Collective Intelligence Report

A Recommender System Implementation

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Contents

1	Introduction Exploratory of the Dataset					
2						
3 Design of the CF implementation						
	3.1 Design with core functions in mind	6				
	3.2 Data repository facilities	7				
	3.3 Evauluation module and Composablity	8				
	3.4 Model class and utilities	8				
	3.5 Details for similarity implementations	10				
4 Evaluation						
	4.1 Evaluation pipeline	11				
	4.2 Evaluation of the result	11				
5	Recap and Future work					
R	References					

1 Introduction

When our daily commerce went online since the turn of the millenia almost 20 years ago, the cost of maintaining information about the goods has drastically reduced, while the interface for shopping, as far as our attention when browsing products go, has not been improved to the same scale as the warehouse storing them (Benn et al. 2015). Take amazon.com for example: the availability of the products on the platform is unproportional to what an average user is able to focus the attention to. Even with the revolutionary search ability introduced through technology, vast majority of the products remain unexplored. Not to mention the economic benefit from tackling the Long Tail problem (Anderson 2006), stores can potentially "sell less for more". This is where recommender systems come in. As a new channel of communication between stores and users, users will also receive recommendation on items that they might be interested in.

Collaborated Filtering (hereafter as **CF**) is one of the 2 major families of recommender system strategy. Unlike content-based strategy, which is focusing on analysing the item's features, CF is focusing on the users' behaviour. One of the reason behind this is **CF** boosts diversity. Simply put, similar items recommended using content-based strategy appeal less in terms of selling point, unless the user are aiming to collect a specific genre, which is not a common shopping behaviour. People are drawn to each other based on liked minds. **CF** can emulate a good friend, should one user's taste is being identified as similar to another within the system.

This report describes the work involved in building a CF based recommender system, using varies methods, including distance metrics based similarity (Shardanand and Maes 1995), prediction formula by (Resnick et al. 1994) etc. Finally with the evalution of the involved methods. The Dataset being targeted in this report is a subset of the *MovieLens* movie rating data. We will first explore the dataset, then introduce the CF techniques with an emphasise on the implementation side. In the later chapters evaluation and metics plots will be shown, with future work being hinted on how to explain the phanomena we observed while evaluating the result.

2 Exploratory of the Dataset

The Dataset is a subset of 100,000 movie ratings from *MovieLens* database. A *rating* is rated by one user on one movie with a rating score from 1 to 5. We expect that not every user has rated every possible movies in the dataset. This implies that there are potentially way smaller number of unique users than the number of ratings. We confirmed this by looking at the dataset by running aggregation statistics, including number of each rating score.

User count: 943 Movie count: 1682 Rating count: 100000

Rating Density: 0.06304669364224531

Rating bin count for 1.0: 6110 2.0: 11370 3.0: 27145 4.0: 34174 5.0: 21201

Mean Rating: 3.52986

Briefly, there are 943 users and 1682 movies in the dataset. Further we can quantise the overall degree how much is the 100,000 ratings being covered in all possible user \times movie combination, namely a density score given by

$$density = \frac{|ratings|}{|possible_pair|}$$

.

As it turns out, the density is merely 6.3%. The average rating overall is 3.5. From figure 1 we can see the involved users have mostly rated high.

As each user might have rated a list of movies, it is also interesting to look at how the rating score looks like from a per user stand point. I calculated basic statistics for each user's rating scores. There are for example only 1 user who have a max rating score of 3, while 26 of them have a max score of 3.

```
table(stats user$max)
```

```
##
## 3 4 5
## 1 14 928
```

table(stats_user\$min)

```
##
## 1 2 3 4
## 723 192 26 2
```

The average variance of rating scores between user is

```
mean(stats_user$sd) ^ 2
```

```
## [1] 1.019751
```

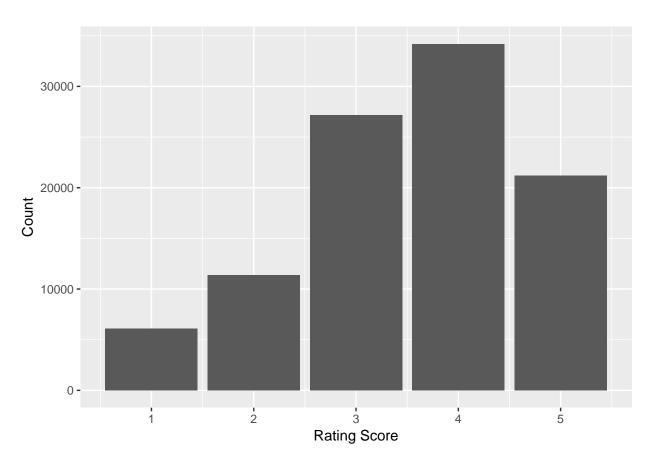


Figure 1: Bin Count

which indicates that even though this batch of users rated mostly higher than 3, the diversity is still significant.

Similar to statistics for each user, I also performed analysis on ratings for each movie.

##	mean	median	sd	max
##	Min. :1.000	Min. :1.000	Min. :0.0000	Min. :1.000
##	1st Qu.:2.660	1st Qu.:3.000	1st Qu.:0.8502	1st Qu.:4.000
##	Median :3.162	Median :3.000	Median :0.9951	Median:5.000
##	Mean :3.076	Mean :3.203	Mean :0.9209	Mean :4.451
##	3rd Qu.:3.653	3rd Qu.:4.000	3rd Qu.:1.1173	3rd Qu.:5.000
##	Max. :5.000	Max. :5.000	Max. :2.0000	Max. :5.000
##	min			
##	Min. :1.000			
##	1st Qu.:1.000			
##	Median :1.000			
##	Mean :1.301			
##	3rd Qu.:1.000			
##	Max. :5.000			

3 Design of the CF implementation

The design is followed along a bottom-up approach, such that core functionalities were first addressed in a composable fashion. The main concerns included how to calculate similarity quickly, how to calculate neighbours based on similarity quickly and how to design data structures that trade memory with speed.

3.1 Design with core functions in mind

As in typical bottom-up strategy, before thinking about optimal data structure for the job, the core functionalities were being specified out first.

In this stage, I looked at similarity calculation between the user pair (User a, User b) first. For both distance and cosine similarity, we have to loop through (in order of O(n)) the common set of movies that both users have rated. This prompted that the **User** class should have method that deliver the common set easily.

Next for the neighbours calculation, I took the assumption that all similarities are calculated and stored to allow O(1) retrieval, given user pair (a,b). The problem of getting neighbours from ranked similarity score is a typical search problem: this hinted at sorting the similarities before doing anything is most efficient approach. Both heap and Red-black tree based datastructure for temporary neighbourhood list for the targeted user can provide this speed up.

Finally for prediction calculation, given the neighbour list, prediction is a simple order O(n) operation.

Pseudo code for compute similarity:

func computeSim(a, b : User) similarity : double

commonMovies := a.getCommonMovies(b)
...
for (m : commonMovies) {
 calculateMetrics(a.getRating(m), b.getRating(m))
}
...

Pseudo code for compute neighbourhood:

return finalMetrics

```
func computeNeighbour(u : User) neighbours : Collection
   init(heap)
   init(neighbours)
   for (v : allUsers) {
      heap.insert( getSimilarity(u,v) )
   }
   while (neighbour.size < desired && heap.notEmpty) {
      neighbour.add( heap.top() )
   }
   return neighbour</pre>
```

Pseudo code for compute prediction:

return finalPrediction

```
func predict(a : User, m : Movie) rating : double
  neighbours := computeNeighbour(a)
  for (n : neighbours) {
      similarity := getSimilarity(a,n)
      calculatePrediction(similarity, a.getRating(m))
  }
  ...
```

High-level pseudo code shown above for the corresponding design suggested that for next stage of design, the focus should be designing correct facilities to make sure the complexity does not go higher.

3.2 Data repository facilities

Following along the rationale from previous section, several data repository are required for the operations. Among which, the similarity repository for allowing O(1) retrieval given a user pair should either be a **User** × **User** matrix, or compounded key based dictionary. While dictionary potentially uses less space than 2-D matrix for users, due to the diagonal value being useless, there is no benefit from using a dictionary. Depending on the language

for the implementation, for this case I'm using **Java**, 2-D matrix for users similarity score is easier to code.

A set of all users were also required in the previous analysis, which is provided by the **IO** facility when doing the dataset **IO** operations.

Because the **IO** facility were providing ratings set and movies set as well, it makes sense to have similarity class take them as argument too, for better encapsulation and prepare for interfacing with **Evaluation** module.

3.3 Evauluation module and Composablity

In the scope of this implementation, the evaluation strategy is leave-out-out. In other words, the evaluation module will essentially loop through all the ratings, such that for each rating we calculate predicted rating for that $\mathbf{User} \times \mathbf{Movie}$ pair and calculate error between the predicted and the actual rating.

Due to the encapsulation of data repositories inside similarity module, the evaluation module is very easy to implement: all it needs is a dependency of similarity. Further, because we know that there are different methods to be benchmarked for the similarity module, it makes sense to extract the previous contract to an interface.

Figure 2 shows the composite between the modules.

3.4 Model class and utilities

It is not common to address the Model class this late, after so many design has been mapped out. The reason behind this is bottom-up imposes a minimal pressure on both the complexity of model class and its dependencies. This however does not mean the Model classes are trivial, as an example the **User** class has 2 special requirement other than normal attributes for data encapsulation:

- for the **User** × **User** similarity matrix in the similarity module contract, indexing using **user_id** makes poor sense. The reason is that **user_id** is depending on the cleaness of the dataset, meanwhile what we need is the consistency for each and every programme execution. Therefore we must introduce an internal **id**. This is achieved using **static** keyword for the class attribute.
- as mentioned before, **User** class need to also have a method for common movies retrieval

Utilities other than **IO** modules are implemented in this stage. Among which the logging facilities were designed to be a generic writer such that object serialisation can be carried out without too much manual formatting.

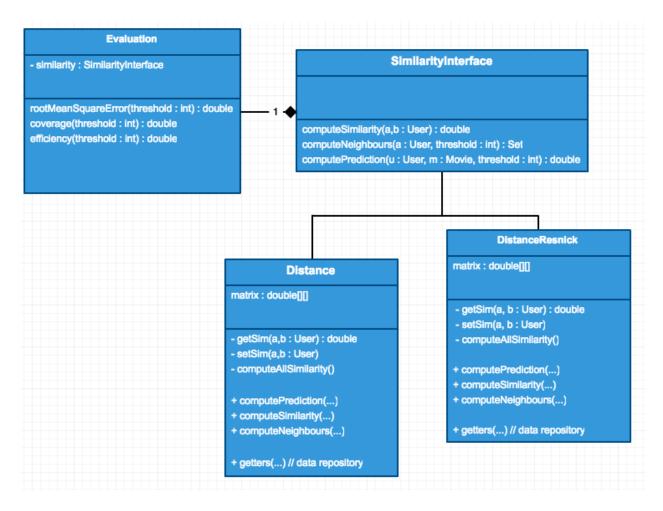


Figure 2: Composite of Evaluation and Similarity

3.5 Details for similarity implementations

4 similarity classes were implemented. The first one is **BaseLine**, which contains only prediction calculation. It exhaustively loop through all users bar from the user being predicted and deliver avarage rating as prediction.

Distance implements (Shardanand and Maes 1995)'s distance based similarity and uses prediction formula detailed in Collective Intelligence lecture. When calculating all similarity for the **User** \times **User** matrix, a fan-out to fan-in (or divide and merge) concurrency pattern was implemented to facilitate speed-up from parallelism. Concretely, the amount of users to be calculated is divided as evenly as possible to available compute cores on the system, and the subset workload is submitted to **ThreadPool** wrapped by **Executor**. Unlike parallelism for matrix multiplication, the $N \times N$ traversal requires slice by row or by column as one side and the other side must remain entire N. Otherwise the pairwise coverage will leave "holes" in the resulting matrix.

$$\sum_{i}^{n} w_{i} \otimes N$$

where w_i is the divided workload and N is the entire set of users.

Resnick's prediction formula is implemented in the **DistanceResnick** and **PearsonResnick**. The latter 3 implementations delivers a prediction rating range of (-1, 1], while if user a and b have no similarity the prediction will be -1.

For all the non **BaseLine** implementation, all similarity scores are computed inside constructor so that when instance object were ready, they are immediately available for prediction and evaluation runs.

4 Evaluation

Result metrics on prediction for **BaseLine**, **Distance** and **DistanceResnick** were evaluated through a pipeline such that methods are parameterised using threshold and runs are repeated to obtain enough datapoints for analysing.

4.1 Evaluation pipeline

The pipeline consists **IO** house-keeping to populate data repositories, after that *similarity* classes are initiated (as mentioned in previous chapter, similarities are calculated at this point). For the **BaseLine** the threshold parameter have no meaning, therefore it's being run first. Followed by loop of runs with increasing threshold value from 10 to 100 in step of 10 and from 100 to 300 in steps of 25.

Due to the nature of measuring runtime (efficiency) requires a full run of **RMSE**, there is no need to run **RMSE** separately in evaluation loop body, the evaluation result, which is augmented by utility classes, is responsible for giving out both run result and timing information.

4.2 Evaluation of the result

4.2.1 Efficiency (Time)

As hinted in previous chapter about the **BaseLine** prediction strategy, **BaseLine** uses by far the longest time to complete due to it exhaustively include all users as neighbours when predicting. The average run time for **BaseLine** on i7 mobile CPU with 8GB ram is 1460788 milliseconds (1460 seconds) or roughly 25 minutes.

Average run time for all other methods shows up heavy correlation as the threshold parameter goes up, matching our expectation. Figure 3 shows the run time plot for threshold between 10 and 100. The high bump on low threshold for resnick is likely due to caching behaviour of the CPU, when running from an older generation chip the bump is gone.

Similarly the time required for completion beyond threshold value of 100 put much pressure on **Resnick**, compared to **MSD** implementation.

4.2.2 Accuracy (RMSE)

For accuracy the **BaseLine** demonstrated a decent value of 1.1268639970237353, this does not necessarily indicate that **BaseLine** is good prediction algorithm, in fact, all it shows is that the mean (expectation) of the spread is a "safe" value.

When comparing MSD with MSD+Resnick a trend of increasing accuracy (decrease of error value) can be clearly observed in the following figure ?? and ??. However here the

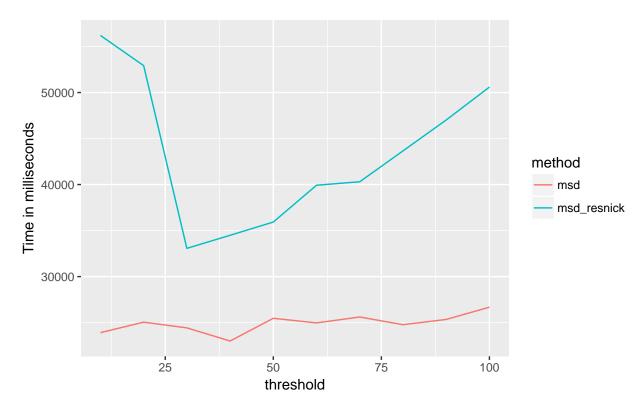


Figure 3: Efficiency Plot for threshold up to 100

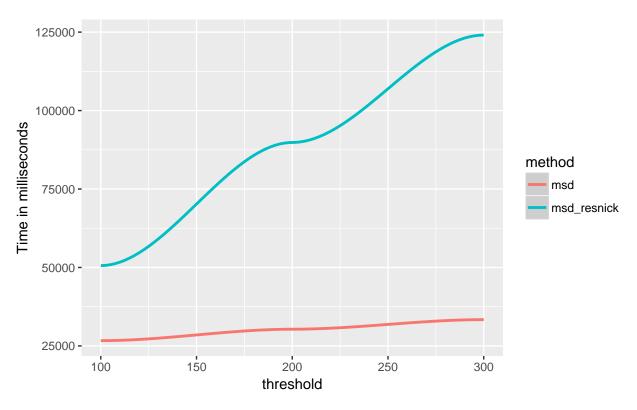
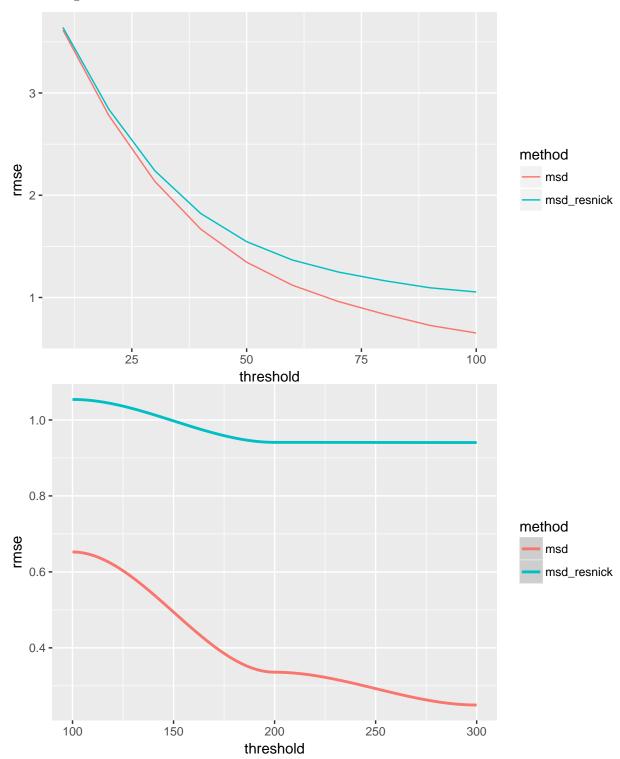


Figure 4: Time Plot for threshold between 100 and 300

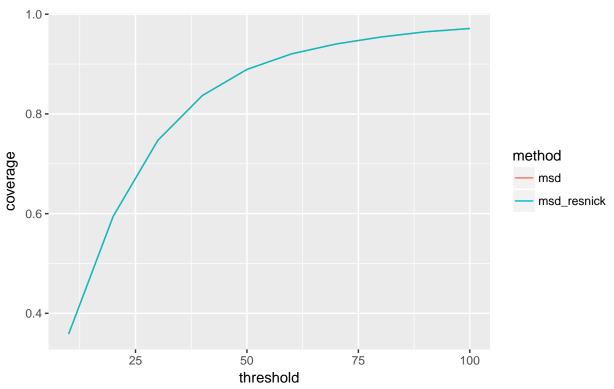
Resnick result shows diminishing growth when threshold value push on towards 100 and beyond. The same phanomena is not observed for \mathbf{MSD} , which performed steadily along the increasing threshold.

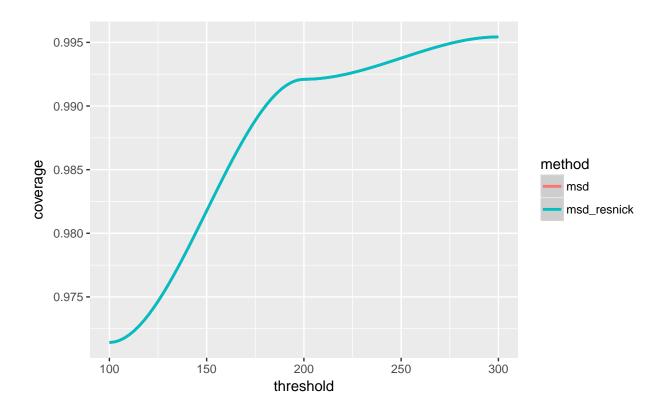


4.2.3 Coverage

Coverage denotes the degree how much ratings can be predicated by the algorithm. The MSD delivers similarity between users based on common movies, and the lack of commonality prevents all further operation, including prediction.

Unlike **BaseLine**, **MSD** and **MSD+Resnick** are affected by the threshold value. The more neighbours are considered for a prediction, the more likely a prediction can be made. Needless to say, **BaseLine** "dominates" this metrics by having a constant 1.0 coverage. Figure ?? and ?? shows the coverage result for both non **BaseLine** methods. Notably, because **MSD+Resnick** uses the same similarity as **MSD** the coverage for both 2 are exactely the same, unlike **Pearson**, which regretably due to limited scope is not being evaluated.





4.2.4 Review

Taking another look at **MSD** and **MSD+Resnick** metrics, I gathered a further metrics by calculating variance of the errors for every rating during prediction. It is expected that the variance will have high correlation in tendency as **RMSE**. Figure 5 and 6 confirms such assumption.

With the previous "strange" diminishing growth of the **RMSE** for **MSD+Resnick** in mind, I was hoping to find the variance would explain the phanomena. However, I failed to find such explanation. According to Figure 7 and 8 the variance do not exhibit obvious cause for the poor performance.

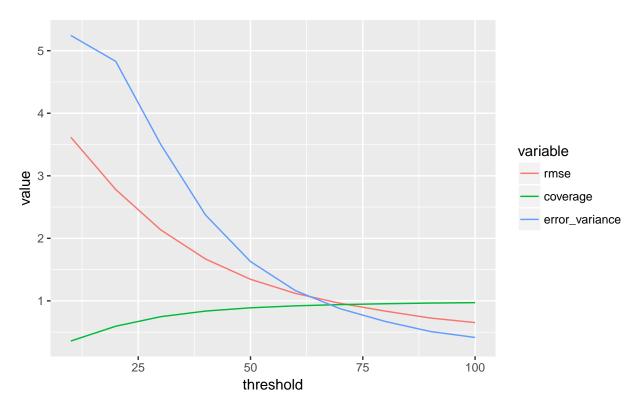


Figure 5: Metrics for MSD Plot for threshold up to 100

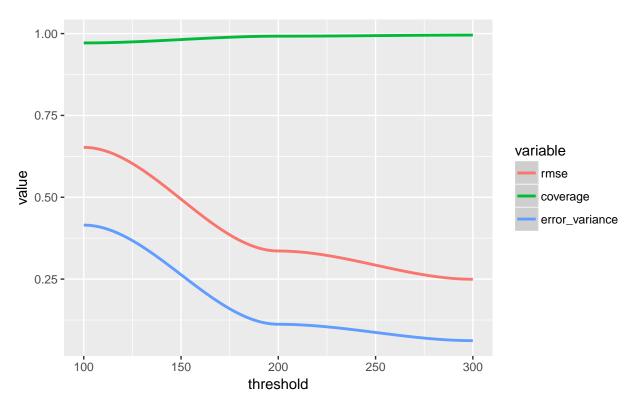


Figure 6: Metrics for MSD Plot for threshold from 100 to 300

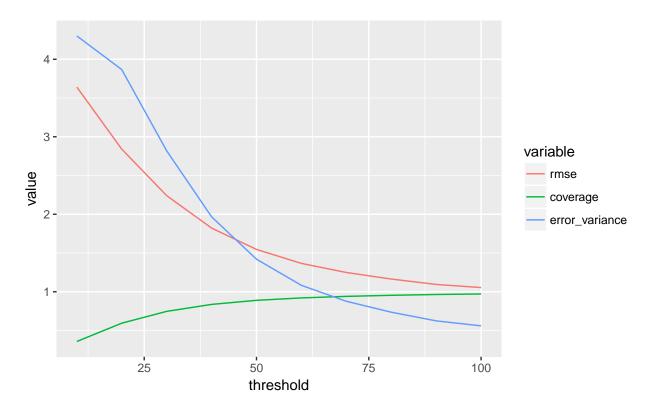


Figure 7: Metrics for MSD + Resnick Plot for threshold up to 100

5 Recap and Future work

In this recommender system report, MSD similarity method and Resnick prediction method are explored by implementation. The importance of choosing suitable data-structure and meaningful architecture design shows great impact on run time efficiency metrics. With the increasing threshold value, MSD based methods outperformed BaseLine in terms of accuracy and efficiency. Future work remains to be done on further evaluating the Pearson similarity's performance, and more anlaysis is needed to explain why Resnick on MSD performs poorly.

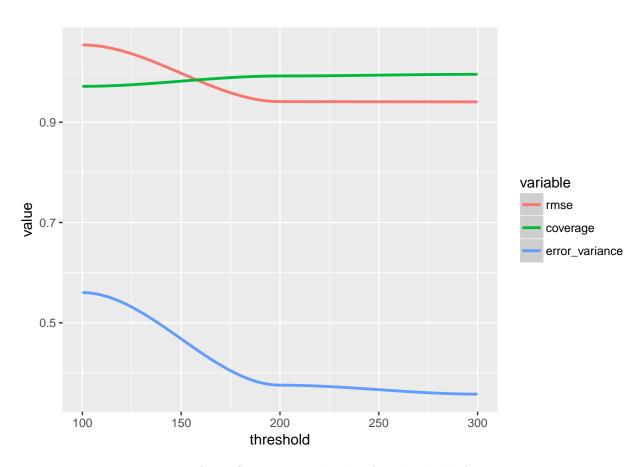


Figure 8: Metrics for MSD + Resnick Plot for threshold from 100 to 300

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