

Capstone

"The first step in the process is to subtract long-term dependencies in light-curve data by subtracting out a third-order polynomial that is fitted on the light curve"

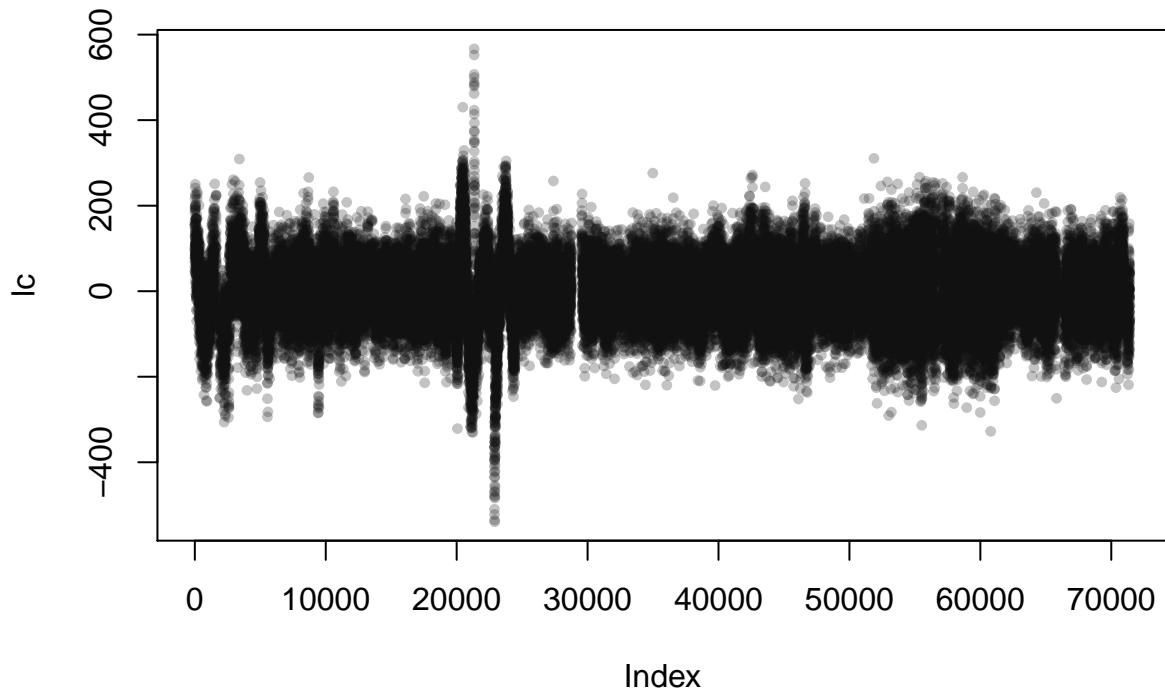
```
## [1] "The first step in the process is to subtract long-term dependencies in light-curve data\n by s

lc = load("000757076_results.Rdat")
lc = results$Lightcurve

summary(lc)

##      Min.   1st Qu.    Median     Mean   3rd Qu.    Max.    NA's
## -539.112 -50.978  -1.478   -1.460   49.054  566.685  17729

#Plot the original, raw light curve
plot(lc, pch=20, cex=0.9, col = "#11111140")
```



```

evaluation_sequence = seq(1, length(lc))

#Fitting a third order polynomial on the raw light curve data
third_order_poly = lm(lc ~ poly(evaluation_sequence, 3), na.action = na.omit)
summary(third_order_poly)

## 
## Call:
## lm(formula = lc ~ poly(evaluation_sequence, 3), na.action = na.omit)
## 
## Residuals:
##      Min       1Q   Median       3Q      Max 
## -538.89  -49.57   -0.06   50.60  566.93 
## 
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)    
## (Intercept)              -1.5212    0.3428  -4.437 9.12e-06 *** 
## poly(evaluation_sequence, 3)1  60.6152   92.9089   0.652  0.51414  
## poly(evaluation_sequence, 3)2 -133.6810   93.6005  -1.428  0.15324  
## poly(evaluation_sequence, 3)3  262.8906   91.9140   2.860  0.00424 ** 
## ---                        
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 
## 
## Residual standard error: 79.31 on 53694 degrees of freedom
##   (17729 observations deleted due to missingness)
## Multiple R-squared:  0.0001891, Adjusted R-squared:  0.0001332 
## F-statistic: 3.385 on 3 and 53694 DF,  p-value: 0.0173

sequence = data.frame(x = seq(1, length(lc)))

predictions = predict(third_order_poly, newdata = sequence)

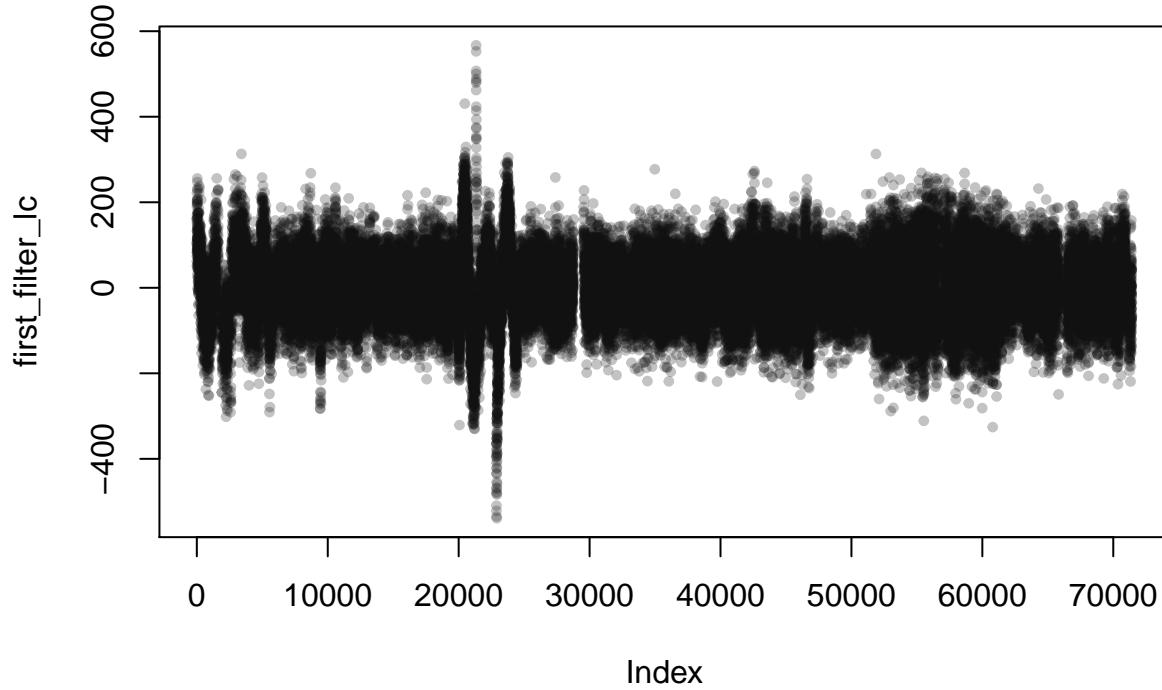
first_filter_lc = lc - predictions

#The first polynomial fitting shows a mean and median that are closer to 0
summary(first_filter_lc)

##      Min.   1st Qu.   Median   Mean   3rd Qu.   Max.   NA's    
## -538.893 -49.572 -0.064   0.000   50.604  566.930  17729 

plot(first_filter_lc, pch=20, cex=0.9, col = "#11111140")

```



```
"Light curve sequences must not contain gaps of more than 0.125 days,
which translates to 6 NA's in a row. In order to be considered for
analysis, a continuous sequence must also be at least 2 days in length"
```

```
## [1] "Light curve sequences must not contain gaps of more than 0.125 days, \n which translates to 6

missing = 0

continuous_sequences = c()

start_of_sequences = c()
new = FALSE

"This for loop keeps track of how many NA's are in a row. If 6 are counted,
then new is set to true and the start of the new sequence is recorded"

## [1] "This for loop keeps track of how many NA's are in a row. If 6 are counted,\n then new is set to

for (i in 1:length(first_filter_lc)){

  if (!is.na(lc[i]) & new == TRUE){
    start_of_sequences = append(start_of_sequences, i)
  }
}
```

```

if (is.na(lc[i])){
  missing = missing + 1
}

else{
  missing = 0
  new = FALSE
}

if (missing == 6){
  new = TRUE
}
}

end_of_sequences = c()
missing = 0

for (i in 1:length(first_filter_lc)){

  if (is.na(lc[i])){
    missing = missing + 1
  }

  else{
    missing = 0
  }

  if (missing == 6){
    end_of_sequences = append(end_of_sequences, i)
  }
}

for (j in 1:length(start_of_sequences)){
  if (j < length(end_of_sequences)-1 & end_of_sequences[j+1]-start_of_sequences[j]>=96){
    continuous_sequences = append(continuous_sequences, j)
  }
}

```

"This step is very similar to the first one, however the fitting and subtraction of the polynomial is done within each continuous sequence of the light curve"

```

## [1] "This step is very similar to the first one, however the fitting and subtraction of the polynomial

#So as to not overwrite the first step's progress
second_filter_lc = first_filter_lc

for (i in 1:length(continuous_sequences)){

  continuous_segment  = second_filter_lc[start_of_sequences[i]:end_of_sequences[i+1]]

```

```

eval_sequence = seq(1, length(continuous_segment))

third_order_poly = lm(continuous_segment ~ poly(eval_sequence, 3, raw = TRUE))

sequence = data.frame(x = seq(1, length(continuous_segment)))
predictions = predict(third_order_poly, newdata = sequence)

second_filter_lc[start_of_sequences[i]:end_of_sequences[i+1]] = lc[start_of_sequences[i]:end_of_sequences[i+1]]
}

## Warning in predict.lm(third_order_poly, newdata = sequence): prediction from a
## rank-deficient fit may be misleading

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## rank-deficient fit may be misleading

summary(second_filter_lc)

##      Min.    1st Qu.     Median      Mean    3rd Qu.      Max.    NA's
## -325.759   -46.238    -1.384   -0.935    44.998   507.022    17729

#It can be seen visually that the values are within a narrower range
plot(second_filter_lc, pch=20, cex=0.9, col = "#11111140")

```

