A comparison of 24-hour urine versus random urine samples for determination and quantification of Bence Jones protein in a South African population

1	B	V
-	_	y

DR Ashandree Reddy

Submitted in partial fulfilment of the requirements for the degree of Master of Medicine in Chemical Pathology in the School of Pathology,

College of Health Sciences,

University of KwaZulu-Natal

April 2019

As the candidate's supervisor, I have approved this dissertation for submission.

Signed: DR V Gounden Name: DR Verena Gounden Date: 21/5/19

Declaration

- I, Dr Ashandree Reddy, declare as follows:
- (i) The research reported in this dissertation, except where otherwise indicated, is my original work.
- (ii) This dissertation has not been submitted for any degree or examination at any other university.
- (iii) This dissertation does not contain other persons' data, pictures, graphs or other information, unless specifically acknowledged as being sourced from other persons.
- (iv) This dissertation does not contain other persons' writing, unless specifically acknowledged as being sourced from other researchers. Where other written sources have been quoted, then:
- a) their words have been re-written but the general information attributed to them has been referenced;
- b) where their exact words have been used, their writing has been placed inside quotation marks, and referenced.
- (v) Where I have reproduced a publication of which I am an author, co-author or editor, I have indicated in detail which part of the publication was actually written by myself alone and have fully referenced such publications.
- (vi) This dissertation does not contain text, graphics or tables copied and pasted from the Internet, unless specifically acknowledged, and the source being detailed in the dissertation and in the References sections.
- (vii) My contribution to the project is as follows:

Literature review, research protocol development, ethical and hospital management approval, data collection and processing, statistical analysis, interpretation of data and journal article first draft write-up.

- (viii) The contributions of others to the project are as follows:
- 1) Supervisor: Dr Verena Gounden: Formation of research idea and literature review, research protocol development, statistical analysis, interpretation of data and review of first draft write-up
- 2) Co-supervisor: Dr Nadine Rapiti: Research protocol review, data collection, review of manuscript

Dr Ashandree Reddy:

ii

Date: 21/5/19

Dedication

Dedicated to my husband and family were strength, wisdom and comfort flow freely.

Acknowledgements

I would like to acknowledge and thank the following:

- 1. National Health Laboratory Service K funding for funding of this study
- 2. Sunitha Sathabridg, a technologist at National Health Laboratory Service based at Inkosi Albert Luthuli Central Hospital for assistance in analysis of the urine protein electrophoresis.
- 3. My supervisor, Dr Verena Gounden, for her guidance and assistance.

Overview of the thesis

Introduction:

Multiple myeloma is a hematological cancer that has a high incidence of relapse despite intensive treatment regiments. The laboratory plays a vital role in treatment response evaluation. Measuring urine Bence Jones protein(BJP) is amongst the tests used to monitor response. The International Myeloma working group (IMWG) and College of American Pathologists recommend a 24-hour collection for BJP. Although a 24-hour urine collection is a definitive means to determine BJP excretion, it has several issues related to sample collection and is prone to inaccuracy. Protein to creatinine ratios have demonstrated good correlation with 24-hour urine. The aim of this study was to compare measured 24-hour urine collection to random urine for the quantitation of BJP in a South African population.

Method:

Known patients with multiple myeloma(MM) collected 24-hour urine as part of their routine clinical assessment for BJP, random urine samples were submitted following completion of the 24-hour collection. The measured 24-hour urine BJP was then compared to 2 estimated 24-hour BJP excretions. The estimated excretions were calculated as follows;

Estimation 1 (E1): Estimated 24-hour BJP (mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) X10,

Estimation 2 (E2): Estimated 24-hour BJP (mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) x 15mg/kg for women or x 20mg/kg for men.

All the 24-hour BJP results were classified according to IMWG treatment response criteria.

Results

When using the Wilcoxon paired test analysis, the measured 24-hour urine BJP was significantly different to both the E1 (p=0.049) and E2 (p=0.049) equations. But analysis following categorization of each patient per IMWG response criteria, indicated no significant difference in classification of treatment response using either the E1 or E2 estimation equations (P=0.69).

Conclusion:

24-hour urine collections are cumbersome. Random urine BJP estimates are simple, rapid and inexpensive. This study demonstrates that both the estimates of 24-hour BJP can be used to monitor response in patients with MM. This can be added to the body of evidence that random samples can be used to monitor patients' treatment response in MM.

Table of Contents

Declaration	i
Dedication	ii
Acknowledgements	iv
Overview of the thesis	
Table of Contents	vi
Part 1: The Review of Literature	1
Part 2: A submission ready manuscript.	
Appendices	XXX
Appendix 1: The Study Protocol	XXXI
Appendix 2: The Guidelines for Authorship for the Journal selected fo manuscript	
Appendix 3: Ethical approvals	LXI
Appendix 4: Data collection tools and consent forms	LXV
Appendix 5: Raw data	LXX

Part 1: The Review of Literature

Introduction

Multiple myeloma is a hematological cancer that develops in bone marrow and has a high incidence of relapse despite intensive treatment regiments. The laboratory plays a vital role in treatment response evaluation. Multiple myeloma has traditionally been based on the evaluation of serum and urine monoclonal protein concentrations via protein electrophoresis and immunofixation for response evaluation. The identification of paraprotein acts as surrogate for the tumour burden. [1]

We are continuously searching for reliable biomarkers in the hope that this could simplify and improve accuracy of medical decisions. This study reviewed the urine protein electrophoresis to determine if the measured 24-hour urine collection is comparable to random urine for the quantitation of Bence Jones protein (BJP) in a South African population. In doing so, we hoped to add to the body of existing knowledge and use random urine samples to monitor treatment response in patients known with multiple myeloma. A random urine sample may improve patient compliance to investigations, management and quality of life whilst on treatment.

The International Myeloma Working Group (IMWG) and College of American Pathologists recommend a 24-hour collection for Bence Jones proteins(BJP). [2-3] Although a 24-hour urine collection is a definitive means to determine BJP excretion, it has several issues related to sample collection and is prone to inaccuracy. [4-7] Protein to creatinine ratios have demonstrated good correlation with 24-hour urine. [4-6]

We intend with this literature review to examine the updated definition of multiple myeloma as well as review updated diagnostic criteria and treatment response criteria. We will also review literature that have performed similar studies in different populations.

Multiple Myeloma

Multiple myeloma is a hematological cancer that develops in bone marrow. It belongs to a group of diseases known as monoclonal gammopathies which are characterized by the neoplastic expansion of a single clone of plasma cells. Plasma cells are responsible for immunoglobulin production. The single clone malignant cell proliferation differs to the normal polyclonal distribution of plasma cells in a healthy individual. Monoclonal gammopathies produce increased amounts of a single clone of immunoglobulins that result in a disproportionate fraction of the number of plasma cells in the body. [2]

Examples of monoclonal gammopathies, also known as plasma dyscrasia's, include; multiple myeloma (MM), monoclonal gammopathy of undetermined significance (MGUS), smoldering multiple myeloma, solitary plasmacytomas, AL amyloidosis (light chain), Waldenstroms macroglobulinaemia and POEMS syndrome. Each of the monoclonal gammopathies have their own specific diagnostic criteria. These monoclonal gammopathies result from an abnormal overproduction of a single abnormal clone of a plasma cell or B lymphocyte resulting in increased production of either the intact immunoglobulin, free light chain component or both. The presence, level and type of the immunoglobulin have vital implications in diagnosis, staging and treatment of these diseases. [2]

Epidemiology

While age-standardized incidence rates differ with ethnicity from 3.9/100 000 in Chinese to 12.7/100 000 in African individuals, the data suggest that MM poses a significant worldwide health problem. [8] It consists of only 1% of all cancers but 10% of hematological malignancies in the United States. [9]

MM was responsible for 0.43% of newly diagnosed cases of malignancies in South Africa in 1999 with the incidence being reported at approximately 0.00054%. [10] While the incidence is highly variable among countries, studies indicate that the incidence of multiple myeloma has increased uniformly since 1990 with the largest increase in middle and low-middle income countries. [8,11]

The prevalence of multiple myeloma is higher in HIV positive compared to uninfected individuals. [12] This increases the disease burden of multiple myeloma in South Africa which has the largest HIV epidemic in the world having 7.1 million people living with HIV. [13] Multiple myeloma is also a disease that is more prevalent in the elderly with a median age of diagnosis being 65-70 years old. [8]

Pathophysiology and Diagnosis

The pathophysiology of multiple myeloma is complex. Briefly, the plasma cells, which originate from post follicular B cells, are characterized by complex chromosomal abnormalities. This results in dysregulation of oncogenes and suppressor genes. There is increasing evidence that suggests that the bone marrow microenvironment of tumor cells also perform a crucial role in the pathogenesis. The imbalances between receptor activator of nuclear factor kappa-B ligand (RANKL) and osteoprotegerin together with osteoclast activity factors significantly contribute to the development of myeloma bone disease. [15]

The pathophysiologic basis for the clinical sequelae of multiple myeloma comprises the skeletal, hematologic, renal, and nervous systems. Multiple myeloma is typically associated with four major dysfunctions i.e. Calcium, Renal, Anemia and Bone lesions, referred to by the acronym CRAB. Patients often have hypercalcemia due to bone lytic lesions caused by plasma cell tumours outgrowing their confined space in bone marrow and invading the bone. The breakdown of bone results in the release of calcium and hence, higher levels of serum calcium. The elevated serum calcium is defined as > 2.75mmol/L or > 0.25mmol/L of the upper reference limit. Patients can present with neurological features due to hypercalcaemia. Renal insufficiency results from deposition of free light chains in renal tubules. The monoclonal free light chains obstruct the renal tubules causing significant renal damage. The creatinine clearance <40 mL per min or serum creatinine >177 µmol/L is diagnostic of renal insufficiency. Anaemia is a consequence of overgrowth of plasma cells leading to crowding out of the cells that produce red blood cells. Erythropoietin production is also reduced in renal failure and contributes to the anaemia in patient with multiple myeloma. Anaemia in multiple myeloma is defined as a haemoglobin value of >2 g/dL below the lower limit of normal, or a haemoglobin value <10

g/dL. Bone lesions result in fractures often in the axial skeleton. The bone lytic lesions are caused by expansion of plasma cells in the bone marrow and is defined as one or more osteolytic lesions on skeletal radiography, computer tomography (CT), or positron emission tomography-CT. [2,15]

Understanding the natural progression of the multiple myeloma is important in order to appreciate the important role that the laboratory plays in the management of a patient with multiple myeloma. It is associated with significant mortality and morbidity and is considered largely incurable and fatal without treatment. With the introduction of new classes of effective drugs for the treatment of multiple myeloma, improved frequencies and degree of patient response has been observed. Many treatments have been shown to significantly prolong survival and simultaneously improve the quality of life for patients. Unfortunately, all patients will ultimately relapse after treatment and will require change to a more responsive therapy. This necessitates regular periodic monitoring of disease in order to detect relapse. Laboratory testing plays a vital role in monitoring response to treatment as well detecting relapse in patient on treatment. [1,2]

The diagnosis and monitoring of multiple myeloma can be accomplished by multiple means and typically includes a thorough clinical examination, history taking and laboratory testing. Laboratory tests include using immunoglobulin studies; serum and urine protein electrophoresis and immunofixation, serum free light chain(SFLC) analysis; bone marrow evaluation, full blood count with differential; serum chemistries: creatinine, calcium together with imaging. Imaging including conventional x-rays, CT, magnetic resonance imaging (MRI) and PET scans. [1,2]

Diagnosis of multiple myeloma is currently determined by the International Myeloma Working Group(IMWG). The IMWG has developed amongst others, guidelines for the diagnosis, management and response criteria for multiple myeloma. These criteria are constantly evolving and updated every few years to reflect our increasing knowledge of the disease. The most recent diagnostic criteria were published in 2014. The updated diagnostic criteria allows for treatment of patients who are at high risk of progression to symptomatic disease. Furthermore, these criteria may assist patients to potentially live longer if they were treated before serious organ damage occurred. [2,16]

The updated diagnostic criteria as per IMWG taken from the Lancet Oncology journal published in 2014 is as follows [2,16]:

Definition of multiple myeloma

Clonal bone marrow plasma cells $\geq 10\%$ or biopsy-proven bony or extramedullary plasmacytoma and any one or more of the following myeloma defining events:

- Myeloma defining events:
- Evidence of end organ damage that can be attributed to the underlying plasma cell proliferative disorder, specifically:
- Hypercalcaemia: serum calcium >0,25 mmol/L higher than the upper limit of normal or >2,75 mmol/L
- • Renal insufficiency: creatinine clearance <40 mL per minor serum creatinine >177 μ mol/L
- Anaemia: haemoglobin value of >2 g/dL below the lower limit of normal, or a haemoglobin value <10 g/dL
- Bone lesions: one or more osteolytic lesions on skeletal radiography, CT, or PET-CT
- Any one or more of the following biomarkers of malignancy:
- Clonal bone marrow plasma cell percentage ≥60%
- Involved: uninvolved serum free light chain ratio ≥100
- >1 focal lesion on MRI studies

One of the new additions to the 2014 IMWG diagnostic criteria was the Myeloma Defining Events (MDE's) to the definition of myeloma. The MDE are associated with inevitable progression to end-organ damage and include: a) clonal bone marrow plasma cell percentage ≥60%. b) Involved: uninvolved serum free light chain ratio ≥100 and c) >1 focal lesion on MRI studies. This addition allows clinicians to start treatment earlier prior to end organ damage. [2,16]

Treatment response guideline

IMWG has guidelines for response to treatment as seen in Table 1. The uniform response criteria for multiple myeloma plays an essential role in disease management. It monitors for response and treatment relapse guides future therapy. The IWMG criteria include the measurement of serum free light chains, serum and urine electrophoresis and immunofixation, bone marrow analysis and imaging. [1,16]

Each laboratory investigation has its advantage and disadvantages in our setting and we have explored a few. Bone marrow examination directly identifies malignant cells, but the heterogeneous nature of the disease makes it problematic to use routinely to follow patients with multiple myeloma as it may not be representative and require repeated sample collection. Being an invasive procedure, bone marrow biopsies pose potential complications to the patient, such as bleeding and infection. They are also expensive and arduous to be performed at a regular interval for treatment monitoring. [17]

Light chains are more challenging to detected than complete immunoglobulins. [18] The serum free light-chain(SFLC) assay has increasingly been used, and in individual patients tracks well with proteinuria. [19-20] The greater sensitivity when compared to urine analysis, has brought forth the widespread use and incorporation of SFLC measurements into multiple guidelines for the management of myeloma, most recently as a myeloma defining event in asymptomatic patients. [1,21] However, the SFLC assay is not readily available in our province of KZN (referred to a NHLS laboratory in another province approximately 600km away). And due to inter-patient variation in the renal metabolism of light chains, quantification of proteinuria cannot be predicted by the SFLC concentration. [22-24] The IMWG states that once a diagnosis of multiple myeloma is made, a 24-hour urine protein electrophoresis(UPEP) and immunofixation should be done for patient monitoring and these measures are not replaceable with SFLC. [1,25]

It is presumed that due to the above-mentioned factors, urine and serum protein electrophoresis are commonly employed in our setting to monitor patients.

Our study focused on urinary monoclonal free light chains as it is part of the response criteria in multiple myeloma and it is measured in the Chemistry department at the National Health Laboratory Service, Inkosi Albert Luthuli Central Hospital. It is an available laboratory test for our population and fairly simple to perform.

Monoclonal free light chains

Monoclonal free light chains (FLCs) appearing in urine, are referred to as Bence Jones proteins (BJP). This was first described by Dr. Henry Bence Jones over 150 years ago. Detection and measurement of BJPs are utilized to aid in the diagnosis and monitoring of monoclonal gammopathies. [2,26] Once renal tubular reabsorption is saturated; BJP is present in urine. In approximately 20% of MM cases, BJP may occur in the absence of a monoclonal band in the serum thus making it a valuable test in the detection of this malignancy. [26, 27]

BJP may be quantified by means of urine protein electrophoresis. This method involves separation of charged proteins in a liquid medium under the influence of an electrical field. Following electrophoresis of the urine specimen and staining of the gel, the size of the BJP peak is measured using densitometry scan of the peak. The percentage area of the peak is then multiplied by the total urine protein concentration of the sample to provide a semi- quantitative value for the BJP. Confirmation of the presence of BJP following urine protein electrophoresis is performed via urine immunofixation electrophoresis(UIFE). The principle of UIFE involves applying antisera to the separated proteins on the gel. Antigen-antibody complexes precipitate and are trapped within the gel matrix. The complexes are then stained and visualized. [18]

Response criteria

The Bence Jones protein quantitated for as a 24-hour urine is used to monitor treatment response in categories described by the IMWG guidelines. The article; International myeloma working group consensus criteria for response and minimal residual disease assessment in multiple myeloma printed in the Lancet Oncology journal in 2016 is presented below, see Table 1. [1,16]

Stringent complete response	Complete response as defined below plus normal FLC ratio and absence of clonal cells in bone marrow biopsy by immunohistochemistry (κ/λ ratio $\leq 4:1$ or $\geq 1:2$ for κ and λ patients, respectively, after counting ≥ 100 plasma cells)
Complete response	Negative immunofixation on the serum and urine and disappearance of any soft tissue plasmacytomas and <5% plasma cells in bone marrow aspirates
Very good partial remission (VGPR)	Serum and urine M-protein detectable by immunofixation but not on electrophoresis or ≥90% reduction in serum M-protein plus urine M-protein level <100 mg per 24 h
Partial response (PR)	≥50% reduction of serum M-protein plus reduction in 24 h urinary M-protein by ≥90% or to <200 mg per 24 h; If the serum and urine M-protein are unmeasurable, a ≥50% decrease in the difference between involved and uninvolved FLC levels is required in place of the M-protein criteria; If serum and urine M-protein are unmeasurable, and serum-free light assay is also unmeasurable, ≥50% reduction in plasma cells is required in place of M-protein, provided baseline bone marrow plasma-cell percentage was ≥30%. In addition to these criteria, if present at baseline, a ≥50% reduction in the size (SPD) of soft tissue plasmacytomas is also required
Minimal response	≥25% but ≤49% reduction of serum M-protein and reduction in 24-h urine M-protein by 50–89%. In addition to the above listed criteria, if present at baseline, a ≥50% reduction in the size (SPD) of soft tissue plasmacytomas is also required
Stable disease	Not recommended for use as an indicator of response; stability of disease is best described by providing the time-to-progression estimates. Not meeting criteria for complete response, very good partial response, partial response, minimal response, or progressive disease
Progressive disease (poor response)	Any one or more of the following criteria: Increase of 25% from lowest confirmed response value in one or more of the following criteria: Serum M-protein (absolute increase must be ≥ 0.5 g/dL); Serum M-protein increase ≥ 1 g/dL, if the lowest M component was ≥ 5 g/dL; Urine M-protein (absolute increase must be ≥ 200 mg/24 h); In patients without measurable serum and urine M-protein levels, the difference between involved and uninvolved FLC levels (absolute increase must be ≥ 10 mg/dL); In patients without measurable serum and urine M-protein levels and without measurable involved FLC levels, bone marrow plasma-cell percentage irrespective of baseline status (absolute increase must be $\geq 10\%$); Appearance of a new lesion(s), $\geq 50\%$ increase from nadir in SPD of ≥ 1 lesion, or $\geq 50\%$ increase in the longest diameter of a previous lesion ≥ 1 cm in short axis; $\geq 50\%$ increase in circulating plasma cells (minimum of 200 cells per μ L) if this is the only measure of disease
Clinical relapse	Clinical relapse requires one or more of the following criteria: Direct indicators of increasing disease and/or end organ dysfunction (CRAB features) related to the underlying clonal plasma-cell proliferative disorder. It is not used in calculation of time to progression or progression-free survival but is listed as something that can be reported optionally or for use in clinical practice; Development of new soft tissue plasmacytomas or bone lesions (osteoporotic fractures do not constitute progression); Definite increase in the size of existing plasmacytomas or bone lesions. A

	definite increase is defined as a 50% (and ≥1 cm) increase as measured serially by the SPD of the measurable lesion; Hypercalcaemia (>11 mg/dL);			
	Decrease in haemoglobin of ≥2 g/dL not related to therapy or other non-myeloma-related conditions;			
	Rise in serum creatinine by 2 mg/dL or more from the start of the therapy			
	and attributable to myeloma;			
	Hyperviscosity related to serum paraprotein			
Relapse from complete response	Any one or more of the following criteria:			
(to be used only if the end point is	Reappearance of serum or urine M-protein by immunofixation or			
disease-free survival)	electrophoresis;			
	Development of ≥5% plasma cells in the bone marrow;			
	Appearance of any other sign of progression (i.e., new plasmacytoma, lytic			
	bone lesion, or hypercalcaemia see above)			
Relapse from MRD negative (to	Any one or more of the following criteria:			
be used only if the end point is	Loss of MRD negative state (evidence of clonal plasma cells on NGF or			
disease-free survival)	NGS, or positive imaging study for recurrence of myeloma); Reappearance			
	of serum or urine M-protein by immunofixation or electrophoresis;			
	Development of ≥5% clonal plasma cells in the bone marrow;			
	Appearance of any other sign of progression (i.e., new plasmacytoma, lytic			
	bone lesion, or hypercalcaemia)			

Table 1. International myeloma working group consensus criteria for response and minimal residual disease assessment in multiple myeloma

SPD=sum of the products of the maximal perpendicular diameters of measured lesions.

CRAB features=calcium elevation, renal failure, anaemia, lytic bone lesions.

Traditional serum and urine assessment of monoclonal proteins and bone marrow assessment was used as response criteria in multiple myeloma. Recent efforts have focused on the identification of residual tumour cells in the bone marrow by means of flow cytometry or gene sequencing. In addition, sensitive imaging techniques can be used to detect the presence of residual disease beyond the bone marrow. Combining these different methods, the International Myeloma Working Group has defined new response categories of minimal residual disease. This permits consistent reporting inside and outside of clinical trials. [1,16]

BJP can be used to differentiate the treatment response categories and guide therapy. The International Myeloma Working Group (IMWG) and the College of American pathology recommend a 24-hour urine collection for quantification of urine BJP. [2,3] The advantage of the 24-hour urine collection is that it defines BJP excretion over the entire 24-hours and directly relates to published data on 24-hour BJP excretion. But it has several issues especially those related to sample collection and is often subject to error and may result in inaccurate BJP quantitation. [4-7]

In particular, the impracticality of a 24-hour collection together with high likelihood of incomplete collections hinder the accuracy of the test. Other disadvantages for the patient include the inconvenience of storage and transport of the samples to the clinic as these are most often in large 5-liter plastic containers. Many of these patients carry their 24-hour urine collections, travelling several hundred kilometers using public transport to reach the haematology clinic. This is not ideal for maintaining sample stability as well as inconvenient and embarrassing for the patient. [28]

It is also more time consuming for the laboratory to supervise as the completed 24-hour urine samples must be accurately weighed for the correct volume. An aliquot of the sample together with the recorded 24-hour volume is sent to the referral laboratory for analysis. Some aliquots are sent without volumes and these samples are unfortunately rejected as the calculation for BJP requires a volume. In these instances, no result can be provided to the clinician for patient management in regard to the recognized response criteria. An important factor affecting 24-hour collections is that total urinary protein has a variation with urine volume and hence, can be variable depending on patients' fluid status or medications (e.g. diuretics). [5,29-30] Random urine collections on the other hand are easy to obtain. There is rapid transfer to laboratory which avoids potential degradation of protein. It can also be collected at any time of day and there is no need for the patient to store the sample and transport it to the laboratory.

Hence the use of random or early morning urine collections have been suggested to avoid the problems associated with 24-hour collections. The main concern with random urine is that the synthesis and release of BJP may be variable throughout the day. But, the clinical utility of urine protein results is improved when expressed as a ratio to urine creatinine. [4-6] As creatinine excretion in urine is fairly constant throughout the 24-hour period, measurement of protein creatinine ratios (PCR) allows correction for variations in urine concentration. The use of PCRs has become widespread for routine urine protein analysis and several studies have demonstrated good correlation with the 24-hour collection. [4-6]

Similar studies

Whilst the use of Bence Jones protein creatinine ratios has emerged as an alternative to the 24-hour collection, there are few studies that have examined its correlation with the 24-hour collection and no reported studies to our knowledge reviewing its utility in an African population. [29-30]

A previous study demonstrated that it may be possible to use the protein/creatinine ratio from random urine samples to estimate the 24-hour BJP excretion. [29] Another study concluded that protein concentrations in the same individual are relatively constant. This group also demonstrated that early morning spot specimens had a linear relation with measured 24-hour BJP collections and were preferred over the random urine collection. [30].

Conclusion

The need for simple, easily available test for treatment response in multiple myeloma in our population can assist and improve patient manage. We decided to compare the use of measured 24-hour urine collection to random urine for the quantitation of BJP in a South African population with this mind. Very few similar studies have been done in different population groups and have shown some comparability.

References

- 1. S. Kumar, B. Paiva, K.C. Anderson, B. Durie, O. Landgren, P. Moreau, et al., International myeloma working group consensus criteria for response and minimal residual disease assessment in multiple myeloma, Lancet Oncol. 17 (2016) e328–e346.
- 2. S.V. Rajkumar, M.A. Dimopoulos, A. Palumbo, J. Blade, G. Merlini, M.-V. Mateos, et al., International myeloma working group updated criteria for the diagnosis of multiple myeloma, Lancet Oncol. 15 (2014) e538–e548.
- 3. Keren DF, Alexanian R, Goeken JA, Gorevic PD, Kyle RA, Tomar RH. Guidelines for clinical and laboratory evaluation of patients with monoclonal gammopathies. Arch Pathol Lab Med 1999; 123:106–7
- 4. Beetham R, Cattell WR. Proteinuria: pathophysiology, significance and recommendation for measurement in clinical practice. Ann ClinBiochem 1993; 30(5):425-34.
- 5. Burtis, Carl A, Edward R. Ashwood, and David E. Bruns. Tietz Textbook of Clinical Chemistry and Molecular Diagnostics. London: Elsevier Health Sciences, 2012;25: 675-676
- 6. Lesley A Inker. et al KDOQI US Commentary on the 2012 KDIGO Clinical Practice Guideline for the Evaluation and Management of CKD, American Journal of Kidney Diseases ,2014;63(5) , 713 735
- 7. Citalia VC,Kthari J, Wells EJ, Livesy JH, Robson RA, Searle M, et al. Cost-benefit analysis and prediction of 24-hour proteinuria from the spot urine protein-creatinine ratio. Clin Nephrol. 2001 Jun;55(6):436-47.
- 8. Surveillance, Epidemiology, and End Results Program. SEER Stat Fact Sheets: Myeloma. National Cancer Institute. Available at http://seer.cancer.gov/statfacts/html/mulmy.html. Accessed: January 11, 2018.
- 9. Ludwig H, Miguel JS et al. International Myeloma Working Group recommendations for global myeloma care. Nature Review, Leukemia 2014 May;28(5):981-92
- 10. Visser HF et al. Retrospective review of multiple myeloma and immunosecretory disorder cases diagnosed in a tertiary setting. SA orthopaedic journal.Summer 2008: 38-43
- 11. Cowan AJ, Allen C, Barac A, et al. Global Burden of Multiple Myeloma: A Systematic Analysis for the Global Burden of Disease Study 2016. JAMA Oncol. 2018;4(9):1221-1227.
- 12. De Groot, J J B Et al. Concomitant HIV infection in newly diagnosed multiple myeloma patients is hard to recognise and should be tested for routinely in areas of high endemicity. South African Medical Journal, 2017;107(9):781-787
- 13. UNAIDS Data 2018, http://www.unaids.org/sites/default/files/media_asset/unaids-data-2018 en.pdf (accessed October 2018),

- 14. Ailawadhi S, Aldoss IT, Yang D, Razavi P, Cozen W, Sher T, et al. Outcome disparities in multiple myeloma: a SEER-based comparative analysis of ethnic subgroups. Br J Haematol. 2012;158(1):91-98
- 15. Christoph Röllig, Stefan Knop, Martin Bornhäuser, Multiple myeloma: Lancet 2015; 385: 2197–208
- 16. International Myeloma Working group www.imwg.myeloma.org
- 17. Amanda D. Tatsas, Madan H. Jagasia, Heidi Chen, Thomas L. McCurley, Monitoring Residual Myeloma: High-Resolution Serum/Urine Electrophoresis or Marrow Biopsy With Immunohistochemical Analysis?, American Journal of Clinical Pathology, Volume 134, Issue 1, July 2010, Pages 139–145, https://doi.org/10.1309/AJCP69TCAVDGSCWI
- 18. Graziani, M., Merlini, G. & Petrini, C. (2005). Guidelines for the Analysis of Bence Jones Protein. Clinical Chemistry and Laboratory Medicine, 41(3): 338-346.
- 19. Abraham RS, Clark RJ, Bryant SC, et al. Correlation of serum immunoglobulin free light chain quantification with urinary Bence Jones protein in light chain myeloma. Clin Chem. 2002;48(4): 655–657.
- 20. Alyanakian MA, Abbas A, Delarue R, Arnulf B. Free immunoglobulin light-chain serum levels in the follow-up of patients with monoclonal gammopathies: correlation with 24-hour urinary light-chain excretion. Am J Hematol 2004; 75: 246–248
- 21. Bradwell AR, Carr-Smith HD, Mead GP, Harvey TC, Drayson MT. Serum test for assessment of patients with Bence Jones myeloma. Lancet 2003; 361: 489–491
- 22. E.L. Jenner, J.A.R. Evans, S.J. Harding, Serum free light chain (FLC) analysis: a guiding light in monoclonal gammopathy management, J. Appl. Lab. Med. 2 (2017) 98–106.
- 23. Dispenzieri A, Zhang L, Katzmann JA et al. Appraisal of immunoglobulin free light chain as a marker of response. Blood 2008; 111: 4908–4915
- 24. Nowrousian MR, Brandhorst D, Sammet C et al. Serum free light chain analysis and urine immunofixation electrophoresis in patients with multiple myeloma. Clin Cancer Res 2005; 11: 8706–8714
- 25. Singhal S, Stein R, Vickrey E, Mehta J. The serum-free light chain assay cannot replace 24-hour urine protein estimation in patients with plasma cell dyscrasias. Blood 2007; 109: 3611–3612
- 26. Beetham R, Detection of Bence-Jones protein in practice, Ann Clin Biochem 2000; 37: 563-570
- 27. Dispenzieri A, Kyle R et al, International Myeloma Working Group (IMWG) Guidelines for Serum Free Light Chain Analysis in Multiple Myeloma and Related Disorders. Leukemia 2009; 23:215–224
- 28. Sebia© HYDRAGEL 7,15 & 30 β1- β2 package insert, 2015-06
- 29. Jennifer S. Kaplan, Gary L Horowitz. Twenty-four-Hour Bence Jones Protein Determinations, can we ensure accuracy? Arch Pathology Lab Med 2011:135: 1048-1051
- 30. Brigden, Malcolm & D. Neal, Evelyn & D. D. Mcneely, Michael & N. Hoag, Gordon. (1990). The Optimum Urine Collections for the Detection and Monitoring of Bence Jones Proteinuria. American journal of clinical pathology.1993; 689-93.

Part 2: A submission ready manuscript.

CHAPTER 2

A comparison of 24-hour urine versus random urine samples for determination and quantification of Bence Jones protein in a South African population

Prepared according to the Instructions for Authors of Clinical Biochemistry

Authors

Ashandree Reddy

Department of Chemical Pathology, Faculty of Health Sciences, University of Kwa-Zulu Natal and National Health Laboratory Service, Durban, South Africa

Nadine Rapiti

Department of Haematology, Faculty of Health Sciences, University of Kwa-Zulu Natal and National Health Laboratory Service, Durban, South Africa

Verena Gounden

Department of Chemical Pathology, Faculty of Health Sciences, University of Kwa-Zulu Natal and National Health Laboratory Service, Durban, South Africa

Corresponding author:

Verena Gounden

Department of Chemical Pathology,

2nd level Pathology Building, Inkosi Albert Luthuli Central Hospital 800 Vuzi Mzimela Road, Cato Manor, 4058, Durban, South Africa

Email:verenagounden@yahoo.com

Number of figures: 2 Number of tables: 3 Word count: 3446

Declaration of interest: none to declare. The authors would like to acknowledge the National Health Laboratory Service K funding for funding of this study

Abstract

Objectives:

The International Myeloma Working Group (IMWG) and College of American Pathologists recommend a 24-hour collection for Bence Jones proteins(BJP). Although a 24-hour urine collection is a definitive means to determine BJP excretion, it has several issues related to sample collection and is prone to inaccuracy. Protein to creatinine ratios have demonstrated good correlation with 24-hour urine. The aim of this study was to compare measured 24-hour urine to random urine collections for the quantitation of BJP in a South African population.

Method:

Known patients with multiple myeloma(MM) collected 24-hour urine as part of their routine clinical assessment for BJP, random urine samples were submitted following completion of the 24-hour collection. The measured 24-hour urine BJP was then compared to 2 estimated 24-hour BJP excretions which were calculated as follows; Estimation 1 (E1): Estimated 24-hour BJP (mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) × 10, Estimation 2 (E2): Estimated 24-hour BJP (mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) × 15mg/kg for women or × 20mg/kg for men. All the 24-hour BJP results were classified according to IMWG treatment response criteria.

Results:

When using the Wilcoxon paired test analysis, the measured 24-hour urine BJP was significantly different to both the E1 (p=0.049) and E2 (p=0.049) equations. But analysis following categorization of each patient per IMWG BJP response criteria, indicated no significant difference in classification of treatment response using either the E1 or E2 estimation equations (P=0.69).

Conclusion:

24-hour urine collections are cumbersome. Random urine BJP estimates are simple, rapid and inexpensive. This study demonstrates that both the estimates of 24-hour BJP can be used to monitor response in patients with MM. This can be added to the body of evidence that random samples can be used to monitor patients' treatment response in MM.

1. Introduction

Plasma dyscrasias (PD) are a group of disorders, which includes multiple myeloma, where a clone of plasma cells or B-lymphocytes have the ability to secrete a homogenous immunoglobulin (Ig) or its components. These may be identified as a monoclonal peak on analysis by serum or urine protein electrophoresis [1]. The presence, level and type of monoclonal Ig have important implications in diagnosis, staging and treatment of disease states [2].

The core diagnostic features of multiple myeloma (MM) include the presence of neoplastic plasma cells on bone marrow aspirate, radiological evidence of osteolytic lesions and detection of monoclonal Igs in serum or urine [1]. It is the second most common haematological cancer accounting for 1% of all malignancies worldwide. MM was responsible for 0.43% of newly diagnosed cases of malignancies in South Africa in 1999 with the incidence being reported at approximately 0.00054%. [3] While the incidence is highly variable among countries, studies indicate that the incidence of MM has increased uniformly since 1990 with the largest increase in middle and low-middle income countries. [4,5] The prevalence of MM is higher in HIV positive compared to uninfected individuals. [6] This increases the disease burden of MM in South Africa which has the largest HIV epidemic in the world having 7.1 million people living with HIV. [7]

Monoclonal free light chains (FLCs) appearing in urine, are referred to as Bence Jones proteins (BJP). This was first described by Dr. Henry Bence Jones over 150 years ago. Detection and measurement of BJPs are utilized to aid in the diagnosis and monitoring of monoclonal gammopathies. [8,9] Once renal tubular reabsorption is saturated; BJP is present in urine. In approximately 20% of MM cases, BJP may occur in the absence of a monoclonal band in the serum thus making it a valuable test in the detection of this malignancy. [8, 10]

BJP may be quantified by means of urine protein electrophoresis. Following electrophoresis of the urine specimen and staining of the gel, the size of the BJP peak is measured using densitometry scan of the peak. The percentage area of the peak is then multiplied by the total urine protein concentration of the sample to provide a semi- quantitative value for the BJP. Confirmation of the presence of BJP following urine protein electrophoresis is performed via immunofixation.

The International Myeloma Working Group (IMWG) and the College of American pathology recommend a 24-hour urine collection for quantification of urine BJP. [9, 11] Although a 24-hour urine is a definitive means to determine renal protein excretion, it has several issues especially those related to sample collection (as summarized in Table 1). In particular, the impracticality of a 24-hour collection together with high likelihood of incomplete collections hinder the accuracy of the test. Hence the use of random or early morning urine collections have been suggested to avoid the problems associated with 24-hour collections. The clinical utility of measured urine protein is improved when expressed as a ratio to urine creatinine. [12-14] As creatinine excretion in urine is fairly constant throughout the 24-hour period, measurement of protein creatinine ratios (PCR) allows correction for variations in urine concentration. The use of PCRs has become widespread for routine urine protein analysis and several studies have demonstrated good correlation with the 24-hour collection. [11-13]

Whilst the use of Bence Jones protein creatinine ratios has emerged as an alternative to the 24-hour collection, there are few studies that have examined its correlation with the 24-hour collection and no reported studies to the authors knowledge reviewing its utility in an African population. [14-17]

	Advantages	Disadvantages
Random Urine	 Easy to obtain Rapid transfer to laboratory which avoids potential degradation Collected any time of day No need for patient to store sample and transport to laboratory 	Synthesis and release of BJP may be variable throughout the day
24-hour Urine	 Defines BJP excretion over the entire 24-hours Directly relates to published data on 24-hour BJP excretion 	 Inconvenient/complex for patient-collection, storage, transport to hospital. Frequently incomplete collections More expensive for laboratory to supervise Total urinary protein has a variation with urine volume

Table 1 Comparison of Random and 24-hour urine collection for BJP [12-15]

The haematology clinic at King Edward VIII Hospital (KEH) is the referral center for the entire province of Kwa-Zulu Natal (KZN), South Africa, for the management of patients with multiple myeloma and other plasma cell dyscrasias. Many of these patients carry their 24-hour collections, travelling several hundred kilometers using public transport to reach the haematology clinic. This is not ideal for maintaining sample stability as well as inconvenient and embarrassing for the patient. [18]

MM is largely incurable and despite new therapy options, most patients relapse and require change in management. Laboratory testing plays a vital role in monitoring response to treatment as well detecting relapse in patient on treatment. [9] This, together with the previously described issues related to 24-hour urine collections prompted us to examine the utility and validity of measured 24-hour urine compared to random urine collections for the quantitation of BJP in a South African population.

2. Material and Methods

Study participants were recruited based on a known diagnosis of PDs from the Haematology clinic at King Edward VIII Hospital, Durban. Samples were collected over a period of 2 years (2016-2018). All participants had the diagnosis of multiple myeloma (per IMWG criteria) and were at different stages of disease and treatment.

Each participant collected a 24-hour urine sample for BJP following a standard protocol as part of the routine clinical assessment. The 24-hour collection was started the day before the clinic visit. On submission of the 24-hour collection, the participants immediately collected a random urine sample as per instructions provided. Thymol was used as the preservative for the 24-hour urine sample and no preservative was utilized for the random sample. Both samples were submitted to the laboratory immediately. The 24-hour collections were analysed as per routine by the chemical pathology laboratory.

The random urine samples were analysed for urine total protein and creatinine. Aliquots of the random urine samples were then frozen at -70°C and stored for a maximum of one month (stability as per manufacturer) until the urine protein electrophoresis (UPEP) was performed. [18]

For both random and 24-hour urine collections, UPEP was performed using the Sebia Hydragel 7 HR kit run on the Sebia Hydrasys (Sebia, Norcross, GA, USA). Quantitation of UPEP fractions was performed using the Sebia Hydrasys densitometer system and Phoresis software. Acid violet staining was used and the sensitivity of this method allows BJP to be detected at concentrations of 15-20 mg/L of the original urine. Urine samples for immunofixation electrophoresis (IFE) analysis were concentrated using BJP concentrators from the Sebia Hydrasys kit for all urine total protein samples measuring < 0.7g/L. [18] Urine total protein(UTP) and urine creatinine were measured using standard spectrophotometric methods on the Siemens Advia 1800 chemistry analyser (Siemens Diagnostics, Tarrytown, NY, USA). A dye binding method using pyrogallol red was used to quantify UTP. [19] Urine creatinine was measured using the modified kinetic Jaffe method. [20]

Only those 24-hour urine sample that were positive for BJP, had their respective random samples analysed to determine comparability. The measured 24-hour BJP excretion was calculated as follows: %BJP peak on densitometer \times UTP (g/L) \times 24-hour urine volume (L) and multiplied by 1000 for mg/24hr. The estimated 24 BJP using the random urine values were calculated as per the two different formulae below:

Estimation 1 (E1):

Estimated 24-hour BJP (mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) ×10 Note: For estimation 1 (E1), a factor of 10 was utilized because while daily excretion of creatinine is dependent on muscle mass, an average daily loss of 10mmol of creatinine can be expected. [13]

Estimation 2 (E2):

Estimated 24-hour BJP (mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) \times 15mg/kg for women or \times 20mg/kg for men (convert mg/kg to mmol/kg by multiplying 0.00884).

Note: The second method includes an estimation of daily creatinine excretion based on body mass (in kilograms). [16]

The estimated 24-hour BJP values were then used to classify patients according to their treatment response based on IMWG criteria. Very good partial response (VGPR) signified a concentration of <100 mg/24 hr, partial response 100-199 mg/24 hr and progressive disease was all those patients with $\geq 200 \text{mg}/24 \text{hr}$. [21]

Demographic details and clinical histories were collected from the patient's clinical records.

The Body mass index (BMI) was calculated as weight/height² (kg/m²) and categorized according to WHO. Statistical analyses were performed using Microsoft® Excel (Microsoft® Office 2016, Microsoft, USA) and MedCalc for Windows, version 10.0 (MedCalc Software, Ostend, Belgium). The Shapiro-Wilk test was used to assess normality. For non-parametric data Spearman rank correlation and Passing Bablock regression analysis was utilized for comparison of different estimated 24-hour BJP equations to the measured 24-hour BJP. Categorical data was compared using the Kruskal Wallis test. Wilcoxon paired sample analysis was used to compare continuous variables. A p value of <0.05 was deemed to be statistically significant. Ethical approval to conduct the study was acquired from the Biomedical Research Ethics Committee (BREC), University of Kwazulu-Natal (ref. no. BE509/15). Written informed consent was taken from each participant in English or isiZulu depending on their requirement.

3. Results

A total of 66 paired 24-hour and random urine samples were collected. Of these, 22 had detectable BJP on 24-hour UPEP and 19 had a quantifiable BJP in g/24hrs. Three patients had faint bands below detectable limit (< 15mg/L) on the measured 24-hour urine with % BJP calculated but, did have a quantifiable BJP peak on their paired random urine sample. The urine TP for those 3 24-hour samples were 0.1g/L, 0.07g/L and 0.05g/L which was much lower than their random paired samples 1.6g/L, 1.1g/L and 1.8g/L. This could account for the non-quantifiable bands in the measured 24-hour urine samples. One sample had a quantifiable 24-hour BJP peak (TP 0.1g/L) with no peak on the random urine sample (TP 4.2g/L) but the monoclonal band was present on UIFE.

The 22 samples were from 19 patients as 3 patients had repeat collections within the study period. There were 10 females and 9 males with 18 of the 19 patients being black African. The remaining one patient was of Indian descent. Table 2 presents the 19 patients with their demographics, immunotyping and other relevant results. Of note, there is only a record of 10 patients tested for HIV, with only one being positive. Serum Free Light Chain's (SFLC) were also only measured in 7 of these patients. Of the 19 patients, 2 did not have UIFE analysis performed despite having detectable BJP on UPEP and the UIFE being suggested by the

reporting pathologist. The mean age was 55,8 years old (SD $\pm 6,6$)) and the mean BMI was 27,5 m²/kg ((SD $\pm 5,4$). Refer to table 3.

No	Age	Gender	Race	SIFE	UIFE	Total Serum Calcium	Albumin	eGFR	ТР	НВ	HIV status	SFLC ratio
1	58	F	В	IgA K and free K	Free K	2,65	42	17	81	5,2	Neg	Nil
2	74	M	В	IgG K and free K	n/a	2,03	22	15	98	5,9	Nil	Nil
3	57	F	В	IgG K and free K	Free K	1,89	17	>60	41	5,7	Nil	Nil
4	61	M	В	IgG K and free K	Free K	2,06	16	>60	108	7,7	Nil	Nil
5	59	F	В	IgG K and free K	Free K	2,2	26	24	82	9,3	Nil	Nil
6	54	F	В	IgA K and free K	IgA K and free K	1,7	20	7	86	7,7	Neg	Nil
7	59	M	В	IgA K	Free K	2,17	29	36	95	8,6	Neg	4.73
8	58	M	В	IgG K	IgG K	3,18	20	18	153	6,4	Neg	Nil
9	63	M	В	IgG K	n/a	2,65	28	37	100	8,8	Neg	Nil
10	47	M	В	Free L	Free L	3,36	34	16	73	5,9	Pos	Nil
11	56	F	В	Free L	Free L	2,6	40	48	73	13,3	Nil	0,01
12	53	F	В	IgG L and free L	IgG L and free L	2,2	35	19	83	9,1	Neg	Nil
13	65	M	В	Free K	Free K	2,28	40	>60	67	7,9	Nil	1,76
14	57	F	I	IgG K	IgG K and free K	1,9	19	9	96	7,4	Nil	Nil
15	47	F	В	IgA K	IgA K and free K	3,49	16	8	140	6,7	Neg	102,23
16	49	M	В	IgA K	IgA K and free K	2,04	21	50	109	9,2	Neg	0,76
17	49	F	В	IgA L	Free L	2,6	21	23	118	6	Nil	Nil
18	55	M	В	Free L	Free L	2,19	38	>60	65	10,3	Nil	0,01
19	56	F	В	Free K	Free K	2,59	37	10	72	6,6	Neg	leaked
Mean (SD)			2,4(0,5)	27,4(8,9)	10,0	91,6(25,9)	7,8(1,9)					

Table 2 Patient characteristics

Notes: Age in years. SIFE; Serum immunofixation electrophoresis, UIFE; Urine immunofixation electrophoresis, M; male, F; female, B; black, I; Indian, K; Kappa, L; Lambda, TP; Total protein, HB; Hemoglobin, HIV; Human Immunodeficiency Virus, Neg; negative, Pos; positive, SFLC; Serum Free Light Chain. eGFR; estimated Glomerular Filtration Rate in ml/min/1.73m². Reference ranges: SFLC; 0.26-1.65, Calcium 2.15 - 2.55mmol/L, Albumin 35-52g/L.

Parameters	Mean (±SD)			
	Median(range)			
Total number of samples	N=22			
Age (years)	55.8 (±6.6)			
BMI (m ² /kg)	27.5(±5.4)			
Measured 24-hour BJP (mg/24hr)	2480 (250-14400)			
E1 (mg/24hr)	1256 (211-7143)			
E2 (mg/24hr)	1403 (267-9597)			

Table 3 Summary data of patient characteristics

[note E1 refers to (E1) Estimated 24-hour BJP (mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) $\times 10$ and E2 refers to Estimated 24-hour BJP (mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) $\times 15$ mg/kg for women or $\times 20$ mg/kg for men (convert mg/kg to mmol/kg by multiplying 0.00884).

Using Wilcoxon paired test analysis, the measured 24-hour urine BJP was significantly different to both the E1 (p=0.049) and for the E2 (p=0.049) equations. Spearman rank correlation for both estimation equations E1 and E2 was 0.893 when compared to the measured 24-hour BJP. On Kruskal Wallis analysis following categorization of each patient per IMWG BJP response criteria, there was no significant difference in classification of treatment response using either the E1 or E2 estimation equations when compared to the measured 24-hour urine BJP results (P=0.69). Results of Passing Bablock regression analysis are shown in Figures 1 and 2. The E2 estimation equation shows a smaller proportional bias with a slope of 0.968 as compared to the E1 estimation equation slope of 0.671 when compared to the measured 24-hour BJP

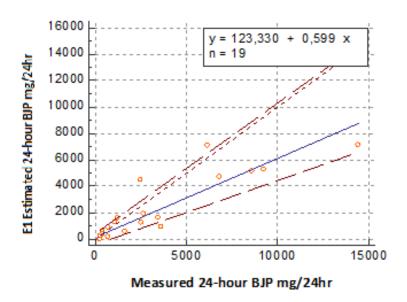


Figure 1
Regression analysis of Measured 24–hour BJP excretion versus E1 estimation for 24 hour BJP

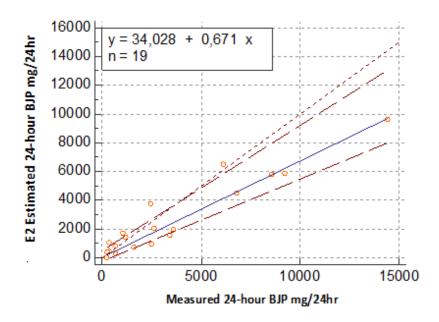


Figure 2
Regression analysis of Measure 24–hour BJP excretion versus E2 estimation for 24-hour BJP

Key for figure 1 and figure 2

—: Regression line

---: 95% Confidence intervals

• : Sample

4. Discussion

The average age of patients in this study was 55,8 years old, which is much younger when compared to the western countries with the average age at diagnosis ranging from 65-70 years. [22] This could be related to the high incidence of HIV in our population. HIV infection is more prevalent among younger than older patients hence, HIV-positive MM patients present at a significantly lower age. [6,22] Ethnicity has been found to effect incidence of MM. [23] Our data showed only 1 HIV positive patient from the 10 patients tested making ethnic disparities an alternative reason for the lower age.

The E2 estimation demonstrations a closer correlation and smaller proportional bias with the measured 24-hour BJP compared to E1 estimation equation. Our study revealed that both methods used to estimate 24-hour BJP were not comparable to the actual measured 24-hour BJP however when the estimated BJPs were used to classify patients according to IMWG treatment response, there was no significant difference with the performance of the measured 24-hour BJP and the estimated BJP using the E1 and E2 equations. This is key with regards to being able to use the random specimens for monitoring of disease. This study indicates that both the estimates of 24-hour BJP can be used to monitor response in patients with MM. This is in keeping with prior findings in other studies performed in different population groups. (16,17)

A previous study demonstrated that it may be possible to use the protein/creatinine ratio from random urine samples to estimate the 24-hour BJP excretion. [16] Another study concluded that protein concentrations in the same individual are relatively constant. This group also demonstrated that early morning spot specimens had a linear relation with measured 24-hour BJP collections and were preferred over the random urine collection. [17]. Because patients travelled long distances and arrived at the Haematology clinic at varying times, early morning specimens where a challenge to collect. Despite this, our study was still able to demonstrate that a random sample can be used to determine an estimate of the measured 24-hour BJP and can be used to monitor disease response to treatment.

As a result of only including patients with densitometrically quantifiable BJP on the measured 24-hour BJP the small sample size was a limitation, however this was also a limitation noted in other studies reviewing use of random urines for BJP estimation. (16,17) Measuring the creatinine on the 24-hour urine collections to verify the accuracy of collection would have been beneficial. [17] Another limitation is the challenges associated with the method to quantify BJP. Different proteins have varying affinities for the dyes used to stain electrophoretic gels, and thus a lack of linearity of the densitometry response may be seen. BJP may also co-migrate with other proteins or present with several bands making it complex to define the BJP peak correctly by densitometry. The measurement of BJP is not standardized and in order to minimize the mentioned analytical variability, it is suggested that patients should be followed up at the same laboratory, which was adhered to in this study. [24] We suggest using the random BJP to monitor

known patients with MM who already have confirmed BJP on immunofixation to minimize the above-mentioned limitations associated with measuring BJP's on electrophoresis.

Light chains are more challenging to detected than complete immunoglobulins. [25] The serum free light-chain(SFLC) assay has increasingly been used and tracks well with proteinuria in individual patients. [26,27] The greater sensitivity when compared to urine analysis, has brought forth the widespread use and incorporation of SFLC measurements into multiple guidelines for the management of myeloma, most recently as a myeloma defining event in asymptomatic patients. [9,28]. All the study participants had a SPEP and UPEP but surprisingly only a few had SFLC's. We found only 7 patients had SFLC's and 1 of the 7 samples had leaked during transit. The SFLC assay is not readily available in our province of KZN. And due to inter-patient variation in the renal metabolism of light chains, quantification of proteinuria cannot be predicted by the SFLC concentration. [29-31] 15 of the 19 patients had GFR's < 60ml/min/1.73m² which may affect the renal metabolism of SFLC's. The IMWG states that once a diagnosis of MM is made, a 24-hour UPEP and immunofixation should be done for patient monitoring and these measures are not replaceable with SFLC. [21,31] MM is associated with significant mortality and morbidity and is considered largely incurable and fatal without treatment. With the introduction of new classes of effective drugs for the treatment of multiple myeloma, improved frequencies and degree of patient response has been observed. Many treatments have been shown to significantly prolong survival and simultaneously improve the quality of life. Unfortunately, all patients will ultimately relapse after treatment and will require change to a more responsive therapy. This necessitates regular periodic monitoring of disease in order to detect relapse. Laboratory testing plays a vital role in monitoring response to treatment as well detecting relapse in patient on treatment. [9,21]

This study is the first to use and to demonstrate the utility of the estimate 24-hour urine BJP in an African population group. Both the E1 and E2 calculations are simple to perform and UTP and urine creatinine measurements are easily available on routine chemistry analysers.

This, together with other studies can be used by IMWG to add to their body of evidence for future use of estimated 24-hour BJP for patient monitoring. [16,17]

5. Conclusion

The random urine BJP estimates are simple, rapid, easily available and an inexpensive method for monitoring known patients with MM including light-chain disease. We have demonstrated that when using the IMWG response classification, the estimating equations, E1 and E2, did not differ from the measured 24-hour BJP. Further studies with larger cohorts can be conducted to validate the constant protein to creatinine ratios in random urine samples throughout the 24-hour period in patients with BJP to verify the accurateness of using a random sample.

Acknowledgements

The authors are grateful to Sunitha Sathabridg, a technologist at National Health Laboratory Service based at Inkosi Albert Luthuli Central Hospital for assistance in analysis of the UPEP.

Conflict of interest:

The authors declare that they have no financial or personal relationship(s) that may have inappropriately influenced them in writing this article.

Authors' contributions:

- 1) Verena Gounden: Formation of research idea and literature review, research protocol development, statistical analysis, interpretation of data and review of first draft write-up 2) Ashandree Reddy: Literature review, research protocol development, ethical and hospital
- management approval, data collection and processing, statistical analysis, interpretation of data and journal article first draft write-up.
- 3) Nadine Rapiti: Research protocol review, data collection, review of manuscript

Disclaimer The views expressed in the submitted article are those of the authors and not an official position of the institution. The authors declare no potential conflicts of interest with respect to the research, authorship, and/or publication of this article. The views expressed in the submitted article are those of the authors and not an official position of the institution.

References:

- 1. Burtis, Carl A, Edward R. Ashwood, and David E. Bruns. Tietz Textbook of Clinical Chemistry and Molecular Diagnostics. London: Elsevier Health Sciences, 2012;25: 547-548
- 2. Jillian Tate, Grahame Caldwell et al. Recommendations for standardized reporting of protein electrophoresis in Australia and New Zealand. Annals of Clinical Biochemistry 2012:49: 242-256.
- 3. Visser HF et al. Retrospective review of multiple myeloma and immunosecretory disorder cases diagnosed in a tertiary setting. SA orthopaedic journal. Summer 2008: 38-43
- 4. Ludwig H, Miguel JS et al. International Myeloma Working Group recommendations for global myeloma care. Nature Review, Leukemia 2014 May;28(5):981-92
- 5. Cowan AJ, Allen C, Barac A, et al. Global Burden of Multiple Myeloma: A Systematic Analysis for the Global Burden of Disease Study 2016. JAMA Oncol. 2018;4(9):1221-1227.
- 6. De Groot, J J B et al. Concomitant HIV infection in newly diagnosed multiple myeloma patients is hard to recognise and should be tested for routinely in areas of high endemicity. South African Medical Journal, 2017;107(9):781-787
- 7. UNAIDS Data 2018, http://www.unaids.org/sites/default/files/media_asset/unaids-data-2018_en.pdf (accessed October 2018),
- 8. Beetham R, Detection of Bence-Jones protein in practice, Ann Clin Biochem 2000; 37: 563-570
- 9. Rajkumar S.V, Dimopoulos M.A, Palumbo A, Blade J, Merlini G, Mateos M.V, et al, International myeloma working group updated criteria for the diagnosis of multiple myeloma, Lancet Oncol. 15 (2014) e538–e548.
- 10. Dispenzieri A, Kyle R et al, International Myeloma Working Group (IMWG) Guidelines for Serum Free Light Chain Analysis in Multiple Myeloma and Related Disorders. Leukemia 2009; 23:215–224
- 11. Keren DF, Alexanian R, Goeken JA, Gorevic PD, Kyle RA, Tomar RH. Guidelines for clinical and laboratory evaluation of patients with monoclonal gammopathies. Arch Pathol Lab Med 1999; 123:106–7
- 12. Beetham R, Cattell WR. Proteinuria: pathophysiology, significance and recommendation for measurement in clinical practice. Ann ClinBiochem 1993; 30(5):425-34.

- 13. Burtis, Carl A, Edward R. Ashwood, and David E. Bruns. Tietz Textbook of Clinical Chemistry and Molecular Diagnostics. London: Elsevier Health Sciences, 2012;25: 675-676
- 14. Lesley A Inker. et al KDOQI US Commentary on the 2012 KDIGO Clinical Practice Guideline for the Evaluation and Management of CKD, American Journal of Kidney Diseases ,2014;63(5) , 713 735
- 15. Citalia VC,Kthari J, Wells EJ, Livesy JH, Robson RA, Searle M, et al. Cost-benefit analysis and prediction of 24-hour proteinuria from the spot urine protein-creatinine ratio. <u>Clin Nephrol.</u> 2001 Jun;55(6):436-47.
- 16. Jennifer S. Kaplan, Gary L Horowitz. Twenty-four-Hour Bence Jones Protein Determinations, can we ensure accuracy? Arch Pathology Lab Med 2011:135: 1048-1051
- 17. Brigden, Malcolm & D. Neal, Evelyn & D. D. Mcneely, Michael & N. Hoag, Gordon. (1990). The Optimum Urine Collections for the Detection and Monitoring of Bence Jones Proteinuria. American journal of clinical pathology.1993; 689-93.
- 18. Sebia© HYDRAGEL 7,15 & 30 β1- β2 package insert, 2015-06
- 19. Siemens ADVIA Chemistry systems, Advia 1800 reagent package insert for Total Protein (Urine) (UPRO), 2007-05
- 20. Siemens ADVIA Chemistry systems, Advia 1800 reagent package insert for Creatinine (Creatinine_2), 2014-06
- 21. S. Kumar, B. Paiva, K.C. Anderson, B. Durie, O. Landgren, P. Moreau, et al., International myeloma working group consensus criteria for response and minimal residual disease assessment in multiple myeloma, Lancet Oncol. 17 (2016) e328–e346.
- 22. Surveillance, Epidemiology, and End Results Program. SEER Stat Fact Sheets: Myeloma. National Cancer Institute. Available at http://seer.cancer.gov/statfacts/html/mulmy.html. Accessed: January 11, 2018.
- 23. Ailawadhi S, Aldoss IT, Yang D, Razavi P, Cozen W, Sher T, et al. Outcome disparities in multiple myeloma: a SEER-based comparative analysis of ethnic subgroups. Br J Haematol. 2012;158(1):91-98
- 24. Graziani, M., Merlini, G. & Petrini, C. (2005). Guidelines for the Analysis of Bence Jones Protein. Clinical Chemistry and Laboratory Medicine, 41(3): 338-346.
- 25. Abraham RS, Clark RJ, Bryant SC, et al. Correlation of serum immunoglobulin free light chain quantification with urinary Bence Jones protein in light chain myeloma. Clin Chem. 2002;48(4): 655–657.
- 26. Alyanakian MA, Abbas A, Delarue R, Arnulf B. Free immunoglobulin light-chain serum levels in the follow-up of patients with monoclonal gammopathies:

- correlation with 24-hour urinary light-chain excretion. Am J Hematol 2004; 75: 246–248
- 27. Bradwell AR, Carr-Smith HD, Mead GP, Harvey TC, Drayson MT. Serum test for assessment of patients with Bence Jones myeloma. Lancet 2003; 361: 489–491
- 28. E.L. Jenner, J.A.R. Evans, S.J. Harding, Serum free light chain (FLC) analysis: a guiding light in monoclonal gammopathy management, J. Appl. Lab. Med. 2 (2017) 98–106.
- 29. Dispenzieri A, Zhang L, Katzmann JA et al. Appraisal of immunoglobulin free light chain as a marker of response. Blood 2008; 111: 4908–4915
- 30. Nowrousian MR, Brandhorst D, Sammet C et al. Serum free light chain analysis and urine immunofixation electrophoresis in patients with multiple myeloma. Clin Cancer Res 2005; 11: 8706–8714
- 31. Singhal S, Stein R, Vickrey E, Mehta J. The serum-free light chain assay cannot replace 24-hour urine protein estimation in patients with plasma cell dyscrasias. Blood 2007; 109: 3611–3612

Appendices

Appendix 1: The Study Protocol

Research Protocol: MMed

Pilot study: A comparison of 24-hour urine versus random urine samples for determination and quantification of Bence Jones protein

Dr Ashandree Reddy 214585580 February 2016

Title of study

A comparison of 24-hour urine versus random urine samples for determination and quantification of Bence Jones protein.

Aims

1 To compare accuracy of random urine Bence Jones:creatinine ratios (mg/mmol) to the gold standard 24-hour Bence Jones protein quantitation (mg/24 hrs)

Secondary objectives

- 1. To assess the accuracy of currently available equations for estimation of 24-hour BJP quantitation from random urine BJP: creatinine ratios
- 2. To develop an equation for estimation of 24-hour BJP values using random urine BJP:creatinine ratio

Background and literature

Plasma dyscrasias are a group of disorders associated with the presence of a monoclonal band (M protein) from malignant or nonproliferative population of cells. Examples of plasma dyscrasias include multiple myeloma (MM), monoclonal gammopathy of undetermined significance (MGUS) and plasmacytoma as well as conditions such as amyloidosis and Waldenstroms macroglobulinaemia. ² These monoclonal gammopathies result from an abnormal overproduction of a single abnormal clone of a plasma cell or B lymphocyte resulting in increased production of one immunoglobulin type (either the intact immunoglobulin, free light chain component or both). ¹The presence, level and type of the immunoglobulin have vital implications in diagnosis, staging and treatment of these diseases. ⁸This study focuses on MM as it is the second most common cancer of blood and accounts for 1% of all malignancies worldwide. ⁶ It is also associated with significant mortality and morbidity and is considered largely incurable. MM accounted for 0.43% of newly diagnosed cases of malignancies in South Africa in 1999 which makes the incidence approximately 0.00054% in our population of 47.8 million. ⁷

The monoclonal immunoglobulin (M protein) is recognized as a band of restricted migration on serum protein electrophoresis (SPEP) or urine protein electrophoresis (UPEP). Bence Jones protein (BJP) refers to the presence of a band of restricted migration on SPEP or UPEP representing a free light chain that is utilised to identify, diagnose and monitor patients with plasma cell dyscrasias in particular multiple myeloma. BJP is also important for the diagnosis of light chain myeloma as it is the only indicator of response to therapy.

The international myeloma working group (IMWG) recommendations provide essential procedures for the diagnosis and follow up of patients with MM. The investigations of interest to this study include SPEP and immunofixation as well as 24-hour urine collection for proteinuria,

electrophoresis and immunofixation. The Bence Jones protein quantitated for as a 24-hour urine is used to monitor treatment response (see Table1) in categories described by the IMWG guidelines.

Near complete remission	Paraprotein visible by IFE but not on electrophoresis of
	serum or urine samples
Very good partial remission	Serum and urine M-protein detectable by immunofixation
(VGPR)	but not on electrophoresis or ≥ 90% reduction in serum M-
	protein plus urine M-protein level < 100 mg/24 h or
	≥ 50% reduction of serum M-protein and reduction in 24-
	hours urinary M-protein by≥90% or to < 200 mg/24 hr.
Partial response (PR)	If serum and urine M-protein are not measurable, and
	serum free light chain assay is also not measurable, ≥50%
	reduction in plasma cells is required in place of M-protein,
	provided baseline bone marrow plasma cell percentage
	$was \ge 30\%$
Progressive disease (poor	24-hour urine Bence Jones protein :25% increase from
response)	nadir of urine M-component (the absolute increase must be
	≥200 mg/24 h)
Relapse from complete response	Reappearance of serum or urine M-protein on
	electrophoresis or immunofixation.

Table 1. IWMG Treatment response categories that include BJP.

Whilst the performance of urine electrophoresis and BJP quantitation is a relatively simple analytical procedure, the collection of urine over a 24-hour period is often subject to error and may result in inaccurate BJP quantitation. The preanalytical errors that can lead to a less reliable result include collection beyond or under the 24hour period is common as well as contamination and improper storage of urine during collection. The process is inconvenient and time consuming for patients and results in delay of sample analysis and ultimately treatment. The 24-hour urine sample is also more expensive for the laboratory to supervise. On the other hand, random urine collections are easy to obtain, allow rapid transfer to the laboratory and may avoid potential contamination and degradation of protein. A random specimen is a non-timed specimen that may be taken at any time of the day. The use of random urine protein/albumin: creatinine ratios have been shown to be equivalent to 24-hour collection for the measurement of urine protein and albumin, hence this study employed a protein(BJP): creatinine ratio.

Key references

- 1. Attaelmannan A and Levison SS. Understanding and Identifying Monoclonal Gammopathies. Clin Chem 2000;46: 1230-1238
- 2. Alexanian R, Weber D, Liu F. Differential diagnosis of monoclonal gammopathies. Arch Pathol Lab Med 1999; 123:108-113
- 3. Jenkins MA. Serum and Urine Electrophoresis for Detection and Identification of Monoclonal Proteins. Clin Biochem Rev 2009; 30:119-122
- 4. Malcolm L, brigden, MD et al. The optimum Urine Collections for the detection and monitoring of Bence Jones Proteinuria, American Journal of Pathology 1990; 93: 689-693
- 5. Jennifer S. Kaplan, Gary L Horowitz. Twenty-four-Hour Bence Jones Protein Deteminations, Can we ensure accuracy? Arch Pathology Lab Med 2011:135: 1048-1051
- 6. Ludwig H, Miguel JS et al. International Myeloma Working Group recommendations for global myeloma care. Nature Review, Leukemia 2014 May;28(5):981-92
- 7. Visser HF et al. Retrospective review of multiple myeloma and immunosecretory disorder cases diagnosed in a tertiary setting. SA orthopaedic journal.Summer 2008: 38-43
- 8. Jillian Tate, Grahame Caldwell et al. Recommendations for standardized reporting of protein electrophoresis in Australia and New Zealand. Annals of Clinical Biochemistry 2012:49: 242-256.

Study design

This is a pilot study as literature on this topic is scarce and previous studies involved very small sample numbers. It is a quantitative observational study

The validity of the study lies in a random urine sample being compared to the gold standard 24-hour urine collection for Bence Jones Protein.

Study population and location

Known patients with plasma dyscrasias enrolled at the Haematology clinic at King Edward VIII Hospital, Durban for which 24-hour urine collections will be performed for Bence Jones protein analysis as part of routine clinical assessment.

Sample size

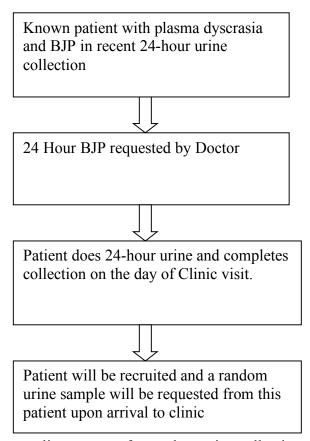
This study will be regarded as a pilot study and the sample number is approximately 100 patient samples. A statistician was consulted with regards to sample size determination

Inclusion/Exclusion criteria

Included are the adult patients that attend the KEH Haematology clinic that have a plasma dyscrasia and a positive BJP urine on most recent UPEP. Excluded are patients less than 18 years.

Sampling strategy

Informed consent will be obtained from all study participants (see appendix 1 Consent form-English and isiZulu version will be made available). Ethical clearance submitted to Biomedical Research Ethics Committee



Flow diagram showing sampling strategy for random urine collection

Data collection strategy and methods

Participants will collect random urine samples as per instructions provided (see appendix 2). Both samples will be submitted to the laboratory, however only the 24-hour collection as requested by the physician will be analysed and resulted by the routine laboratory for clinical patient care. Aliquots of both the random specimen and 24-hour urine sample will be analysed for total protein and creatinine (to ensure complete collection). The random sample will then be

frozen and stored at -70 C for a maximum of one month (stability as per manufacturer- Sebia) until UPEP is performed. Participants mass and height will also be recorded.

For both random and 24 hr urine collections: Urine protein electrophoresis will be performed using the Sebia BJP kits run on the Sebia Hydrasys. Quantitation of the M spike on UPEP (BJP) will be performed by densitometric scan of the UPEP gel. Total urine protein quantitation will be performed on the Siemens Advia 1800 chemistry analyser using a dye binding Pyrogallol red, which complexes with proteins in anacid environment containing molybdate ions. The resulting blue-colored complex is read via spectrophotometric method. Urine creatinine will be analysed on the Siemens Advia 1800 chemistry analyser using the modified kinetic Jaffe method. BJP: creatinine ratio will be calculated and the 24-hour BJP estimation will then be calculated using each of the following equations:

Twenty-four-hour BJP excretion calculated in 3 different ways:

- 1. Submitted BJP (mg) = %BJP X UTP (mg/dL) X 24-hour urine volume (dL)
- 2. Normalized BJP (mg) = $%BJP \times UTP (mg/dL)/UCR (mg/dL) \times patient's mean 24-hour creatinine excretion (mg)$
- 3. Estimated BJP (mg) = % BJP X UTP (mg/dL)/UCR (mg/dL) X patient's weight-based expected 24-hour creatinine excretion (mg) 5

Data will be collected from the patients' files and will be captured into an electronic database example Windows Excel and subsequently transferred to a statistical program for analysis. These programs will be password protected.

Clinical variables extracted from the charts include: hospital number, date of birth, height (cm), weight (kg), diagnosis, gender, race and age.

Statistical analyses

BJP quantitation values for 24-hour verse estimations from the random urine collections will then be compared using regression analysis and other relevant statistical analyses (Students t test)

The BJP results for each equation for the random urine specimens will be compared to the 24-hour urine BJP results also using regression analysis and Bland Altman plots.

Study period

The study period will be approximately 1 year January 2016-December 2016.

Limitations

Random specimens in comparison to timed or early morning urine collections are more prone to dilution of the specimen when collection occurs soon after the patient has consumed fluids. However, using a random sample is more convenient for the patient. Literature available on this topic is restricted.

Ethical considerations

This study has minimal ethical issues as it involves a non-invasive procedure of urine collection. This study will have no direct immediate impact on patients and will not affect their treatment. Patient confidentiality will be maintained at all times. Data will be collected on a password protected computer and the primary investigator will be the only person with access to it. Patients will be identified by hospital numbers and their identities will not be revealed.

Supervision and collaboration:

This research project will be performed under the supervision of Dr Verena Gounden, consultant chemical pathologist at the department of chemical pathology, and in collaboration with Dr Nadine Rapiti, Head of Department of Haematology at King Edward.

Appendix 2: The Guidelines for Authorship for the Journal selected for submission of the manuscript

CLINICAL BIOCHEMISTRY

Official Journal of the Canadian Society of Clinical Chemists

TABLE OF CONTENTS

- Description p.1
- Audience p.1
- Impact Factor p.1
- Abstracting and Indexing p.1
- Editorial Board p.2
- Guide for Authors p.4

DESCRIPTION

AUTHOR INFORMATION PACK

ISSN: 0009-9120

Clinical Biochemistry publishes articles relating to clinical chemistry, molecular biology and genetics, therapeutic drug monitoring and toxicology, laboratory immunology and laboratory medicine in general, with the focus on analytical and clinical investigation of laboratory tests in humans used for diagnosis, prognosis, treatment and therapy, and monitoring of disease.

Manuscripts are categorized as Analytical or Clinical Investigations and may be offered as Full Papers, Short Communications, or Letters. Opinion pieces and Special Reports are welcome, but contributors are encouraged to contact the Editor-in-Chief to avoid conflict with other forthcoming publications.

AUDIENCE

Clinical chemists, laboratory directors, physicians, as well as other laboratory professionals including, hematologists, geneticists, microbiologists, pathologists, biochemists, toxicologists, immunologists, analytical chemists, and molecular biologists.

IMPACT FACTOR

2017: 2.584 © Clarivate Analytics Journal Citation Reports 2018

ABSTRACTING AND INDEXING

MEDLINE® BIOSIS Chemical Abstracts Elsevier BIOBASE Reference Update Current Contents/Life Sciences EMBASE

Scopus

AUTHOR INFORMATION PACK 16 Apr 2019

www.elsevier.com/locate/clinbiochem 1

EDITORIAL BOARD

Editor-in-Chief:

Loralie Langman, Mayo Clinic, Rochester, Minnesota, USA Review Editor:

Damien Gruson, Université Catholique de Louvain (Cliniques Universitaires St-Luc), Brussels, Belgium *Special Issue Editor:*

Edgard Delvin, University of Montreal, Montreal, Quebec, Canada Associate Editors:

Peter Hickman, Australian National University, Canberra, Australian Capital Territory, Australia Patricia Jones, University of Texas Southwestern Medical Center, Dallas, Texas, USA Leslie Lai, Gleneagles Medical Centre, Kuala Lumpur, Malaysia Nathalie Lepage, University of Ottawa, Ottawa, Ontario, Canada

Erin McElvania, NorthShore University Health System, Evanston, Illinois, USA Amy Saenger, University of Minnesota, Minnesota, USA Hao Sun, The Chinese University of Hong Kong, Hong Kong, China Pierre E. Wallemacq, Université Catholique de Louvain (Cliniques Universitaires St-Luc), Brussels, Belgium

Editorial Board:

Abdolamir Allameh, Tarbiat Modares University, Tehran, Iran
Borros Arneth, Technische Universität Dresden, Dresden, Germany
Hikmet Aydin, Ege University, Izmir, Turkey
Hassan Azzazy, American University in Cairo, Cairo, Egypt
Tony Badrick, Bond University, Gold Coast, Queensland, Australia
Vipin Bhayana, Western University, London, Ontario, Canada
Ronald Booth, University of Ottawa, Ottawa, Ontario, Canada
Rossa Chiu, The Chinese University of Hong Kong, Hong Kong, China
Paul Collinson, St. George's Hospital Medical School, University of London, London, UK Eleftherios
Diamandis, Mount Sinai Hospital CA, Toronto, Ontario, Canada
Dennis Dietzen, Washington University School of Medicine, St Louis, Missouri, USA
James Donnelly, Quotient, Eysins, Switzerland

K.C. Dooley, Life Labs, Victoria, BC, Canada

Ozcan Erel, Ataturk Education Hospital, Ankara, Turkey

Jiri Frohlich, University of British Columbia, Vancouver, British Columbia, Canada

Ronda Greaves, RMIT University, Melbourne, Victoria, Australia

Ming Guan, Fudan University, Shanghai, China

Ola Hammarsten, Göteborg University (Sahlgrenska University Hospital), Göthenburg, Sweden Neil

Selwyn Harris, University of Florida College of Medicine, Gainesville, Florida, USA

Trefor Higgins, Dynacare Kasper Medical Laboratories, Edmonton, Alberta, Canada

Stephen Hill, McMaster University, Hamilton, Ontario, Canada

Barry Hoffman, Mount Sinai Hospital CA, Toronto, Ontario, Canada

Allan S. Jaffe, Mayo Clinic, Rochester, Minnesota, USA

Petr Jarolim, Brigham and Women's Hospital, Boston, Massachusetts, USA

Atholl Johnston, Queen Mary, University of London (QMUL), London, UK

Jawahar Kalra, University of Saskatchewan, Saskatoon, Saskatchewan, Canada

Morten Karsdal, Nordic Bioscience, Herlev, Denmark

Alexey Katrukha, HyTest Ltd., Turku, Finland

Denis Lehotay, University of Saskatchewan, Regina, Saskatchewan, Canada

Andrew Lyon, University of Saskatchewan, Saskatoon, Saskatchewan, Canada

Jacques Massé, Cité de la Santé de Laval, Québec, Quebec, Canada

Qing Meng, University of Texas, Houston, Texas, USA

Mathias Müller, Austrian Society of Quality Assurance and Standardisation, Wien, Austria Michael

Oellerich, Georg-August-Universität Göttingen, Göttingen, Germany

Mario Plebani, University Hospital of Padova, Padova, Italy

Joseph Queraltó, Hospital de la Santa Creu i Sant Pau, Barcelona, Spain

Nader Rifai, Harvard University, Boston, Maryland, USA

Robert Salvayre, INSERM, Toulouse, France

Hans Schneider, AlfredHealth, Victoria, Victoria, Australia

Christoph Seger, Medizinische Universität Innsbruck, Innsbruck, Austria

Grazyna Sypniewska, Nicolaus Copernicus University, Bydgoszcz, Poland

Gregory J. Tsongalis, Dartmouth Hitchcock Medical Center, Lebanon, New Hampshire, USA Sam

Vasikaran, Royal Perth Hospital, Perth, Western Australia, Australia

Naoki Watanabe, Sapporo Medical University, Sapporo, Japan

Eberhard Wieland, Universitätsklinikum Stuttgart - Katharinenhospital, Stuttgart, Germany

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 2

Edward Young, David Bradley Research Institute, Hamilton, Ontario, Canada **Martina Zaninotto**, University Hospital of Padova, Padua, Italy

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 3

GUIDE FOR AUTHORS

Your Paper Your Way

We now differentiate between the requirements for new and revised submissions. You may choose to submit your manuscript as a single Word or PDF file to be used in the refereeing process. Only when your paper is at the revision stage, will you be requested to put your paper in to a 'correct format' for acceptance and provide the items required for the publication of your article.

To find out more, please visit the Preparation section below. INTRODUCTION

Clinical Biochemistry publishes articles relating to clinical chemistry, molecular biology and genetics, therapeutic drug monitoring and toxicology, laboratory immunology and laboratory medicine in general, with the focus on analytical and clinical investigation of laboratory tests in humans used for diagnosis, prognosis, treatment and therapy, and monitoring of disease.

Manuscripts are categorized as Analytical or Clinical Investigations and may be offered as Full Papers, Short Communications, or Letters. Opinion pieces and Special Reports are welcome, but contributors are encouraged to contact the Editor-in-Chief to avoid conflict with other forthcoming publications.

Types of submission and criteria

•Original Research Communications (designated as one of two categories: Analytical or Clinical Investigation) may be offered as Full Papers or as Short Communications. The latter format is recommended for presenting technical evaluations and short clinical notes, comprising up to 1,500 words of text, 15 references, and two illustrative items (Tables and/or Figures). • Case Reports will be accepted only where they provide novel insight into disease mechanisms or diagnostic applications. • Reviews will be welcome but prospective authors are strongly advised to seek authorization from the Editor-in-Chief to avoid conflict with scheduled reviews invited by the Editorial Board. They should address new topics or trends in clinical biochemistry or related fields. •Consensus recommendations or guidelines on the use of laboratory test for clinical practice will be considered if they are compiled by a recognized organization or expert panel (e;g. IFCC, IUPAC, AACC, etc.) Please contact the Editor-in-Chief for consideration. The responsibility for such material remains with the originating body. •Editorial and opinion pieces. Please contact the Editor-in-Chief for consideration.

Contact details for submission

Papers should be submitted using the *Clinical Biochemistry* online submission system, http:// ees.elsevier.com/clb.

Submission checklist

You can use this list to carry out a final check of your submission before you send it to the journal for review. Please check the relevant section in this Guide for Authors for more details.

Ensure that the following items are present:

One author has been designated as the corresponding author with contact details: ullet E-mail address

Full postal address

All necessary files have been uploaded:

Manuscript:

- Include keywords
- All figures (include relevant captions)
- All tables (including titles, description, footnotes)
- Ensure all figure and table citations in the text match the files provided Indicate clearly if

color should be used for any figures in print *Graphical Abstracts / Highlights files* (where applicable)

Supplemental files (where applicable)

Further considerations

- Manuscript has been 'spell checked' and 'grammar checked'
- All references mentioned in the Reference List are cited in the text, and vice versa

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 4

- Permission has been obtained for use of copyrighted material from other sources (including the Internet)
- A competing interests statement is provided, even if the authors have no competing interests to declare
- Journal policies detailed in this guide have been reviewed
- Referee suggestions and contact details provided, based on journal requirements

For further information, visit our Support Center. **BEFORE YOU BEGIN**

Ethics in publishing

Please see our information pages on Ethics in publishing and Ethical guidelines for journal publication.

Policy and ethics

The work described in your article must have been carried out in accordance with *The Code of Ethics of the World Medical Association (Declaration of Helsinki) for experiments involving humans* http://www.wma.net/en/30publications/10policies/b3/index.html; *Recommendations for the Conduct, Reporting, Editing, and Publication of Scholarly Work in Medical Journals* http://www.icmje.org, published by the International Committee of Medical Journal Editors. This must be stated at an appropriate point in the article

Please note: Clinical Biochemistry does not accept submission of papers based on animal studies.

STARD initiative

Clinical Biochemistry supports the STARD initiative on reporting of diagnostic accuracy. http://www.stard-statement.org

Declaration of interest

All authors must disclose any financial and personal relationships with other people or organizations that could inappropriately influence (bias) their work. Examples of potential competing interests include employment, consultancies, stock ownership, honoraria, paid expert testimony, patent applications/registrations, and grants or other funding. Authors must disclose any interests in two places: 1. A summary declaration of interest statement in the title page file (if double-blind) or the manuscript file (if single-blind). If there are no interests to declare then please state this: 'Declarations of interest: none'. This summary statement will be ultimately published if the article is accepted. 2. Detailed disclosures as

part of a separate Declaration of Interest form, which forms part of the journal's official records. It is important for potential interests to be declared in both places and that the information matches. More information.

Submission declaration and verification

Submission of an article implies that the work described has not been published previously (except in the form of an abstract, a published lecture or academic thesis, see 'Multiple, redundant or concurrent publication' for more information), that it is not under consideration for publication elsewhere, that its publication is approved by all authors and tacitly or explicitly by the responsible authorities where the work was carried out, and that, if accepted, it will not be published elsewhere in the same form, in English or in any other language, including electronically without the written consent of the copyright- holder. To verify originality, your article may be checked by the originality detection service Crossref Similarity Check.

Use of inclusive language

Inclusive language acknowledges diversity, conveys respect to all people, is sensitive to differences, and promotes equal opportunities. Articles should make no assumptions about the beliefs or commitments of any reader, should contain nothing which might imply that one individual is superior to another on the grounds of race, sex, culture or any other characteristic, and should use inclusive language throughout. Authors should ensure that writing is free from bias, for instance by using 'he or she', 'his/her' instead of 'he' or 'his', and by making use of job titles that are free of stereotyping (e.g. 'chairperson' instead of 'chairman' and 'flight attendant' instead of 'stewardess').

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 5

Authorship

All authors should have made substantial contributions to all of the following: (1) the conception and design of the study, or acquisition of data, or analysis and interpretation of data, (2) drafting the article or revising it critically for important intellectual content, (3) final approval of the version to be submitted.

Changes to authorship

Authors are expected to consider carefully the list and order of authors **before** submitting their manuscript and provide the definitive list of authors at the time of the original submission. Any addition, deletion or rearrangement of author names in the authorship list should be made only **before** the manuscript has been accepted and only if approved by the journal Editor. To request such a change, the Editor must receive the following from the **corresponding author**: (a) the reason for the change in author list and (b) written confirmation (e-mail, letter) from all authors that they agree with the addition, removal or rearrangement. In the case of addition or removal of authors, this includes confirmation from the author being added or removed.

Only in exceptional circumstances will the Editor consider the addition, deletion or rearrangement of authors **after** the manuscript has been accepted. While the Editor considers the request, publication of the manuscript will be suspended. If the manuscript

has already been published in an online issue, any requests approved by the Editor will result in a corrigendum.

Article transfer service

This journal is part of our Article Transfer Service. This means that if the Editor feels your article is more suitable in one of our other participating journals, then you may be asked to consider transferring the article to one of those. If you agree, your article will be transferred automatically on your behalf with no need to reformat. Please note that your article will be reviewed again by the new journal. More information.

Copyright

Upon acceptance of an article, authors will be asked to complete a 'Journal Publishing Agreement' (see more information on this). An e-mail will be sent to the corresponding author confirming receipt of the manuscript together with a 'Journal Publishing Agreement' form or a link to the online version of this agreement.

Subscribers may reproduce tables of contents or prepare lists of articles including abstracts for internal circulation within their institutions. Permission of the Publisher is required for resale or distribution outside the institution and for all other derivative works, including compilations and translations. If excerpts from other copyrighted works are included, the author(s) must obtain written permission from the copyright owners and credit the source(s) in the article. Elsevier has preprinted forms for use by authors in these cases.

For gold open access articles: Upon acceptance of an article, authors will be asked to complete an 'Exclusive License Agreement' (more information). Permitted third party reuse of gold open access articles is determined by the author's choice of user license.

Author rights

As an author you (or your employer or institution) have certain rights to reuse your work. More information.

Elsevier supports responsible sharing

Find out how you can share your research published in Elsevier journals.

Role of the funding source

You are requested to identify who provided financial support for the conduct of the research and/or preparation of the article and to briefly describe the role of the sponsor(s), if any, in study design; in the collection, analysis and interpretation of data; in the writing of the report; and in the decision to submit the article for publication. If the funding source(s) had no such involvement then this should be stated.

Funding body agreements and policies

Elsevier has established a number of agreements with funding bodies which allow authors to comply with their funder's open access policies. Some funding bodies will reimburse the

author for the gold open access publication fee. Details of existing agreements are available online.

Elsevier journals comply with current NIH public access policy.

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 6

Open access

This journal offers authors a choice in publishing their research:

Subscription

- Articles are made available to subscribers as well as developing countries and patient groups through our universal access programs.
- No open access publication fee payable by authors.
- The Author is entitled to post the accepted manuscript in their institution's repository and make this public after an embargo period (known as green Open Access). The published journal article cannot be shared publicly, for example on ResearchGate or Academia.edu, to ensure the sustainability of peer- reviewed research in journal publications. The embargo period for this journal can be found below. **Gold open access**
- Articles are freely available to both subscribers and the wider public with permitted reuse.
- A gold open access publication fee is payable by authors or on their behalf, e.g. by their research funder or institution.

Regardless of how you choose to publish your article, the journal will apply the same peer review criteria and acceptance standards.

For gold open access articles, permitted third party (re)use is defined by the following Creative Commons user licenses:

Creative Commons Attribution (CC BY)

Lets others distribute and copy the article, create extracts, abstracts, and other revised versions, adaptations or derivative works of or from an article (such as a translation), include in a collective work (such as an anthology), text or data mine the article, even for commercial purposes, as long as they credit the author(s), do not represent the author as endorsing their adaptation of the article, and do not modify the article in such a way as to damage the author's honor or reputation.

Creative Commons Attribution-NonCommercial-NoDerivs (CC BY-NC-ND)

For non-commercial purposes, lets others distribute and copy the article, and to include in a collective work (such as an anthology), as long as they credit the author(s) and provided they do not alter or modify the article.

The gold open access publication fee for this journal is **USD 2400**, excluding taxes. Learn more about Elsevier's pricing policy: https://www.elsevier.com/openaccesspricing.

Green open access

Authors can share their research in a variety of different ways and Elsevier has a number of green open access options available. We recommend authors see our open access page for further information. Authors can also self-archive their manuscripts immediately and enable public access from their institution's repository after an embargo period. This is the version that has been accepted for publication and which typically includes author-incorporated changes suggested during submission, peer review and in editor-author communications. Embargo period: For subscription articles, an appropriate amount of time is needed for journals to deliver value to subscribing customers before an article becomes freely available to the public. This is the embargo period and it begins from the date the article is formally published online in its final and fully citable form. Find out more.

This journal has an embargo period of 12 months.

Elsevier Researcher Academy

Researcher Academy is a free e-learning platform designed to support early and mid-career researchers throughout their research journey. The "Learn" environment at Researcher Academy offers several interactive modules, webinars, downloadable guides and resources to guide you through the process of writing for research and going through peer review. Feel free to use these free resources to improve your submission and navigate the publication process with ease.

Language (usage and editing services)

Please write your text in good English (American or British usage is accepted, but not a mixture of these). Authors who feel their English language manuscript may require editing to eliminate possible grammatical or spelling errors and to conform to correct scientific English may wish to use the English Language Editing service available from Elsevier's WebShop.

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 7

Informed consent and patient details

Studies on patients or volunteers require ethics committee approval and informed consent, which should be documented in the paper. Appropriate consents, permissions and releases must be obtained where an author wishes to include case details or other personal information or images of patients and any other individuals in an Elsevier publication. Written consents must be retained by the author but copies should not be provided to the journal. Only if specifically requested by the journal in exceptional circumstances (for example if a legal issue arises) the author must provide copies of the consents or evidence that such consents have been obtained. For more information, please review the Elsevier Policy on the Use of Images or Personal Information of Patients or other Individuals. Unless you have written permission from the patient (or, where applicable, the next of kin), the personal details of any patient included in any part of the article and in any supplementary materials (including all illustrations and videos) must be removed before submission.

Submission

Our online submission system guides you stepwise through the process of entering your article details and uploading your files. The system converts your article files to a single PDF

file used in the peer-review process. Editable files (e.g., Word, LaTeX) are required to typeset your article for final publication. All correspondence, including notification of the Editor's decision and requests for revision, is sent by e-mail.

Referees

A minimum of six suitable potential reviewers must be supplied (please provide their name, email addresses, and institutional affiliation). When compiling this list of potential reviewers please consider the following important criteria: they must be knowledgeable about the manuscript subject area; must not be from your own institution; at least two of the suggested reviewers should be from another country than the authors'; and they should not have recent (less than four years) joint publications with any of the authors. However, the final choice of reviewers is at the editors' discretion.

PREPARATION

NEW SUBMISSIONS

Submission to this journal proceeds totally online and you will be guided stepwise through the creation and uploading of your files. The system automatically converts your files to a single PDF file, which is used in the peer-review process.

As part of the Your Paper Your Way service, you may choose to submit your manuscript as a single file to be used in the refereeing process. This can be a PDF file or a Word document, in any format or lay- out that can be used by referees to evaluate your manuscript. It should contain high enough quality figures for refereeing. If you prefer to do so, you may still provide all or some of the source files at the initial submission. Please note that individual figure files larger than 10 MB must be uploaded separately.

References

There are no strict requirements on reference formatting at submission. References can be in any style or format as long as the style is consistent. Where applicable, author(s) name(s), journal title/ book title, chapter title/article title, year of publication, volume number/book chapter and the article number or pagination must be present. Use of DOI is highly encouraged. The reference style used by the journal will be applied to the accepted article by Elsevier at the proof stage. Note that missing data will be highlighted at proof stage for the author to correct.

Formatting requirements

There are no strict formatting requirements but all manuscripts must contain the essential elements needed to convey your manuscript, for example Abstract, Keywords, Introduction, Materials and Methods, Results, Conclusions, Artwork and Tables with Captions. If your article includes any Videos and/or other Supplementary material, this should be included in your initial submission for peer review purposes.

Divide the article into clearly defined sections.

Figures and tables embedded in text

Please ensure the figures and the tables included in the single file are placed next to the relevant text in the manuscript, rather than at the bottom or the top of the file. The corresponding caption should be placed directly below the figure or table.

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 8

Peer review

This journal operates a single blind review process. All contributions will be initially assessed by the editor for suitability for the journal. Papers deemed suitable are then typically sent to a minimum of two independent expert reviewers to assess the scientific quality of the paper. The Editor is responsible for the final decision regarding acceptance or rejection of articles. The Editor's decision is final. More information on types of peer review.

REVISED SUBMISSIONS

Use of word processing software

Regardless of the file format of the original submission, at revision you must provide us with an editable file of the entire article. Keep the layout of the text as simple as possible. Most formatting codes will be removed and replaced on processing the article. The electronic text should be prepared in a way very similar to that of conventional manuscripts (see also the Guide to Publishing with Elsevier). See also the section on Electronic artwork.

To avoid unnecessary errors you are strongly advised to use the 'spell-check' and 'grammar-check' functions of your word processor.

Article structure

Subdivision - numbered sections

Divide your article into clearly defined and numbered sections. Subsections should be numbered 1.1 (then 1.1.1, 1.1.2, ...), 1.2, etc. (the abstract is not included in section numbering). Use this numbering also for internal cross-referencing: do not just refer to 'the text'. Any subsection may be given a brief heading. Each heading should appear on its own separate line.

Introduction

State the objectives of the work and provide an adequate background, avoiding a detailed literature survey or a summary of the results.

Material and methods

Provide sufficient details to allow the work to be reproduced by an independent researcher. Methods that are already published should be summarized, and indicated by a reference. If quoting directly from a previously published method, use quotation marks and also cite the source. Any modifications to existing methods should also be described.

Experimental

Provide sufficient details to allow the work to be reproduced by an independent researcher. Methods that are already published should be summarized, and indicated by a reference. If quoting directly from a previously published method, use quotation marks and also cite the source. Any modifications to existing methods should also be described.

Theory/calculation

A Theory section should extend, not repeat, the background to the article already dealt with in the Introduction and lay the foundation for further work. In contrast, a Calculation section represents a practical development from a theoretical basis.

Results

Results should be clear and concise.

Discussion

This should explore the significance of the results of the work, not repeat them. A combined Results and Discussion section is often appropriate. Avoid extensive citations and discussion of published literature.

Conclusions

The main conclusions of the study may be presented in a short Conclusions section, which may stand alone or form a subsection of a Discussion or Results and Discussion section.

Glossary

Please supply, as a separate list, the definitions of field-specific terms used in your article.

Appendices

If there is more than one appendix, they should be identified as A, B, etc. Formulae and equations in appendices should be given separate numbering: Eq. (A.1), Eq. (A.2), etc.; in a subsequent appendix, Eq. (B.1) and so on. Similarly for tables and figures: Table A.1; Fig. A.1, etc.

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 9

Essential title page information

- *Title.* Concise and informative. Titles are often used in information-retrieval systems. Avoid abbreviations and formulae where possible.
- Author names and affiliations. Please clearly indicate the given name(s) and family name(s) of each author and check that all names are accurately spelled. You can add your name between parentheses in your own script behind the English transliteration. Present the authors' affiliation addresses (where the actual work was done) below the names. Indicate all affiliations with a lower- case superscript letter immediately after the author's name and in front of the appropriate address. Provide the full postal address of each affiliation, including the country name and, if available, the e-mail address of each author.

- **Corresponding author.** Clearly indicate who will handle correspondence at all stages of refereeing and publication, also post-publication. This responsibility includes answering any future queries about Methodology and Materials. **Ensure that the e-mail address is given and that contact details are kept up to date by the corresponding author.**
- **Present/permanent address.** If an author has moved since the work described in the article was done, or was visiting at the time, a 'Present address' (or 'Permanent address') may be indicated as a footnote to that author's name. The address at which the author actually did the work must be retained as the main, affiliation address. Superscript Arabic numerals are used for such footnotes.

Highlights

Highlights are mandatory for this journal. They consist of a short collection of bullet points that convey the core findings of the article and should be submitted in a separate editable file in the online submission system. Please use 'Highlights' in the file name and include 3 to 5 bullet points (maximum 85 characters, including spaces, per bullet point). You can view example Highlights on our information site.

Abstract

A concise and factual abstract is required. The abstract should state briefly the purpose of the research, the principal results and major conclusions. An abstract is often presented separately from the article, so it must be able to stand alone. For this reason, References should be avoided, but if essential, then cite the author(s) and year(s). Also, non-standard or uncommon abbreviations should be avoided, but if essential they must be defined at their first mention in the abstract itself.

Maximum length of submissions

Full length articles should not exceed 4000 words (maximum 40 references), and up to 6 tables and/or figures with short communications comprising up to 1500 words of text, maximum 15 references, and two illustrative items (Tables and/or Figures). Letters will be classified as Case Reports (provide novel insight into disease mechanisms or diagnostic applications). Laboratory Notes (technical evaluation or important insight into analytical methodology), or Letters to the Editor (focused on a specific article that has appeared in Clinical Biochemistry within 4 weeks of print issue date of article). For all 3 types of letters listed above, the text should not exceed 500 words, with no abstract, a maximum of 1 table or figure and up to 5 references. Review Articles and Special Reports may exceed the word and reference limit for Full length articles as per the comprehensive nature of these articles. However, both of these articles (Reviews and Special Reports) will still require an abstract (unstructured, 250 word maximum). Editorials and Opinion pieces will not require an abstract and will be limited to 2000 words and up to 20 references.

Graphical abstract

Although a graphical abstract is optional, its use is encouraged as it draws more attention to the online article. The graphical abstract should summarize the contents of the article in a concise, pictorial form designed to capture the attention of a wide readership. Graphical abstracts should be submitted as a separate file in the online submission system. Image size: Please provide an image with a minimum of 531×1328 pixels (h \times w) or

proportionally more. The image should be readable at a size of 5×13 cm using a regular screen resolution of 96 dpi. Preferred file types: TIFF, EPS, PDF or MS Office files. You can view Example Graphical Abstracts on our information site.

Authors can make use of Elsevier's <u>Illustration Services</u> to ensure the best presentation of their images and in accordance with all technical requirements.

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 10

Keywords

Immediately after the abstract, provide a maximum of 6 keywords, using American spelling and avoiding general and plural terms and multiple concepts (avoid, for example, 'and', 'of'). Be sparing with abbreviations: only abbreviations firmly established in the field may be eligible. These keywords will be used for indexing purposes.

Abbreviations

Define abbreviations that are not standard in this field in a footnote to be placed on the first page of the article. Such abbreviations that are unavoidable in the abstract must be defined at their first mention there, as well as in the footnote. Ensure consistency of abbreviations throughout the article.

Acknowledgements

Collate acknowledgements in a separate section at the end of the article before the references and do not, therefore, include them on the title page, as a footnote to the title or otherwise. List here those individuals who provided help during the research (e.g., providing language help, writing assistance or proof reading the article, etc.).

Formatting of funding sources

List funding sources in this standard way to facilitate compliance to funder's requirements:

Funding: This work was supported by the National Institutes of Health [grant numbers xxxx, yyyy]; the Bill & Melinda Gates Foundation, Seattle, WA [grant number zzzz]; and the United States Institutes of Peace [grant number aaaa].

It is not necessary to include detailed descriptions on the program or type of grants and awards. When funding is from a block grant or other resources available to a university, college, or other research institution, submit the name of the institute or organization that provided the funding.

If no funding has been provided for the research, please include the following sentence:

This research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

Nomenclature and units

Follow internationally accepted rules and conventions: use the international system of units (SI). If other quantities are mentioned, give their equivalent in SI. You are urged to consult IUB: Biochemical Nomenclature and Related Documents for further information.

Math formulae

Please submit math equations as editable text and not as images. Present simple formulae in line with normal text where possible and use the solidus (/) instead of a horizontal line for small fractional terms, e.g., X/Y. In principle, variables are to be presented in italics. Powers of e are often more conveniently denoted by exp. Number consecutively any equations that have to be displayed separately from the text (if referred to explicitly in the text).

Footnotes

Footnotes should be used sparingly. Number them consecutively throughout the article. Many word processors build footnotes into the text, and this feature may be used. Should this not be the case, indicate the position of footnotes in the text and present the footnotes themselves separately at the end of the article.

Artwork

Image manipulation

Whilst it is accepted that authors sometimes need to manipulate images for clarity, manipulation for purposes of deception or fraud will be seen as scientific ethical abuse and will be dealt with accordingly. For graphical images, this journal is applying the following policy: no specific feature within an image may be enhanced, obscured, moved, removed, or introduced. Adjustments of brightness, contrast, or color balance are acceptable if and as long as they do not obscure or eliminate any information present in the original. Nonlinear adjustments (e.g. changes to gamma settings) must be disclosed in the figure legend.

Electronic artwork General points

Make sure you use uniform lettering and sizing of your original artwork.

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 11

- Preferred fonts: Arial (or Helvetica), Times New Roman (or Times), Symbol, Courier. Number the illustrations according to their sequence in the text.
- Use a logical naming convention for your artwork files.
- Indicate per figure if it is a single, 1.5 or 2-column fitting image.
- For Word submissions only, you may still provide figures and their captions, and tables within a single file at the revision stage.
- Please note that individual figure files larger than 10 MB must be provided in separate source files. A detailed guide on electronic artwork is available.

You are urged to visit this site; some excerpts from the detailed information are given here.

Formats

Regardless of the application used, when your electronic artwork is finalized, please 'save as' or convert the images to one of the following formats (note the resolution requirements for line drawings, halftones, and line/halftone combinations given below):

EPS (or PDF): Vector drawings. Embed the font or save the text as 'graphics'.

TIFF (or JPG): Color or grayscale photographs (halftones): always use a minimum of 300 dpi.

TIFF (or JPG): Bitmapped line drawings: use a minimum of 1000 dpi.

TIFF (or JPG): Combinations bitmapped line/half-tone (color or grayscale): a minimum of 500 dpi is required.

Please do not:

- Supply files that are optimized for screen use (e.g., GIF, BMP, PICT, WPG); the resolution is too low. Supply files that are too low in resolution.
- Submit graphics that are disproportionately large for the content.

Color artwork

Please make sure that artwork files are in an acceptable format (TIFF (or JPEG), EPS (or PDF), or MS Office files) and with the correct resolution. If, together with your accepted article, you submit usable color figures then Elsevier will ensure, at no additional charge, that these figures will appear in color online (e.g., ScienceDirect and other sites) regardless of whether or not these illustrations are reproduced in color in the printed version. For color reproduction in print, you will receive information regarding the costs from Elsevier after receipt of your accepted article. Please indicate your preference for color: in print or online only. Further information on the preparation of electronic artwork.

Illustration services

Elsevier's WebShop offers Illustration Services to authors preparing to submit a manuscript but concerned about the quality of the images accompanying their article. Elsevier's expert illustrators can produce scientific, technical and medical-style images, as well as a full range of charts, tables and graphs. Image 'polishing' is also available, where our illustrators take your image(s) and improve them to a professional standard. Please visit the website to find out more.

Figure captions

Ensure that each illustration has a caption. A caption should comprise a brief title (**not** on the figure itself) and a description of the illustration. Keep text in the illustrations themselves to a minimum but explain all symbols and abbreviations used.

Tables

Please submit tables as editable text and not as images. Tables can be placed either next to the relevant text in the article, or on separate page(s) at the end. Number tables consecutively in accordance with their appearance in the text and place any table notes below the table body. Be sparing in the use of tables and ensure that the data presented in them do not duplicate results described elsewhere in the article. Please avoid using vertical rules and shading in table cells.

The Editor-in-Chief, on accepting a manuscript, may recommend that additional tables containing important backup data, too extensive to be published in the article, may be published as supplementary material (see below) or deposited with the National Auxiliary Publications Service or made available by the author(s). In that event, an appropriate statement will be added to the text. Submit such tables for consideration with the manuscript.

References

Citation in text

Please ensure that every reference cited in the text is also present in the reference list (and vice versa). Any references cited in the abstract must be given in full. Unpublished results and personal communications are not recommended in the reference list, but may be mentioned in the text. If these

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 12

references are included in the reference list they should follow the standard reference style of the journal and should include a substitution of the publication date with either 'Unpublished results' or 'Personal communication'. Citation of a reference as 'in press' implies that the item has been accepted for publication.

Web references

As a minimum, the full URL should be given and the date when the reference was last accessed. Any further information, if known (DOI, author names, dates, reference to a source publication, etc.), should also be given. Web references can be listed separately (e.g., after the reference list) under a different heading if desired, or can be included in the reference list.

Data references

This journal encourages you to cite underlying or relevant datasets in your manuscript by citing them in your text and including a data reference in your Reference List. Data references should include the following elements: author name(s), dataset title, data repository, version (where available), year, and global persistent identifier. Add [dataset] immediately before the reference so we can properly identify it as a data reference. The [dataset] identifier will not appear in your published article.

References in a special issue

Please ensure that the words 'this issue' are added to any references in the list (and any citations in the text) to other articles in the same Special Issue.

Reference management software

Most Elsevier journals have their reference template available in many of the most popular reference management software products. These include all products that support Citation Style Language styles, such as Mendeley. Using citation plug-ins from these products, authors only need to select the appropriate journal template when preparing their article,

after which citations and bibliographies will be automatically formatted in the journal's style. If no template is yet available for this journal, please follow the format of the sample references and citations as shown in this Guide. If you use reference management software, please ensure that you remove all field codes before submitting the electronic manuscript. More information on how to remove field codes from different reference management software.

Reference formatting

There are no strict requirements on reference formatting at submission. References can be in any style or format as long as the style is consistent. Where applicable, author(s) name(s), journal title/ book title, chapter title/article title, year of publication, volume number/book chapter and the article number or pagination must be present. Use of DOI is highly encouraged. The reference style used by the journal will be applied to the accepted article by Elsevier at the proof stage. Note that missing data will be highlighted at proof stage for the author to correct. If you do wish to format the references yourself they should be arranged according to the following examples:

Reference style

Text: Indicate references by number(s) in square brackets in line with the text. The actual authors can be referred to, but the reference number(s) must always be given. Example: '.... as demonstrated [3,6]. Barnaby and Jones [8] obtained a different result' List: Number the references (numbers in square brackets) in the list in the order in which they appear in the text.

Examples:

Reference to a journal publication:

- [1] J. van der Geer, J.A.J. Hanraads, R.A. Lupton, The art of writing a scientific article, J. Sci. Commun. 163 (2010) 51–59. https://doi.org/10.1016/j.Sc.2010.00372.
- Reference to a journal publication with an article number:
- [2] Van der Geer, J., Hanraads, J.A.J., Lupton, R.A., 2018. The art of writing a scientific article. Heliyon. 19, e00205. https://doi.org/10.1016/j.heliyon.2018.e00205. Reference to a book:
- [3] W. Strunk Jr., E.B. White, The Elements of Style, fourth ed., Longman, New York, 2000. Reference to a chapter in an edited book:
- [4] G.R. Mettam, L.B. Adams, How to prepare an electronic version of your article, in: B.S. Jones, R.Z. Smith (Eds.), Introduction to the Electronic Age, E-Publishing Inc., New York, 2009, pp. 281–304. Reference to a website:
- [5] Cancer Research UK, Cancer statistics reports for the UK.
- http://www.cancerresearchuk.org/ aboutcancer/statistics/cancerstatsreport/, 2003 (accessed 13 March 2003).

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 13

Reference to a dataset:

[dataset] [6] M. Oguro, S. Imahiro, S. Saito, T. Nakashizuka, Mortality data for Japanese oak wilt disease and surrounding forest compositions, Mendeley Data, v1, 2015. https://doi.org/10.17632/xwj98nb39r.1.

Journal abbreviations source

Journal names should be abbreviated according to the List of Title Word Abbreviations.

Abbreviations and units

Standard abbreviations as listed in the *Council of Biology Editors Style Manual* may be used without definition. Use non-standard abbreviations sparingly, preceding their first use in the text with the corresponding full designation. Use units in conformity with the standard International System (SI) of units.

Video

Elsevier accepts video material and animation sequences to support and enhance your scientific research. Authors who have video or animation files that they wish to submit with their article are strongly encouraged to include links to these within the body of the article. This can be done in the same way as a figure or table by referring to the video or animation content and noting in the body text where it should be placed. All submitted files should be properly labeled so that they directly relate to the video file's content. . In order to ensure that your video or animation material is directly usable, please provide the file in one of our recommended file formats with a preferred maximum size of 150 MB per file, 1 GB in total. Video and animation files supplied will be published online in the electronic version of your article in Elsevier Web products, including ScienceDirect. Please supply 'stills' with your files: you can choose any frame from the video or animation or make a separate image. These will be used instead of standard icons and will personalize the link to your video data. For more detailed instructions please visit our video instruction pages. Note: since video and animation cannot be embedded in the print version of the journal, please provide text for both the electronic and the print version for the portions of the article that refer to this content.

Data visualization

Include interactive data visualizations in your publication and let your readers interact and engage more closely with your research. Follow the instructions here to find out about available data visualization options and how to include them with your article.

Supplementary material

Supplementary material such as applications, images and sound clips, can be published with your article to enhance it. Submitted supplementary items are published exactly as they are received (Excel or PowerPoint files will appear as such online). Please submit your material together with the article and supply a concise, descriptive caption for each supplementary file. If you wish to make changes to supplementary material during any stage of the process, please make sure to provide an updated file. Do not annotate any corrections on a previous version. Please switch off the 'Track Changes' option in Microsoft Office files as these will appear in the published version.

Supplementary material captions

Each supplementary material file should have a short caption which will be placed at the bottom of the article, where it can assist the reader and also be used by search engines.

Research data

This journal encourages and enables you to share data that supports your research publication where appropriate, and enables you to interlink the data with your published

articles. Research data refers to the results of observations or experimentation that validate research findings. To facilitate reproducibility and data reuse, this journal also encourages you to share your software, code, models, algorithms, protocols, methods and other useful materials related to the project.

Below are a number of ways in which you can associate data with your article or make a statement about the availability of your data when submitting your manuscript. If you are sharing data in one of these ways, you are encouraged to cite the data in your manuscript and reference list. Please refer to the "References" section for more information about data citation. For more information on depositing, sharing and using research data and other relevant research materials, visit the research data page.

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 14

Data linking

If you have made your research data available in a data repository, you can link your article directly to the dataset. Elsevier collaborates with a number of repositories to link articles on ScienceDirect with relevant repositories, giving readers access to underlying data that gives them a better understanding of the research described.

There are different ways to link your datasets to your article. When available, you can directly link your dataset to your article by providing the relevant information in the submission system. For more information, visit the database linking page.

For supported data repositories a repository banner will automatically appear next to your published article on ScienceDirect.

In addition, you can link to relevant data or entities through identifiers within the text of your manuscript, using the following format: Database: xxxx (e.g., TAIR: AT1G01020; CCDC: 734053; PDB: 1XFN).

Mendeley Data

This journal supports Mendeley Data, enabling you to deposit any research data (including raw and processed data, video, code, software, algorithms, protocols, and methods) associated with your manuscript in a free-to-use, open access repository. During the submission process, after uploading your manuscript, you will have the opportunity to upload your relevant datasets directly to *Mendeley Data*. The datasets will be listed and directly accessible to readers next to your published article online.

For more information, visit the Mendeley Data for journals page.

Data in Brief

You have the option of converting any or all parts of your supplementary or additional raw data into one or multiple data articles, a new kind of article that houses and describes your data. Data articles ensure that your data is actively reviewed, curated, formatted, indexed, given a DOI and publicly available to all upon publication. You are encouraged to submit your article for *Data in Brief* as an additional item directly alongside the revised version of your manuscript. If your research article is accepted, your data article will automatically be

transferred over to *Data in Brief* where it will be editorially reviewed and published in the open access data journal, *Data in Brief*. Please note an open access fee of 500 USD is payable for publication in *Data in Brief*. Full details can be found on the *Data in Brief* website. Please use this template to write your Data in Brief.

Data statement

To foster transparency, we encourage you to state the availability of your data in your submission. This may be a requirement of your funding body or institution. If your data is unavailable to access or unsuitable to post, you will have the opportunity to indicate why during the submission process, for example by stating that the research data is confidential. The statement will appear with your published article on ScienceDirect. For more information, visit the Data Statement page.

AFTER ACCEPTANCE

Online proof correction

Corresponding authors will receive an e-mail with a link to our online proofing system, allowing annotation and correction of proofs online. The environment is similar to MS Word: in addition to editing text, you can also comment on figures/tables and answer questions from the Copy Editor. Web-based proofing provides a faster and less error-prone process by allowing you to directly type your corrections, eliminating the potential introduction of errors.

If preferred, you can still choose to annotate and upload your edits on the PDF version. All instructions for proofing will be given in the e-mail we send to authors, including alternative methods to the online version and PDF.

We will do everything possible to get your article published quickly and accurately. Please use this proof only for checking the typesetting, editing, completeness and correctness of the text, tables and figures. Significant changes to the article as accepted for publication will only be considered at this stage with permission from the Editor. It is important to ensure that all corrections are sent back to us in one communication. Please check carefully before replying, as inclusion of any subsequent corrections cannot be guaranteed. Proofreading is solely your responsibility.

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 15

Offprints

The corresponding author will, at no cost, receive a customized Share Link providing 50 days free access to the final published version of the article on ScienceDirect. The Share Link can be used for sharing the article via any communication channel, including email and social media. For an extra charge, paper offprints can be ordered via the offprint order form which is sent once the article is accepted for publication. Both corresponding and co-authors may order offprints at any time via Elsevier's Webshop. Corresponding authors who have published their article gold open access do not receive a Share Link as their final published version of the article is available open access on ScienceDirect and can be shared through the article DOI link.

AUTHOR INQUIRIES

Visit the Elsevier Support Center to find the answers you need. Here you will find everything from Frequently Asked Questions to ways to get in touch.

You can also check the status of your submitted article or find out when your accepted article will be published.

© Copyright 2018 Elsevier | https://www.elsevier.com

AUTHOR INFORMATION PACK 16 Apr 2019 www.elsevier.com/locate/clinbiochem 16

Appendix 3: Ethical approvals



05 April 2016

Dr A Reddy (214585580) Discipline of Chemical Pathology School of Laboratory Medicine and Medical Sciences Ashandree.reddy@nhls.ac.za

Protocol: Comparison of 24 hour urine verse a random urine sample for determination and quantification of Bence Jones protein.

Degree: MMed

BREC reference number: BE509/15

EXPEDITED APPLICATION

The Biomedical Research Ethics Committee has considered and noted your application received on 14 December 2015.

The study was provisionally approved pending appropriate responses to queries raised. Your responses dated 01 April 2016 to queries raised on 16 March 2016 have been noted and approved by a subcommittee of the Biomedical Research Ethics Committee. The conditions have now been met and the study is given full ethics approval.

This approval is valid for one year from 05 April 2016. To ensure uninterrupted approval of this study beyond the approval expiry date, an application for recertification must be submitted to BREC on the appropriate BREC form 2-3 months before the expiry date.

Any amendments to this study, unless urgently required to ensure safety of participants, must be approved by BREC prior to implementation.

Your acceptance of this approval denotes your compliance with South African National Research Ethics Guidelines (2015), South African National Good Clinical Practice Guidelines (2006) (if applicable) and with UKZN BREC ethics requirements as contained in the UKZN BREC Terms of Reference and Standard Operating Procedures, all available at http://research.ukzn.ac.za/Research-Ethics/Biomedical-Research-Ethics, aspx.

BREC is registered with the South African National Health Research Ethics Council (REC-290408-009). BREC has US Office for Human Research Protections (OHRP) Federal-wide Assurance (FWA 678).

The sub-committee's decision will be RATIFIED by a full Committee at its meeting taking place on 10 May 2016.

We wish you well with this study. We would appreciate receiving copies of all publications arising out of this study.

Yours sincerely

Professor J Tsoka-Gwegweni

Chair: Biomedical Research Ethics Committee

cc supervisor: goundenv1@ukzn,ac.za cc postgrad: dudhrajhp@ukzn,ac.za

Biomedical Research Ethics Committee Professor J Tsoka-Gwegweni (Chair) Westville Campus, Govan Mbeki Building Postal Address: Private Bag X54001, Durban 4000

ne: +27 (0) 31 260 2486 Facelmile: +27 (0) 31 260 4609 Email: breodukan.ac.za

Website: http://research.ukzn.ac.za/Research-Ethics/Biomedical-Research-Ethics.aspr

1910 - 2010 AND 100 YEARS OF ACADEMIC EXCELLENCE

Fourting Compuses: Se Edgewood Howard College Medical School Pleasmarksburg Westwille



DIRECTORATE:

330 Langalibalele street, Private Bag X9051 PMB, 3200 Tel: 033 395 2805/3189/3123 Fax: 033 394 3782 Email: hrkm@kznhealth.gov.za www.kznhealth.gov.za

Health Research & Knowledge Management (HKRM)

Reference: HRKM88/16 KZ_2016RP19_462

01 April 2016

Dear Dr A Reddy

(University of KwaZulu-Natal/ National Health Laboratory Service)

Subject: Approval of a Research Proposal

 The research proposal titled 'Comparison of 24 hour urine verse a random urine sample for determination and quantification of Bence Jones protein' was reviewed by the KwaZulu-Natal Department of Health (KZN-DoH).

The proposal is hereby approved for research to be undertaken at King Edward VIII Hospital.

- 2. You are requested to take note of the following:
 - Make the necessary arrangement with the identified facility before commencing with your research project.
 - Provide an interim progress report and final report (electronic and hard copies) when your research is complete.
- Your final report must be posted to HEALTH RESEARCH AND KNOWLEDGE MANAGEMENT, 10-102, PRIVATE BAG X9051, PIETERMARITZBURG, 3200 and e-mail an electronic copy to hrkm@kznhealth.gov.za

For any additional information please contact Ms G Khumalo on 033-395 3189.

Yours Sincerely

Dr E Lutge

Chairperson, Health Research Committee

Date: 01/04/16.

Fighting Disease, Fighting Poverty, Giving Hope

Appendix 4: Data collection tools and consent forms

Procedure to collect random urine sample

The 24-hour urine collections are usually completed on the day the patient comes to haematology clinic. On arrival, a sterile jar will be provided to patients to collect the sample of urine following the last void for the 24-hour urine collection.

Instruction on how to collect the urine:

Step 1

Wash your hands with soap and water.

Step 2

Open the small sterile specimen jar

Avoid contamination by not touching the inside of the jar or the jar lid.

Step 3

Void urine into the container; ensure that the container is at least half filled with urine.

Step 4

Give both the 24-hour and random urine samples to the nurse.

isiZulu transltion

Isithasiselo 2

Ukuthathwa komchamo emahoreni angama-24 kuvame ukuba kuphothulwe ngosuku isiguli esifika ngalo emtholampilo we-haematology. Uma zifika iziguli, zizonikwa ujeke ongenamagciwane ukuba zifake umchamo emva kokuthathwa komchamo wokugcina wamahora angama-24.

Iimiyalelo yokuthathwa komchamo:

Okokuqala

Hlamba izandla ngamanzi nensipho.

Okwesibili

Vula ujeke omncane wokuthatha umchamo

Ungalithinti ingaphakathi noma isivalo sikajeke.

Okwesithathu

Chamela esitsheni; qinisekisa ukuthi umchamo uba, okungenani uhhafu esitsheni.

Okwesine

Nika umhlengikazi onke amasampula omchamo.

ACADEMIC COMPLEX BUSINESS UNIT



Chemical Pathology Department, NHLS Inkosi Albert Luthuli Central Hospital 800 Vusi Mzimela (Bellair) Road, Mayville, 4091 Private Bag X03, Mayville, 4058

Tel: +27 (0)31 240 2557/68/69 Fax: +27 (0)31 240 2576 Practice Number: 5200296



Informed Consent form for early morning spot urine collection for Bence Jones protein

This Informed Consent Form is for men and women who attend the haematology clinic at King Edward V11 Hospital, who we are inviting to participate in research on plasma dyscrasias. The title of our research project is "Spot the Bence Jones"

I am Dr Ashandree Reddy, working for the National Health Laboratory Services. We are doing research on plasma dyscrasias. I am going to give you information and invite you to be part of this research.

There may be some words that you do not understand. Please ask me to stop as we go through the information and I will take time to explain. If you have questions later, you can ask them of me, the study doctor or the staff.

Plasma dyscrasias include diseases like multiple myeloma and amyloidosis which are serious and life-long diseases. A 24-hour urine collection for a protein called Bence Jones is done for patients with these types of disease to help the doctor to monitor a patient on treatment. Collecting a 24-hour urine sample, example from 6am to 6am the next morning is difficult. We are doing this research to see if we can use just one early morning urine sample in place of a 24-hour urine collection.

This research will involve an early morning spot (one sample) urine sample on the day that you are doing your 24-hour urine collection.

Your participation in this research is entirely voluntary. Whether you choose to participate or not, all the services you receive at this clinic will continue and nothing will change. You may change your mind later and stop participating even if you agreed earlier.

We will collect your early morning urine specimen each time you do your routine 24-hour urine collection for Bence Jones protein over a period of approximately 3 months. At the end of the research, in approximately 1 year, any leftover urine samples will be destroyed. The samples will only be used for this test.

The samples will be collected on the day of your usual clinic visit so there is no extra visits or time needed for this project.

Your participation is likely to help us find the answer to the research question and may benefit future generations in that they may have to simply collect a single early morning urine sample rather than a 24-hour collection.

The information that we collect from this research project will be kept confidential. Any information about you will have a number on it instead of your name. Only the researchers will know what your number is and we will lock that information.

You do not have to take part in this research if you do not wish to do so. You may also stop participating in the research at any time you choose. It is your choice and all of your rights will still be respected.

You can ask me any more questions about any part of the research study, if you wish to. Do you have any questions?

PART II: Certificate of Consent

Print Name of Participant

I have read the foregoing information, or it has been read to me. I have had the opportunity to ask questions about it and any questions that I have asked have been answered to my satisfaction. I consent voluntarily to participate as a participant in this research.

Signature of Participant		
Date		
Date Day/month/year		
If illiterate A literate witness must sign (if possible, this pe no connection to the research team). Participants		
I have witnessed the accurate reading of the individual has had the opportunity to ask questreely.		
Print name of witness	_ AND	Thumb print of participant
Signature of witness		
Date Day/month/year		
Statement by the researcher/person taking collinate accurately read out the information sability made sure that the participant unders 1. An additional early morning spot urine collection.	theet to the potentiands that the following the desired to be done	ollowing will be done: ne on the same day as the 24-hour urin
I confirm that the participant was given an the questions asked by the participant have be confirm that the individual has not been coer freely and voluntarily.	been answered c	correctly and to the best of my ability.
A copy of this ICF has been provided to the print Name of Researcher/person taking the		
Signature of Researcher /person taking the co	onsent	
Date Day/month/year		

ACADEMIC COMPLEX BUSINESS UNIT



Chemical Pathology Department, NHLS Inkosi Albert Luthuli Central Hospital 800 Vusi Mzimela (Bellair) Road, Mayville, 4091 Private Bag X03, Mayville, 4058 Tel: +27 (0)31 240 2557/68/69

Fax: +27 (0)31 240 2576 Practice Number: 5200296



Isithasiselo 1

Ifomu lemvume yokuthatha umchamo ukuze kuhlolwe iphrotheyini i-Bence Jones

Leli fomu lemvume elabesilisa nabesifazane abaya emtholampilo we-haematology esiBhedlela i-King Edward VIII abamenywa ukuba babe yingxenye yocwaningo lwe-*plasma cell dyscrasias*. Sanibona, igama lami nginguDkt. Ashandree Reddy, osebenza e-National Health Laboratory Services. Senza ucwaningo lwe-*plasma cell dyscrasias* (uhlobo lomdlavuza wegazi). Ngizoninika ulwazi bese nginimema ukuba nibe yingxenye yocwaningo.

Kungahle kube namagama eningawaqondi. Ngiyocela ningimise uma kukhona lapho nisala khona, ngiyobe sengicacisa lapho. Uma uba neminye imibuzo kamuva, ukhululekile ukungibuza.

I-Plasma dyscrasias ibandakanya izifo ezinjenge-multiple myeloma kanye ne-amyloidosis okuyizifo ezibucayi futhi ezihlalayo. Ukuthathwa komchamo isikhathi esingamahora angama-24 kutholwa iphrotheyini ebizwa nge-Bence Jones kwenziwa ezigulini ezinalezi zifo ukusiza udokotela ukuba akwazi ukubheka iziguli ezelashwayo. Ukuthathwa kwesampula lomchamo wamahora angama-24 kunzima, isibonelo nje, kusukela ngehora lesi-6 ekuseni kuye ehoreni lesi-6 ekuseni ngosuku olulandelayo. Senza lolu cwaningo ukuze sithole ukuthi singakwazi yini ukuba sithathe isampula elilodwa lomchamo esikhundleni sokuthatha umchamo wamahora angama-24. Lolu cwaningo luzobandakanya ukuthathwa komchamo ngelanga osuke uzothathwa ngalo umchamo wamahora angama-24.

Siyothatha umchamo wakho zikhathi zonke uma uzothathwa umchamo wamahora angama-24 wephrotheyini i-*Bence Jones* esikhathini esiyizinyanga ezi-3. Ekupheleni kocwaningo, okuyoba emva konyaka, ayolahlwa onke amasampula omchamo ayobe esele. La masampula ayosetshenziselwa lolu cwaningo kuphela. Amasampula ayothathwa ngosuku osuke uze ngalo emtholampilo ukuze ungabi nezinsuku ezengeziwe ozozizela le phrojekthi.

Ukuzibandakanya kwakho kungase kusisize ukuba sithole izimpendulo zombuzo wocwaningo futhi kungasiza izizukulwane ezizayo ngokuthi seziyothathwa isampula lomchamo elilodwa kunokuthi uthathwe isikhathi esingamahora angama-24.

Ulwazi esiluqoqa kulolu cwaningo luyohlala luyimfihlo. Yonke imininingwane yakho iyoba nenombolo esikhundleni segama lakho. Abacwaningi kuphela abayokwazi inombolo yakho futhi siyoluvalela lolo lwazi.

Uyazikhethela ukubamba iqhaza kulolu cwaningo. Noma ungakhetha ukuzibandakanya nocwaningo noma ukhethe ukungazibandakanyi nocwaningo, akukho lutho oluzoshintsha noma ozokwephucwa khona ngokuza kulo mtholampilo. Ungabuye ushintshe umqondo wakho uyeke ukuba yingxenye yocwaningo. Kuyilungelo lakho okumele lihlonishwe.

Uma ufisa ungangibuza noma yimuphi umbuzo omayelana nocwaningo. Unayo imibuzo?

Isitifiketi Sencwadi Yemvume Ngilufunde lonke ulwazi olungenhla, noma ngifundelwe lona. Ngibe nethuba lokub	บเรล
imibuzo ngalo, yonke imibuzo engiyibuzile ngenelisekile ngokuphendulwa kwa	
	ıyu.
Ngiyavuma ukubamba iqhaza kulolu cwaningo.	
Igama Lombambiqhaza	
Isiginesha Yombambiqhaza	
· ————	

Usuku	Usuku/inyanga/unyaka
Kongakwazi ukufunda n	okubhala
Bengikhona kufundwa	n ngobunyoninco incwadi yesicelo semvume kongaba unikeziwe ithuba lokubuza imibuzo. Ngiyaqinisekisa ukuthi
	ıkubamba iqhaza ocwaningweni.
	KANYE nokucindezela isithupha kombambiqhaza
	Usuku/inyanga/unyaka
Isitatimende somcwaning	
•	azi lowo ongaba ngumbambiqhaza, ngawo onke amandla ami
	ımbambiqhaza ukuqonda konke ukuthi kuzokwenziwa lokhu
okulandelayo:	
1. Konke ukuthathwa	komchamo okwengeziwe kuyokwenziwa ngosuku olufanayo
lokuthathwa komchamo	
	9
	umbambiqhaza wanikwa ithuba lokubuza imibuzo mayelana
	imibuzo ayibuza yaphendulwa ngokufanelekile. Ngiyaqinisekisa
ukuthi umbambiqhaza	akaphoqwanga ukunikeza imvume kepha uzivumele yena
ngokwakhe. Ikhophi vale	ncwadi yemvume unikeziwe naye umbambiqhaza.
	i/umuntu ocela imvume
	umuntu ocela imvume
Usuku	
Usuku/inyanga	/unyaka
Lolu cwaningo lubukiwe lwase luv	unywa yi-UKZN Biomedical research Ethics Committee (inombolo yokuvunywa).Uma
kuba khona izinkinga noma imibu	zo ungaxhumana nomcwaningi ku-0312402558 noma i-UKZN Biomedical Research Ethics
Committee, imininingwane yokuxh	umana imi kanje: BIOMEDICAL RESEARCH ETHICS ADMINISTRATION
Research Office, Westville Campus	
Govan Mbeki Building	
University of KwaZulu-Natal	
Private Bag X 54001, Durban, 4000	
KwaZulu-Natal, SOUTH AFRICA	

Tel: 27 31 2602486 - Fax: 27 31 2604609 Email: BREC@ukzn.ac.za

Appendix 5: Raw data

	Summary	Summary Data Sheet											
	Study No	Diagnosis	Age (years)	Gender	Race	BMI kg/m2	BJP mg/24hr	MWG response	% BJP	JP: creat ratio (mg/mmol)	E 1 mg/24hr	E 2 mg/24hr	IMWG response
1	MM001	MM	28	Н	В	22,5	0252	2	8′29	125,6	1256,3	949,5	2
2	MM002	MM	74	Μ	В	23,1	1080	2	20	128,6	1286,2	1705,5	2
3	MM004	MM	57	F	В	24,6	3430	2	82,3	163,6	1636,4	1562,3	2
4	MM010	MM	61	Μ	В	15,4	069	2	53	0006	9'668	795,3	2
2	MM011	MM	59	F	В	36,7	רסר	0	40,6	35,1	350,8	437,3	2
9	MM014	MM	54	Н	В	24,9	0089	2	6′29	471,3	4712,5	4499,2	2
7	MM020	MM	59	Μ	В	23,5	רסר	0	15,5	46,1	460,5	447,8	2
8	MM021	MM	58	Μ	В	33,1	380	2	26,2	59,8	298,0	1046,6	2
6	MM022	MM	63	Μ	В	28,7	רסר	0	25,2	39,7	397,1	575,8	2
10	MM024	MM	47	Δ	В	26,3	14400	2	52,2	714,3	7142,9	7,7656	2
11	MM025	MM	99	Н	В	29,8	0858	2	98	513,7	5137,1	5790,0	2
12	MM028	MM	99	F	В	32,8	9240	2	81,6	528,6	5286,4	5888,2	2
13	MM030	MM	53	ч	В	33,3	380	2	53	39,4	393,5	412,3	2
14	MM031	MM	92	Μ	В	22,3	7660	2	57,1	191,8	1917,8	2034,4	2
15	650MM	MM	22	Ь	-	35,4	250	2	35	0,0	0′0	0'0	0
16	MM060	MM	47	F	В	20,3	2480	2	81,9	448,2	4481,6	3743,9	2
17	MM061	MM	49	Μ	В	30,4	089	2	37	21,2	211,8	269,7	2
18	MM062	MM	49	F	В	27,4	1200	2	55	162,8	1628,2	1403,4	2
19	MM063	MM	55	Μ	В	35,5	3600	2	69	22,7	2'956	1945,2	2
70	MM064	MM	48	Μ	В	26,2	290	2	34,5	29,7	297,0	383,4	2
21	MM065	MM	47	F	В	56,6	1610	2	75,9	54,9	549,5	728,6	2
22	MM066	MM	26	F	В	27,0	6150	2	77	708,8	7087,6	6484,8	2
		Summary Data Sheet	ta Sheet										
		NOTES:	MM: Multiple myeloma	myeloma									
			M:Male F: Female	male									
			B: Black I: Indian	an									
			E1: Estimated 24-hour BJP	24-hour BJP	(mg/24hou	ır)= Urine BJ	P/Creatinine ration	(mg/24hour) = Urine BJP/Creatinine ratio (mg/mmol) X10					
			E2: Estimated 24-hour BJP			ır) = Urine BJP,	/Creatinine ratio	(mg/mmol) x 15mg,	/kg for women or	(mg/24hour) = Urine BIP/Creatinine ratio (mg/mmol) x 15 mg/kg for women or x 20mg/kg for men (convert mg/kg to mmol/kg by multiplying 0.00884;	cg to mmol/kg by multiplyi	ng 0.00884)	
			LDL: Lower than Detectab	an Detectable	le limit								
			Red: Random urine	urine									