## Collaborative solving in a human computing game

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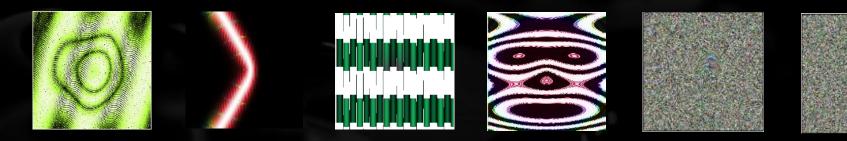
McGill School of Computer Science Seminar, October 11th

## Outline

- Introduction to human computing
- Collaborative solving with a market, skills and challenges

- Human computing is about combining the strengths of the human brain and those of the computer
- Can be used to solve specific problems
  - Perception (image labeling, sound recognition)
  - Knowledge (collecting common-sense facts)
  - Reasoning (puzzle solving)
  - Coordination (robotics)

 Deep Neural Networks are Easily Fooled (Nguyen *et al.* 2014)



 Deep Neural Networks are Easily Fooled (Nguyen *et al.* 2014)



#### Confidence score > 99%

 Typically, participants (usually called "workers") perform the task voluntarily or in exchange of money

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 Mechanical
 Turk :

amazon mechanical turk					<u>Sign I</u>	
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## Human computing games

- It is also possible to embed the tasks into games
- Human computing games (games with a purpose):
  - ESP
  - Foldit
  - Phylo
  - Ribo

## Human brain power available

Top	Games By Current F				Next page
	Name	Current Players	Last 30 Days	Peak Players	Hours Played
	Dota 2	531,393			
2.	Counter-Strike: Global Offensive	325,786			
3.	Sid Meier's Civilization V	47,907			
4.	Team Fortress 2	43,942			
	Rocket League	42,995			
	Grand Theft Auto V	38,181			
	ARK: Survival Evolved	35,715			
8.	Rust	34,233			
	Garry's Mod	29,275			
10	Paladins	27,434			
11	Mafia III	26,180			
12	Football Manager 2016	25,550			
13	Arma 3	24,770			
14	The Elder Scrolls V: Skyrim	22,523			
15	Fallout 4	22,051			
16	Warframe	21,586			
17	Unturned	21,162			
18	H1Z1: King of the Kill	18,361			
19	Terraria	14,783			
20	Path of Exile	14,727			

In the last 30 days, for the 20 most popular PC games on Steam alone:

- At least 2,586,602 distinct players in total
- A total of 995,605,427 hours played

## Human brain power available

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995,605,427 hours played in 30 days x 12 months =

#### http://steamcharts.com/

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995,605,427 hours played in 30 days Х 12 months 11,947,265,124 12 billion hours

per year of brain power available

http://steamcharts.com/

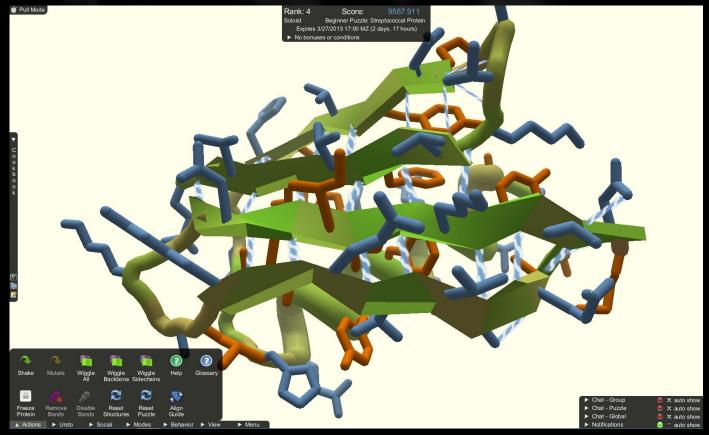
## ESP game

0	0	R The ESP Game - Neticape			
		O:11 Tame Left	The I	ESP Game	2 100
				Taboo Words MAN BEARD	Your Guesses HAT
Player 1 guesses: purse Player 1 guesses: bag Player 1 guesses: brown	Player 2 guesses: handbag	Type your	r next gues	st	
Success! Agreement on "purse"	Player 2 guesses: purse Success! Agreement on "purse"	a 1000	(18) Caraga Hah	n inicality at 1920 second Pasi	P.P.otra

#### "It turns out that the string on which the two players agree is typically a good label for the image"

## Foldit

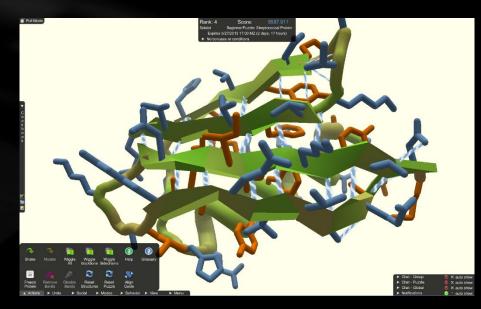
 First human computing game on molecular biology



http://fold.it/portal/

## Foldit

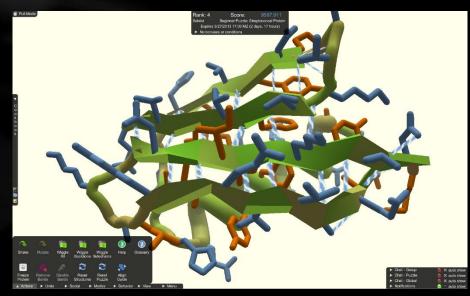
 Has been used to solve the crystal structure of a retroviral protease (Khatib *et al.* 2011)



13

## Foldit

- Has been used to solve the crystal structure of a retroviral protease (Khatib *et al.* 2011)
- It is not a casual game:
  - Complex interface
  - Requires a certain level of scientific knowledge



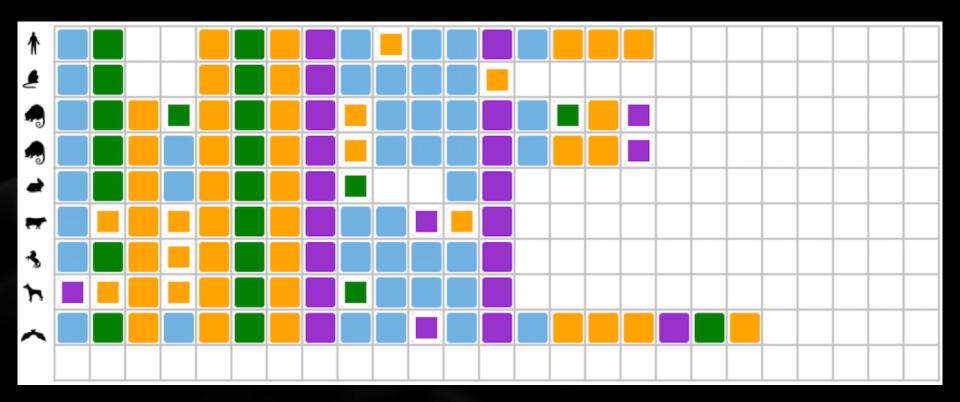
## Phylo

- In 2010, Phylo was released
- It is tackling a fundamental problem in comparative genomics: Multiple Sequence Alignment (MSA)
- It converts the MSA problem into a casual game that can be played by ordinary web users

### **Comparative Genomics - MSA**

Gaps human getcaaccaaccaacaatttctttaeca<mark>acgt</mark>gaataactaattattegtctaecaaccattte ΤG TG baboon GGTCAACGAGCAAGAATTTCTTTAGCAAGGTGAA--+-TAATTATTGGTCTAACAAGCATTTG macaque GGTCAACGAGCAAGAATTTCTTTAGCAAGGTGAA---+-TAATTATTGGTCTAACAAGCA ТG vervet GGTCAACGAGCAAGAATTTCTTTAGCAAGGTGAA----TAATTATTAGTCTAACAAGCATT dusky titi ggtcaacgaggaagaatttctttagga<mark>aggt</mark>gaataactgattattggtctaggaagdat TG тG mouse lemur getcaaccaecaacaatttctttaccaaccatcaataatccattattgctccactcaccatttc 1emur GGTCAACGAGCAAGAATTTCTTTAGCAAGGTGAATAACGGATTA TG TTGGTCCAGTGAGCA CA 1000 GOT CACCOACCAACAATCTCTTTACCAACCTCAATAATCCCCCATACACTCCACTCCCACTCCCCCTTTC rabbit GGTCAACGAGCAAGAATTTCTTTAGCAAGGTGAGTATCTGATTATTGGTCTACCAAGCAT TG PAT GGTCAACGTGCAAGAATTTCTTTAGCAAGGTAAACGTTCAACTGTTGGTTTGCTGAGAA CTTG MOUSE GGTCAGCGTGCAAGGATTTCTTTAGCAAGGTAAATAT TTAACTG GIGAGCA TG TTGGTCTT COM GETCAGEGAGEGAGAATTTETTAGEAAGGTGAATATETGEETA TGGGTT CAGCAAGCA Sheep GGTCAGCGAGCAAGAATTTCTTTAGCAAGGTAAATATCTGCTTA TEGTCCAGCAAGCA TG TG pic GETCACCEACEARGARTTTCTTACCA<mark>ACCT</mark>CAATATCACTTATTGGTCCACCAACCG TG horse GGTCAGCGAGCAAGGATTTCTTTAGCAAGGTGAATAGCTGATT TEGTTCAGTGAGCC TTG CAT GETCAGEGAGEAAGAATTTEETTAGEAAGGTGAATATETEATTAGET CAGTGAGC TTG dog GGGCAAAGAGGAAGAATTTCCCTAGCA<mark>AGGT</mark>GAATATCCCGACGATTGGTTTGGCGAGGA GG TG CEPAT GEOCACCOACCOACAATTTCTTACCAACGTCAATATCTCATTATTCGT-CACTCACCA TTG hedgehoo GGTCAACGAGCAAGAATTTCATTAGCAAGGTGAATA AGCATTTG armad 110 GGTCAACGAGCAAGAATTTCTTTAGCAAGGT-ATTATCTCACTATTGGACCAGTTGACA CTTG TG monode 1ph is GGTCAACGAGCAAGAATTTCTTTAGCAAGGTTAATATTTTGGTA GTT CAG ΤG Wallaby GGTCAACGAGCAAGAATTTCTTTAGCAAGGTTAATATTTTGGTAT GTT CAGT TGARA dunnart GGTCAACGAGCAAGAATTTCCTTAGCAAGGTTAATA TTTTGATAT GT A G platypus GGTCAGCGGGCCAGAATTTCATTAGCCAGGTGAGTA TCAGGTGGCGTTTG chicken GGCCAGCGAGCACGAATCTCACTAGCGAGGTGAGCATTTTGCTAT torto ise GGTCAACGGGCTAGAATCTCACTAGCTAGGTGAATATTTTATCAT zeprafish GGTCAGAAGGCACGCGTGGCGTCTGGCCAGGT--ATG+TCACAC ---TTTCACAG tetra GGTCAGAGGGGCACGTCTGGGTTTGGGCC<mark>AGGT</mark>--ACT+TCTCTCACAC ---CTTCAGCACACC FUSU GGTCAAAGGGCACGCCTGGGTTTGGCCCAGGT-ACT+TCTGCCACAC ---TTCAGCGCACC ю

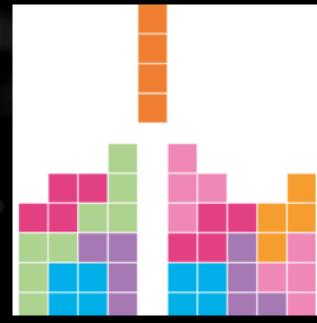
## Phylo DNA Puzzles



Turn the multiple sequence alignment problem into a casual tile-matching game.

## The 3 Pillars of Phylo

Why? Fundamental problem in molecular biology.
What? Well-defined problem proven difficult for computers.
How? Humans are good at matching colors.



http://thegoddamn90s.com

#### Whole-genome multiple alignment calculated with computers

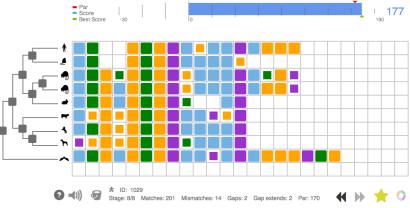
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с	GCC C C	C GCCC GCA	AGGC AGAA G A CA C AGA G CCAA	A CAGAAAG G CAG CCA CCA ACCACC AGGAGC A CAA G GGGC GCAGC A GGA C	AAA
	G CACAA	AGGAAG GCCAC GGCC	C AGAG GG AGAA GA CAGGGA GC AA AA CA CCCACO	CA CC ACAG GC CAGAACAGCACCCC ACCC CACCCCA CAACAAAGAA A CCAGCCCAAAAA GCCAA A G GCCC	CAGA
CA	G CACAG	AGGGGG AC AC GGCA	C A CG GG GGAG GA AGGGA AC GA AA C	A C ACAG GCACAGGAC <mark>AG ACCCC AC CACCCCACAA CAAAGAA A CCAGCCCAAAA GCCAACA G GC C</mark>	CAGA
G	G CACAA	GGGGGA AC AC GGCA	C AA G GG AGAG GA CAGGGA AC GA AA	GC ACAG GCACAGGACAGCACCC A C CACCCCAAAAGCAAAG A A CCAGCCCCAAA GCCAA G G GC C	CAGA
GEHOG	G CA AG	GA A A GGGC	C AG A GACAAAGAAA AAGA G C GG AG C	A C GC CCA A GA AGCAC CCCA C CAC CCAAAA AAGAG CA CA AC CAG G GCCAA A G GCCC	CAGA
	G CACAG	GGAGGA G AC GACA	C AGAG AG AGAC AAAGA AC GA AG C	ACCCCA G GCAC C CCAACAA AA GGC CA CGAAACC AAA GCCAA C GCCAA A G C C	CAG
A	G CACAA	GGAGGA G AC GGCA	C AGAG AG AGAC AAGGACAC GA AA C	A AC A GC GCAC CCAACAA AA GGC CA C AGACC AAA ACCAA C GCCAA A A CC	CAG
A	A CACAA	GGGGAACACCAC GGCA	C CGGG AGCAGGC CAGGCA GC GG AA	A AC ACAG GCACAG ACAG CCCCACA CCCGCACCAACAACA GG A GC GCCCAAAG GCCAG G GC CC	CACG
	A CACAA	GGGGG GCCACGG CC	C CAG G GG AGAG AA CAGGGAGGC GA AACC	ACCC GCAG GCACAGGGCAG GCC CCAC CCCACCACAACAA GGAGAA A GGGCCCCCAAA GCCAA A G GCCC	CAAG
	A CACAG	GGGGGA GCCAC GGCC	C AAG G GG AGAG AA CAGGGAGGC GAAAACC	ACCC GCAGAGCACGGGGCAG GCC CACCACCAC CCAACAACGGAGAA A GGG CCCAAA GCCAA A G GCCC	CAGG
	G CAGAA	GGGGGA GC C GGC C	AC G GG AGAG AAACAGGGA GC A AA C	A CC ACAG GCACAGGACAG ACCCCCCCCCCCCCCCC	
ACA	G CAGAA	GGGGGA GC C GGC C	AC G GG AGAG AAACAGGAA GC A AA C	A CC ACAG GCACAGG CAG ACCCCCCCCCCCCCCCC	
A	G CAGAA	GGGGGA GC C GGC C	AC G GG AGAA AAACAGGGA GC A AA C	A CC ACAG GCACAGGACAG ACCCCCCCCCCCCCCCC	
ANGUTA	G CACGA	GGGAGA GC C GGC C	GAC G GG AGAG AAGCGGGGA GC A AA C	A CCAACAG GCACAGGACAG ACCCCCCACCCCACAC CCAG AA GAAGAA CAC GGACCCAAAA G AA G G G CC	CAGG
GORILLA	G CACGA	GGGGGA GC C GGC C	A AC G GG AGAG AAG GGGGA GC A AC C	A CC ACAG GCACAGGACAG ACCCCCACCCACAC CCAG AA GAAGAA CA AGACCGAAAA G AA G G G CC	CAGG
С	G CACGA	GGGGGA GC C GGC C	A AC G GG AGAG AAGCGGGGA GC A AA C	A CC ACAG GCACAGGACAG ACCCCCACCCACAC CCAG AA GAAGAA CA AGACCGAAAA G AA G G G CC	CAGA
A	G CACGA	GGGGGA GC C GGC C	A AC G GG AGAG AAGCGGGGA GC A AA C	A CC ACAG GCACAGGAC <mark>A</mark> G ACCCCCACCACCACAC CCAG AA GAAGAA CA AGACC AAAA G AA G G G CC	

## Extract dubious alignment region

CATAGAG-CC-CAGGGGT	GCTGC!
ACAAAGAATTAACTAGCCC	CAA TAA'
GTT	TGA:
GTCTTAAAATGCACAGTGTAGCC( CCC	TGA
ATCAGAAAGTGTTCAG CCCC	TGG
ATCCTACAGTGCTCAGAACAGCA(	TGA!
ATTCTACAGTGCACAGGACAGTA( CTC	TGA!
GCTTTACAGTGCACAGGACAGCA( ACC	TGG!
ATTCTGCTTTCCATATGATAGCA( ACC	TGA!
ACCCCATTGTGCACCCCC	TGA
ATACTATGCTGCACCCCC	TTA:
ATACTACAGTGCACAGTACAGTT( TCC	TTA
ACCCTGCAGTGCACAGGGCAGTG( CCC	TTA
ACCCTGCAGAGCACGGGGCAGTG	TTA
ATCCTACAGTGCACAGGACAGTA(	TTA
ATCCTACAGTGCACAGGTCAGTA(	TTA
ATCCTACAGTGCACAGGACAGTA( CCG	TTA
ATCCAACAGTGCACAGGACAGTA( CCG	
ATCCTACAGTGCACAGGACAGTA( CCT)	
ATCCTACAGTGCACAGGACAGTA(	
ATCCTACAGTGCACAGGACAGTA(	



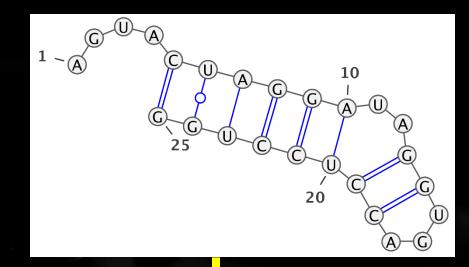
#### Reinsertion into original alignment + Evaluation



#### Database of interesting puzzles

http://phylo.cs.mcgill.ca 19

## **RNA sequence & structure**

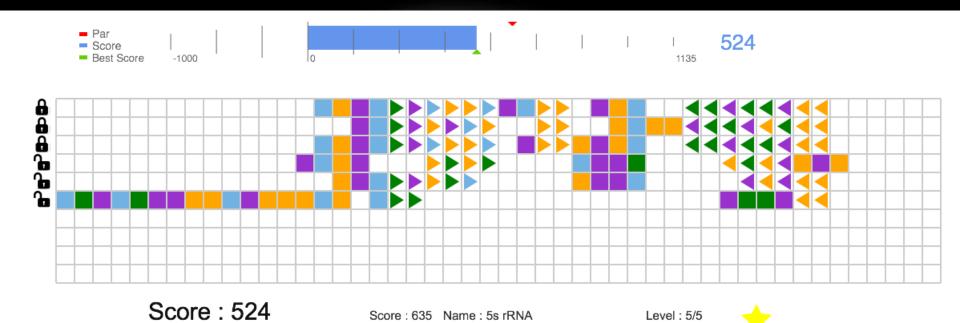


2D structure:

# Bracket notation:

Game representation:

## Ribo



http://ribo.cs.mcgill.ca

## Still not convinced about the potential of human computing?

## Still not convinced about the potential of human computing?



- Using a human computing game to solve a problem that has a large search space is not straightforward:
- 1. Overwhelming for a single player
- 2. Impossible to find an optimal solution without considering all the data

• How to deal with a large search space?

- How to deal with a large search space?
- A. Decompose the problem, distribute small tasks, aggregate the answers (AMT, Crowdcrafting):
  - No interactions
  - Trying to limit groupthink
  - Cannot benefit from collective intelligence

- How to deal with a large search space?
- Build a collaborative A. Decompose the problem, B. distribute small tasks, environment: aggregate the answers Interactions allowed but (AMT, Crowdcrafting): controlled
  - No interactions
  - Trying to limit groupthink
  - Cannot benefit from collective intelligence

- Promote cooperation
- Allow exchange of information
- Improve the solutions of others

- Before transitioning to collaborative models in games, we need to:
  - Estimate the potential gains in productivity
  - Quantify the usefulness of different mechanisms

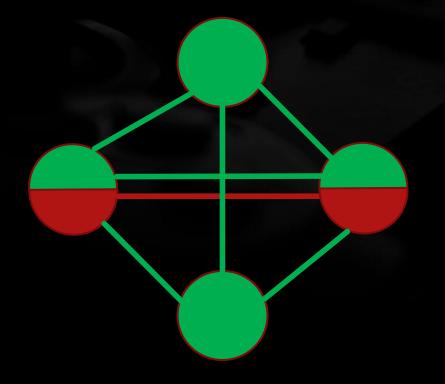
- Before transitioning to collaborative models in games, we need to:
  - Estimate the potential gains in productivity
  - Quantify the usefulness of different mechanisms
- We propose a formal framework to study human collaborative solving in a video game with:
  - A market
  - Skills
  - Challenges

## Hypotheses

- 1. A market system will help players build longer solutions
- 2. Skills are useful to guide the players
- 3. A challenge system can encourage players to do certain actions
- 4. Collected solutions are better when all the three features are present

## Problem

 Problem we want to solve is equivalent to finding maximal cliques in a colored multigraph



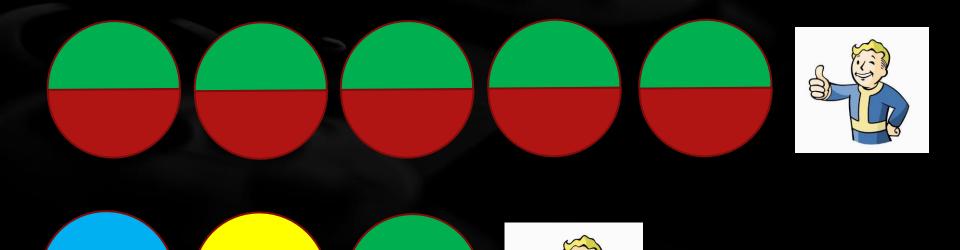
Exact solution:  $O(|V|^2)^{|C|}$ 

## Problem

- The problem can actually be solved quickly by algorithms
- Allows us to find the exact solutions and evaluate the performance of players

## Goal of the game

 Goal: build long sequences with many colors in common

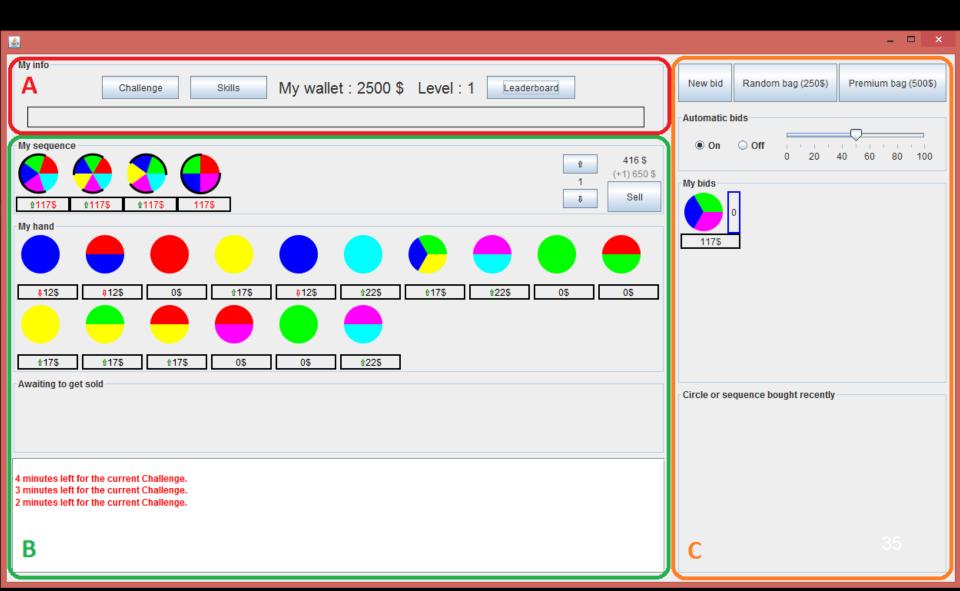


## Parameters

• Scoring function: baseScore, \* seqLength<sup>2</sup>

Number of colors	baseScore <sub>n</sub>
0	0
1	5
2	14
3	26
4	40
5	55
6	72

## Game interface



## Features

- Market:
  - Selling/buying individual circles
  - Buying sequences that have been sold to the system (buyout)

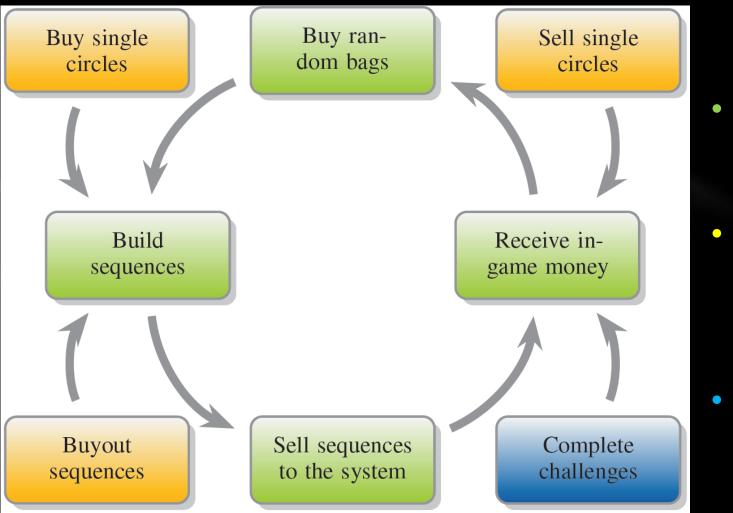
#### Features

- Skills:
  - Buyout King: lowers the price of buying a sequence from another player
  - Color Expert: gives a bonus to selling sequences that have more than one color in common
  - Sequence Collector: gives an additional sequence slot
  - Master Trader: gives a bonus to selling circles to other players

#### Features

- Challenges:
  - Sell/buy circles
  - Buyout sequences
  - Minimum number of colors
  - Minimum sequence length
  - Specific colors in common

### Gameplay loop



 Normal actions

 Marketrelated actions

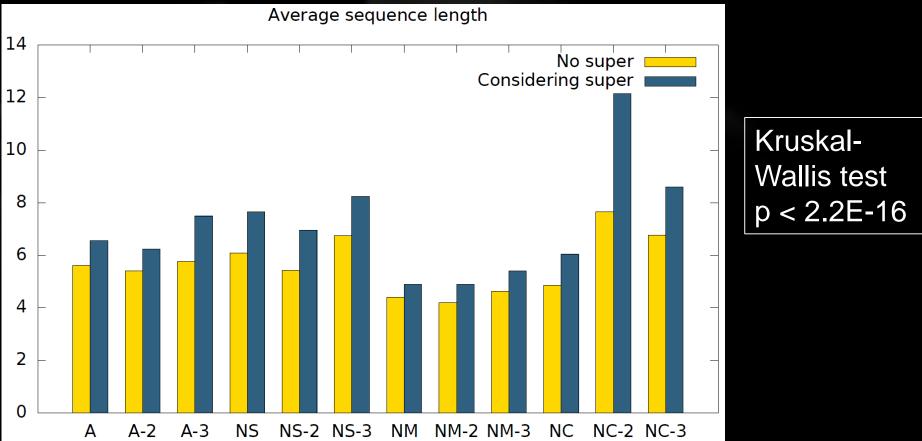
Challenges

#### Experiments

- Generated a graph of 300 vertices and 6 colors
- We recruited 12 groups of 10 people
- We tested 4 conditions (3 times each):
  1) All (skills + market + challenges)
  2) No skills (market + challenges)
  3) No market (skills + challenges)
  4) No challenges (skills + market)

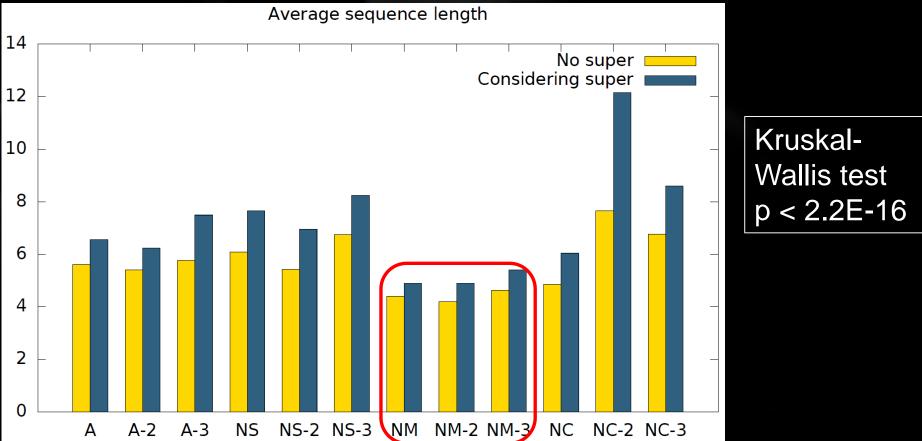
## Testing hypothesis 1 (market)

Presence of the market and sequence length



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Presence of the market and sequence length



# Testing hypothesis 1 (market)

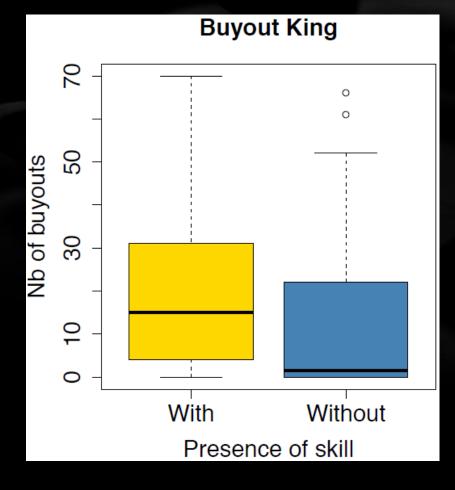
Presence of the market and sequence length

#### Similar groups (Dunn's test):

	A-2	A-3	NS	NS-2	NS-3	NM	NM-2	NM-3
Α	n/s	n		n/s				
A-2		n		n/s				
A-3			n/s					
NC						n		n/s
NC-3					n/s			
NM							n/s	n

- The skills have an effect on the players' strategies:
  - Buyout King (affects nb buyouts)
  - Master Trader (increases the nb of circles sold)
  - Color Expert (increases the proportion of sequences with many colors)
  - Sequence collector (increases the sequence length and nb of colors)

Buyout King skill and number of buyouts



Median values: With = 15 Without = 1.5

<u>Mann-Whitney's U test:</u> p = 0.004, effect size r = 0.28

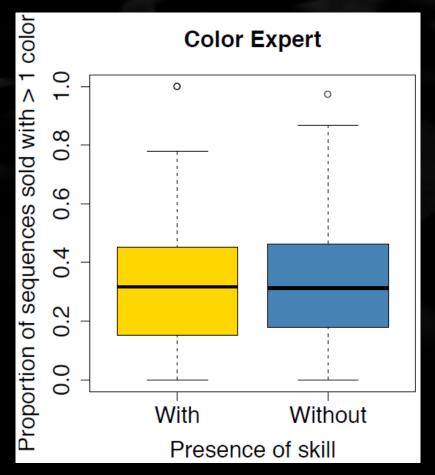
Master Trader skill and number of circles sold



Median values: With = 73 Without = 21.5

<u>Mann-Whitney's U test:</u> p = 7.2E-4, effect size r = 0.33

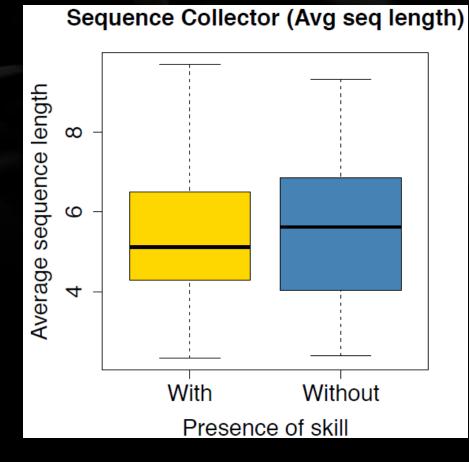
 Color Expert skill and proportion of multicolored sequences



 $\frac{\text{Median values:}}{\text{With} = 0.317}$ Without = 0.313

<u>Mann-Whitney's U test:</u> p = 0.89 -> failure to reject the null hypothesis

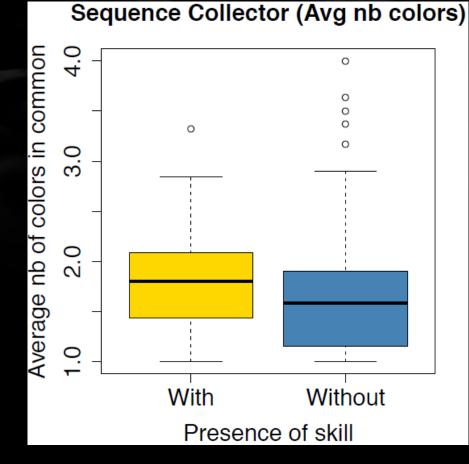
 Sequence Collector skill and average sequence length



Median values: With = 5.12 Without = 5.63

<u>Mann-Whitney's U test:</u> p = 0.69 -> failure to reject the null hypothesis

 Sequence Collector skill and average number of colors

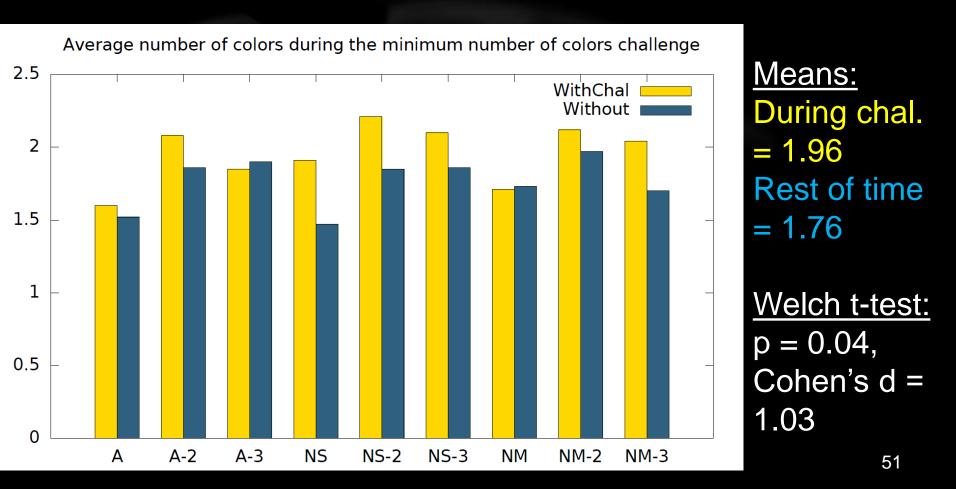


Median values: With = 1.80 Without = 1.58

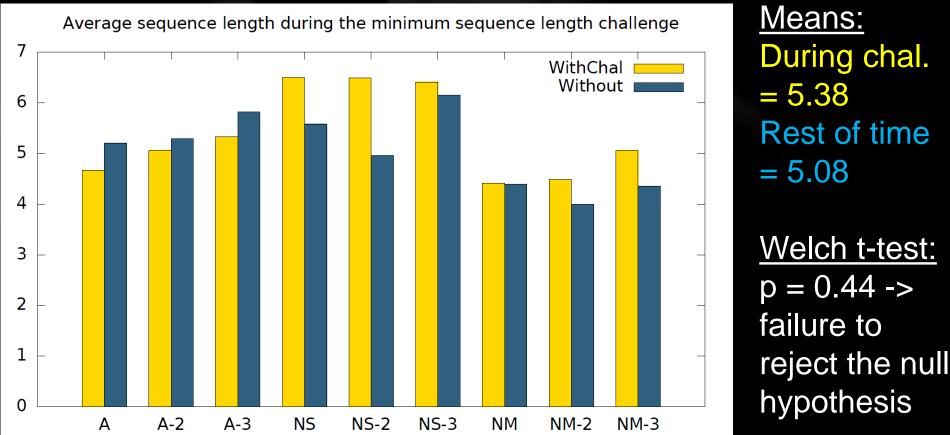
<u>Mann-Whitney's U test:</u> p = 0.01, effect size r = 0.21

- Challenges encourage players to do a certain action
  - Minimum number of colors challenge
  - Minimum sequence length challenge
  - Sell/buy challenge
  - Buyout challenge
  - Specific colors in common challenge

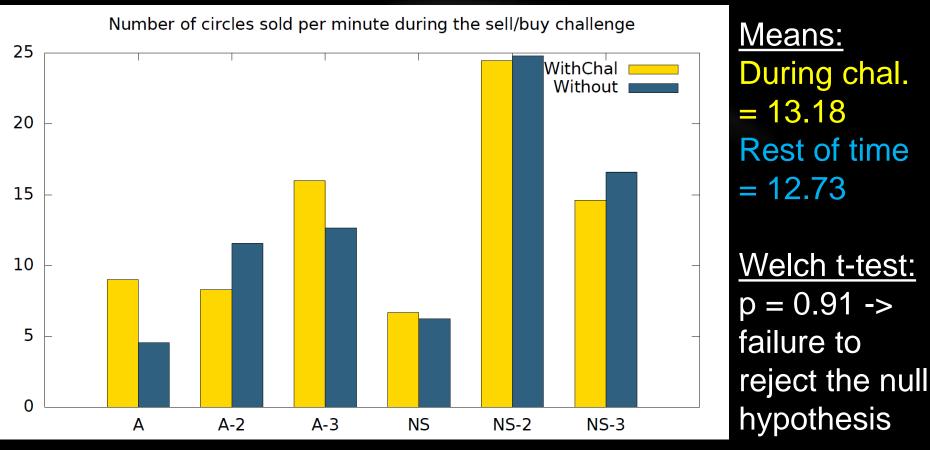
• Minimum number of colors challenge



• Minimum sequence length challenge



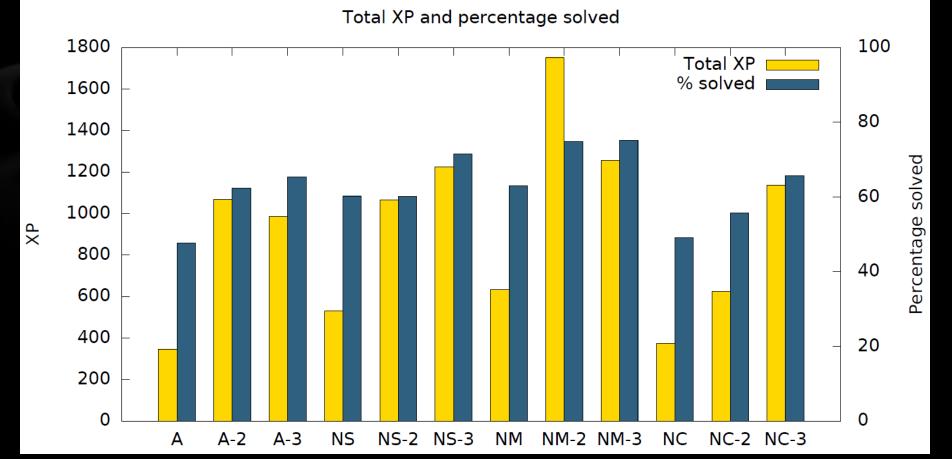
#### Sell/buy challenge



- Buyout challenge: appeared only once in all the game sessions
- Specific set of colors in common challenge:
  - Appeared 11 times in total
  - Completed only 8 times

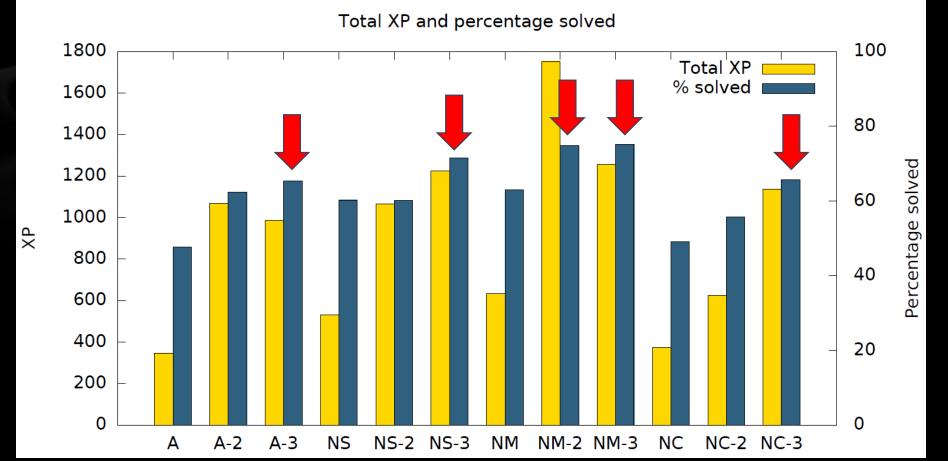
## Testing hypothesis 4 (% solved)

Is the percentage solved better when all the features are present?



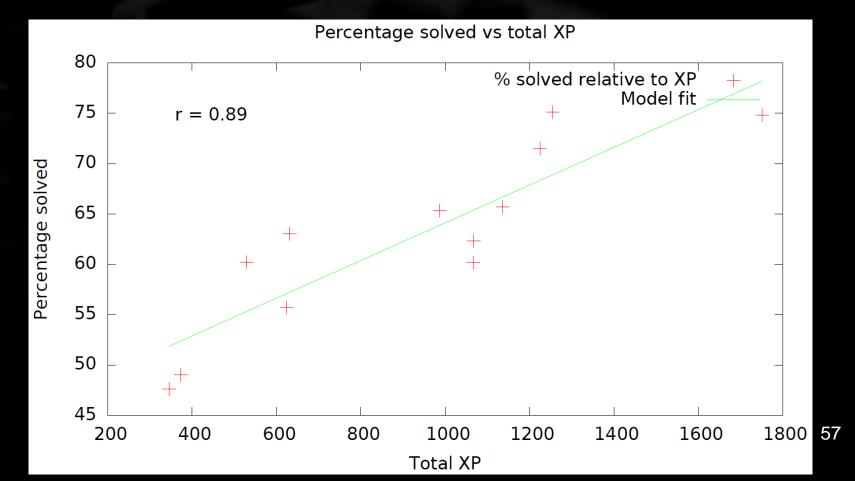
## Testing hypothesis 4 (% solved)

Is the percentage solved better when all the features are present?



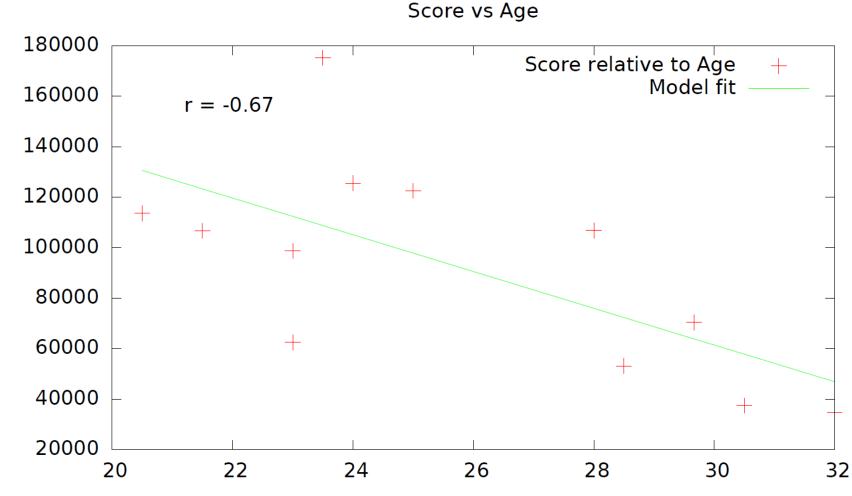
## Testing hypothesis 4 (% solved)

• Is the percentage solved better when all the features are present?



• Top 12 players VS the others:

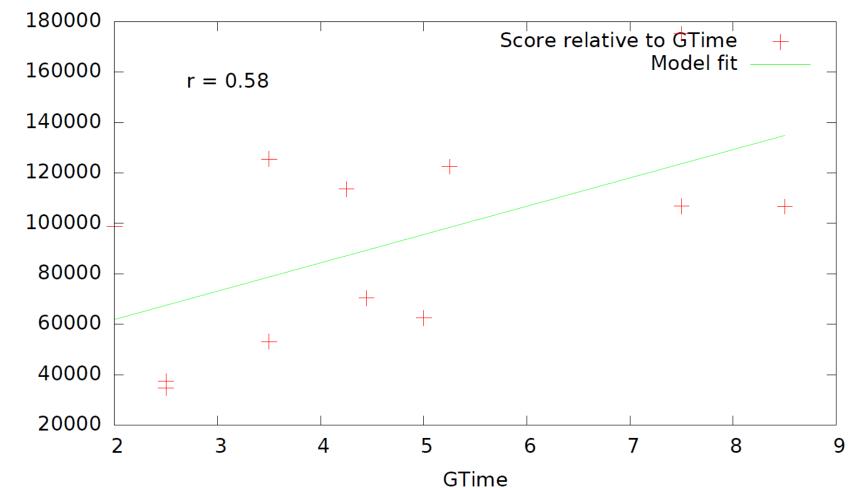
	Тор 12	Others
Age	23.42	25.99
Self evaluation	3.67	2.90
Game time / week	10.00	4.11



Age

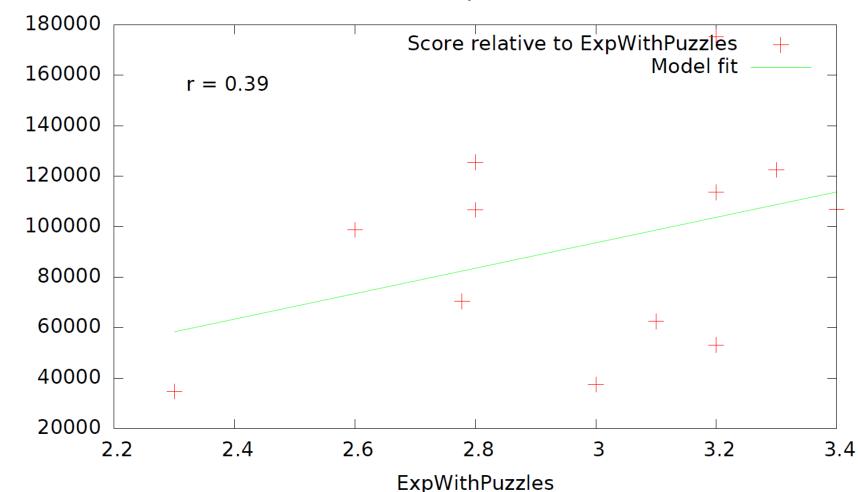
Score





Score

Score vs ExpWithPuzzles



Score

### Conclusions

- A market system is a useful tool to help players build longer solutions
- Skills are helpful to guide the players into doing specific actions in the game
- Well-balanced challenges in terms of difficulty can promote a certain action from the players
- Percentage of the problem solved depends not only on the features present in the game, but also on the players' skills

### Conclusions

- Players' satisfaction:
  - Average score of 7.16/10
  - Very addictive
  - Leaderboard = powerful motivation
  - Some participants found the game too easy / hard

#### Conclusions

- Scalability:
  - Would we get similar results with more players?
  - Probably: in the tests during development, we noticed that having more players only helped
  - Future work: development of an online version that would be available 24/7

### More future work

- Verify if the results are task dependent
- Build bots (AI) and test their performance
- Define an optimal crowd of players with different strategies
- Switch to a problem that makes a better use of human skills

## Crowdsourcing genomic databases

- Development of a web and mobile crowdsourcing platform for curating genomic databases
- Why: more and more data, manual curation is necessary, but funding is limited

## Crowdsourcing genomic databases

- Develop crowdsourcing systems for curating, maintaining and updating genomic databases
- 2. Develop learning/teaching interface
- Explore new technologies: virtual reality (3D genome browser)

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- Shu Hayakawa
- All the players

