[Introduction to mass collaboration], [Human computation], [Open call], [Distributed data collection], [Fragile Families Challenge]

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- 1) Introduction
- 2) Observing behavior
- 3) Asking questions
- 4) Running experiments
- 5) Mass collaboration
- 6) Ethics
- 7) The future

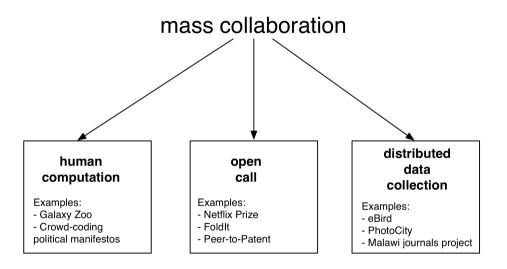


Fig 5.4 (Salganik 2018)

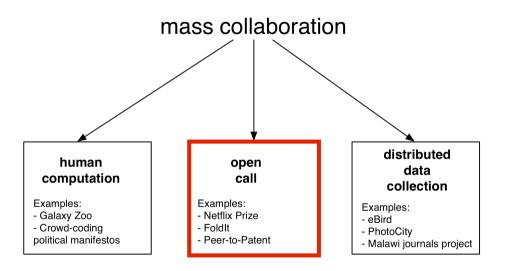


Fig 5.4 (Salganik 2018)

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- ightharpoonup Released a **training set** of \sim 100 million ratings
- ▶ Held-back a **test set** of \sim 1.5 million ratings

	Movie 1	Movie 2	Movie 3	 Movie 20,000
User 1	2	5		?
User 2		?	2	3
User 3				4
:				
User 500,000	?		2	1

Goal is clear, but path is not clear.

- ▶ Open it up to the world and offer a prize of \$1,000,000
- ▶ 44,014 valid submissions from 5,169 different teams
- ► Fortunately, they were easy to check

$$RMSE = \sqrt{\frac{\sum_{i} \sum_{j} (\widehat{r_{ij}} - r_{ij})^{2}}{n}}$$



So, in other words, if we take the rank-40 singular value decomposition of the 8.5B matrix, we have the best (least error) approximation we can within the limits of our user-movie-rating model. I.e., the SVD has found our "best" generalizations for us. Pretty neat, eh?

Only problem is, we don't have 8.5B entries, we have 100M entries and 8.4B empty cells. Ok, there's another problem too, which is that computing the SVD of ginormous matrices is... well, no fun. Unless you're into that sort of thing.

But, just because there are five hundred really complicated ways of computing singular value decompositions in the literature doesn't mean there isn't a really simple way too: Just take the derivative of the approximation error and follow it. This has the added bonus that we can choose to simply ignore the unknown error on the 8.4B empty slots.

So, yeah, you mathy guys are rolling your eyes right now as it dawns on you how short the path was.

Sifter aka Simon Funk

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Moved him instantly into fourth place, and was later used by all serious competitors.



Foldit

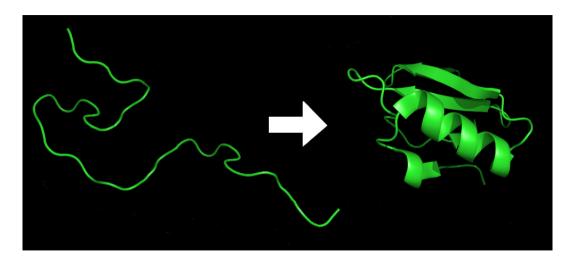
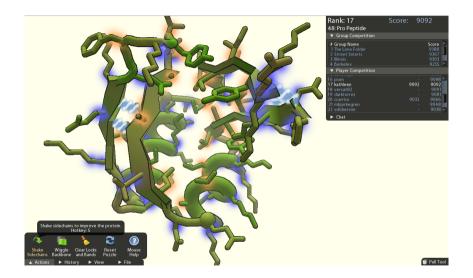
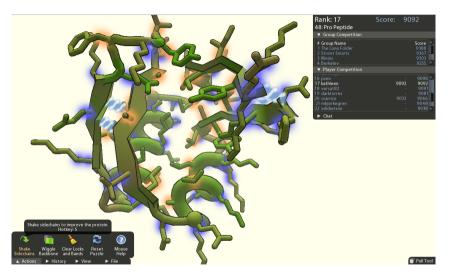


Fig 5.7 (Salganik 2018)

FoldIt



FoldIt



Gamers outperformed best known computational algorithms on 5 out of 10 proteins of unknown structure (Cooper et al., 2010)

Fragile Families Challenge

Measuring the predictability of life outcomes with a scientific mass collaboration

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What to read next:

- ► Longitude, Sobel (1996)
- ► "Statistical Significance of the Netflix Challenge", Feurverger et al. (2012)

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