\Delta_{ij}|i]) \\ & (\mathbb{E}_{j \sim p_B^t}[\Delta_{ij}|i] = \text{expected} \\ & \text{differential fitness of a} \\ & \text{genotype with allele i for gene A} \\ & \text{at time t} \end{aligned}$$

$$weak$$
Selection
$$w_{ij} \in [1 - s, 1 + s]$$

$$p_{ij}^{t} \approx p_{A}^{t}(i) \cdot p_{B}^{t}(j)$$
allele frequencies
$$p_{A}^{t+1}(i) \approx \frac{1}{Z^{t}} p_{A}^{t}(i)(1 + s \cdot \mathbb{E}_{j \sim p_{B}^{t}}[\Delta_{ij}|i])$$

$$p_{B}^{t+1}(j) \approx \frac{1}{Z^{t}} p_{B}^{t}(j)(1 + s \cdot \mathbb{E}_{i \sim p_{A}^{t}}[\Delta_{ij}|j])$$