



www.bioinformatics.net
Volume 20(10)

Research Article

Received October 1, 2024; Revised October 31, 2024; Accepted October 31, 2024, Published October 31, 2024

DOI: 10.6026/9732063002001418

BIOINFORMATION 2022 Impact Factor (2023 release) is 1.9.

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Citation: Seth *et al.* Bioinformatics 20(10): 1418-1423 (2024)

Evaluation of pre-analytical errors using six sigma metrics towards quality of laboratory performance in an Indian tertiary care hospital

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Abstract:

Six Sigma methodologies is one of the standard management system adopted to check the quality and improve laboratory performance continuously by assessing the laboratory process using six sigma metrics. A one-year prospective study from January 2019 to December 2019 was conducted in the laboratory medicine department of a tertiary care hospital. A total of 162974 samples were received from both inpatient and outpatient departments, and total laboratory errors were found to be 1144, with the maximum errors identified in the pre-analytical phase coming out to be 978 (0.6%). The types of errors in the pre-analytical phase after being identified and categorized were hemolyzed samples, clotted samples, phlebotomy errors, *etc.* The most typical error identified was clotted sample error (0.29%) followed by haemolysed sample error (0.20%). After calculating the error rate or defects per million error samples on a Sigma calculator, the value obtained was between 4 and 5, an average class quality. Our study concludes that continuous and repeated recognition of errors and evaluating them using Six Sigma metrics can help achieve overall laboratory procedure quality and better clinical diagnosis.

Keywords:

Pre-analytical errors, six sigma metrics, laboratory quality, total laboratory testing, laboratory errors & quality indicators

Aim:

The main aim of our study is to identify the pre-analytical errors and laboratory quality assessment based on these errors using the Six Sigma methodology.

Abbreviations:

EDTA = Ethylene diamine tetraacetic acid

TTP = Total testing process

DPM = Defects per million

LIS = Laboratory Information System

HINAI = Software of LIS

Background:

Total Laboratory testing (TTP) is a process that starts with raising a test request of an individual by a clinician, followed by sample collection at the laboratory end for testing and ending with test reports of that individual after testing [1]. As laboratory results play a crucial role in disease diagnosis, it is vital to guarantee the reliability of the results furnished by the clinical laboratory [2]. Errors in this process can disprove the test results concerning the patient's health status [1]. According to many studies, 70% of medical decisions rely on the accuracy of laboratory tests, and the whole laboratory testing cycle has three phases: Pre-analytical, Analytical, and Post-analytical [3-4]. Over the years, the considerable changes leading to advancements in automation, sample collection, transport, and dispatch of results have led to a substantial change in the working and quality of performance in the laboratories with a decline in the errors to a remarkable level [2]. These advancements have provided enormous help in clinical decision-making by supporting, preventing, diagnosing, and therapeutic monitoring of human disorders [5]. However, achieving 100% accuracy and precision is far beyond the approachability [2]. Any defect from ordering tests to reporting results and appropriately interpreting and reacting to these defects is known as Laboratory error. These errors, which arise at any phase, can lead to misdiagnosis, mistreatment, *etc.*, [5]. A remarkable decrease is seen in the

analytical phase due to the advanced instrument technology compared to the pre-analytical phase error [6], which is reported to be around 46 - 68%, according to some studies [4]. Some other studies showed that 61.9% of errors arose in the pre-analytical phase, while 15% of errors were in the analytical phase, and 23.1% were in the post-analytical phase [6]. As evident, the laboratory bears the maximum burden of errors arising in the pre-analytical phase; its regulation and continuous monitoring arise because of unmanageable pre-analytical variables such as too many medical professionals involved, including physicians, nursing, transport staff, phlebotomists, *etc.* is crucial [3]. Therefore, it is of much concern to identify the viable areas of errors in this phase and take corrective steps at repeated intervals to reduce them and, in turn, provide high-quality, reliable test results and safe health care [6]. The areas of errors identified further need to be checked on quality by measuring the defects based on the Six Sigma scale. Six Sigma methodologies were introduced into the industry and business as early as the 1980s and were developed by Motorola, Inc. This methodology measures the error rate or defects expressed as defects per million (DPM) in the pre-analytical phase. It thus monitors the outcome process by converting the errors or defects to sigma metrics using a standard table available in any six-sigma text. Laboratory processes with poor outcomes are

counted as errors. Higher value validates the good quality laboratory report results. Six Sigma metrics from 6.0-3.0 represent the range from “best to worst case quality. Six Sigma is a world-class quality process, with the 4 Sigma value considered the average class quality performance [2, 7]. The objective of the present study is to enumerate the types, estimate the frequency, and evaluate the quality of the identified pre-analytical phase errors observed during the 1- year study period in the Department of Laboratory Medicine.

Materials and Methods:

The present descriptive study was conducted in the central Laboratory, Department of Laboratory Medicine, Shri Mata Vaishno Devi Narayana Super Specialty Hospital, and a tertiary Care Hospital. A total of 162974 samples were received randomly, and 761955 tests were performed from both outpatient and inpatient departments for pre-defined pre-analytical errors during the 1-year duration from January 2019 to December 2019. Phlebotomists collected outpatient samples, and inpatient samples were collected by nursing staff and transported to the laboratory by supporting staff. At the sample receiving desk, the person assigned for sample receiving checks for errors and makes entries in the sample rejection register as per standard operating procedure. This data was evaluated monthly using percentages and Six Sigma scale methodology for laboratory testing performance quality checks.

Six sigma values were calculated by first calculating the DPM rate using the following formula:

$DPM = \text{number of errors} \times 1,000,000 / \text{total number of samples}$

After calculating DPM, the rate was converted to a sigma value based on the sigma score calculators available online at <http://www.westgard.com/six-sigma-calculators.htm>.

Depending upon the Six Sigma value, the performance level of the laboratory in the pre-analytical phase was evaluated.

Results:

The present study was conducted in the Central Laboratory during the 1- year duration from January 2019-December 2019. A total of 162974 samples were received, and 761955 tests were performed from both outpatient and inpatient departments. Out of 162974 samples received, as per the LIS system of the hospital, the total number of samples found to have laboratory errors is 1144 (0.7%), with 978 (0.6%) errors identified in the pre-analytical phase along with 150 (0.10%) and 16 (0.01%) in both analytical and post-analytical phases (Figure 1).

Table 1: The various pre-analytical errors observed in the present study were summarized as follows:

S. No.	Type of Pre-analytical error(s)	Frequency
1	Clotted samples	325 (0.29%)
2	Hemolysed samples	282 (0.20%)
3	Phlebotomy error	128 (0.18%)
4	Wrong sample	85 (0.10%)
5	Wrong Collection	82 (0.10%)

6	Wrong billing	61 (0.04%)
7	Lipemic samples	15 (0.01%)

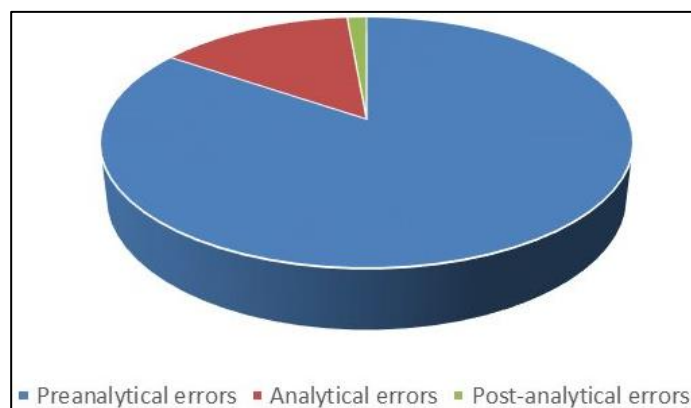


Figure 1: Error percentage in different phases of total laboratory process

The hospital's LIS system (HINAI) made it easy to trace the sample right from the start, *i.e.*, from when the clinician ordered the test until receiving it. The errors identified and included as quality indicators for the improvement of total laboratory quality were: -

- [1] Hemolysed sample
- [2] Clotted sample
- [3] Phlebotomy error
- [4] Wrong sample
- [5] Wrong billing
- [6] Lipemic
- [7] Wrong collection

Month-wise percentage data of pre-analytical errors per error (Figure 2) and their Six Sigma value (Figure 3) were calculated and significant changes were seen in the month of July (12.2% and 4.7).

The calculated percentage and Six Sigma value of the quality indicators identified and rejected in our laboratory were 28.8% (4.5) of hemolysed samples, and 33.2% (4.4) were clotted due to improper mixing or delayed transport. 13.1% (4.7) errors were due to incorrect phlebotomy, 8.7% & 8.4% (4.8 each) samples were rejected because of wrong samples (tube and cap interchange or samples not matching with advised test), and wrong collection (biochemistry sample collected in EDTA tube or vice versa and saliva instead of sputum in case of microbiology tests). 6.24% (4.9) of wrong billing errors were either due to missing tests as requested by the doctor or test requests were wrongly billed. 1.5% (5.3) of lipemic samples was rejected over one year of the study. (Table 1) (Figure 4 & 5). The month-wise distribution of different pre-analytical errors identified is given in Table 2.



Figure 2: The percentage of pre-analytical errors per error observed distribution month-wise

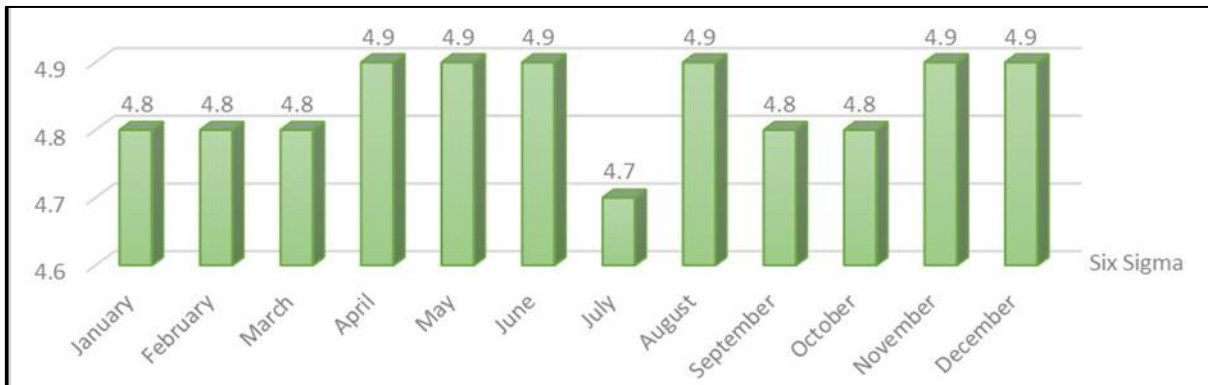


Figure 3: The six-sigma value for the pre-analytical errors calculated month-wise

Table 2: Further, the distribution of different pre-analytical errors month-wise was presented as follows:

Type of Pre-analytical Errors	Month											
	Jan	Feb	Mar	Apr	May	Jun	Jul	Aug	Sep	Oct	Nov	Dec
HEMOLYSED	24	24	29	22	22	23	35	29	21	26	11	16
CLOTTED	29	11	28	27	27	32	48	3	31	17	36	36
PHLEBOTOMY ERROR	15	22	9	16	3	5	12	14	10	19	1	2
WRONG SAMPLE	4	17	1	2	2	7	11	5	6	8	7	15
WRONG BILLING	6	7	6	0	0	1	1	5	10	11	10	4
LIPEMIC	2	2	2	1	1	2	0	1	0	2	0	2
WRONG COLLECTION	7	5	7	4	4	5	12	11	5	9	10	3

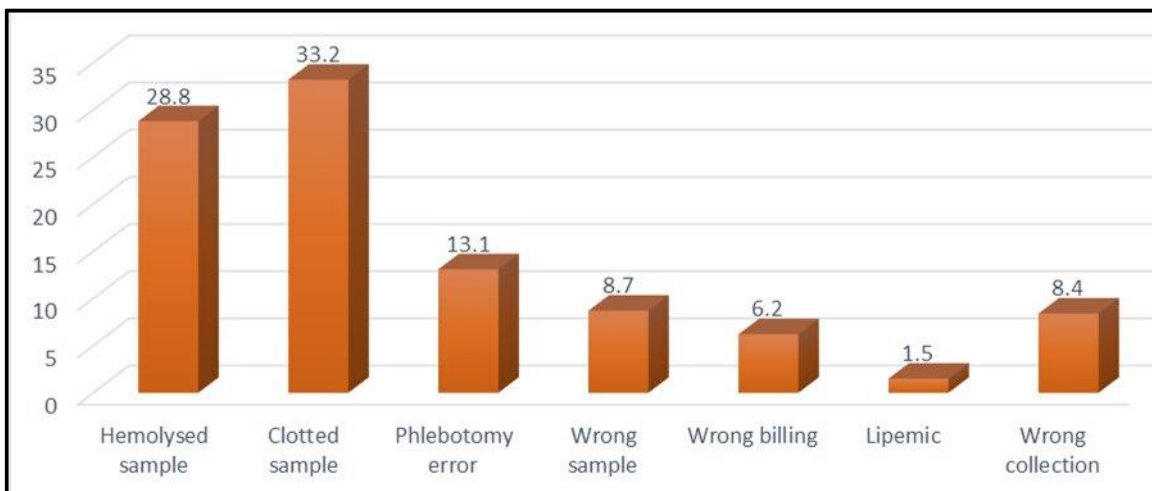


Figure 4: The percentage of each of the pre-analytical errors per total pre-analytical errors over 12 months was calculated and shown in the figure

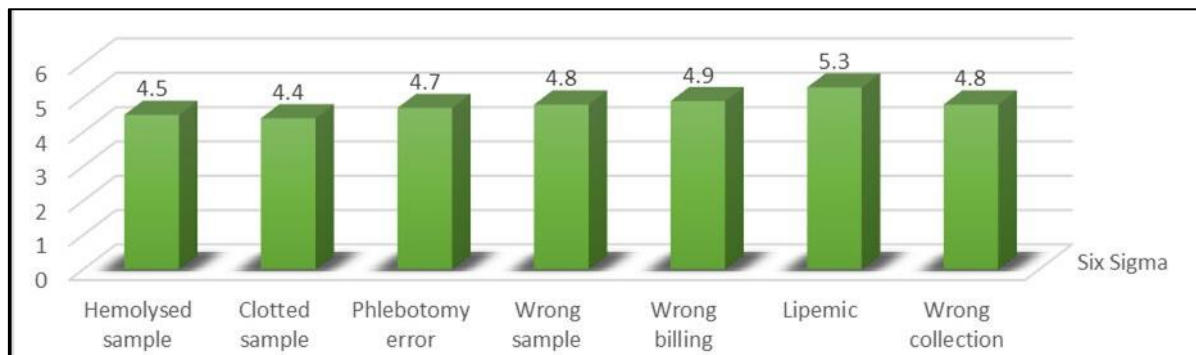


Figure 5: The six-sigma value calculated for each pre-analytical error over 12 months was calculated and shown in the figure.

Discussion:

Though the laboratories adopt state-of-the-art facilities, primarily concerned with a fully automated system to give appropriate patient care decisions, there are still laboratory diagnostics errors [8]. The total error rate in the present study was 0.7%, which is in the range of 0.1 – 9.3%, as suggested by Carraro and Plebani for good laboratory care [4]. In the present study, the pre-analytical error was found to be 85.5%, an analytical error was 13%, and the post-analytical error was 1.4% concerning total errors. A study on laboratory errors showed that pre-analytical errors were 81% and analytical errors 10%, primarily human and technical errors. [9]. In the current location, we first conducted a study pointing out the types and frequencies of pre-analytical errors with a high occurrence of pre-analytical errors in concordance with the previous studies [10, 11]. Most pre-analytical errors were due to clotted samples, hemolysed samples, and phlebotomy errors. Hemolysis might be caused by temperature variations, incorrect sample collection (forceful dispensing of the sample through fine needle, vigorous shaking, and improper centrifugation), untrained staff, and improper storage and transport. It is very pertinent to remind phlebotomists and lab staff that hemolysis is one of the most common causes of pre-analytical errors, causing considerable harm to the accuracy of analytical tests [12, 13]. A similar study done in a laboratory of a tertiary care hospital also showed a higher percentage of hemolyzed sample errors (33.2%) [14]. It is evident from the study's findings that medical and paramedical professionals play a substantial role in the causation of pre-analytical errors, demanding collaboration as key to improving laboratory results quality [15]. Clotted samples accounted for rejection of 0.29% in the present study, possibly due to improper sample mixing with clot activator. This is in accordance with the study done by Arul *et al.* which reported that clotted samples (0.12%) were the second most common error, which might be primarily due to inappropriate mixing of samples after collection [16]. Improper mixing and/or under filled/overfilled EDTA tubes might introduce pre-analytical errors as the latter changed the sample to additive proportion and consequently impacted the results' quality [17, 18].

Wrong sample and collection accounted for a rejection of 0.1% each, and wrong billing accounted for a rejection of 0.04% in the

present study. A survey by Dudani mentioned that though collecting samples from the wrong patient accounted for a small proportion of sample collection errors, it was a considerable concern for laboratory settings. Wrong billing or data entry non-conformances might arise due to incorrect name and age spelling and incomplete contact information. Inappropriate investigation requests occurred because of unreadable writing on the request slips and deficient medical knowledge linked with insufficient training [19]. Laboratory experts should conduct sensitizing programs for treating clinicians and urge the bedside phlebotomist to take care of the appropriate filling of test request forms to help laboratory workers better validate. Adopting computerized test requests by clinicians, the lab lean process, and introducing Six Sigma rules could reduce the frequency of pre-analytical errors [20]. Again, lipemic samples accounted for rejection in the present study was 0.01%. In a study done by Chawla *et al.* it was reported that the rejection of samples due to lipemia was 0.03% and 0.11% in the admitted patients and outpatients, respectively [5]. Lipemia samples might be due to interference of heavy metals, or a patient might suffer from hyperlipoproteinemia disorder. It became mandatory for both the clinician and phlebotomist to ensure that proper patient preparation should be introduced before sample collection [21]. However, lipemia could be avoided by taking overnight fasting samples [5]. Several studies also reported that the significant interference of lipemia in the laboratory analysis and ultracentrifugation procedure might help reduce lipemia in samples [22, 23]. In the present study, we calculated the Six Sigma value for 1-year study and different pre-analytical variables. We did our best in this quality check by adopting Six Sigma metrics of value 4.0 as average class quality performance and less than 3.0 as unsatisfactory. As our Six Sigma values of most of the parameters were between 4 – 5 sigma, which is average class quality, there is a necessity for rigorous quality checks in a pre-analytical zone by identifying and taking care of the gaps in the pre-analytical stage and eventually refining the quality and performance of clinical laboratory by achieving world-class performance of 6.0 sigma value. Various studies adopted Six Sigma Metrics to point out pre-analytical errors and improve the quality and performance of the laboratory [24, 25]. Training the laboratory technicians, nursing staff, and interns frequently on sample collection, transport, and appropriate

filling of test request forms with all required information is necessary. Making mistakes is human nature, and those who correct the errors are good humans [26].

Conclusion:

The quality of laboratory performance for identifying, categorizing the pre-analytical errors as quality indicators and for interpreting the Six Sigma value is of concern. This process helped us take remedial actions needed like training technicians, nurses, and doctors to avoid and reduce the number of errors and eventually improving the total quality management in the laboratory.

Acknowledgments:

I would also like to extend my heartfelt thanks to all the laboratory technicians and non-technical staff, especially Ms. Asha, for their understanding of our research work and their constant help in compiling and completing the work. Their contributions are greatly appreciated. Lastly, I want to express my sincere thanks and deep gratitude to my colleagues, Dr. Neetika and Dr. Rita, for their endless help and support throughout my research work. Their guidance and encouragement have been invaluable. I would like to take this opportunity to express my gratitude to the Almighty for his boundless blessings, which have enabled me to successfully complete my research work. I would also like to extend my heartfelt appreciation to the laboratory technicians and non-technical staff, particularly Ms. Asha, for their unwavering support and understanding of our research work. Their invaluable contributions in terms of compiling and completing the work are deeply appreciated. Furthermore, I would like to extend my sincere thanks and profound gratitude to my colleagues, Dr. Neetika and Dr. Rita, for their continuous assistance and encouragement throughout my research work. Their unwavering support, guidance and input have been instrumental in the successful completion of my research work. Once again, thank you all for your invaluable support and cooperation.

References:

[1] Kaplan LA & Pesce AJ. *Clinical Chemistry: Theory, analysis, correlation*. (2009) P:1200, Location: USA, Publisher: Mosby

- [2] Aysenur A *et al. Biochemia Medica*. 2014 **24**:376[PMID: 25351356]
- [3] Patra SMD *et al. International Journal of Bioassays*. 2013 **2**:551
- [4] Dereen N. *PLoS One*. 2017 **12**:e0170211.[PMID: 28107395]
- [5] Chawla R *et al. LABMEDICINE*. 2010 **41**:89.[DOI:10.1309/LM9JXZBMLSVJT9RK]
- [6] Neogi SS *et al. J Clin Sci Res*. 2016 **5**:171[DOI:10.15380/2277-5706.JCSR.15.062.]
- [7] Burtis CA *et al. Tietz Textbook of Clinical Chemistry and Molecular Diagnostics*. W B Saunders Co Ltd; 2448 pages (2005) New Delhi
- [8] Sakyi AS *et al. Annals Med Health Sci Res*. 2015 **5**:8.[PMID: 25745569]
- [9] Carraro P & Plebani M. *Clin Chem*. 2007 **53**:1338.[PMID: 17525103]
- [10] Abdollahi A *et al. N. Am. J. Med. Sci*. 2014 **6**:224.[PMID: 24926448]
- [11] Plebani M. *Clin Biochem Rev*. 2012 **33**:85.[PMID: 22930602]
- [12] Jay D & Provasek D. *Clin Chem*. 1993 **39**:1804[PMID: 8375051]
- [13] Carraro P *et al. Clin Chem*. 2000 **46**:306.[PMID: 10657399]
- [14] Wadhwa N. *Int J Clin Biochem Res*. 2020 **7**:354[DOI: 10.18231/j.ijcbr.2020.076]
- [15] Valenstein P & Meier P. *Arch Pathol Lab Med*. 1999 **123**:1145[PMID: 10583917]
- [16] Arul P *et al. J Lab Physicians*. 2018 **10**:237.[PMID: 29692594]
- [17] Gupta M *et al. Biochem Physiol*. 2015 **4**:181.[DOI: 10.4172/2168-9652.1000181]
- [18] Da Rin G. *Clin Chim Acta*. 2009 **404**:68[PMID: 19302988]
- [19] Dudani S. *Int J Med Lab Res*. 2018 **3**:27.
- [20] Munilakshmi. U *et al. Int J Clin Biochem Res*. 2018 **5**:467.[DOI:10.18231/2394-6377.2018.0099]
- [21] Naz S *et al. Pak J Med Res*. 2012 **51**:27.
- [22] Jabbar J *et al. J Pak Med Assoc*. 2006 **56**:59.[PMID: 16555635]
- [23] Calmarza P & Cordero J. *Biochem Med*. 2011 **21**:160.[PMID: 22135856]
- [24] Shah S *et al. Int J Med Res Rev*. 2014 **2**:140.[DOI:10.17511/ijmrr.2014.i02.20]
- [25] Mukhopadhyay T *et al. Int J Health Sci Res*. 2021 **11**:171.
- [26] Kulkarni S *et al. Indian J Clin Biochem*. 2018 **33**:102.[PMID: 29371778]