# Michaelis-Menten

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## Question:

The below data represents the substrate concentration (S) and the observed velocity (v) based on an enzymology experiment.

- 1)Fit a non linear regression model that relates velocity to concentration using Michaelis-Menten equation .
- 2)Analyse and examine whether you can fit a simple linear regression model that relates velocity and substrate concentration by using any suitable transformation.

## Variable of Interest:

- S: Concentration of the chemical substance
- V: Velocity of the chemical substance.

## Import dataset

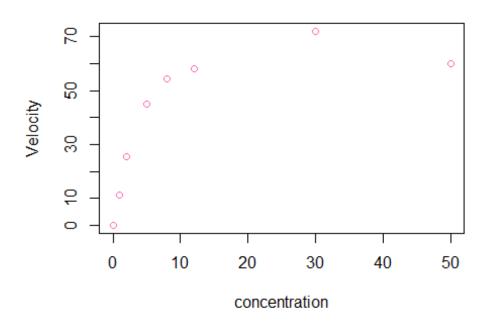
```
library(readxl)

rate <- read_excel("C:/Users/mayur/Desktop/Mstat/Semesters/Tri-sem2/Regression/Dataset
/rate.xlsx")

View(rate)
attach(rate)

plot(rate$S,rate$V,col="hotpink",xlab="concentration",ylab="Velocity",main="substrate
concentration and velocity")
```

# substrate concentration and velocity



```
model1=nls(V\sim Vmax*S/(k_m+S), start=c(Vmax=72, k_m=5), data=rate)
model1
## Nonlinear regression model
## model: V \sim Vmax * S/(k_m + S)
   data: rate
##
## Vmax k m
## 73.261 3.437
## residual sum-of-squares: 156.4
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 6.202e-06
summary(model1)
##
## Formula: V \sim Vmax * S/(k_m + S)
##
```

the value for k and  $v_max$  values are not zero and hence a significant value.  $r^2$  not a good metric to study the fit since x and y are not linear . thus the value of  $v_max$  is 73.26 and  $k_m$  is 3.43

```
confint(model1)

## Waiting for profiling to be done...

## 2.5% 97.5%

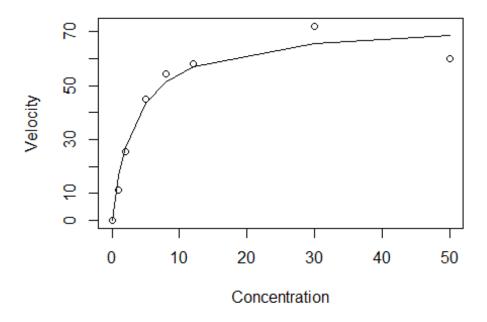
## Vmax 63.302914 84.889668

## k_m 1.958589 5.794501
```

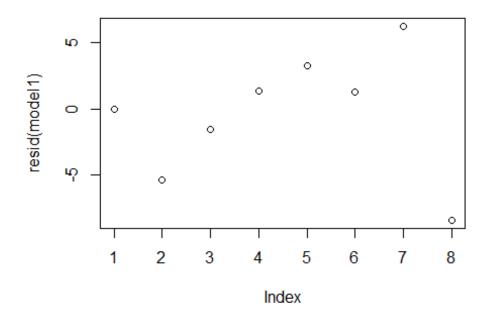
95% of the time the true value will lie within this range

Vmax 63.302914 84.889668 k\_m 1.958589 5.794501

```
plot(S,V,xlab="Concentration", ylab="Velocity")
lines(S,predict(model1))
```



```
summary(model1)
##
## Formula: V \sim Vmax * S/(k_m + S)
##
## Parameters:
##
     Estimate Std. Error t value Pr(>|t|)
## Vmax 73.2614 4.5825 15.987 3.8e-06 ***
## k_m 3.4372 0.8173 4.205 0.00565 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5.106 on 6 degrees of freedom
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 6.202e-06
plot(resid(model1))
```



```
resid(model1)

## [1] 0.000000 -5.410856 -1.548379 1.384119 3.255559 1.250639 6.269496

## [8] -8.449099

## attr(,"label")

## [1] "Residuals"
```

# normality

```
shapiro.test((resid(model1)))

##

## Shapiro-Wilk normality test

##

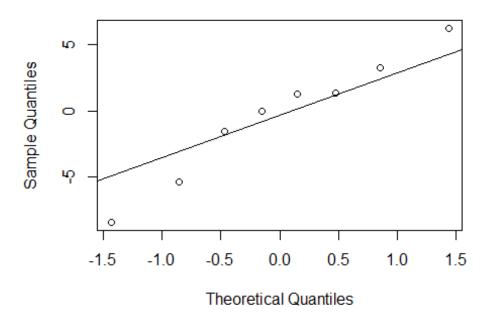
## data: (resid(model1))

## W = 0.96435, p-value = 0.8504

qqnorm(resid(model1))

qqline(resid(model1))
```

# Normal Q-Q Plot



since the p value is greater than alpha(0.05) thus it follows normality condition. We can also infer the same from the above plot.

## Transformation to linear model.

To perform linear regression, we need to have a function of V be linearly related to a function of [S]. This is achieved via taking the reciprocal of both sides of this equation.

$$\frac{1}{V} = \frac{[S] + K_m}{V_{\mathsf{max}}[S]} = \frac{1}{V_{\mathsf{max}}} + \frac{K_m}{V_{\mathsf{max}}} \frac{1}{[S]}$$

Thus, we have a linear relationship between 1/V ,the response variable, and 1/S ,the explanatory variable

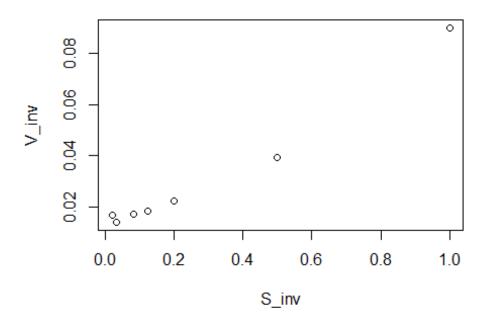
library(readxl)

rate1 <- read\_excel("C:/Users/mayur/Desktop/Mstat/Semesters/Tri-sem2/Regression/Datas et/rate1.xlsx")

```
View(rate1)
attach(rate1)

S_inv=1/(s)

V_inv=1/(v)
plot(S_inv,V_inv)
```



```
model2=lm(V_inv~S_inv)
model2

##

## Call:

## lm(formula = V_inv ~ S_inv)

##

## Coefficients:

## (Intercept) S_inv

## 0.00996 0.07551

summary(model2)
```

```
##
## Call:
## lm(formula = V_inv \sim S_inv)
##
## Residuals:
      1
            2
                  3
                                         7
## 0.0046231 -0.0083435 -0.0027401 -0.0010499 0.0009298 0.0014119 0.0051687
##
## Coefficients:
##
        Estimate Std. Error t value Pr(>|t|)
## S inv
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.005083 on 5 degrees of freedom
## Multiple R-squared: 0.9712, Adjusted R-squared: 0.9655
## F-statistic: 168.7 on 1 and 5 DF, p-value: 4.824e-05
```

# Normality and Constant Variance Assumption

```
shapiro.test(resid(model2))

##

## Shapiro-Wilk normality test

##

## data: resid(model2)

## W = 0.93643, p-value = 0.6068

library(Imtest)

## Loading required package: zoo

##

## Attaching package: 'zoo'
```

```
## The following objects are masked from 'package:base':
##
## as.Date, as.Date.numeric

bptest(model2)
##
## studentized Breusch-Pagan test
##
## data: model2
## BP = 1.2317, df = 1, p-value = 0.2671
```

since for both p values are greater than alpha, we accept null hypothesis and say that it follows normal distribution and has constant variance

Conclusion: Thus, we have demonstrated Michaelis-Menten model for the given data-set and have demonstrated that an inverse transformation to the model can make it a linear model.