

Estimating deep web data source size by capture—recapture method

Information Retrieval

February 2010, Volume 13, Issue 1, pp 70-95

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Article

First Online:

13 August 2009

Received:

09 April 2009

Accepted:

27 July 2009

DOI (Digital Object Identifier): 10.1007/s10791-009-9107-y

Cite this article as:

Lu, J. & Li, D. Inf Retrieval (2010) 13: 70. doi:10.1007/s10791-009-9107-y

- 14 Citations
- 409 Downloads

Abstract

This paper addresses the problem of estimating the size of a deep web data source that is accessible by queries only. Since most deep web data sources are non-cooperative, a data source size can only be estimated by sending queries and analyzing the returning results. We propose an efficient estimator based on the capture—recapture method. First we derive an equation between the overlapping rate and the percentage of the data examined when random samples are retrieved from a uniform distribution. This equation is conceptually simple and leads to the derivation of an estimator for samples obtained by random queries. Since random queries do not produce random documents, it is well known that the traditional methods by random queries underestimate the size, i.e., those

estimators have negative bias. Based on the simple estimator for random samples, we adjust the equation so that it can handle the samples returned by random queries. We conduct both simulation studies and experiments on corpora including Gov2, Reuters, Newsgroups, and Wikipedia. The results show that our method has small bias and standard deviation.

Keywords

Deep web Estimators Capture—recapture <u>Download</u> fulltext PDF

1 Introduction

The deep web (Bergman 2001) is the web that is dynamically generated from data sources such as databases or file systems. Unlike the surface web where data are available through URLs, data from deep web data sources are guarded by search interfaces.

Estimating the size of a deep web data source is an important component of data source sampling which obtains a profile or summary of a data source (Callan and Connell 2001; Caverlee et al. 2004; Ipeirotis et al. 2001; Si and Callan 2003; Wu et al. 2003). Data source sampling has attracted much attention in numerous contexts, such as distributed or integrated information retrieval, data source selection and categorization, and peer-to-peer information retrieval. Given the proliferation of web data sources and web services, quite often there are many similar data sources serving the same purpose. There is a need to select the one that is most comprehensive. While data providers do have the details such as the size of the data source, the same information may not be available to a third party.

Estimating the data source size is also an indispensable step when crawling a deep web data source for the purpose of indexing, backup, and downloading. There has been extensive research on deep web crawling and data extraction by issuing queries (Lu et al. 2008; Ntoulas et al. 2005; Barbosa and Freire 2004; Liddle et al. 2002; Raghavan and Garcia-Molina 2001; Shestakov et al. 2005; Knoblock et al. 2000; Crescenzi et al. 2001; Nelson et al. 2006), and all of them need to decide as to when most of the data have been harvested. Without knowledge on the data source size, it is difficult to decide when to stop the crawling process, and how to evaluate the performance of the data extractors.

Almost all organizations have web presence, and most of them provide a keyword based search interface. The size of these deep web data sources can reflect the size of the organization, or the intensities of activities in these organizations. For example, by estimating the size of the data sources, we can learn the number of products being sold by an e-commerce company, the number of books in a university library, the number of articles published by a newspaper, or the number of blogs posted in a social networking web site. Such information is of interest to general public, sometimes vital for attracting advertisements from which many web sites live upon.

Since most of the deep web data sources can be accessed only by queries, the estimation is based on the queries issued and the results (either complete documents or document ids) obtained. This kind of query based data size estimation has been widely studied (Bar-Yossef and Gurevich 2006, 2007; Bharat and Broder 1998; Bolshakov and Galicia-Haro 2003; Broder et al. 2006; Caverlee et al. 2004; Shokouhi et al. 2006; Si and Callan 2003; Wu et al. 2003; Xu et al. 2007). One of the basic techniques is the traditional capture—recapture method (Amstrup et al. 2005; Chao and Lee 1992; Darroch 1958; Schumacher and Eschmeyer 1943) in ecology. The idea is that if one takes several samples from a population, there would be some overlapping between the samples. According to the overlapping information, various estimation methods are proposed and applied in areas including data collection size estimation.

Our first *contribution* in this paper is the derivation of a simple equation between the overlapping rate and the percentage of the data examined. The equation holds only when many queries are fired, which is common in query based sampling, but may not be feasible for estimation problems in ecology. Although the equation itself can not be directly used in size estimation due to the assumption of the availability of uniformly distributed random samples, the simplicity of the equation leads to the derivation of a real estimator for samples obtained by queries.

One challenge in data size estimation is the difficulty in obtaining random samples from a data source, partially because documents have unequal probabilities of being retrieved. It is well known that random queries do not return random documents, and that sampling by random queries will consistently underestimate the size (Bar-Yossef and Gurevich 2007; Broder et al. 2006; Shokouhi et al. 2006; Xu et al. 2007). Shokouhi et al. observed that there is a fixed relationship between the estimation and the actual size, and proposed the use of regression to adjust the estimation result obtained by an estimator for random samples (Shokouhi et al. 2006). However, this estimator is not consistent—it will overestimate when the sample size is large.

Based on our simple estimator for random samples, our second *contribution* is to adjust our estimator so that it can handle the samples returned by random queries. When documents have unequal probabilities of being captured, the degree of heterogeneity can be used to modify the estimation. Our simulation study shows that our estimator works very well if the degree of heterogeneity is known. We also conducted experiments on various corpora and show that our method compensates for this kind of query bias (Bharat and Broder 1998) very well.

Many data sources, especially large search engines such as Google, rank the matching results and return only top k elements. This kind of rankbias (Bharat and Broder 1998) can be tackled by downloading and analyzing the documents (Bar-Yossef and Gurevich 2007). Since our method does not download the documents, it can't overcome the rank bias. For ranked data sources, it works well only when there are not many overflowing queries, i.e., the queries match more than k documents.

Our method is not intended for estimating the size of web (Dobra and Fienberg 2004), or huge databases such as general purpose search engines (Bharat and Broder 1998), where ranking and overflowing queries are common and can not be ignored. Rather, it is for unranked data sources, or ranked data sources with low density of overflowing queries. These data sources are typically of a smaller size. Compared with the methods for ranked data sources, our estimator is rather efficient—we do not need to download and analyze the documents.

Another restriction of this paper is that we focus on textual data sources that contain plain text documents only. This kind of data source usually provides a simple keywords-based query interface, instead of multiple attributes as studied in (Wu et al. 2006).

2 Related work

2.1 Basic concepts of capture-recapture method

Capture—recapture method was originally developed in ecology and used to estimate the size of an animal population (Amstrup et al. 2005; Pollock et al. 1990). In the estimation process, animals are captured, marked, and released in several trapping occasions. The data collected during the process, including the number of capture occasions, the recaptured animals, and the distinct animals captured, allow one to estimate the total population. This estimation process corresponds nicely to the query based estimation of data collection sizes, where a trapping occasion corresponds to sending a query and retrieving a set of documents from a data source.

If all the documents have an equal probability of being matched by a query, and all the matched documents are returned, we have the simplest model for which many estimators have been developed. The classic estimator is the famous Petersen estimator (Pollock et al. 1990) that can be applied only to two capture occasions:

$$\hat{n}_{Petersen} = n_2 U_2/d_2,$$

(1)

where n_2 is the number of documents retrieved by the second query, U_2 is the number of distinct documents retrieved just before the second query, and d_2 is the number of duplicate documents that are captured after the second query. Here $U_2 = n_1$, i.e., it is the same as the number of documents retrieved by query one. We use U_2 instead of n_1 in order to show the connection between Eqs. 1 and 2 introduced below.

This estimator can be derived using maximum likelihood method. The problem of this estimator is that d_2 could be zero when n_2 and U_2 are not large enough. According to the birthday paradox, in general n_2 , U_2 should be greater than \sqrt{n} , where n is the actual size of the data source, in order to have overlaps between the results of two queries. Unfortunately, many queries do not have that many matches.

One approach to solving the problem is by obtaining two large samples, each is produced by many queries instead of just one query (Broder et al. <u>2006</u>).

Another approach is expanding the estimator to multiple capture occasions or queries, and taking the weighted average of the estimations. i.e.,

$$\hat{n} = rac{\sum_{i=2}^{q} w_i n_i U_i / d_i}{\sum_{i=2}^{q} w_i}$$

where q is the number of queries issued, w_i is the weight for the i-th query, n_i is the number of documents, including the duplicates, that are retrieved by i-th query; U_i is the number of distinct documents that are collected just before i-th query; d_i is the number of duplicates that are retrieved by the i-th query.

When weight $w_i = d_i$, it is the classical Schnabel estimator (Pollock et al. <u>1990</u>) for multiple captures:

$$\hat{n}_{Schnabel} = rac{\sum_{i=2}^{q} n_i U_i}{\sum_{i=2}^{q} d_i}$$

When weight $w_i = d_i U_i$, it is the Schumacher estimator (Schumacher and Eschmeyer 1943) that is used in Shokouhi et al.'s method (Shokouhi et al. 2006):

$$\hat{n}_{Schumacher} = rac{\sum_{i=2}^{q} n_i U_i^2}{\sum_{i=2}^{q} d_i U_i}$$

Unlike two capture occasions, the MLE (Maximum Likelihood Estimator) for multiple capture model does not have a closed form solution. Without surprise, both Schnabel and Schumacher estimators are approximate estimators. However, they are widely accepted as good estimators with very small bias and variance when animals (or documents) are captured with equal probability.

In reality, individuals, be it documents or animals, seldom have equal capture probability. For animals, young animals may be easier to be captured because they are more active. For documents, large documents may be easier to be retrieved by a query because there are more words in those documents.

For this kind of heterogeneous population where each individual has an unequal catachability, the estimation is notoriously difficult (Amstrup et al. 2005). MLE technique can no longer be used to derive an estimator because there can be as many as n+1 parameters: n and capture probabilities $p_1, p_2,...,p_n$. Estimating this many parameters from the capture data is not possible. Although there are several empirical estimators proposed for this model, including the Jacknife estimator (Pollock et al. 1990) and Chao (Chao and Lee 1992) method, both can be only applied to small population with hundreds of elements, and require large sample size.

2.2 Estimators for data collections

Data source size estimation methods can be classified into two categories. One relies on the sample set(s) of documents and the lexical analysis of those documents (Bar-Yossef and Gurevich 2006, 2007; Broder et al. 2006), while the other analyzes the document ids only (Bharat and Broder 1998; Gulli and Signorini 2005; Shokouhi et al. 2006; Wu et al. 2003; Xu et al. 2007).

2.2.1 Methods based on document analysis

Estimation methods based on document analysis date back to a sample—resample method proposed by Si and Callan <u>2003</u>. More recently fairly sophisticated methods such as (Bar-Yossef and Gurevich <u>2006</u>, <u>2007</u>; Broder et al. <u>2006</u>) are proposed.

Taking Broder et al's work (Broder et al. 2006) for example, they first try to establish the sizes of two subsets of the corpus. A subset is the documents that are indexed by a query pool, which may contain millions of queries such as all the eight digit numbers. Since the query pool is rather large, it is impractical to fire all the queries to decide the number of documents that can be retrieved using the query pool. Hence, from the query pool a set of sample queries are randomly selected and sent to the data source. All the matched documents are downloaded and analyzed to obtain the weight (Broder et al. 2006) of each query. Based on the average weight of the queries from the sample, the number of the documents that can be obtained by the query pool can be estimated by multiplying the average weight by the query pool size. Once two such subsets of the documents are obtained, the Petersen estimator (Eq. 1) is used to estimate the total corpus size.

Obviously this approach is rather inefficient and sometimes infeasible because

- Some deep web data sources may not support the downloading of the documents. For example, Amazon book search web service will return the basic information of the books, instead of the entire books. For this kind of data sources, it is impossible to use the returned documents to obtain the source profile.
- Even when the documents are downloadable, the estimation process is very expensive. In addition to the downloading of the documents, obtaining random documents may incur further costs. For example, one of Bar-Yossef et al's methods (Bar-Yossef and Gurevich 2006) needs to fire 2,000 queries in order to obtain one random document.
- The approach is not stable. Quite often the links provided by a deep web data source may not be active, and this may disrupt the estimation process.

2.2.2 Methods based on document ids

This paper focuses on another category of estimation methods, which rely on the returning documents ids instead of the entire documents. Document ids can be in various forms such as ISBN numbers for books in Amazon, URLs for web pages indexed in Google, or file names in our experiments. Compared with the bulk of the documents downloaded, this type of estimator only needs to know the document identifiers, which will save network traffic tremendously.

This approach originated from the Capture–Recapture method, which is extensively studied and widely used in estimating the population of wild animals (Amstrup et al. 2005). Liu et al. proposed using the Capture–Recapture method to estimate a data source size (Liu et al. 2002). More recently, Carverlee et al. (Caverlee et al. 2004) discussed the problem in the setting of a distributed environment. The estimator they used is the traditional Petersen estimator. Lu (2008) briefly introduced a method based on overlapping information.

Shokouhi et al. (2006) proposed to use multiple capture—recapture method, or Capture with History (hereafter *CH*) method, to estimate a data source size. Using regression method, they developed a new estimator based on the traditional Schumacher and Eschmeyer estimator (Schumacher and Eschmeyer 1943) as shown in Eq. 4.

Bharat and Broder used a large query pool to estimate the sizes of search engines and the web (Bharat and Broder <u>1998</u>). In particular, they identified various biases, especially the *querybias* and *rankingbias*, during the estimation process. Gulli and Signorini (Gulli and Signorini <u>2005</u>) improved the method described in (Bharat and Broder <u>1998</u>).

2.3 Compensate the bias

Capture—recapture based methods tend to underestimate the size of a data source due to several reasons. One is the assumption that documents (or the wild animals in the case of animal population estimation) are of the same probability of being captured. When using random queries to capture documents, the retrieved documents are actually not randomly selected.

Various methods have been proposed to compensate this kind of bias. In the traditional capture—recapture research, people proposed various estimators (Amstrup et al. 2005; Chao and Lee 1992) to cope with the population that have unequal catchability. In data source estimation, Bar-Yossef and Gurevich strive to obtain random samples (Bar-Yossef and Gurevich 2006, 2007) by downloading and analyzing the texts.

Shokouhi et al. (2006) corrected the bias of the CH method using regression (CH_{reg} hereafter). They conjecture that there is a fixed relationship between the initial estimation, which is obtained by capture—recapture methods, and the actual data size. Based on this hypothesis, they use a training data set to obtain an equation between the initial estimation and the actual size using non-linear regression.

Xu et al. (2007) tried to compensate the bias by modeling the document capture probabilities with logistic regression.

3 An estimator for random samples

When random samples are obtained from a uniform distribution, the collection size can be estimated fairly accurately using various estimators such as Eq. <u>4</u>. However, it is difficult to adjust Eq. <u>4</u> for samples obtained by queries due to its complexity.

Following the setting of capture—recapture method introduced in Sect. <u>2.1</u>, suppose that q number of queries are issued in sequence. Matched documents of various sizes are taken, with replacement, allowing documents to be seen in multiple capture occasions. In this case let $t = \sum_{i=1}^q n_i$ be the total of all documents including repetition, while $u = U_{q+1} = \sum_{i=1}^q (n_i - d_i)$ the number of distinct items. Let P = u/n denote the percentage of the sample items obtained from the data source, OR = t/u (Overlapping Rate) is the ratio between the accumulative total number of data items and the distinct data items in the samples. Then, we derive a simple equation as follows: $P = 1 - OR^{-2.1}$

3.1 Simulation study

Before deriving Eq. 5, we run a simulation study in the context of a variation of urn model (Holst <u>1979</u>) with replacement.

Urn model: Consider an urn that contains n balls, each is labeled with a distinct number from 1 to n. Randomly select k balls, where $k \ll n$, from the urn, record the ball numbers that are drawn, then put back the balls into the urn. Repeat the process for i times, in which stage u(i) distinct balls are selected, which constitutes P(i) percent of all the balls in the urn. The overlapping rate OR(i) is t(i)/u(i).

The question is: What is the relationship between P and the overlapping rate OR, if there is any?

We first run an urn model simulation to reveal the relation.

Urn model simulation: In this experiment, n is the total number of data items in the data source, and k is the size of each sample. In each step, we generate k random numbers that are uniformly distributed within the range between 1 and n. Add those numbers to the result set and record the overlapping rate OR and hit rate P. Repeat the process until the cardinality of the result set is close to n.

Figure $\underline{1}$ draws a relationship between P and OR for step lengths ranging from 1 to 40 where n, the total of number data items, is 5,000. We tested other values of n and observed similar results.

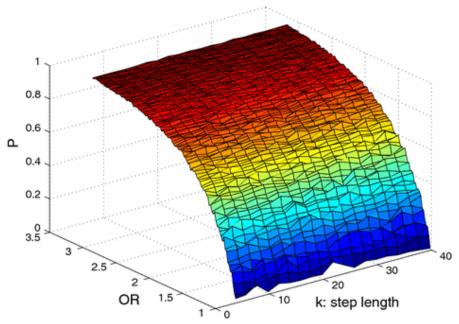


Fig. 1

Urn model simulation. n = 5,000

The simulation indicates that the hit rate P is a function of the overlapping rate OR, independent of the step length k. This prompts us to derive the equation between P and OR.

3.2 Derivation

P(i), the percentage of the data obtained up to i-th iteration, can be also interpreted as the probability of one particular document that is captured in any of the iterations. We start with:

$$P(1) = \frac{k}{n}$$

(6

$$P(2) = rac{k}{n} + rac{k}{n} - rac{k}{n} * rac{k}{n}$$

• ••

$$P(i) = P(i-1) + \frac{k}{n} - P(i-1) * \frac{k}{n}$$

(7)

Solving the Eqs. 6 and 7 we obtain the following:

$$P(i) = 1 - \left(1 - \frac{k}{n}\right)^i$$

$$= 1 - \left(1 - \frac{u}{n}\frac{ik}{u}\frac{1}{i}\right)^i$$

$$= 1 - \left(1 - \frac{P(i) * OR(i)}{i}\right)^i$$

Moving OR(i) to the left hand side of the equation we have:

$$OR(i) = \frac{i * \left(1 - (1 - P(i))^{\frac{1}{i}}\right)}{P(i)}$$

The question we want to ask is: whether OR is a function of P? For example, given a predetermined P = 10%, what is the OR at that point? Since i is assumed to be a large number and P is fixed, the corresponding OR can be derived as follows:

$$\begin{split} OR &= \lim_{i \to \infty} \frac{i * (1 - (1 - P)^{\frac{1}{i}})}{P} \\ &= \frac{1}{P} \lim_{i \to \infty} \frac{1 - (1 - P)^{\frac{1}{i}}}{\frac{1}{i}} \\ &= \frac{1}{P} \lim_{i \to \infty} \frac{\frac{\delta(1 - (1 - F)^{\frac{1}{i}})}{\delta(i)}}{\frac{\delta(\frac{1}{i})}{\delta(i)}} \\ &= \frac{1}{P} \lim_{i \to \infty} \frac{\frac{\delta(1 - (1 - F)^{\frac{1}{i}})}{\delta(i)}}{\frac{\delta(\frac{1}{i})}{\delta(i)}} \times \frac{\frac{\delta(\frac{1}{i})}{\delta(i)}}{\frac{\delta(\frac{1}{i})}{\delta(i)}} \\ &= \frac{1}{P} \lim_{i \to \infty} \frac{\frac{\delta(1 - (1 - P)^{\frac{1}{i}})}{\delta(i)}}{\frac{\delta(\frac{1}{i})}{\delta(i)}} \\ &= \frac{1}{P} \lim_{i \to \infty} \frac{\delta(1 - (1 - P)^{\frac{1}{i}})}{\frac{\delta(\frac{1}{i})}{\delta(i)}} \\ &= \frac{1}{P} \lim_{i \to \infty} \left(-(1 - P)^{\frac{1}{i}} \times ln(1 - P) \right) \\ &= \frac{-ln(1 - P)}{P} \lim_{i \to \infty} \left(-(1 - P)^{\frac{1}{i}} \right) \\ &= -\frac{ln(1 - P)}{P} \end{split}$$

Hence we have Eq. 9:

$$OR = -\frac{ln(1-P)}{P}$$

(9)

Note that although both OR and P are dependent on i, k, and n, what Eq. 9 shows is that when i is large, no matter what values i, k, and n take, OR solely depends on P.

Another view to understand Eq. 9 is that in the sampling process there are various ways, either by increasing i or k, to reach a certain value of P. When i is large, Eq. 9 states that no matter how P is obtained, OR is a function of P only.

Equation 9 can be also illustrated by drawing Eq. 8 as in Fig. 2, where OR is a function of both P and i. It shows that while the relation between P and OR is volatile when i is small, it tends to be stable with the increase of i. We can see that the threshold value for i is dependent on P. Although in the above derivation i in theory should be a large number, our empirical experiments showed that 100 is good enough.

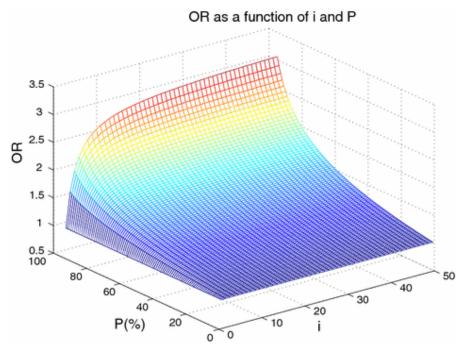


Fig. 2

Plot for Eq. $\underline{8}$ with i as X-axis, P as Y-axis, and OR as Z-axis. With the increase of i, the relation between P and OR becomes stable

To estimate the value of P based on OR, we need to inverse Eq. $\underline{9}$. Since P is smaller then 1, we can postulate the regression equation as follows:

$$P = 1 - \alpha OR^{\beta}$$

$$ln(1-P) = ln\alpha + \beta ln(OR)$$

By running linear regression on the above equation with data generated from Eq. 9 when P is smaller than 0.5, we have

$$\ln(\widehat{1-P}) = 0.005448767 - 2.108591182ln(OR),$$

$$R^2 = 0.999791835$$

i.e., $\hat{\alpha} = 1.005463639$, and $\hat{\beta} = 2.108591182$.

Here R-square is very close to one, and the standard errors for the intercept and slope are close to zeroes (0.000748673 and 0.00453560, respectively). Thus, we derive the following approximation for the relation between P and OR when P is smaller than 0.5: $P = 1 - OR^{-2.1}$ (5)

One of the reasons to restrict P to be smaller than 0.5 is that in estimation applications, the samples in general do not need to be very large. Another reason is that when P is

small, Eq. $\underline{9}$ and 5 can fit very well as illustrated in Fig. 3. However, when P (and OR) becomes larger, there is a small discrepancy between these two equations as shown in Fig. 3a.

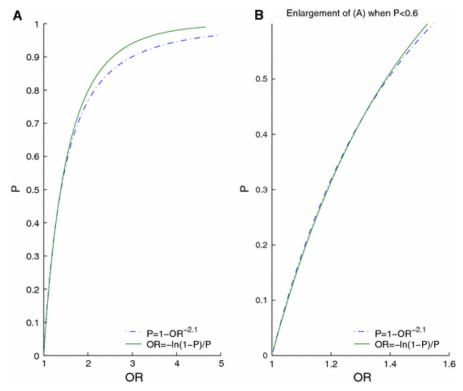


Fig. 3

Plots for Eqs. 5 and 9. **b** is an enlargement of **a** when *OR* is small

3.3 Varying k

In the derivation of Eq. 5 we assume that all the captures have a fixed size k. We need to show that Eq. 5 also holds for varying k size as long as $k \ll n$. This will make the equation more applicable in practice, since in many cases each capture may return varying number of elements.

We conducted another simulation which is the same as the previous one described in Sect. 3.1 except that now in each capture the number of elements is a random number, i.e., in each step we generate a random number *stepSize* within range 1 and k, then we generate *stepSize* number of random numbers within [1, n]. The relationship between P and QR is similar to the study in Sect. 3.1. In the next section more simulations on varying step lengths are conducted to confirm this result. Figure 4 shows the results of two runs of the simulation with exactly the same condition, and the comparison with Eq. 5.

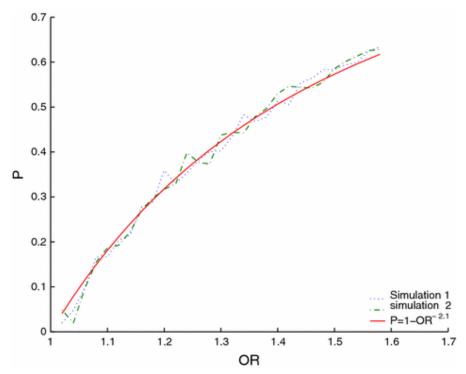


Fig. 4

Simulation with random step length and the comparison with Eq. 5. Two simulations (Simulation 1 and Simulation 2) are run on the same condition, i.e., n = 5,000, stepSize is a random number between 1 and 100

3.4 A naive estimator

Based on Eq. 5, a native estimator is given below:

$$\hat{n} = \frac{a}{1 - OR^{-2.1}}$$

(10)

where u is the number of distinct documents retrieved.

Although the estimator works only for random samples from a uniform distribution, which is rare in practical applications, it leads to the derivation of the estimator for unequal catch probabilities as described in the next section.

Figure 5 shows the results of the estimation using Eq. 10, where the elements are selected randomly with uniform distribution in the range of [1, 5,000]. Here the actual size n is 5,000. When both i and k are small (i < 20, k < 20), many of the estimates are infinite and are not shown in the chart. This can be explained by the Birthday Paradox that the sample size should be larger than \sqrt{n} to produce collisions, or make overlapping rates greater than 1. With the increase of k and i, the estimate becomes stable and accurate.

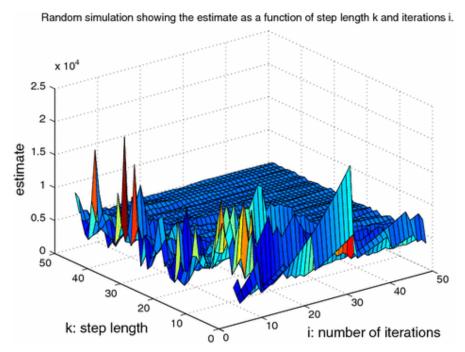


Fig. 5 Estimation result using Eq. 10 with different but fixed step lengths and iterations. n = 5,000. The estimate becomes stable and accurate when k and i increase

3.5 Comparison with Schumacher and Eschmeyer method

We also compared our method (OR method) described in Eq. 10 with the traditional Schumacher and Eschmeyer method (Schumacher and Eschmeyer 1943) described in Eq. 4. Table 1 explains the details of one simulation where n = 2,000. k(i) is the number of elements captured in the i-th step, d(i) is the duplicate elements in the i-step, u(i) is the distinct elements captured up to the i-th step. Figure $\underline{6}$ is the plot of the data in Table 1.

Table 1

Details of the simulation and the stepwise estimation result by SE (Schumacher & Eschmeyer) method and OR method. n = 2,000, step size ranges between 1 and 40

i k(i) d(i) u(i) \hat{n}_{SE} \hat{n}_{OR}

- 1 3 0 3 ∞ ∞
- 2 6 0 9 ∞ ∞
- 3 37 0 46 ∞ ∞
- 4 23 2 67 1,359 1,119
- 5 8 1 74 1,086 924
- 6 10 0 84 1,425 1,182
- 7 35 0 119 3,808 2,336
- 8 38 0 157 8,311 3,051
- 9 10 1 166 5,359 2,748
- 10 31 3 194 3,317 2,384
- 11 38 0 232 5,456 3,376
- 12 32 3 261 4,253 3,143
- 13 14 2 273 3,693 2,933
- 14 28 1 300 4,240 3,285
- 15 18 1 317 4,400 3,426
- 16 32 10 339 2,614 2,442
- 17 27 6 360 2,360 2,261

i k(i) d(i) u(i)

18	34	9	385	2,097	2,054
19	39	3	421	2,420	2,235
20	32	7	446	2,347	2,193
21	24	7	463	2,221	2,109
22	17	5	475	2,155	2,064
23	39	8	506	2,204	2,076
24	21	7	520	2,125	2,024
25	6	0	526	2,181	2,066
26	6	2	530	2,160	2,053
27	18	3	545	2,217	2,097
28	28	6	567	2,258	2,112
29	38	13	592	2,164	2,039
30	19	3	608	2,232	2,092

•••

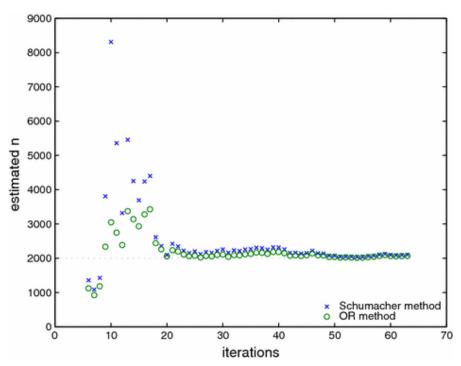


Fig. 6

Estimation result using our *OR* method (i.e. Eq. <u>10</u>) and Schumacher and Eschmeyer method. n = 2,000. In each iteration k elements are selected, where k is a random number between 1 and 40

Table $\underline{2}$ summarizes the Mean Squared Error (MSE), variance, and bias for various sample sizes ranging between 20 and 50% of n. For each sample size 50 simulations are done and the MSE et al. are calculated from the 50 simulations. It can be seen that our OR method is systematically better than Schumacher and Eschmeyer method.

Table 2

Comparison of SE (Schumacher and Eschmeyer) method and *OR* method. The actual size is 2,000, and the step size is a random number between 1 and 40. The data is obtained by 50 runs

	i	\hat{n}	MS	SE	V	ar	Bi	ias
Sample size (%)	SE	OR	SE	OR	SE	OR	SE	OR
20	2239	2000	138,909	61,335	81,582	61,334	239.43	0.87
30	2105	1959	48,645	28,579	37,612	26,975	105.04	-40.04
40	2070	1990	18,323	10,507	13,286	10,415	70.97	-9.64
50	2052	2017	11,789	7,106	9,031	6,782	52.52	17.99

4 Estimator when documents are selected by queries

Equation <u>10</u> is derived on the assumption that documents are selected randomly from uniform distribution. Although we can select random queries, picking out a set of random documents is by no means an easy task (Bar-Yossef and Gurevich <u>2006</u>; Bharat and Broder <u>1998</u>; Thomas and Hawking <u>2007</u>). When samples are not randomly selected with a uniform distribution, there is a tendency that the estimation is negatively biased in traditional capture—recapture method (Amstrup et al. <u>2005</u>) and in data size estimation (Shokouhi et al. <u>2006</u>; Bar-Yossef and Gurevich <u>2007</u>; Bharat and Broder <u>1998</u>). Almost all the methods underestimate the size consistently.

Prompted by Eq. 5, we conjecture that there is also a fixed relation between *P* and *OR* in real data sources, with a modified equation:

$$P = 1 - OR^{\alpha}$$
,

(11)

where α is a value, between 0 and -2.1, that is to be determined by the corpus under investigation. The previous section has shown that if the documents can be sampled with a uniform distribution then $\alpha = -2.1$. Before describing the method to calculate α for documents obtained by queries, we need to understand the non-uniform sampling first.

4.1 The unequal catching probability

To reveal the cause of the negative bias, we conducted an experiment to show how documents are captured when random queries are issued. In this experiment 20,000 documents in 20 Newsgroups (Lang 1995) are sorted according to their file sizes, and are numbered through 1–20,000 in a decreasing order. Then we retrieve documents by

firing single word queries that are randomly selected from the Webster dictionary. Figure 7a depicts the 13,600 documents that are selected by queries, and includes duplicates. To give a better view of 7a, b depicts the first 100 documents that are captured.

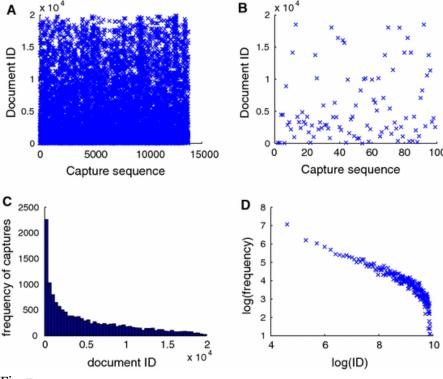


Fig. 7

Capture frequency of news groups documents by queries: $\bf a$ is the scatter plot when documents are selected by queries. In total 13,600 documents are retrieved. $\bf b$ is the first 100 captures in $\bf a$. $\bf c$ is the histogram of $\bf a$. $\bf d$ is the log-log plot of $\bf c$

To understand the plot better, we draw the histogram of retrieved documents as in Figure 7c, where the bin size is 100. It shows that the largest 1% of the documents (i.e., the largest 200 documents) are captured around 2,300 times. The second largest 200 documents are retrieved around 1,000 times. Many small documents, i.e., the documents with larger IDs, are rarely retrieved by queries. The histogram demonstrates that the capture frequency follows the power law, as its log—log plot in Fig. 7d shows.

Since the capture frequency follows the power law, in the following simulation studies, we will generate random numbers following Pareto distribution to simulate the capture frequency.

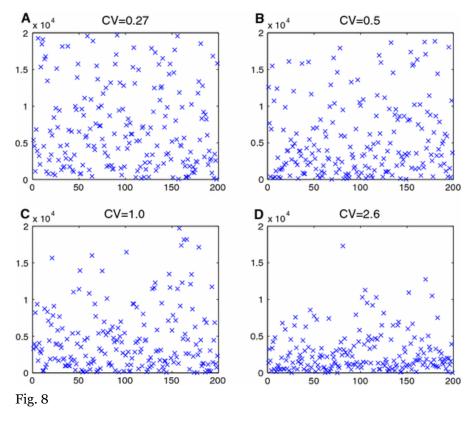
4.2 Measure the heterogeneity

One way to measure the degree of heterogeneity of the capture probability distribution is coefficient of variation (CV) (Chao and Lee $\underline{1992}$). Assume that the documents in the data

source have different but fixed probabilities of being captured, i.e., $p = \{p_1, p_2, ..., p_n\}$, $\sum_{j=1}^n (p_j) = 1$. Let $p_1, p_2, ...$, and p_n have mean $\bar{p} = \sum p_i/n = 1/n$. The coefficient of variation of the probabilities p is defined as

$$\gamma = rac{\sqrt{\sum_{i=1}^{\overline{n}}(\overline{p_i} - ar{p})^2/n}}{ar{p}}$$

Figure <u>8</u> shows how documents are captured when CV varies, where x-axis is the capture sequence, and y-axis is the document ID. If a marker appears at the position (10, 1,000), it means that the document labeled 1,000 is retrieved in the 10th capture occasion. Here we suppose that in each capture we will retrieve only one document. Figure <u>8</u>a has a small CV, hence almost all the documents are retrieved with similar probability. With the increase of CV, certain documents, most probably large ones, are retrieved more frequently, hence resulting in higher overlapping between the query results. We can see that when CV = 2.6, among the 50% of the small documents with document IDs greater than 10,000, only a few of them are retrieved in 200 captures. If we continue to use the estimator in Eq. <u>10</u>, a higher CV will induce a larger negative bias. Hence we need to adjust α in Eq. <u>11</u> according to the value of γ .



Scatter plots for various CVs. 200 random numbers within the range of 1 and 20,000 are generated in Pareto distribution

4.3 Estimate α using CV

To reveal the relationship between α and γ , we conducted a simulation study that produces a set of pairs (γ , α). We first generate random numbers that follow Pareto distributions with different exponents and cutoff values. Each distribution will have a different value for γ .

For each distribution, i.e., each γ , we randomly select some elements from the population, and keep record of the values for P and OR. After certain amount of elements are selected from the population, we obtained $\alpha = ln(1-P)/ln(OR)$ according to Eq. 11.

Figure 9 shows the relationship between α and γ . From the figure it is obvious that α is dependent on γ .

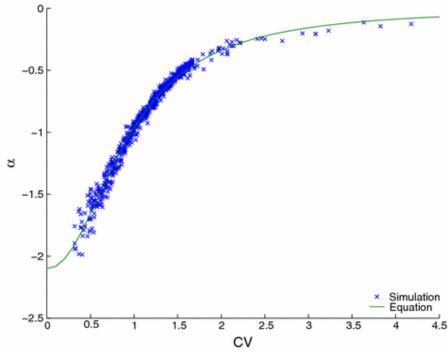


Fig. 9

Relationship between $CV(\gamma)$ and α

Next we need to find a formula to describe the relation. Note that α ranges between 0 and -2.1. When $\gamma = 0$, it is a uniform distribution, hence $\alpha = -2.1$. When γ increases, α will become closer to zero. Given those observations, we postulate that the following relation between α and γ holds, where a and b are to be determined.

$$lpha = rac{-2.1}{1+a\gamma^b}$$

To find the values for a and b, we run linear regression again on $ln(-2.1/\alpha - 1) = lna + bln\gamma$.

with the values for α and γ that are obtained in the simulation. The regression produces $ln(-2.1/\alpha-1)=0.1644+2.1408ln\gamma, R^2=0.9743$ i.e., $\hat{a}=1.1786, \hat{b}=2.1408.$

Thus we derived the equation
$$\alpha=\frac{-2.1}{1+1.1786\gamma^{2.1408}}$$
 Hence,
$$P=1-OR^{-2.1/(1+1.1786\gamma^{2.1408})}$$
 (12) and
$$\hat{n}=u\Big(1-OR^{-2.1/(1+1.1786\gamma^{2.1408})}\Big)^{-1}$$
 (13)

4.4 Simulation study

To evaluate the behavior of the estimator described in Eq. 13, we carried out a simulation study that includes 30 combinations of the following

- Six different CVs, i.e., 6.04, 3.83, 2.86, 2.32, 1.24, and 0.66.
- Five different sample sizes in terms of the fraction of data source size *n*, i.e., 0.05, 0.25, 0.45, 0.65, and 0.85.

For each combination, we run 100 trials, and record in Table 3 the mean estimation of the data size \hat{n} and its standard deviation. In each trial, we randomly generate numbers in the Pareto distribution within the range 1 to n, where n = 100,000. Other values for n are also tested and similar results are obtained. We adjust the exponent of the Pareto distribution and cutoff value so that various CVs are produced. For each CV and sample size we run 100 trials and obtained the estimation \hat{n} . Table 3 gives the simulation result.

Table 3

Simulation study for Eq. 13 with various CV and sample size. Mean \hat{n} and standard deviation (SD) are based on 100 runs. n = 100,000

Sample size (/n)

 CV^2

0.05 0.25 0.45 0.65 0.85

mean $\hat{n}(\times 10^5)$ 1.0481 0.9935 1.0086 1.0367 1.0613 6.04 *SD*(×10⁴) 1.1102 0.9952 1.0073 1.0340 1.0582 Mean $\hat{n}(\times 10^5)$ 1.0669 1.0220 1.0198 1.0327 1.0509 3.83 $SD(\times 10^4)$ 1.1355 1.0243 1.0180 1.0300 1.0471 Mean $\hat{n}(\times 10^5)$ 1.0747 1.0284 1.0234 1.0300 1.0423 2.86 $SD(\times 10^4)$ 1.1958 1.0317 1.0227 1.0272 1.0389 Mean $\hat{n}(\times 10^5)$ 1.0589 1.0262 1.0231 1.0275 1.0370 2.32 $SD(\times 10^4)$ 1.1809 1.0267 1.0218 1.0252 1.0337 Mean $\hat{n}(\times 10^5)$ 1.0447 1.0205 1.0188 1.0203 1.0243 1.24 $SD(\times 10^4)$ 1.2247 1.0248 1.0177 1.0169 1.0206 Mean $\hat{n}(\times 10^5)$ 1.0116 1.0068 1.0080 1.0140 1.0176 0.66 *SD*(×10⁴) 1.2388 1.0159 1.0068 1.0113 1.0143

From the table we can see that both the bias and standard deviation are rather small for all of those combinations, although in general smaller CVs demonstrate a smaller bias. In particular, small sample size such as 5% of n can also produce a rather accurate estimation, although the standard deviation is larger than that of the bigger sample sizes.

4.5 Estimate CV

One estimator for CV based on capture history is given by Chao et al. (1992):

$$\hat{\gamma}^2 = \hat{n_1} rac{\sum_{i=1}^n i(i-1)f_i}{t(t-1)} - 1,$$
 (14)

where \hat{n}_1 is the initial estimation for the data source size, f_i is the number of documents that are captured exactly i times, and t is the total occurrences of captured documents. Given that α is a value ranging between 0 and -2.1, we set the initial estimation for n_1 as: $\hat{n}_1 = u/(1 - OR^{-1.1})$. (15)

5 Experiments

This section evaluates our estimator in Eq. 13 using standard corpora. Experiments are implemented using Lucene (Hatcher and Gospodnetic 2004) on local data. We did not estimate the size of the real deep web data sources because the total number of documents in a real web data source is usually unknown or inaccurate, hence it is impossible to evaluate the estimation methods. An online demo of our estimation method is also available at http://cs.uwindsor.ca/~jlu/estimat (http://cs.uwindsor.ca/~jlu/estimat).

5.1 Data

We run our experiments on a variety of data collected from various domains. They are of different sizes ranging between 30 thousands to 1.4 millions. The corpora are Reuters, Gov2, Wikipedia, and Newsgroups, which are summarized in Table 4. These are standard test data used by many researchers in information retrieval. The file sizes of the corpora roughly follow the power law distribution. The vast majority of the files have very small sizes. On the other hand, there are also very large files, albeit the number of such files is small. Figure 10 shows the log—log plots of the occurrences against the file sizes of the documents in the corpora.

- *Wikipedia* is the corpus provided by wikipedia.org which contains 1.4 millions of English documents.
- Gov2 is a TREC test data collected from .gov domain during 2004, which contains 25 million documents. We used two subsets of the data for efficiency consideration.
- *Newsgroups* corpus includes posts in various newsgroups. We used two subsets of the corpus, named NG_1 and NG_2, respectively.
- Reuters is a TREC data set that contains 806,790 news stories in English (http://trec.nist.gov/data/reuters/reuters.html)
 (http://trec.nist.gov/data/reuters/reuters.html)).

Table 4Summary of test corpora. 'Number of words' is the number of distinct words that appear in both the documents and the Webster dictionary

Corpus name	Docs	Size in MB	Avg size (KB)	Mean #words	SD of #words
Wiki	1,475,022	1950	1.35	284	285
Gov2_1	1,077,019	5420	5.15	396	409
Gov2_2	1,075,270	5241	4.99	389	407
NG_1	1,372,911	1023	0.76	294	223
NG_2	30,000	22	0.73	290	180
Reuters	806,791	666	0.85	125	82

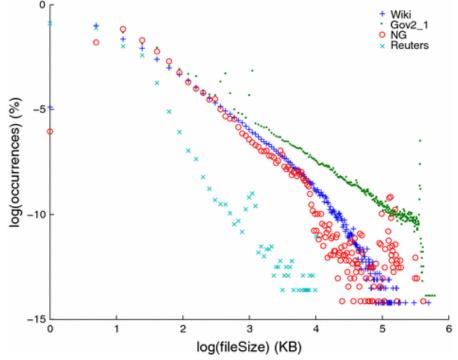


Fig. 10

File size distributions of the four corpora

5.2 Evaluation metrics

The method is evaluated in terms of relative bias, i.e., how far away the estimation is from the actual value, and relative standard deviation, i.e., how close those estimations are with each other.

Let n denote the actual size of the collection, \hat{n} the estimation. Suppose that there are m number of estimations. The expectation of \hat{n} , denoted as $E(\hat{n})$, represents the mean of m estimations, i.e., $E(\hat{n}) = (\hat{n_1} + \hat{n_2} + \cdots + \hat{n_m})/m$. Relative bias (RB) $(E(\hat{n}) - n)/n$ is to measure how close the estimations are to the actual size of the data source. Relative standard deviation (RSD) characterizes the variations of estimations, i.e., how close the estimations are to the center of the estimations. It is defined as

$$RSD = rac{1}{E(\hat{n})} \sqrt{rac{1}{m} \sum_{i=1}^{m} (\hat{n_i} - E(\hat{n}))^2}$$

Another important measurement is the cost of the estimation, including the number of queries sent, and the total number of document Ids retrieved. We call the number of examined document IDs the sample size.

5.3 Experimental setup

In the experiments, each query is a single word randomly selected from the Webster dictionary, which consists of 60,000 entries. For each query we will count all the matched documents. When a popular word, e.g., a stop word, is selected, it may match with most of the documents and make the overlapping rate extremely low, especially when the number of queries are small. Hence, we remove the top two percent of the most popular words from the query pool.

Our first experiment recorded in Table 5 studies 30 combinations of six data collections and five sample sizes in terms of query numbers that include 50, 100, 200, 400, and 800.

Table 5

Bias and standard deviation of the estimation over 100 runs. In each run queries are randomly selected from the Webster dictionary. Sample size is the fraction of n

Number of queries

Corpus

50 100 200 400 800

RB -0.09 -0.15 -0.11 -0.02 0.11

RSD 0.28 0.14 0.12 0.08 0.05

Wiki

Sample size 0.07 0.11 0.18 0.31 0.47

Mean $\hat{\alpha}$ -1.08 -1.05 -0.99 -0.91 -0.83

RB 0.09 -0.18 -0.22 -0.13 0.05

RSD 0.99 0.48 0.28 0.21 0.18

Gov2_1

Sample size 0.07 0.12 0.19 0.31 0.49

Mean $\hat{\alpha}$ -1.09 -1.06 -1.00 -0.90 -0.81

RB 0.19 -0.12 -0.17 -0.09 0.03

RSD 1.33 0.50 0.29 0.24 0.17

Gov2_2

Sample size 0.09 0.13 0.21 0.33 0.49

Mean $\hat{\alpha}$ -1.10 -1.06 -0.99 -0.89 -0.82

RB 0.30 0.08 0.13 0.21 0.28

RSD 0.64 0.28 0.16 0.10 0.05

 NG_1

Sample size 0.09 0.12 0.23 0.39 0.57

Mean $\hat{\alpha}$ -1.09 -1.06 -1.01 -0.95 -0.90

Number of queries

α			~
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C	orpus					
		50	100	200	400	800
	RB	0.45	0.14	0.06	0.12	0.21
NC o	RSD	0.78	0.30	0.13	0.10	0.06
NG_2	Sample size	0.06	0.11	0.18	0.30	0.49
	Mean \hat{lpha}	-1.1	-1.09	-1.05	-1.00	-0.95
	RB	0.74	0.03	0.00	-0.04	-0.06
Doutons	RSD	3.08	0.27	0.18	0.12	0.06
Reuters	Sample size	0.03	0.04	0.07	0.13	0.23
	Mean \hat{lpha}	-1.11	-1.11	-1.10	-1.09	-1.08

For each combination 100 trials are run, each with randomly selected queries from the Webster dictionary. For each set of 100 estimates generated, bias and standard deviation are computed. To have a clear comparison among corpora of different sizes, we record the bias and standard deviation relative to the data collection size.

Table 5 shows that smaller sample sizes (when the number of queries is 50) cause significant overestimation. This is in accordance with our simulation study as illustrated in Figs. 5 and $\underline{6}$. When sample size is too small, some queries may not have overlapping matches at all, hence resulting in overestimation. Also, the variance is large in both simulation studies and real data experiments. However, when the number of queries is equal or greater than 100, the estimation results are in general rather accurate. In particular, when the sample size increases, the standard deviation becomes smaller as expected.

In the file size distributions of the corpora experimented, some corpora have heavier tails than others, as shown in Fig. 10. For example, Gov2 has a flatter slope and heavier tail than Reuters, indicating that Gov2 has more large files than Reuters. A greater portion of large files will lead to higher degree of heterogeneity for capture probability, because those large files will be sampled repeatedly while the small files, albeit they are the

majority of the corpus, are seldom sampled. Thus flatter slope will cause lower values of $|\hat{\alpha}|$, which is confirmed by our experiment. In Table 5, Gov2 consistently has a smaller value of $|\hat{\alpha}|$ than that of Reuters.

5.3.1 Impact of sample size

Our second experiment reported in Table $\underline{6}$ focuses on Reuters corpus with more sample sizes. Again, each data item is obtained from 100 trials, each with a different set of words randomly selected from the Webster. In addition to bias and standard deviation, we also record the estimated values for γ and α . We can see that more queries will produce better results, and converge to the actual n.

Table 6Impact of sample size on Reuters corpus. Data are obtained from 100 trials

Queries	Mean sample size	Mean \hat{n}	RB	SD	Mean $\hat{\gamma}$	Mean $\hat{\alpha}$
50	22,335	1,476,418	0.830	1,757,016	0.81	-1.11
100	37,884	902,351	0.118	301,139	0.81	-1.11
200	67,503	816,023	0.011	170,302	0.82	-1.10
400	119,537	755,362	-0.064	82,819	0.84	-1.09
800	237,623	754,529	-0.065	55,199	0.85	-1.08
1600	478,303	778,159	-0.035	40,524	0.89	-1.06
3200	930,729	796,311	-0.013	25,271	0.93	-1.04

5.3.2 Impact of vocabulary

Previous experiments use the words selected from the Webster dictionary. Our next experiment tests the impact of vocabulary. We build three other vocabularies that consist of the words from very different domains:

- newsgroups: about 20k newsgroups postings;
- wsdl: about 2k web service description files;

• OHSUMED: about 50k medical documents.

Table <u>7</u> shows that four different vocabularies have little influence on the estimation result. In addition to the data reported here, we also experimented with other vocabularies from different areas, and observed similar behaviors.

Table 7

Impact of vocabulary on Reuters corpus. 200 query terms are randomly selected from different domains. Data are obtained from 100 trials

Vocabulary	$E(\hat{n})$	RB	SD
Webster	803,578	-0.004	156,320
Newsgroups	860,436	0.066	46,134
wsdl	931,883	0.155	94,160
OHSUMED	870,843	0.079	18,286

5.4 Comparison with other methods

Compared with Broder et al.'s method (Broder et al. 2006), which is computationally much more expensive, most of our results are better than the best reported 0.22 relative bias. Broder et al.'s method also suffers from large variance.

For the logistic method described briefly in (Xu et al. <u>2007</u>), they tested on data collections with smaller sizes, which in general will result in a smaller bias than larger data collections. Even with those small data sets, the mean absolute error ratio (relative bias) reported in the paper is 0.15, which is higher than our method in most cases.

Shokouhi et al.'s CH_{reg} method is the closest to ours in that their method is also based on the multiple capture—recapture method. In addition, they also use document ids only, hence saving the effort of document downloading and parsing. Table $\underline{8}$ shows the comparison of the two methods. For OR method, 100 queries are issued and all the matched documents are accounted for. More queries will produce more accurate estimations. For CH_{reg} method, the parameters are set as described in the paper (Shokouhi et al. $\underline{2006}$), i.e., 5,000 queries are issued and only the top k documents, where k=10, are selected. We find that both the value of k and the ranking method affect the estimation result greatly. We used the ranking method that produces the best

estimation, i.e., the default ranking by Lucence search engine. Larger k will result in larger positive bias.

Table 8

Comparison of OR and CH_{reg} methods. Each data item in the table is obtained from 100 runs

Corp	us	CH_{reg}	OR
Wiki	RB	0.91	-0.15
WIKI	RSD	0.03	0.14
Gov2_1	RB	-0.41	-0.18
G0V2_1	RSD	0 0.03 -0.41 0 0.01 -0.41 0 0.01 1.35 0 0.04 0.08	0.48
Gov2_2	RB	-0.41	-0.12
G0V2_2	RSD	0.01	0.50
NG_1	RB	1.35	0.08
110_1	RSD	0.04	0.28
NG_2	RB	0.08	0.14
110_2	RSD	0.01	0.30
Reuters	RB	0.56	0.03
_1041010	RSD	0.03	0.27

From Table $\underline{8}$ it can be seen that our *OR* method has much smaller bias in most of the cases. CH_{reg} works well only when the data source is small, i.e., corpus NG_2 which contains 30,000 documents. The standard deviation of CH_{reg} method is smaller because in each run it uses a large portion of the words (5,000) out of a vocabulary of size 60,000. In the contrary *OR* method uses only 100 queries.

In addition to the accuracy, our method is also more efficient than Shokouhi et al.'s method in terms of both the number of queries and the sample size, i.e., the total number of documents retrieved. We tabulate the correspondence of number of queries and sample size in the case of Reuters in Table <u>6</u>. It shows that in average 100 queries result in 37,884 number of documents in average, which is smaller than 50,000 number of documents retrieved in Shokouhi et al's method.

We also tried Chao's classical estimation method in ecology (Chao and Lee <u>1992</u>). As noted in the literature and confirmed by experiments on our data, the method works well only when the sample size is large and CV square is small. For example, when CV is around one, the sample size needs to be $4 \times n$. Obviously this is too costly for data source size estimation.

6 Discussions and conclusions

This paper proposes a novel method to estimate the size of deep web data sources by sending random queries and analyzing returned document ids.

Our first estimator in Eq. <u>10</u> is meant for uniformly distributed samples. It is analytically derived and empirically verified by simulation. The simulation study shows that it has smaller bias and variation than the classical Schumacher et al.'s estimator (Schumacher and Eschmeyer <u>1943</u>; Darroch <u>1958</u>).

Our second estimator in Eq. 13 is derived for samples obtained by queries. It is verified by extensive simulation studies with various degrees of heterogeneity of catching probability and different sample sizes. In applications on various data collections, the estimator also demonstrates a small bias and variance compared with other estimation methods for data collection size.

In addition to the accuracy of the estimation, one salient feature of our method is its efficiency. We only need to issue random queries, instead of carefully selected ones. In most of the cases, only 100 queries are needed, although more queries will result in higher accuracy.

One limitation of our method is the assumption that all the matched document IDs can be retrieved. Very large data sources, such as general-purpose search engines, usually impose a limit on the maximal number of matched results that can be returned for practical consideration. When a query matches more documents than the limit k, i.e., when the query overflows, the matched documents are ranked and only the top-k are returned. This strategy will cause the rank bias. While our method solves the query bias quite well, the rank bias is not tackled in this paper. Nonetheless, our method has a wide area of applications such as (1) data sources that do not impose a return limit; (2) data sources that are not very large, hence most queries will match less than k number of documents; and (3) data sources that do not rank the returning results.

Our study shows that when there are not many overflowing queries, our method still works well by ignoring the overflowing queries. When the density of overflowing queries is high, filtering out those queries will induce another bias, because rare words usually introduce higher overlapping rate. Hence our method is only applicable to ranked data sources with low overflowing density. From this perspective, our method is suitable for relatively small text databases, depending on the result limit. To estimate the size of large search engines such as Google, throwing away overflowing queries is no longer an option since most of the random queries will match more pages than the 1,000 limit Google sets.

On the other hand, with the proliferation of web services and programmable web interfaces, which typically contain data sources of relatively small size, there is a growing demand for probing those data sources to learn their profiles, such as their size. We also expect that the method can be applied in other areas where ranking bias does not exist, such as the estimation of the number of values of an attribute in a database relation (Haas et al. 1995).

It is not easy to obtain the detailed performance comparison with the methods based on document analysis (Broder et al. 2006, Bar-Yossef and Gurevich 2006, 2007; Bharat and Broder 1998). All those works target very large data sources such as the size of the major search engines and the entire web, while we provide a convenient estimator for relatively small data sources. Compared with this group of work, the main advantage of our method is its efficiency-it does not need to download and analyze the documents; does not need to carefully select the uncorrelated queries; and does not need to send out many queries.

Future work on document id based estimators may focus on correcting the rank bias so that our method can be applied to very large data sources. In addition, we need to investigate biases caused by other sources, such as duplicate documents and documents that do not contain English words.

Acknowledgments

We would like to thank reviewers for their insightful comments, and Jie Liang for providing the query interface for the corpora. The research is supported by NSERC (Natural Sciences and Engineering Research Council Canada), SSHRC (Social Sciences and Humanities Research Council Canada), and State Laboratory for Novel Software Technology, Nanjing University.

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