

Scientific Breakthroughs From Siggaard–Andersen Nomogram To Latest Developed Graphical Tool Representation For Arterial Blood Gas (ABG) Interpretation

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ABSTRACT

Arterial Blood Gas (ABG) interpretation plays an immense role in critical care medicine. Various approaches are available to interpret ABG data. The most commonly used approaches are the physiological approach using the bicarbonate-carbon dioxide buffer system and the base excess approach developed by Astrup and Siggaard Anderson. Physico-chemical approach is used to explain the causative mechanism of metabolic acid base disturbances. The graphical representation may serve as an useful tool for easier and quicker interpretation of ABG data. But only few graph methods are proposed and devised for depicting the various respiratory and metabolic acid–base disturbances. They are used for teaching purposes only and these are not practically convenient to use in clinical settings. Graphical methods for ABG interpretation like Siggaard- Andersen chart (S-A chart), Davenport or Bicarbonate-pH diagram and Grogono diagram are not used frequently in the clinical practice due to certain practical difficulties and providing inaccurate diagnosis in few of the ABG data. A newer graphical method using 4 quadrant method was constructed and proposed by the current author. The purpose of this review is to discuss in detail the construction, interpretation, advantages, clinical utility and the limitations of the various proposed graphical tools.

KEY WORDS: Graphical Tool, Arterial Blood Gas, Siggaard-Andersen chart, Davenport diagram, Grogono diagram, 4 quadrant Graphical Tool

INTRODUCTION:

Blood Gas Analyzer remains an indispensable medical equipment in emergency department and critical care medicine. **Arterial Blood Gas (ABG)** interpretation plays an immense role in emergency health conditions. Various approaches are available to interpret ABG data.[1] The commonly used approaches are the **physiological approach** using bicarbonate and the **base excess** approach developed by Astrup and Siggaard Anderson.

The most commonly used physiological approach is based on the bicarbonate-carbon dioxide buffer system(based on $p\text{CO}_2$ / carbonic acid/bicarbonate equilibrium).[2,3]

The contribution to the advancement of clinical acid-base chemistry by **Siggaard-Anderson** is significant. He implemented a method based on the **Van Slyke equation** which emphasizes the use of [base excess](#) (BE) or

deficit. In 1948, **Singer and Hastings** proposed the term '**buffer base**' to define the sum of HCO_3^- and the non-volatile weak acid buffers. A change in buffer base directly reflects the changes in the metabolic component. The methods for calculating the **change in buffer base** were later refined by many investigators to yield the base excess (BE) methodology.[4,5]

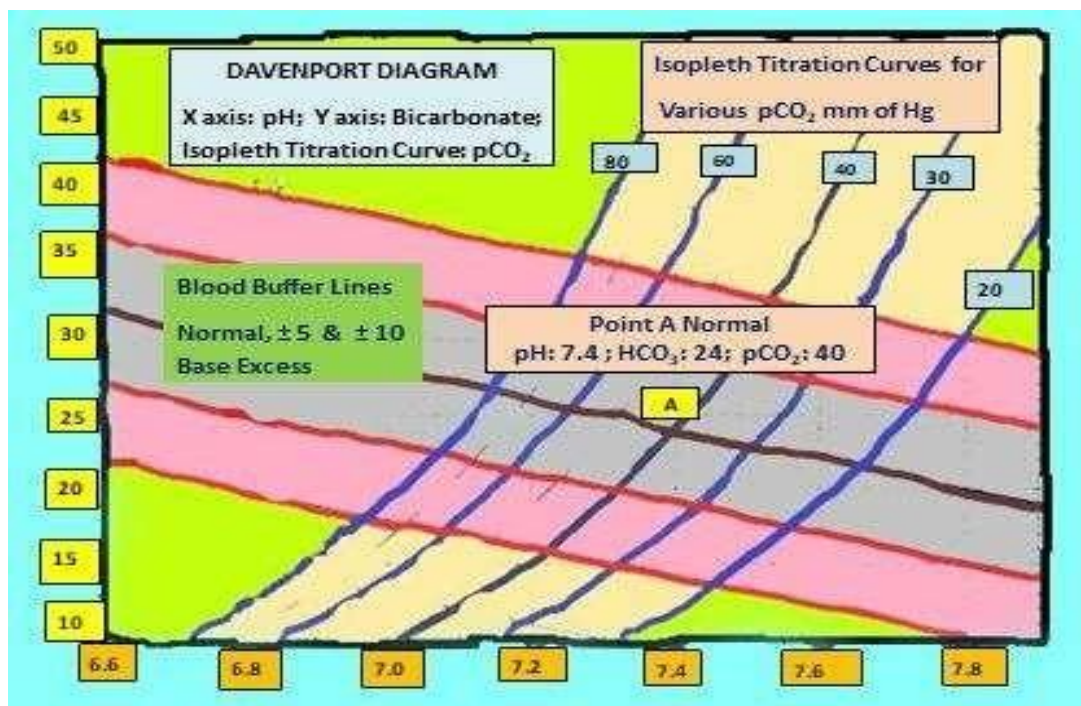
The **base excess calculation** was modified to '**standardize**' the effect of haemoglobin. A blood sample **diluted threefold** with its own plasma may serve as a **model of the extracellular fluid**.[4,5] **Standard base excess (SBE)** or extracellular base excess was developed to represent the in-vivo base excess which is the base excess at haemoglobin concentration of **5g/dl (or 3.1 mmol/L)**.[6,7] *Thus the **Base excess** is defined as the amount of strong acid that must be added to each litre of fully oxygenated blood to return the pH to 7.40 at a temperature of 37°C and a pCO_2 of 40 mmHg.*[8,9,10]

The graphical representation may serve as a useful tool for easier and quicker interpretation of ABG data. But only few graph methods are proposed and devised for depicting the various respiratory and metabolic acid–base disturbances. They are used for teaching purposes only and these are not practically convenient to use in clinical settings.[11,12] Graphical methods for ABG interpretation like **Siggaard-Andersen chart (S-A chart)**, **Davenport or Bicarbonate-pH diagram** and **Grogon diagram** are not used frequently in the clinical practice due to certain practical difficulties and providing inaccurate diagnosis in few of the ABG data. A newer graphical method using **4 quadrant method** was constructed and proposed by the **current author (Rajini Samuel)**.[11-15] The knowledge of the scientific breakthroughs from Siggaard–Andersen nomogram to latest developed graphical tool representation for Arterial Blood Gas (ABG) interpretation may help us to understand the contribution of various researchers in the field of acid base clinical chemistry. The purpose of this review is to discuss in detail the construction, interpretation, advantages, clinical utility and the limitations of the various proposed graphical tools.

DAVENPORT DIAGRAM:

The plot of both the Henderson-Hasselbach Equation and the Van Slyke equation has been called ““Davenport or Bicarbonate-pH diagram”. In the davenport diagram, **x-axis** represents the **pH** values, **y-axis** represents the plasma **bicarbonate** values and the **curved lines (isopleths)** denote the **pCO_2** values derived using the Henderson Hasselbach Equation(depicted in **figure 1**). These titration curves are called **isopleths**, because they are generated at a fixed partial pressure of carbon dioxide (shown in the **figure 1**). Each isopleth pCO_2 curved line has a **fixed pCO_2 value** for a given variable pH and calculated bicarbonate concentration.[16,17]

FIGURE 1: DAVENPORT DIAGRAM



The **blood-buffer line** is the relationship between HCO_3^- and pH as carbonic acid is added to whole blood. This **negative slope line** denotes the **buffering capacity** of blood which is achieved by varying pCO_2 through variation in alveolar ventilation.

CONSTRUCTION OF THE DAVENPORT DIAGRAM:

The **normal blood buffer line** goes through **point A** ($\text{pH} = 7.4$, $[\text{HCO}_3^-] = 24$, $\text{pCO}_2 = 40 \text{ mm Hg}$) on the Davenport diagram (depicted in **figure 1**). If HCO_3^- is added or eliminated while maintaining the pCO_2 fixed at **40 mmHg**, it will move **along the 40 mmHg isobar away from point A**.

The blood sample from a healthy patient is collected and then placed in a chamber in which the partial pressure of carbon dioxide (pCO_2) is held at 40 mmHg. The pH and bicarbonate concentration are measured once equilibrium is reached, and then they are plotted on a diagrammatic chart. Then the pH of the blood sample is modified, first by adding a **strong acid** and then by adding a **strong base** but pCO_2 is kept constant in the chamber. A **titration curve** for a **given pCO_2 value** is obtained by the changes in the pH and bicarbonate concentration for a given pCO_2 value (**40 mm Hg**). Similarly with a new, identical blood sample from the same patient the experiment is performed with the chamber reset to a pCO_2 of **60 mmHg**. After equilibration, a new point is reached, indicating a new pH and a new bicarbonate concentration (shown in **figure 1**). The **bicarbonate** concentration at the newer, **higher pCO_2** will be **higher** than the **value at pCO_2 of 40** and the pH will be lower. [16,17] A **titration curve** for a **given pCO_2 value** is obtained by the changes in the pH and bicarbonate concentration for a given pCO_2 value (**60 mm Hg**). A series of isobars can be constructed depending on the pCO_2 values.[16]

The central **point A** (**figure 1**) is the normal value of pCO_2 (40mm Hg), pH (7.40) and bicarbonate (24 mmol/L). If this experiment is repeated at **various partial pressures of carbon dioxide**, with a new identical blood sample from the same patient **a series of points** will be obtained. A **line is drawn** through these points connecting the central point with other points representing the **buffer line**(clearly depicted in **figure 1**) which shows the possible values of pH and bicarbonate achieved by changing pCO_2 through changes in minute ventilation.

A series of **parallel blood buffer lines** can be constructed by changing the **Base excess values**. The normal buffer line passes through the base excess value of zero. The other parallel blood buffer lines drawn above and below the normal buffer lines depend on the **positive** and **negative** values of the base excess respectively (clearly shown in **figure 1**).

As pCO_2 is increased, the magnitude of the resulting change in pH is dependent on the buffering power of the non-bicarbonate buffers present in the solution. As pCO_2 increases, it reacts with water molecules to form carbonic acid which dissociates into hydrogen and bicarbonate ions. The albumin, hemoglobin, and phosphate buffer system help in buffering of hydrogen ions. So, the concentration of bicarbonate increases as pCO_2 also increases. The generation of a bicarbonate molecule is accompanied with release of a proton, so the pH is decreased. If **strong non-bicarbonate buffers** are present, then they will quickly absorb the vast majority of protons released by the generation of bicarbonate, and pH will change very little for a given rise in bicarbonate concentration. This results in a **buffer line** with a **very steep slope**. On the other hand, if **only weak non-bicarbonate buffers** are present (or **no non-bicarbonate buffer**), then a much larger change in pH will be observed for a given change in bicarbonate concentration, and the **buffer line** will have a **slope closer to zero**.[16]

The slope of the line will **never** actually reach **zero (horizontal)** under equilibrium conditions, even in the complete absence of non-bicarbonate buffers because the production of protons resulting from an increase in pCO_2 is concomitant with the production of bicarbonate ions. Thus, a decrease in pH resulting from an increase in pCO_2 always occurs with some minimal increase in bicarbonate concentration. Similarly, an increase in pH resulting from a decrease in pCO_2 must occur with some minimal decrease in bicarbonate

concentration.[16]

INTERPRETATION OF THE DAVENPORT DIAGRAM:

A **series of isobars** depending on the **pCO₂** values and a **series of parallel blood buffer lines** depending upon the **Base Excess (BE)** values are constructed in the davenport diagram. The **respiratory status** is assessed by the location on the **pCO₂ isobar** and the **metabolic status** is evaluated by the location on the **blood buffer line**. A **negative BE** (base deficit) indicates metabolic acidosis and a **positive BE** (base excess) denotes the presence of metabolic alkalosis. The concentration of bicarbonate corrected to pH of 7.4 is called the **Van Slyke Standard Bicarbonate (VSSB)** or **predicted bicarbonate value** (term used in some research articles). The difference between observed or actual bicarbonate (calculated from Henderson equation) and the predicted bicarbonate value (or Van Slyke Standard Bicarbonate) denotes the **delta Vanslyke Standard Bicarbonate (ΔVSSB)** which is numerically equal to the **base excess value**. [16,18,19]

An acid-base disturbance is characterized by two parameters namely the **delta Vanslyke Standard Bicarbonate (ΔVSSB)** and **delta beta (Δβ)**. **ΔVSSB** denotes the **change in the position** of the Van Slyke curve and **delta beta** indicates the **change in the slope** of the Van Slyke curve. Delta VSSB measures the magnitude of a metabolic acidosis or alkalosis and is numerically equal to base excess. Delta beta indicating the buffer strength measures the change in the plasma's ability to resist perturbations of pH.

The observed [HCO₃⁻] can be calculated from the pH and pCO₂ values. The predicted [HCO₃⁻] is the value predicted from the normal blood buffer line and the observed pH which can be calculated using the given below equation.[16]

$$\text{Predicted [HCO}_3^-] = 24 + \beta \cdot (7.4 - \text{pH})$$

The buffer value β depends on the concentration of haemoglobin, albumin and phosphate values. Multiple formulas exist to calculate the same. Base Excess is defined as the difference between the observed and predicted [HCO₃⁻].

$$\text{BE} = \text{observed [HCO}_3^-] - \text{predicted [HCO}_3^-]$$

$$= [\text{HCO}_3^-] - \{24 + \beta \cdot (7.4 - \text{pH})\}$$

$$= [\text{HCO}_3^-] - 24 - \beta \cdot (7.4 - \text{pH}) \text{ or}$$

$$\text{BE} = [\text{HCO}_3^-] - 24 + \beta \cdot (\text{pH} - 7.4)$$

{(7.4 - pH) is changed to (pH - 7.4), so the sign is changed}

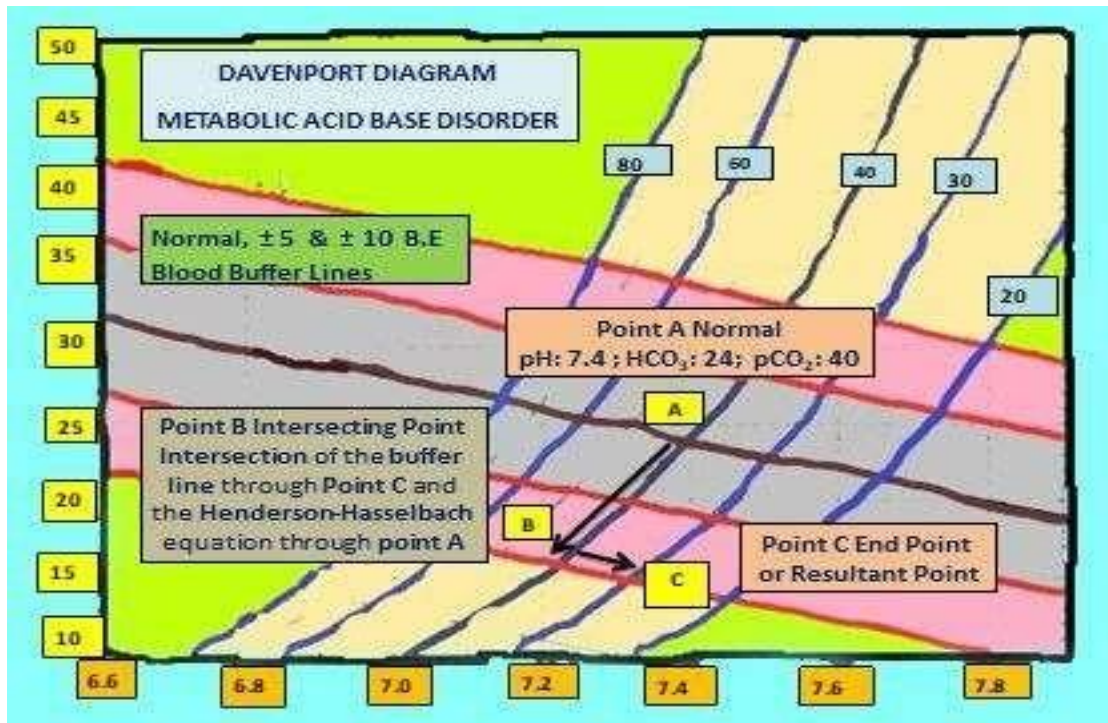
The above equation can be compared with the commonly used formulae to calculate the base excess values.

With a **pure metabolic disturbance** the point remains on the **normal pCO₂** (40 mm of Hg) isobar, but can develop **either a negative BE or a positive BE** by **moving away from the normal point along this isobar** (seen in **figure 1**). With a **pure respiratory disturbance**, the point remains on the **normal blood-buffer line**, but can **move along it from one isobar to another**. Movement along this line represents respiratory acidosis (decreasing pH) or respiratory alkalosis (increasing pH) which is clearly seen in **figure 1**. However, acid-base disturbances are rarely pure. A primary metabolic disturbance is compensated by respiratory mechanisms and vice versa. Also, it is not uncommon for two primary disturbances (mixed disorder) to coexist in the same patient at the same time. The components defining the acid base status can be graphically represented using the **relevant isobar**, as determined by the **pCO₂**, and the **relevant blood buffer line**, as determined by the value for **BE**. [16,17]

If a respiratory disturbance is superimposed on a non-respiratory (metabolic) one, movement from **one isobar(pCO₂) to another** along a blood buffer line that is **displaced from the normal line** by a distance equal

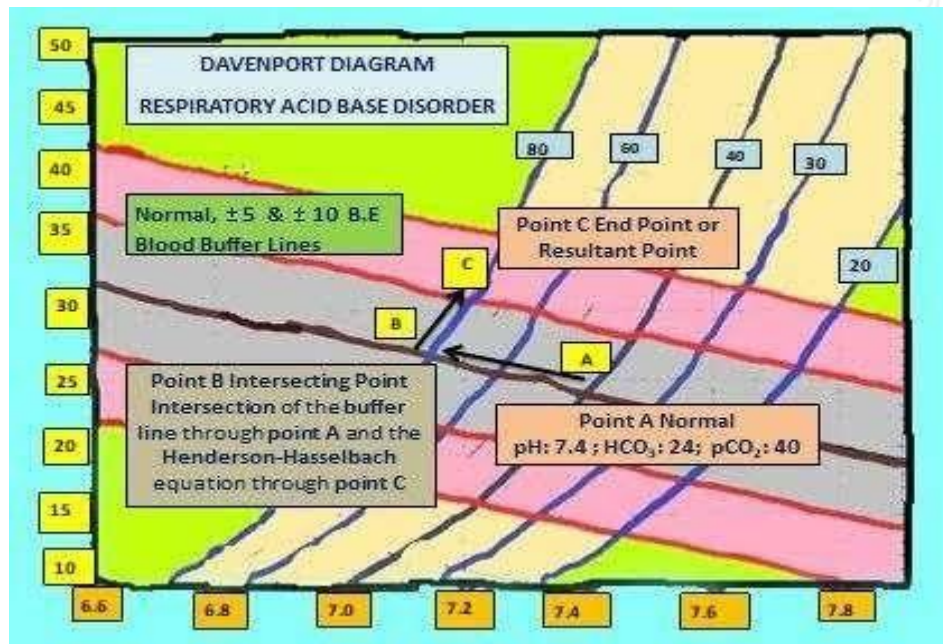
to the value for **BE**. The Henderson-Hasselbalch Curve of pCO_2 40 mm of Hg(**normal pCO_2 isobar**) and the **normal buffer line** passes through the **point A** which is the **starting point** with **normal pH**, pCO_2 and bicarbonate level. The **point C** is the **final result** of the measured pH, pCO_2 and bicarbonate values. (shown in **figure 2**) The **intersection** of the buffer line (**Van Slyke Curve**) with the **isopleth pCO_2 curve** (Henderson Hasselbalch Equation curve) yield the possible combinations of pH and bicarbonate concentration measured in the blood achievable by changes in alveolar ventilation **denoted** by the **point B**. [16,17,20] In case of **metabolic acid base disorder**, Point B is the intersection of the **buffer line** through **Point C** and the **Henderson- Hasselbach** equation through **point A** (clearly depicted in the **figure 2**).

FIGURE 2: Davenport Diagram-Metabolic Acid Base Disorder



In case of **Respiratory acid base disorder** **Point B** is the **intersection** of the buffer line through **point A** and the **Henderson-Hasselbach** equation through **point C** which is clearly shown in the **figure 3**. The resultant point C is the result of the **initial displacement** from the **normal point A** to the point B (**intersecting point**) and the **final displacement** from this intersecting point B to the end point C. The **initial displacement** indicates the **primary acid base disorder** and the **final displacement** represent the **compensatory mechanism** either respiratory compensations in metabolic acid base disorders or metabolic compensations in respiratory acid base disorders. [17]

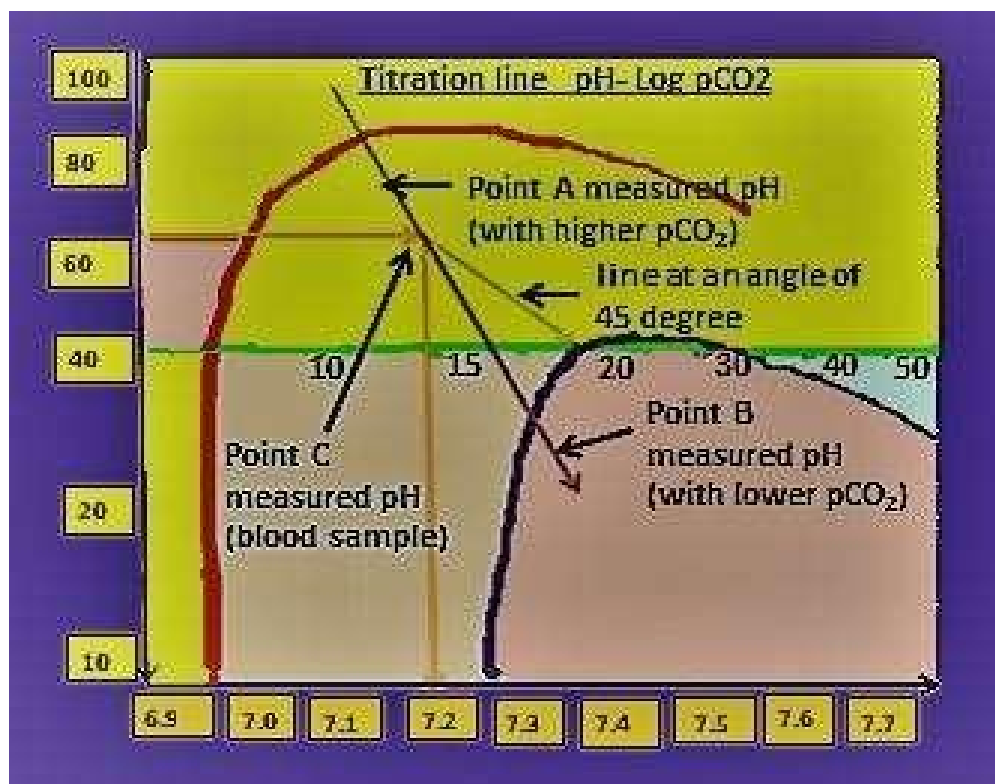
FIGURE 3: Davenport Diagram-Respiratory Acid Base Disorder



THE SIGGAARD-ANDERSEN CURVE NOMOGRAM:

This nomogram has **pCO₂** plotted on a **log scale** on the **vertical axis** and **pH** on the **horizontal axis**. Siggaard-Andersen used the procedure of equilibrium titration curves to determine the values of Buffer base and base excess. The two curves namely the **buffer base curve** and the **base excess curve** are constructed in this coordinate system using the determination of its **pH- log pCO₂ line** (depicted in the figure 4). The **pH** of the blood sample is measured **without** any **equilibration** in vitro. Then the pH of the blood sample is measured **after equilibration** with gas mixtures containing known values of **higher** and **lower pCO₂** values.[21]

FIGURE 4: pH-logpCO₂ line



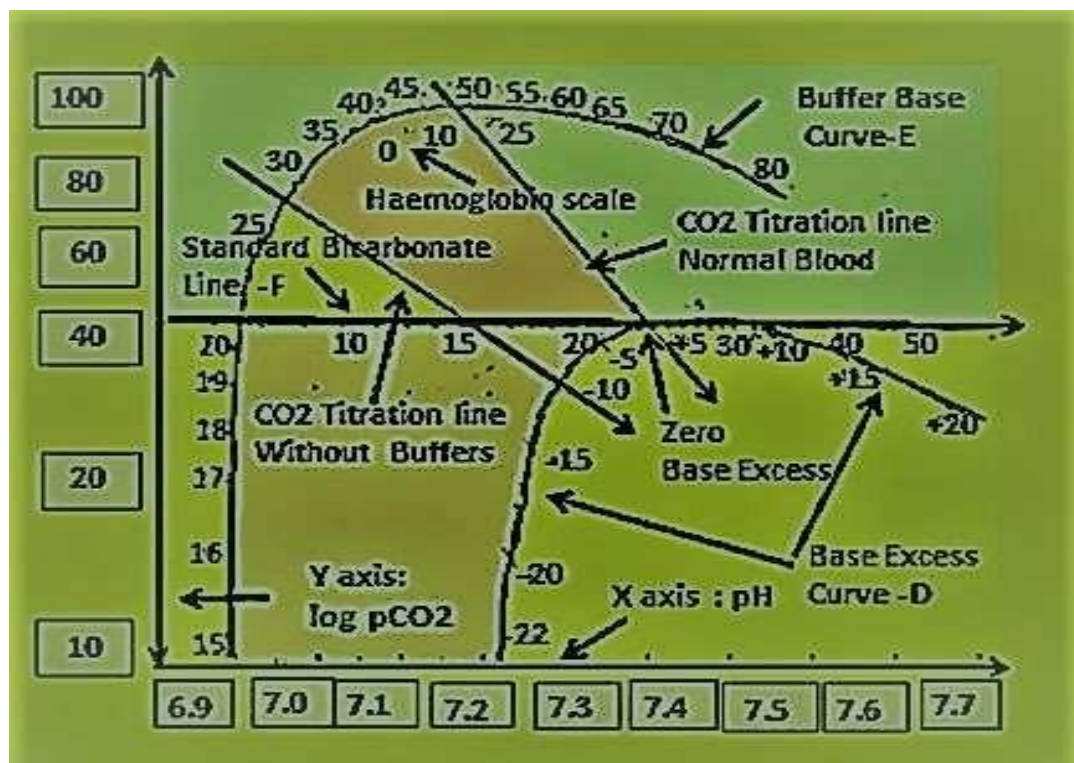
A – denote the measured pH after equilibration with higher pCO_2

B- denote the measured pH after equilibration with lower pCO_2

C- denote the measured pH of the blood sample (without in vitro equilibration)

A line is drawn connecting the points A and B. From the measured pH of the blood sample(without in vitro equilibration), a perpendicular line is drawn that meet the line connecting the points A and B. The value of pCO_2 of the blood sample is read off the ordinate.[21] This is the **Astrup method** of estimation of pCO_2 . The **titration curves** (lines in the semi-logarithmic scale) have **different slopes** after blood titration with carbon dioxide, depending on **haemoglobin** concentrations. If **buffers** were present, the **slope** of the line would be **steeper**. For normal blood containing 15 g of haemoglobin/dL, the CO_2 titration line passes through the 15g/dL mark on the haemoglobin scale (on the underside of the upper curved scale) and the point where the pCO_2 equal to 40 mm Hg and pH equal to 7.40 lines intersect.

FIGURE 5: CO_2 TITRATION LINE WITH BUFFERS AND WITHOUT BUFFERS



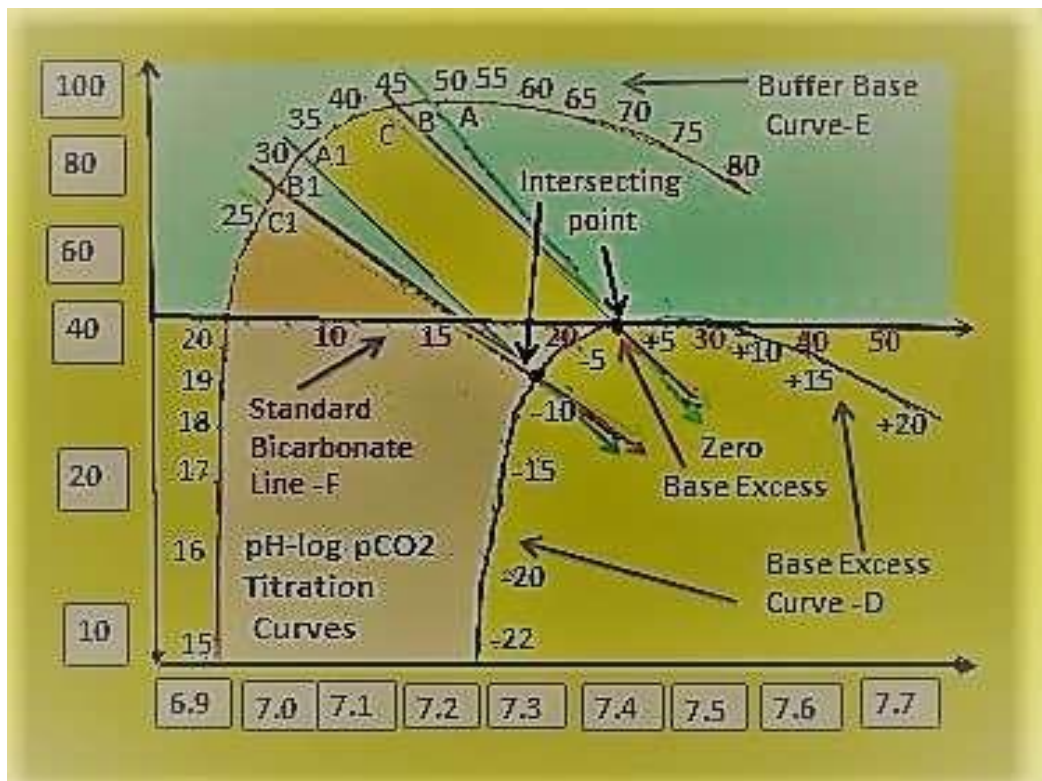
When the haemoglobin content of the blood is low, there is significant loss of buffering capacity, and the slope of the CO_2 titration line diminishes. However, blood of course contains buffers in addition to haemoglobin, so that even the line drawn from the zero point on the haemoglobin scale through the normal pCO_2 -pH intercept is steeper than the curve for a solution containing no buffers. (depicted in the **figure 5**)

CONSTRUCTION OF THE BASE EXCESS AND BUFFER BASE CURVE:

Let **A** denote the **whole blood** , **B** denote the blood with **haemoglobin** concentration of **5g/100ml** and **C** denote the plasma(**blood without haemoglobin**).Then the equilibrium titration was done and the results were plotted in the axis in $\log pCO_2$ /pH coordinates.The **pCO_2 -pH plot** can be constructed by **plotting several pH's while altering pCO_2** . Two points are usually enough because the graph is approximately a straight line

on log $p\text{CO}_2$ -pH coordinates. The titration curves (being lines in the semi-logarithmic coordinates) of the **blood samples with various haematocrit** and the **same BE** always **crossed in the same points**. (shown in the **figure 6**) Similarly, the titration curves of the samples with various haematocrit concentrations (and various BE), but with the **same BB** crossed in the **same points**, too. The **intersection of the points** were the base for the **experimental determination** of the base excess and buffer base curve.[21]

FIGURE 6: Intersecting Points - Determination of Base Excess



The **base excess** values are **changed** by adding defined amounts of **strong acids** or **bases** to blood samples with various haematocrit concentrations. BB and BE change after addition of a strong acid (or strong base) or bicarbonates to the blood sample. Addition of one millimole of a strong acid to one litre of blood results in BE fall by one millimol; addition of one millimol of bicarbonates (or withdrawal of one millimol of hydrogen ions by a reaction with a strong base) results in BB and BE increased by one millimol. The experiment is repeated.

The titration curves of the samples with the **same BE** crossed in the **same points**. By connecting all the intersecting points(of the different base excess values) the base excess curve is constructed(depicted in the **figure 6**). Similarly, the **sample with the same BB** crossed in the **same points**. The buffer base curve is also constructed by changing the concentration of buffer base and connecting all the intersecting points.[21]

Increased partial pressure of carbon dioxide($p\text{CO}_2$) causes it to freely diffuse into the erythrocyte where it reacts with **water molecules** to form **carbonic acid**. The **bicarbonate** and **hydrogen ions** are dissociated from carbonic acid catalysed by the enzyme **carbonic anhydrase** present inside the red blood cells(RBC). Haemoglobin is an important intracellular protein buffer present inside the red blood cells. Bicarbonates are transported into plasma by the concentration gradient (by exchange for chloride ions). Thus, the increase in CO_2 concentrations is associated with the **decrease** in the value of **Buffer Base** concentrations **inside erythrocytes**

(BB_e) or **increase** in the value of **Buffer Base** in the **plasma** (BB_p). The **equilibration titration** with carbon dioxide helps to achieve pCO₂ level at which the Buffer Base (BB) concentrations in erythrocytes and plasma equilibrate (**BB_e = BB_p**). With this level of pCO₂ at BB_p=BB_e, this value determines the point where the titration curves with the same total BB and various haematocrit (Hct) of the blood samples will cut each other in the same point on Siggaard-Andersen nomogram.[21]

$$\begin{aligned}\text{As: } BB &= BB_p (1 - \text{Hct}) + BB_e \text{ Hct} \\ &= BB_p + \text{Hct} (BB_e - BB_p)\end{aligned}$$

The term Hct (BB_e – BB_p) is zero if BB_e = BB_p, so the **whole blood BB** does **not depend on haematocrit**. Thus, the Buffer Base(BB) curve on Siggaard-Andersen nomogram is a **geometric site** of the points where **plasma** and **erythrocytes** have the **same buffer base concentrations**, as at BB_e is equal to BB_p the whole blood BB does not depend on haematocrit (Hct).

Similarly this can be applied for the Base Excess(BE) curve, too.

$$\begin{aligned}\text{As: } BE &= BE_p (1 - \text{Hct}) + BE_e \text{ Hct} \\ &= BE_p + \text{Hct} (BE_e - BE_p)\end{aligned}$$

The term Hct (BE_e – BE_p) is zero at BE_e=BE_p then and the whole blood BE does not depend on haematocrit (Hct) or haemoglobin concentration. Thus, the **BE curve** on Siggaard-Andersen nomogram is a **geometric site** of the points with the **same BE** in the whole blood and plasma, as the whole blood BE does not depend on haematocrit at proper pCO₂ and pH, when BE_e is equal to BE_p.[21]

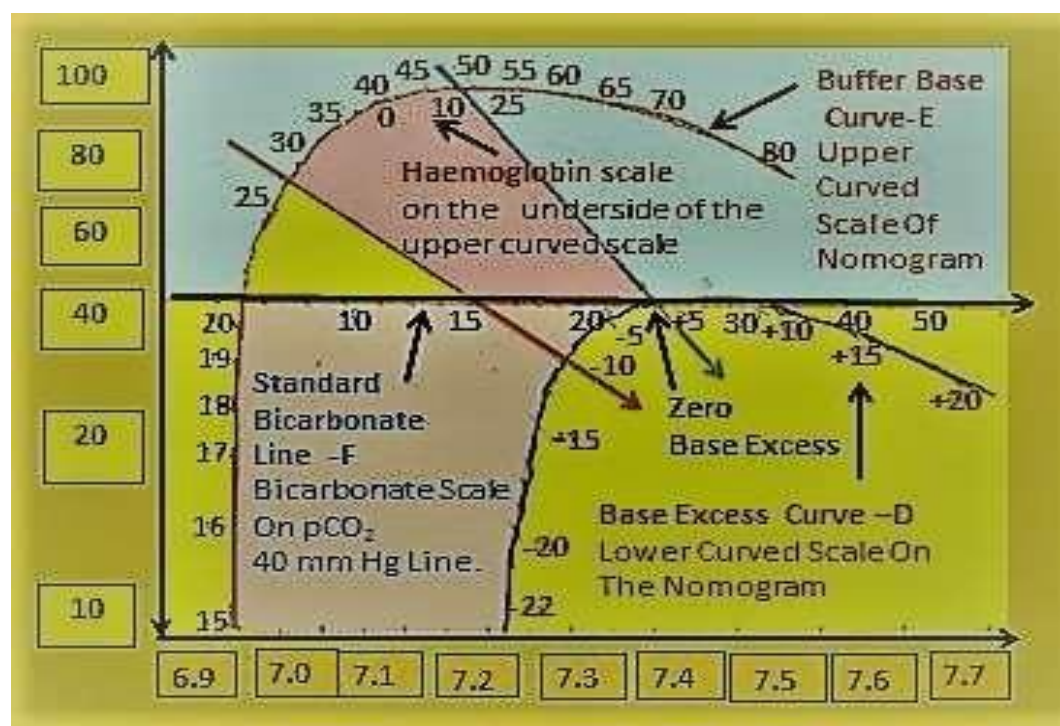
Thus, a nomogram with BE and BB curves with semi-logarithmic coordinates was obtained and these curves enable the determination of BE and BB in the samples to be tested. Siggaard-Andersen used the **mixture of O₂**

- **CO₂** for blood titration with fully oxygen-saturated blood , so the BE curves are those for fully oxygenated blood

$$BE = BE_{ox} + 0.2 \text{ cHB} (1-sO_2)$$

cHb denotes the haemoglobin concentration [g/100ml] and sO₂ denotes the haemoglobin oxygen saturation.[21]

FIGURE 7: DIFFERENT CURVES OF THE NOMOGRAM



Buffer Base Curve (E)-

upper curved scale of nomogram shows the mEq/L of buffer base in the sample

Base Excess Curve(D)-

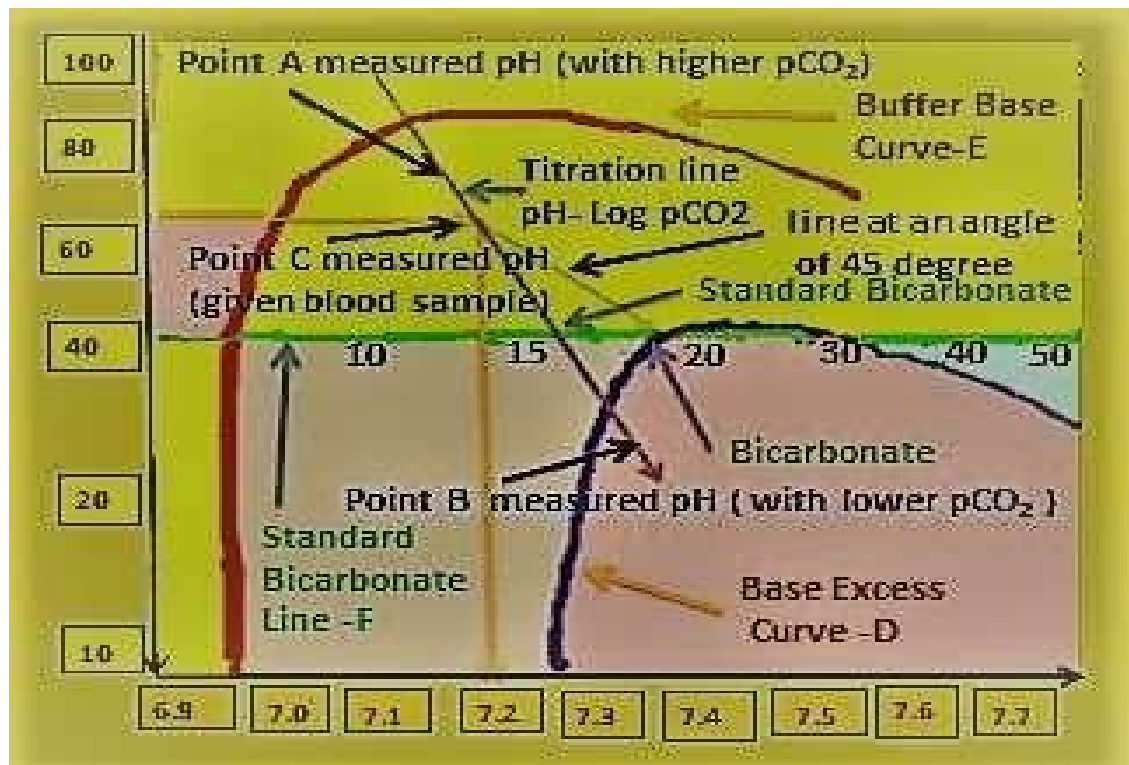
a lower curved scale on the nomogram indicates the base excess.

Standard Bicarbonate(F) Line Scale-

bicarbonate scale on the $p\text{CO}_2$ equal to 40 mm Hg line.

The **point of intersection** of the CO_2 calibration line of the arterial blood sample with the base excess curve(D), buffer base curve(E) and the standard bicarbonate(F) line scale **indicate the values** of base excess, buffer base and standard bicarbonate values respectively.(depicted in **figure 7 and 8**) The **actual bicarbonate** is determined in the graph by drawing a line at an **angle of 45 degree** from the point which denotes the **measured pH** of the blood sample (**without in-vitro equilibration**) to the **standard bicarbonate scale**. This intersecting point denotes the value of the actual bicarbonate.[21,22]

FIGURE 8: Siggaard-Andersen nomogram



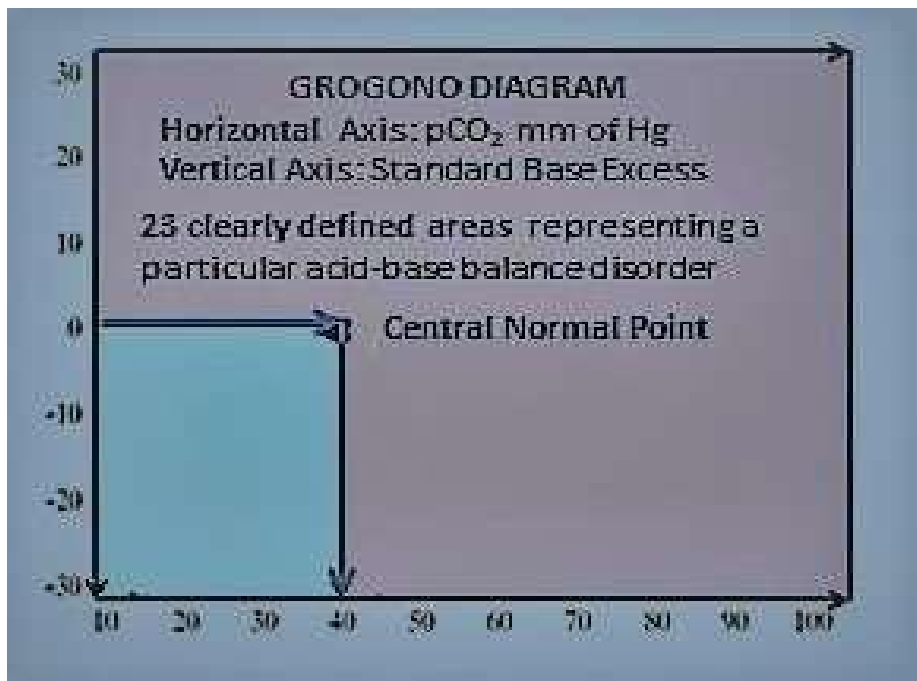
THE OXYGEN-STATUS-ALGORITHM (OSA) SOFTWARE:

This software was developed by the Radiometer-Copenhagen Company and installed in all blood gas analyzers manufactured by Radiometer-Copenhagen. This is based on the diagram and workdone by Siggaard-Andersen.[23]

THE GROGONO DIAGRAM: The initial concept for the construction of the Grogono diagram was based on the previously proposed Siggaard- Andersen and J. Severinghaus diagrams. It has two axes where the $p\text{CO}_2$

representing the **respiratory** component of the acid base disorder is plotted in the **horizontal** axis and the **standard base excess** denoting the **metabolic** component of the acid base disorder is plotted in the **vertical** axis. This diagrammatic representation tool consists of **23 clearly defined areas**, each one representing a particular acid-base disorder. (depicted in **figure 9**) This new diagrammatic approach proposed by A. Grogono is superior to the S- A chart, but it cannot be utilized for the interpretation of acid-base disorders, because it has not provided accurate interpretation in at least 25% of the ABG data. Despite its higher diagnostic agreement compared to the classic Siggaard-Andersen diagram, Grogono diagram is not superior to the Oxygen Status Algorithm software program. [23,24]

FIGURE 9: Grogono Diagram



A NOVEL FOUR QUADRANT GRAPHICAL METHOD:

The great technological advance in laboratory methods and instruments that has occurred in the last decades has not brought a respective development in the field of diagnosis of various acid base disturbances. [23] A new graphical tool was developed by the **current author** for ABG interpretation using **standard base excess** in the **x axis** and **ratios** derived using bicarbonate, standard bicarbonate and carbonic acid values in the **y axis**. The various acid-base disturbances are plotted and analysed in a four quadrant graph method. [25,26,27]

Modified Henderson equation is applied to calculate the concentration of bicarbonate. The concentration of bicarbonate varies with $p\text{CO}_2$, so Standard bicarbonate is utilized which denotes the concentration of bicarbonate with a normal PaCO_2 (40 mmHg) and a normal $p\text{O}_2$ (over 100 mmHg) at a normal temperature (37°C). [28,29] Under normal ventilatory conditions both are similar but in abnormal ventilation (either hypo or hyperventilation) both the actual bicarbonate and the standard bicarbonate concentrations deviate from each other depending on the changes in magnitude and direction of variation in $p\text{CO}_2$. [13,14,30]

Ratio 1 = $\text{HCO}_3 / \text{Std HCO}_3$

Ratio 2 = $(\text{HCO}_3 / \text{H}_2\text{CO}_3) - (\text{Std HCO}_3 / \text{H}_2\text{CO}_3)$ **OR**

$$\text{Ratio 2} = (\text{HCO}_3 - \text{Std HCO}_3) / \text{H}_2\text{CO}_3$$

$$\text{Modified Ratio 2} = (\text{Std HCO}_3/1.2) - (\text{HCO}_3/\text{H}_2\text{CO}_3)$$

At pCO_2 40 mmHg, H_2CO_3 concentration is 1.2 mmol/L. So **ratio 2 is modified here** to correlate Std bicarbonate with H_2CO_3 concentration of 1.2 mmol/L. The **Compensation bedside rules** using the **Boston Method** (6 rules) involving bicarbonate can be applied for the assessment of compensation and to identify the presence of compensations or a mixed acid base disorder in clinical practice. [13,14,29,30] The graphical relationship between pCO_2 and **ratio 1**, **ratio 2** and **modified ratio 2** were clearly depicted in the **figure 10**. A **4 quadrant graphical tool model constructed** using **standard base excess** and either (pCO_2 - 40 mm of Hg) parameter or **modified ratio 2** is shown in **figure 11** which clearly shows the plot for various acid base disturbances.

FIGURE 10: NOVEL DERIVED RATIOS

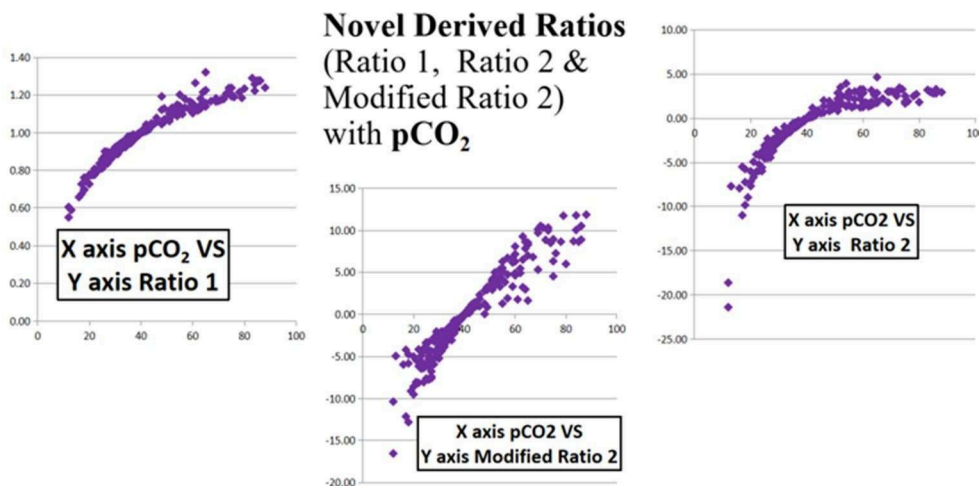
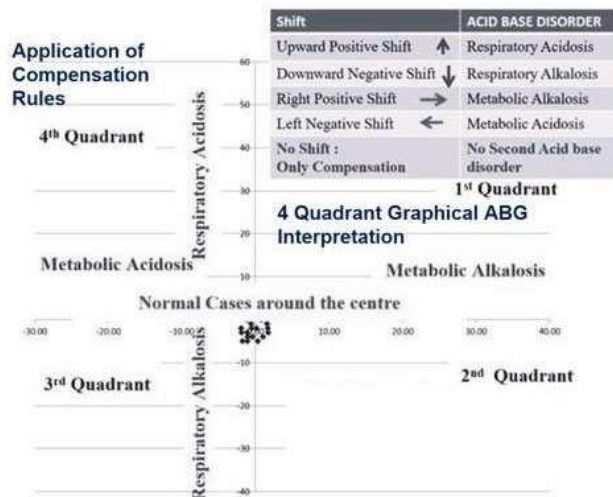


FIGURE 11: 4 Quadrant Graphical Tool Model for ABG Interpretation



If the value of **standard base excess** is $> +2$ mmol/L then it denotes **metabolic alkalosis** and if the value is < -2 mmol/L then it denotes **metabolic acidosis**. The normal range for **pCO₂** is **35 to 45** mm Hg. **Higher pCO₂** values in **respiratory acidosis** and **lesser pCO₂** values in **respiratory alkalosis** are seen. [13,14,29,30]The **modified ratio 2** is **greater positive** for respiratory acidosis and **greater negative** for respiratory alkalosis. Using the **concept of shift** in the **plotted point position** after the application of compensation rules, the mixed acid base disorders can be identified in this graphical tool. [14,30] The scientific breakthroughs from Siggaard-Andersen nomogram to latest developed graphical tool representation for Arterial Blood Gas (ABG) Interpretation may help us to observe the transition, transformation and evolution in the field of diagnosis of acid base clinical chemistry through graphical representation methods.

CONCLUSION:

The detailed discussion of the construction, clinical interpretation and the limitations of the various proposed graphical tools may help us in teaching the students to gain in-depth knowledge in acid base chemistry. The latest proposed four quadrant graph method which involves the application of compensation rules looks very simple, easy and user friendly to overcome the arduous task of complex acid base disorders involving various compensations and mixed disorders.

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