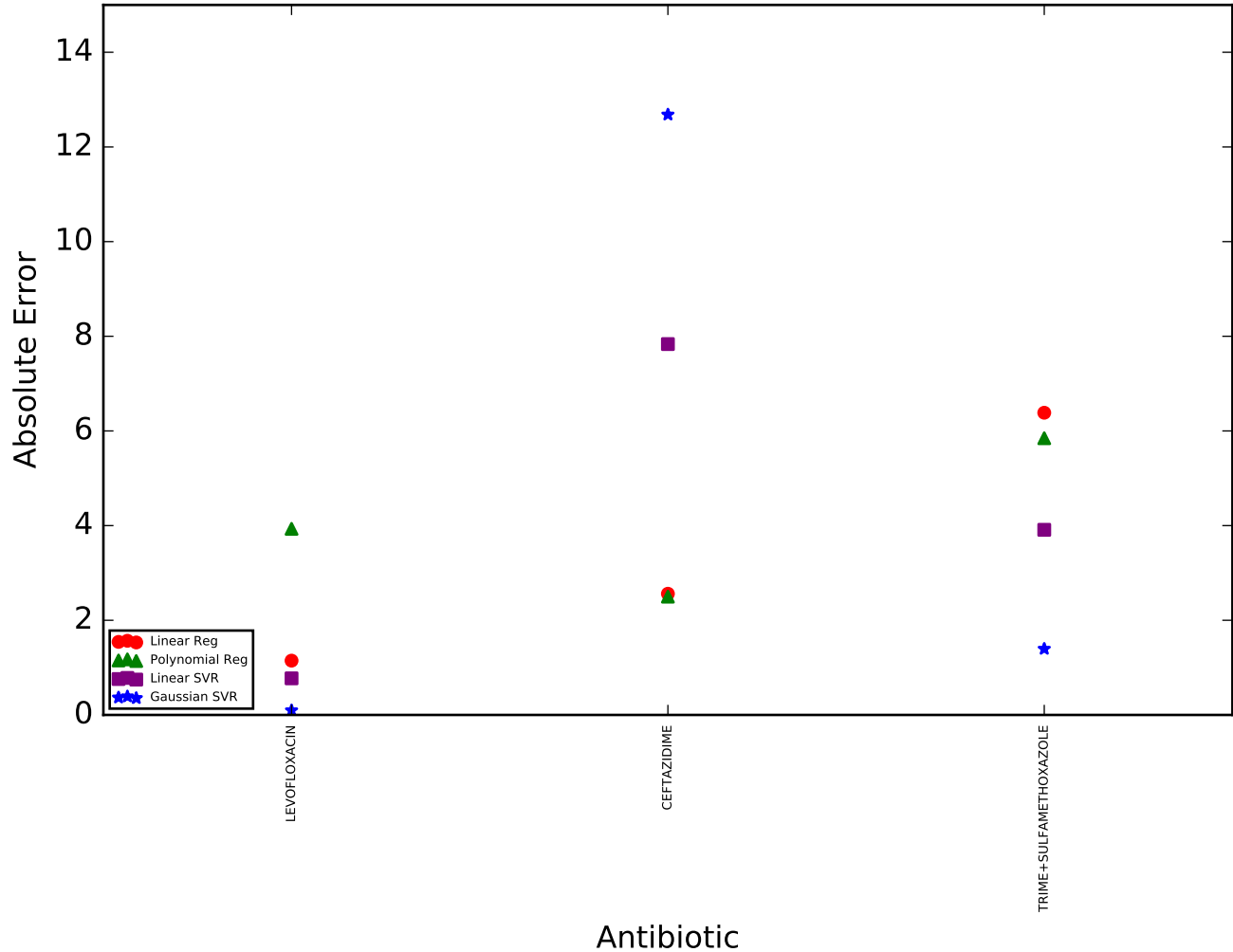


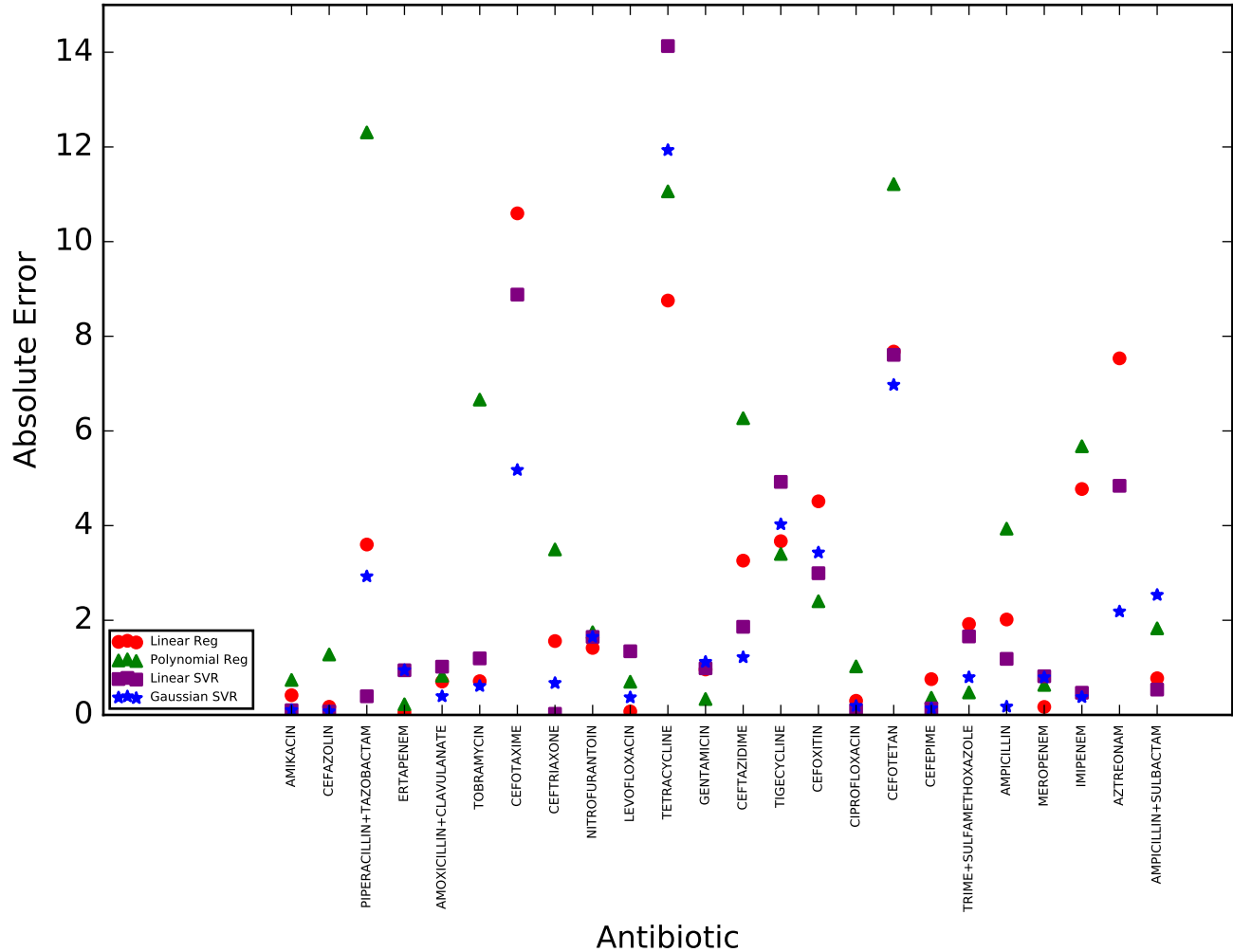
# Stenotrophomonas maltophilia

using data from 2006-2014 to predict 2015

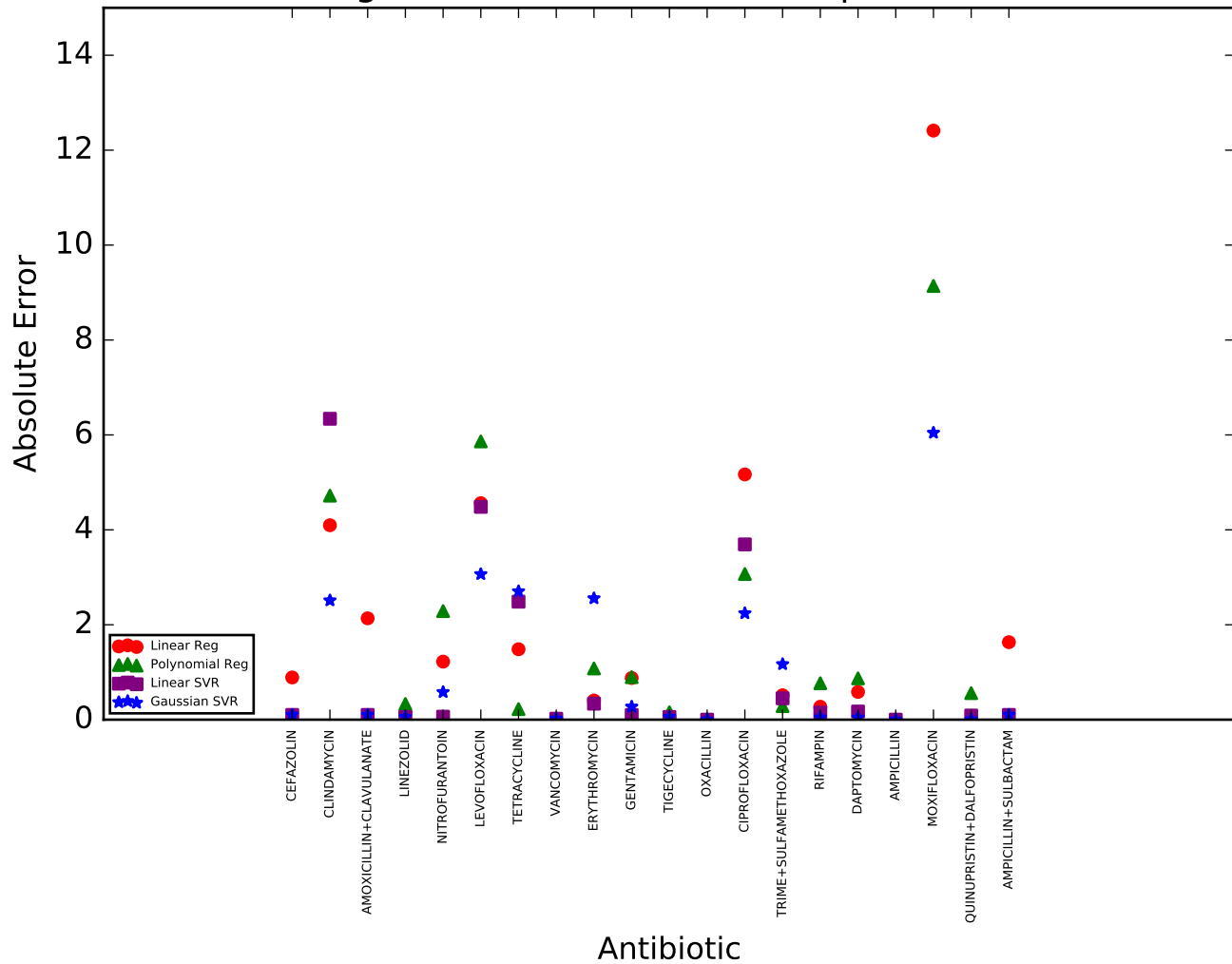


# Serratia marcescens

## using data from 2006-2014 to predict 2015

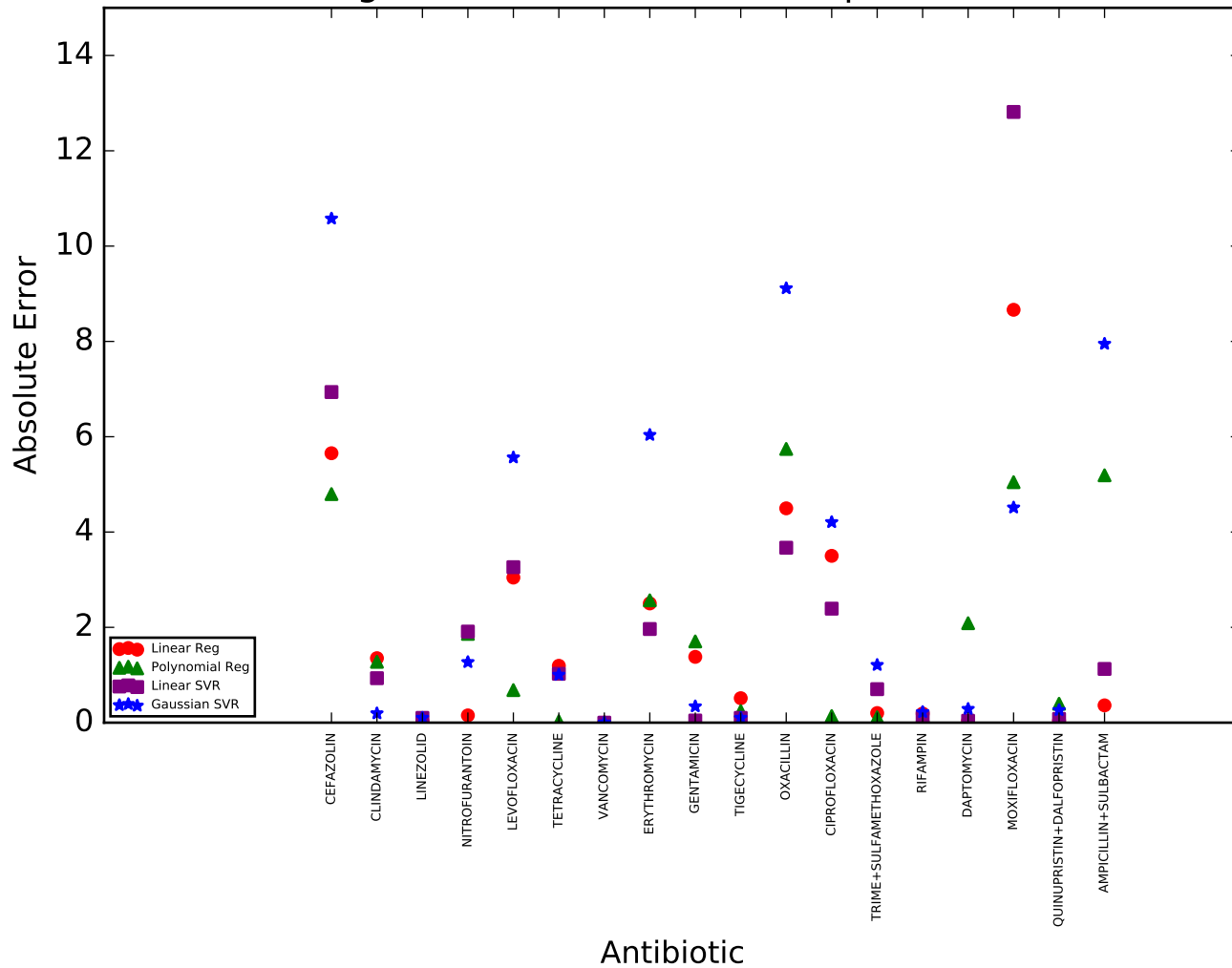


using data from 2006-2014 to predict 2015



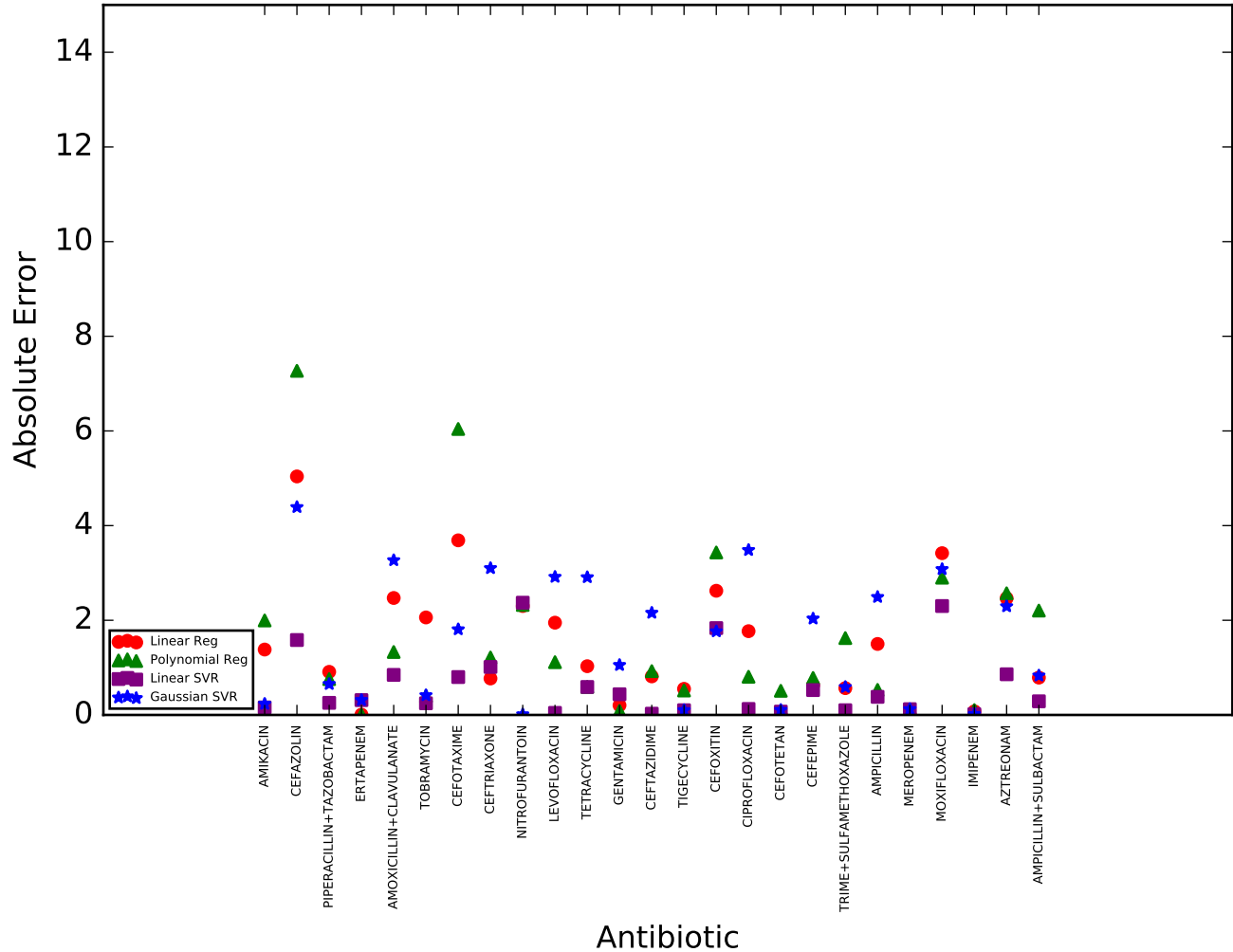
# Staphylococcus aureus

## using data from 2006-2014 to predict 2015



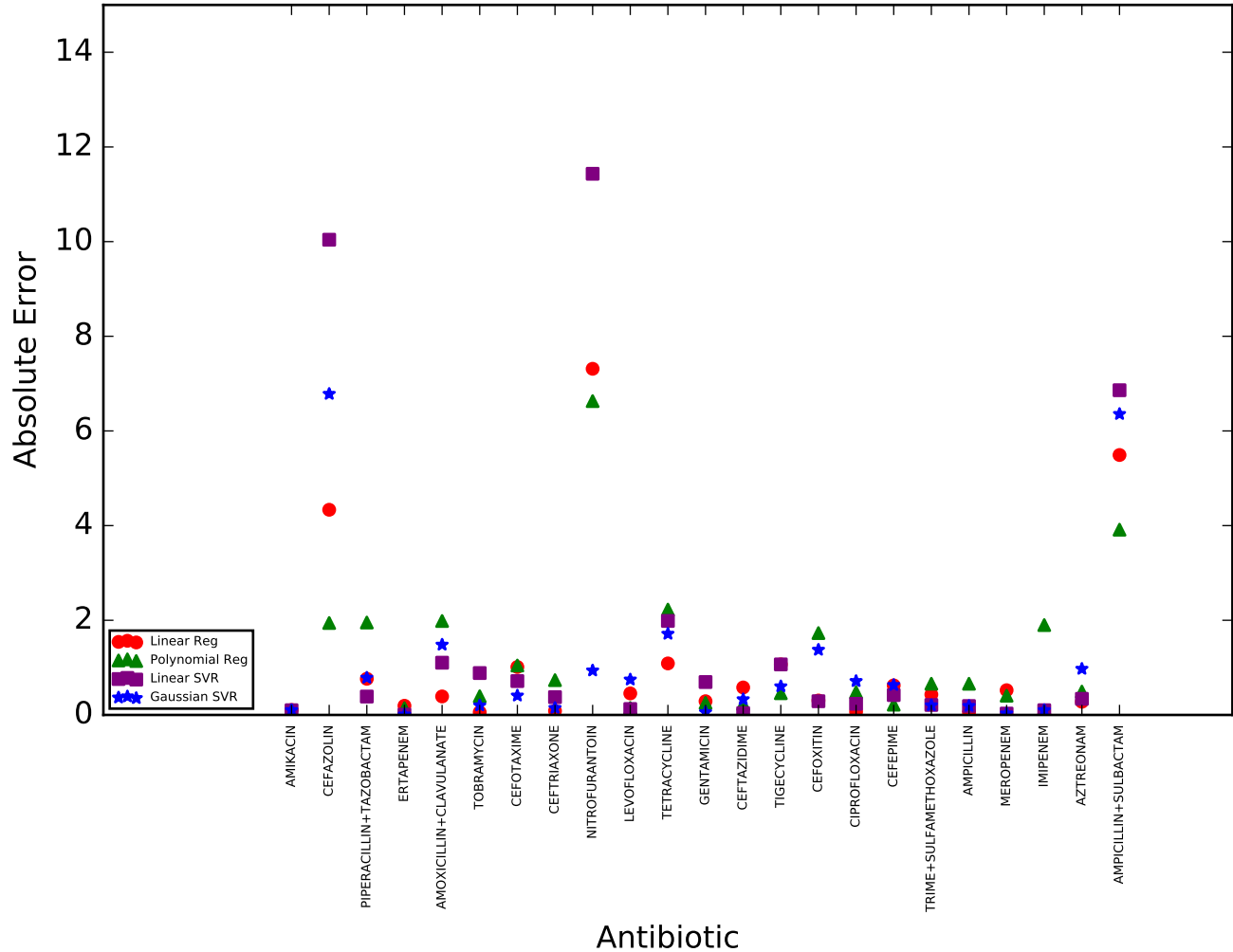
# E. coli

## using data from 2006-2014 to predict 2015



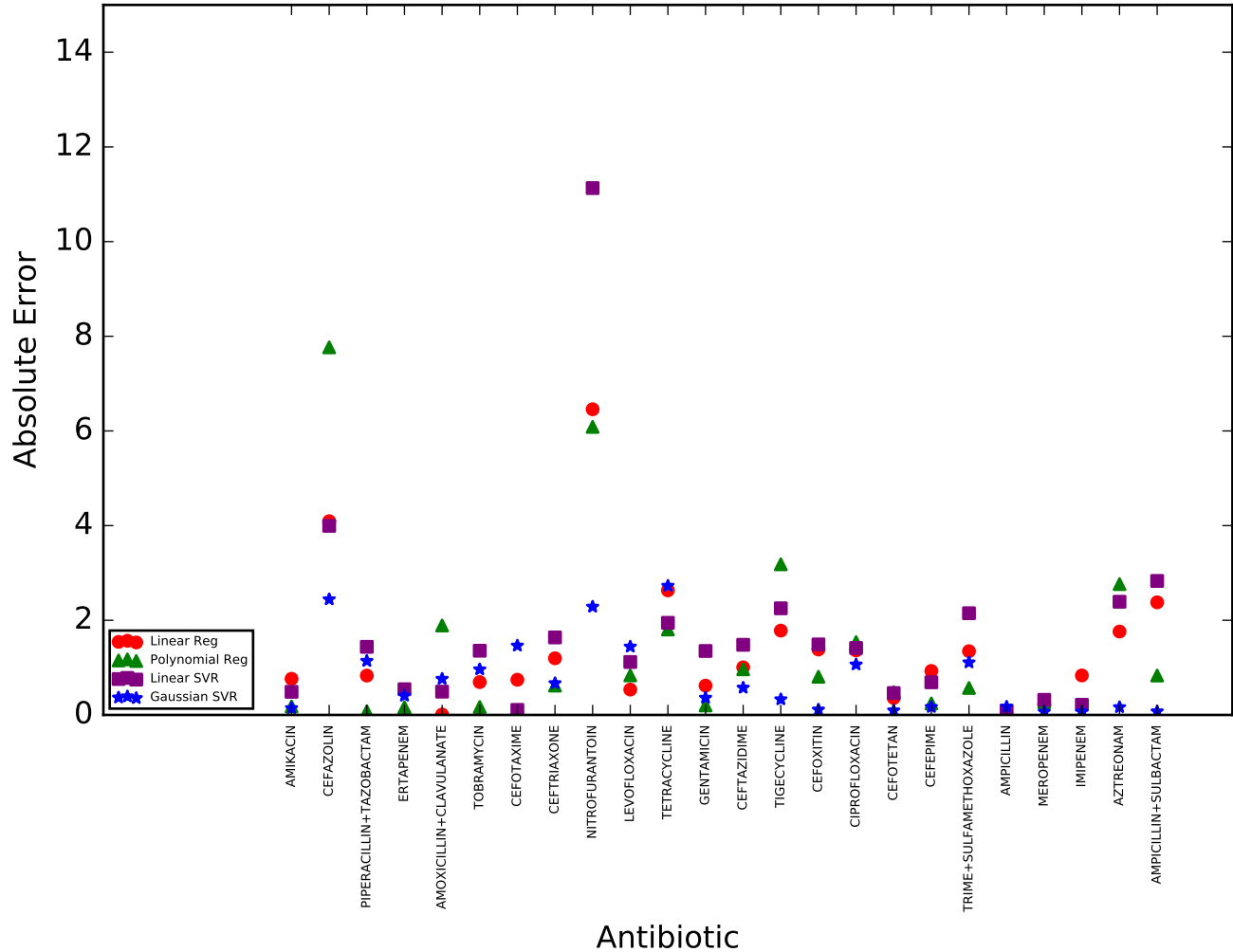
# Klebsiella oxytoca

## using data from 2006-2014 to predict 2015

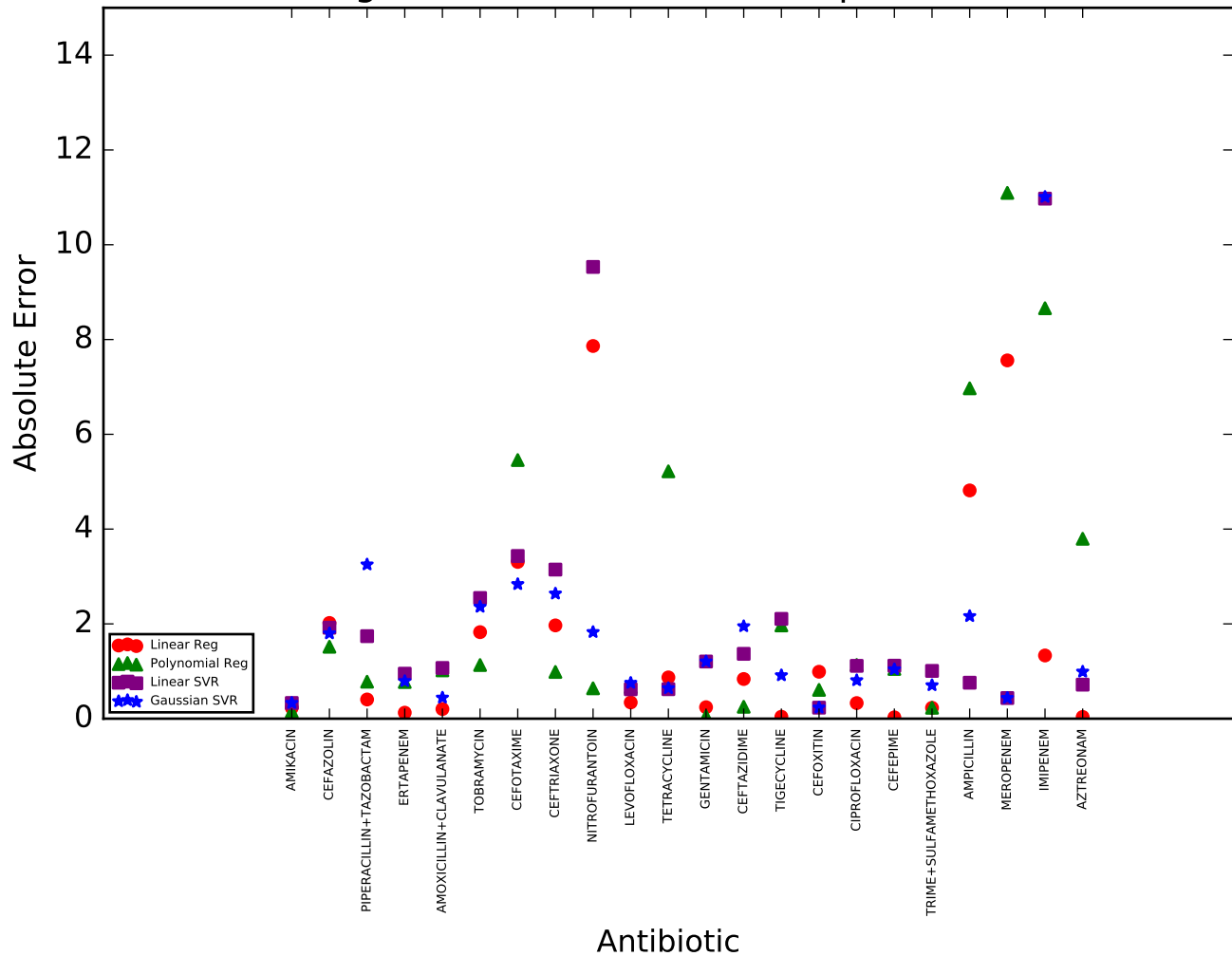


# Klebsiella pneumoniae

## using data from 2006-2014 to predict 2015



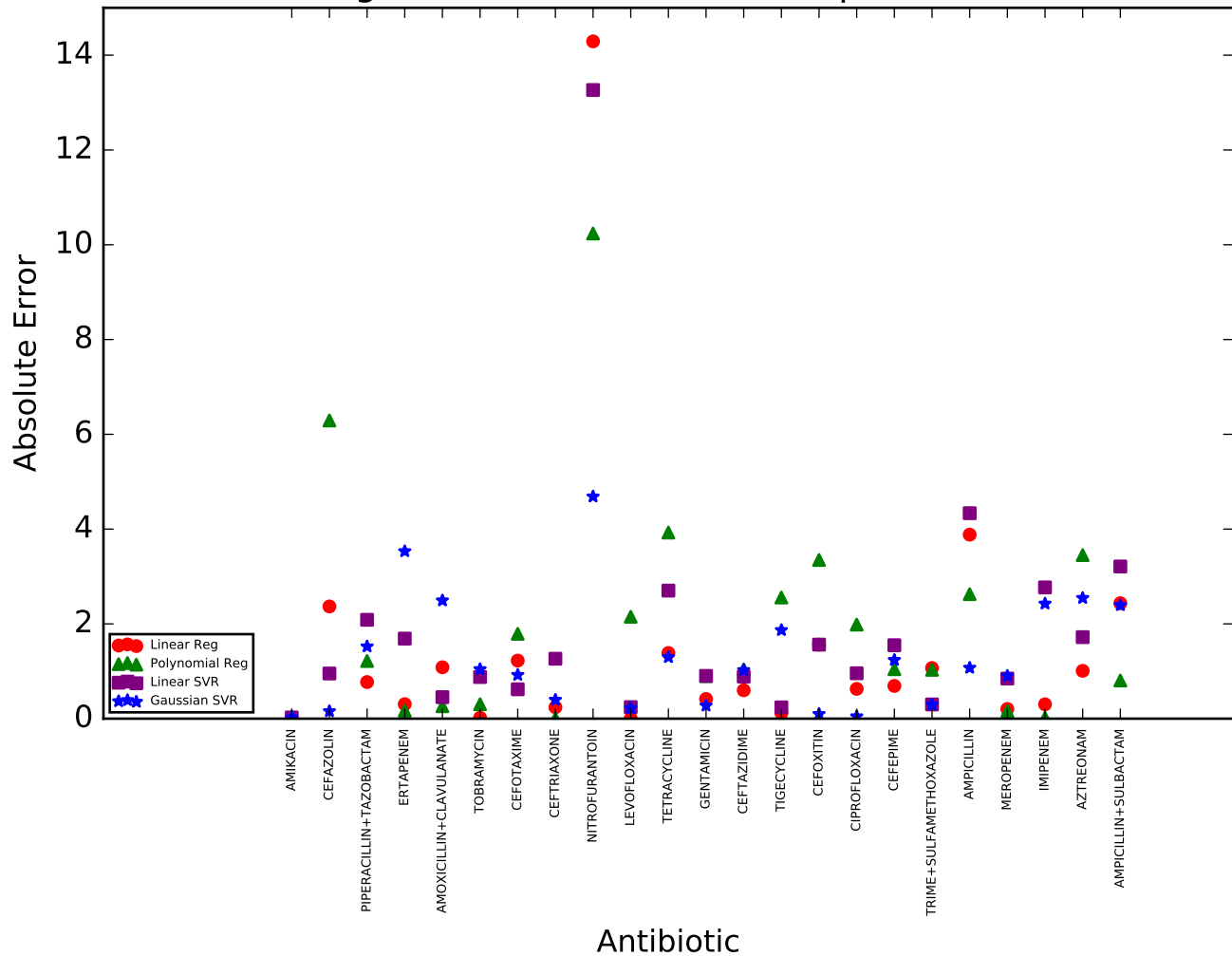
# Enterobacter aerogenes using data from 2006-2014 to predict 2015



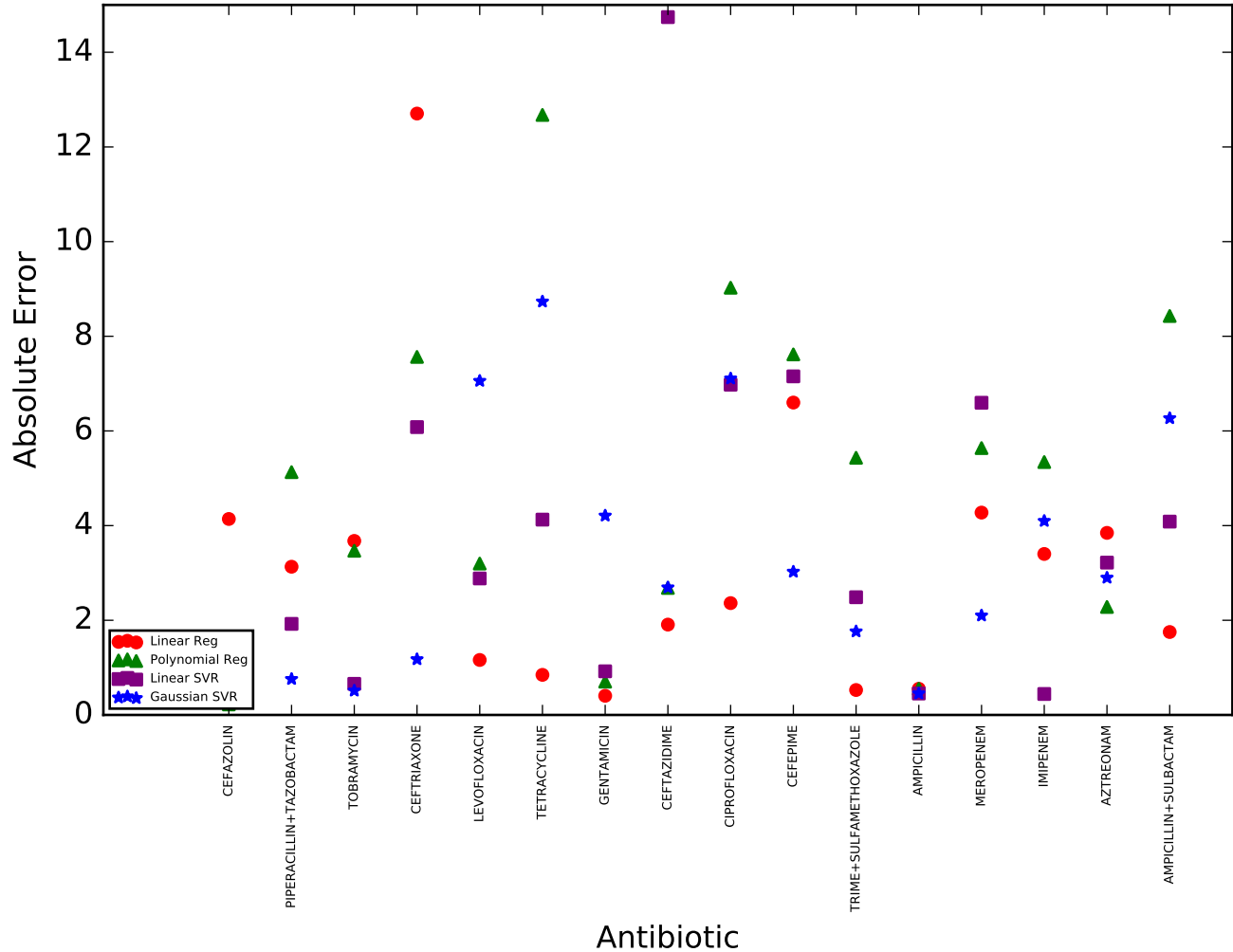


# Enterobacter cloacae

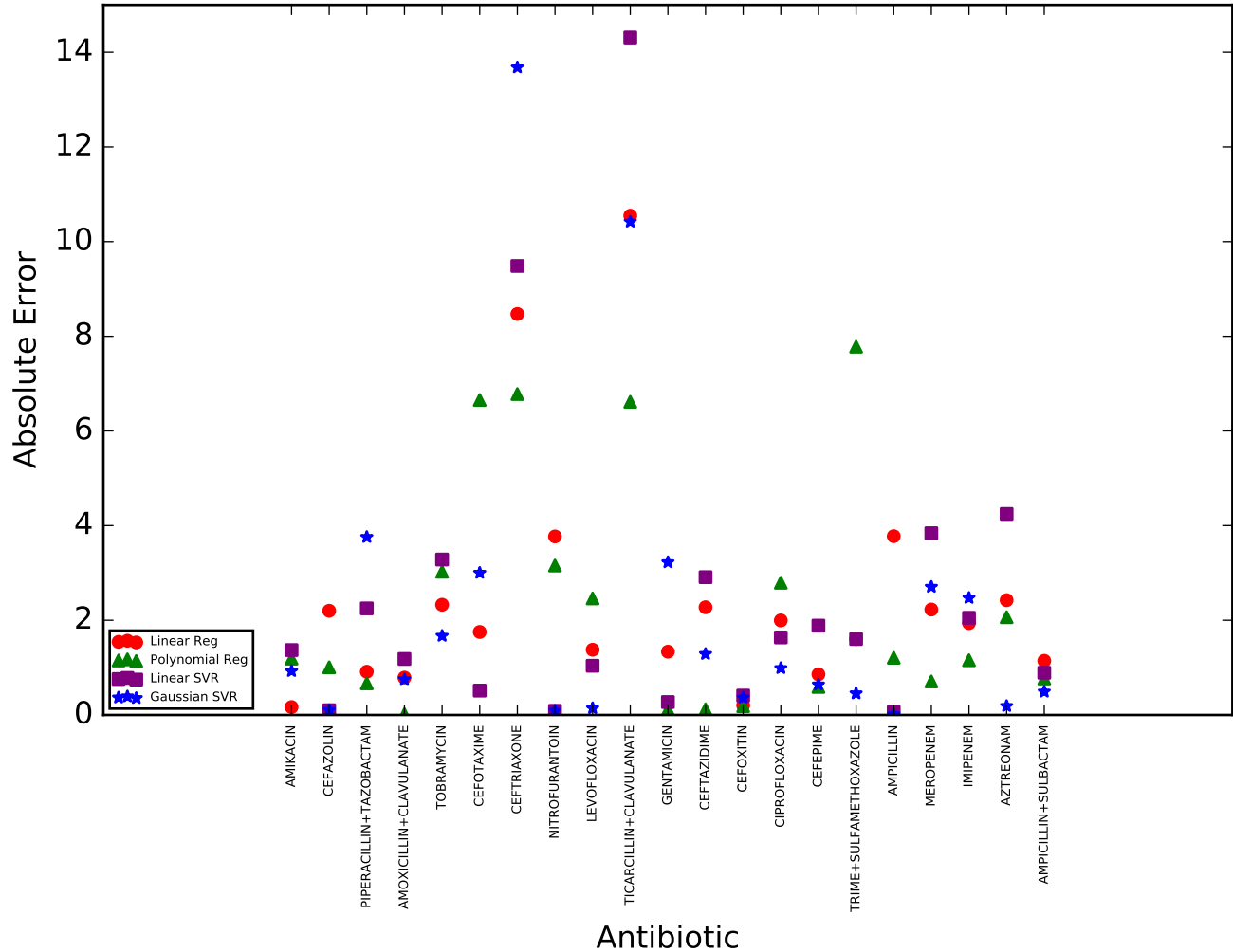
## using data from 2006-2014 to predict 2015



# Acinetobacter baumannii using data from 2006-2014 to predict 2015

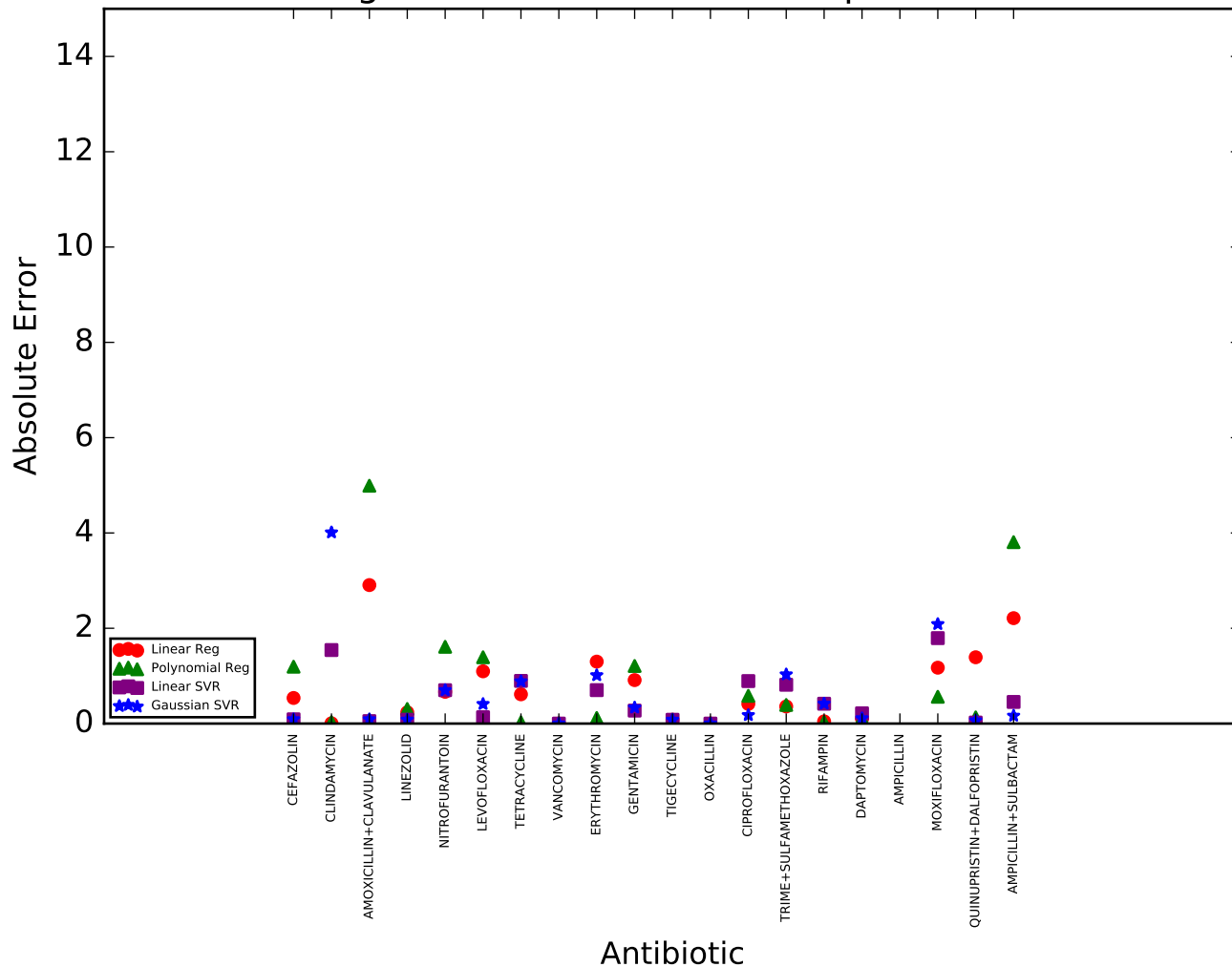


# *Pseudomonas aeruginosa* using data from 2006-2014 to predict 2015



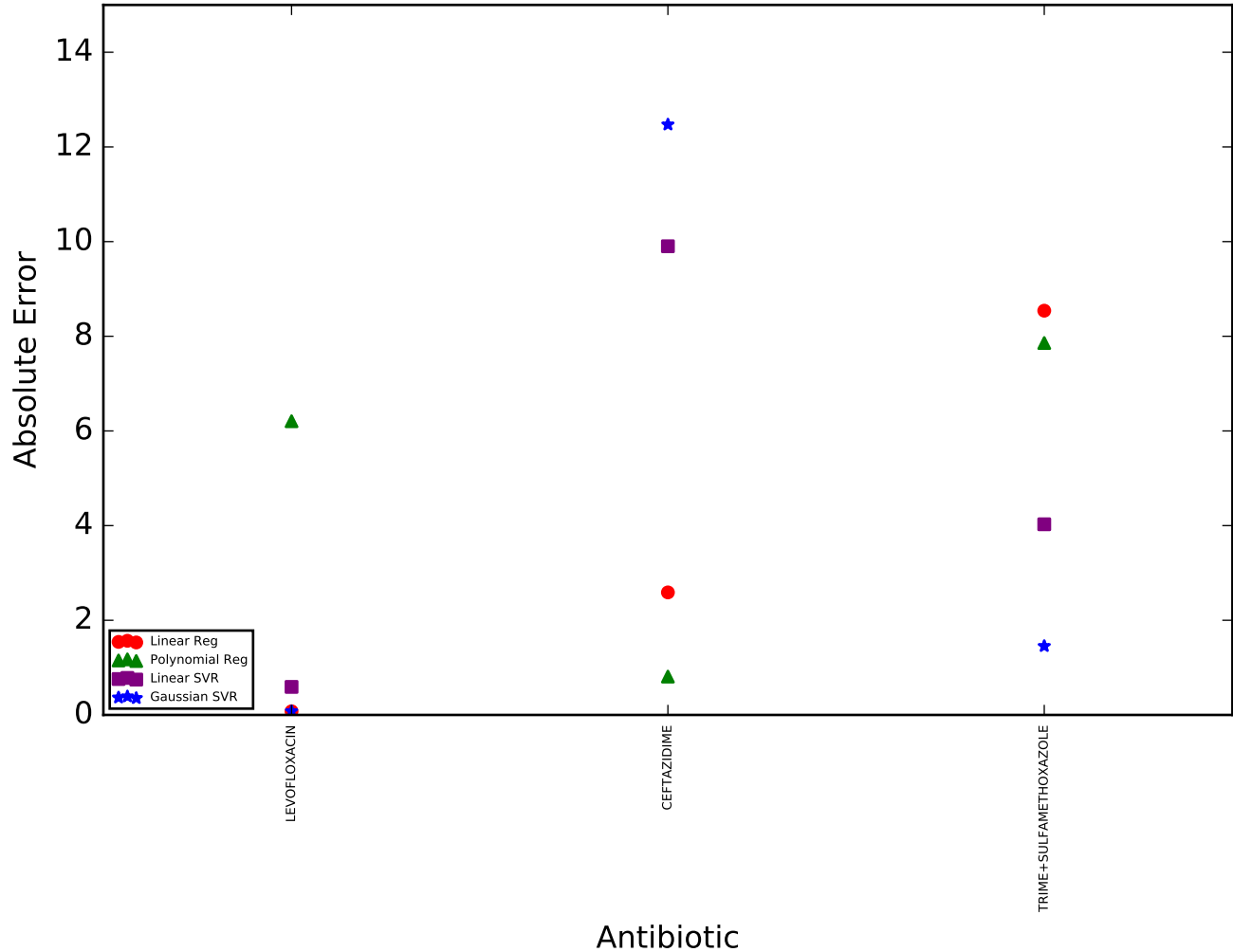
# MSSA

## using data from 2006-2014 to predict 2015



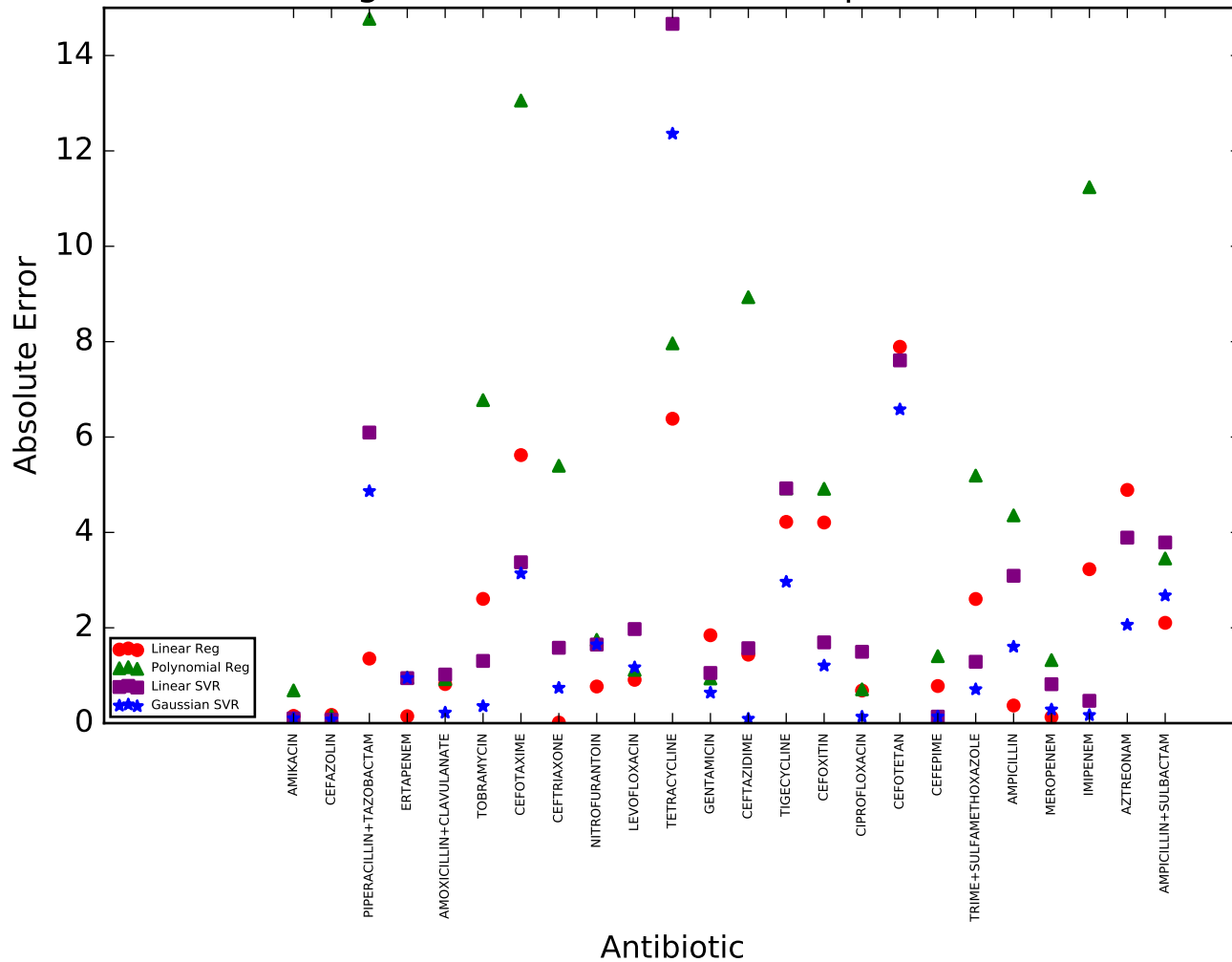
# Stenotrophomonas maltophilia

using data from 2005-2013 to predict 2015



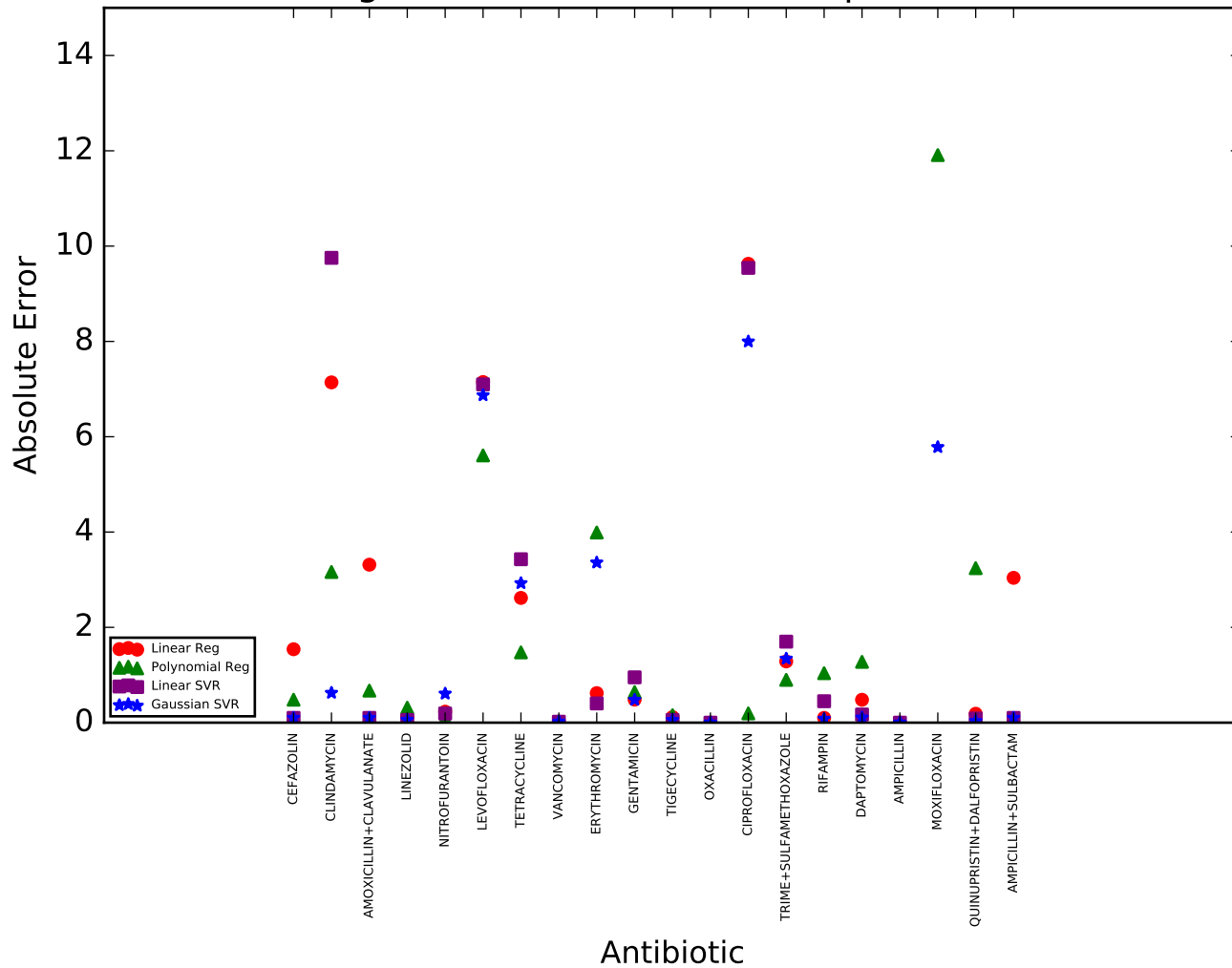
# Serratia marcescens

## using data from 2005-2013 to predict 2015



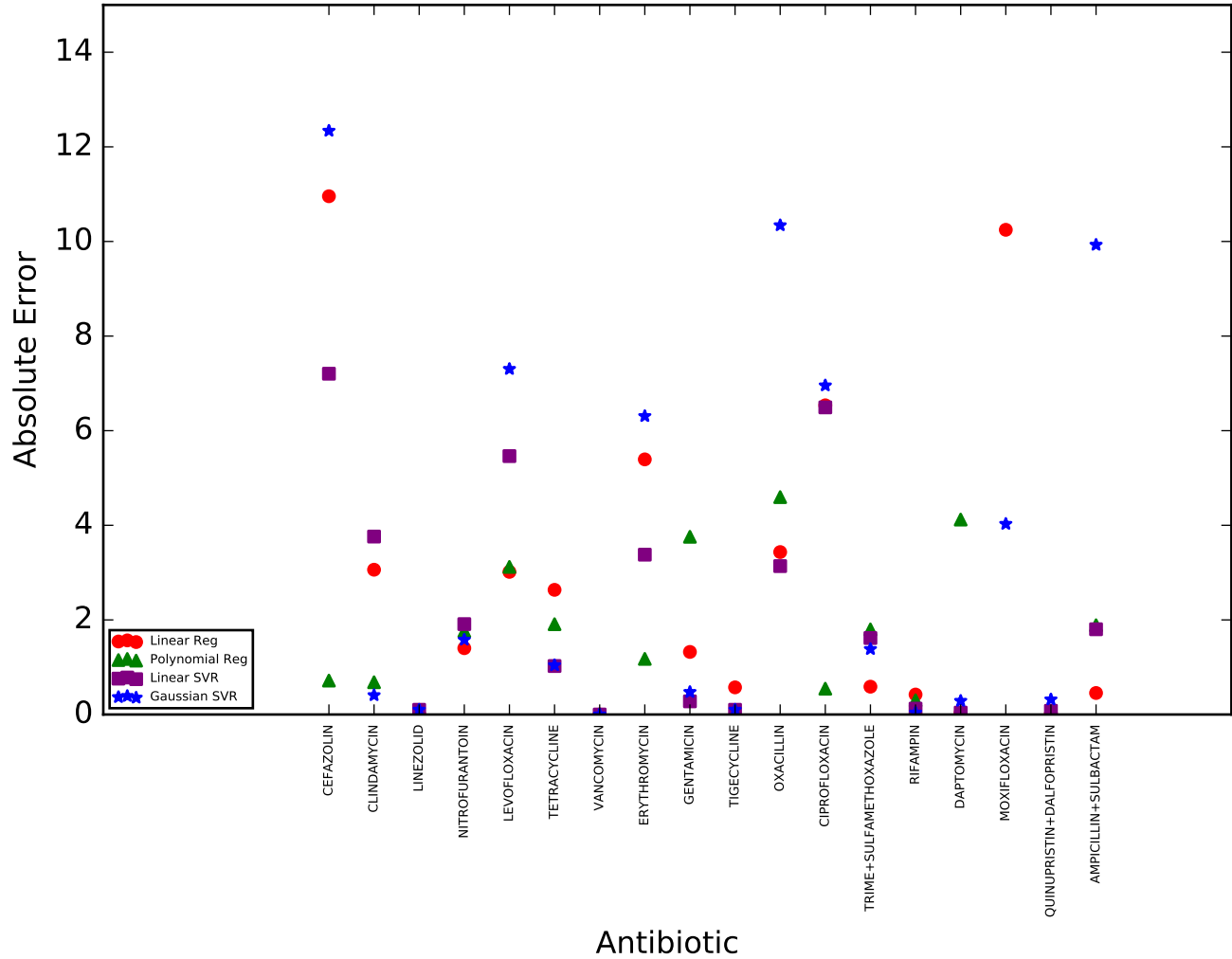
# MRSA

using data from 2005-2013 to predict 2015



# Staphylococcus aureus

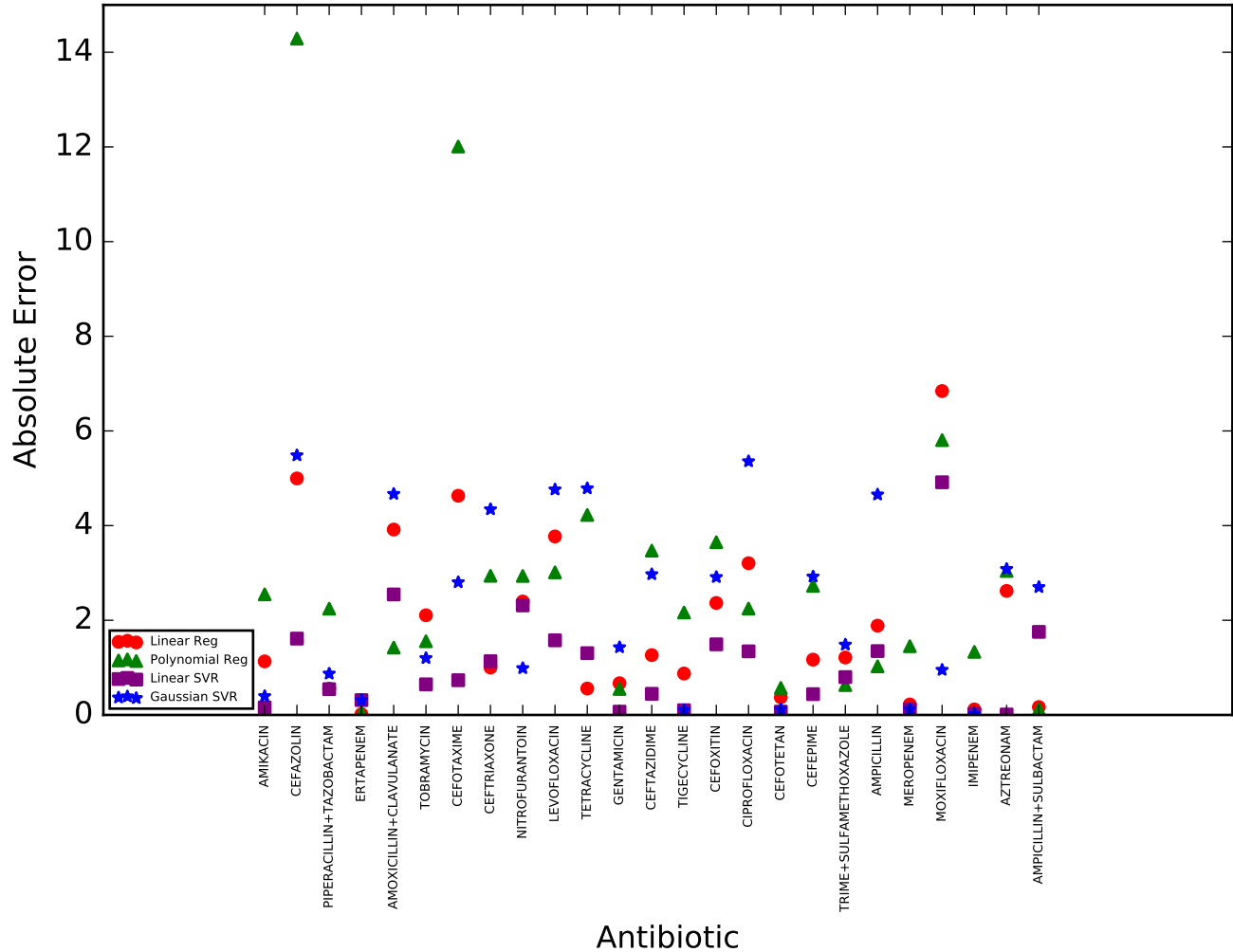
## using data from 2005-2013 to predict 2015





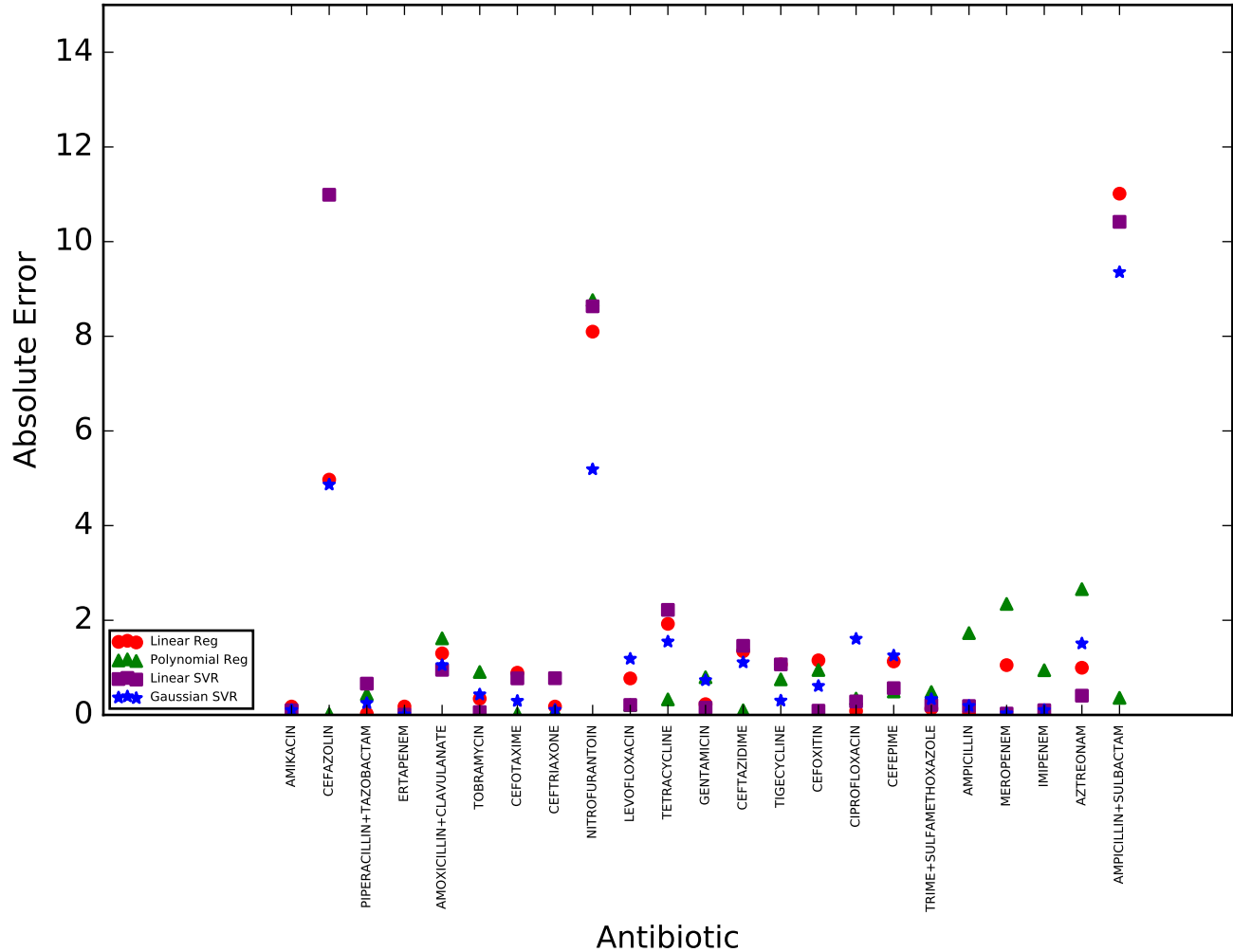
# E. coli

## using data from 2005-2013 to predict 2015



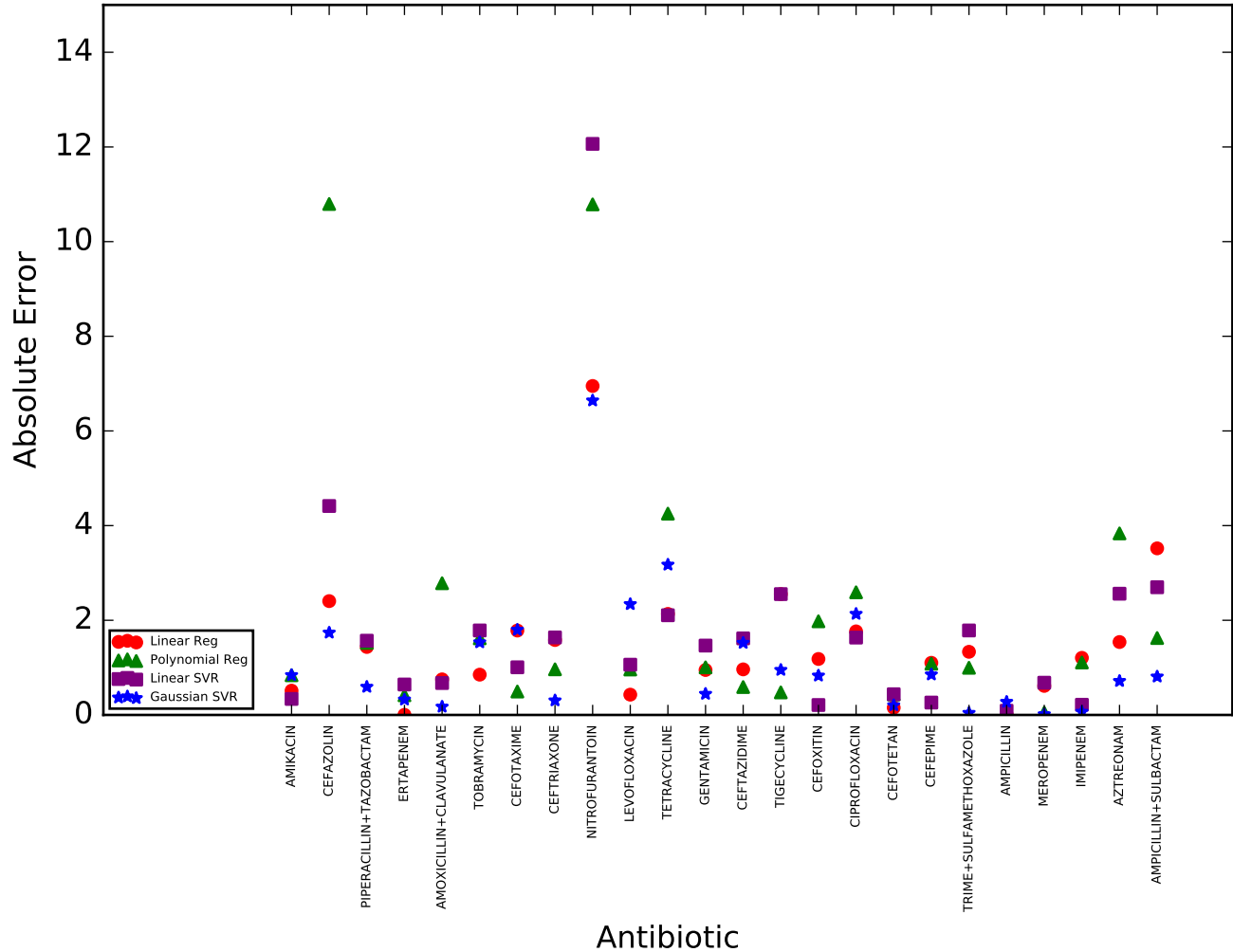
# Klebsiella oxytoca

## using data from 2005-2013 to predict 2015



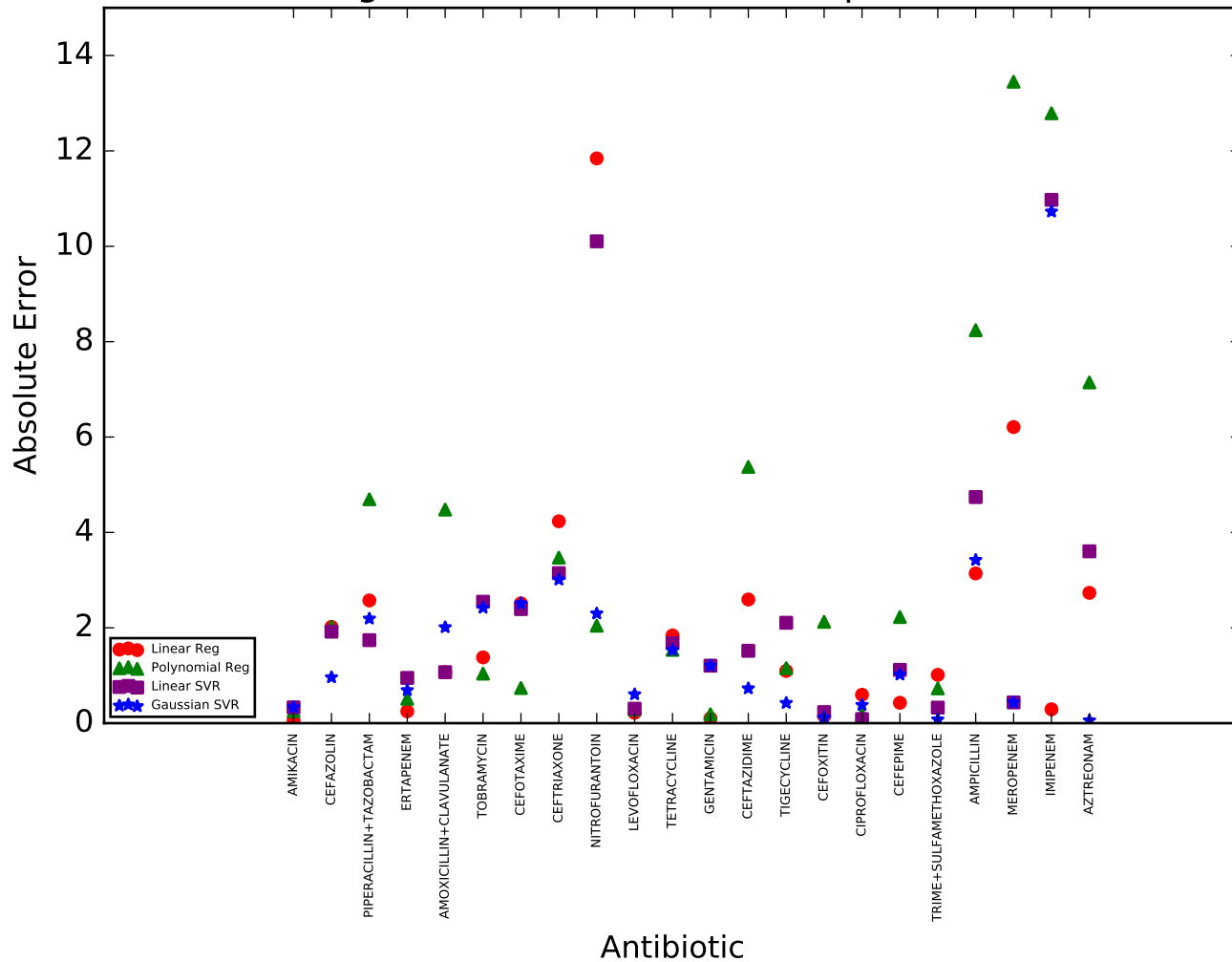
# Klebsiella pneumoniae

## using data from 2005-2013 to predict 2015



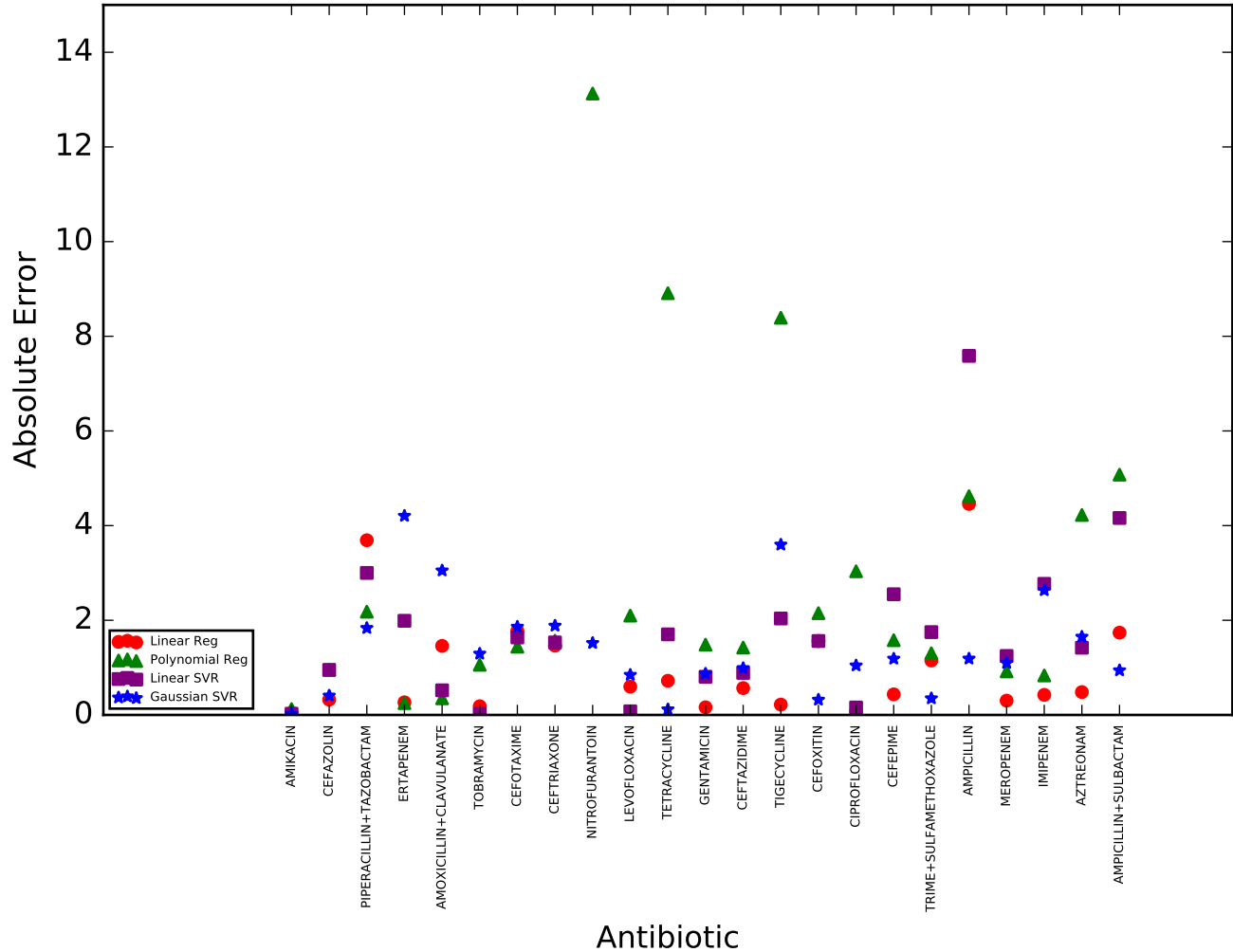
# Enterobacter aerogenes

## using data from 2005-2013 to predict 2015

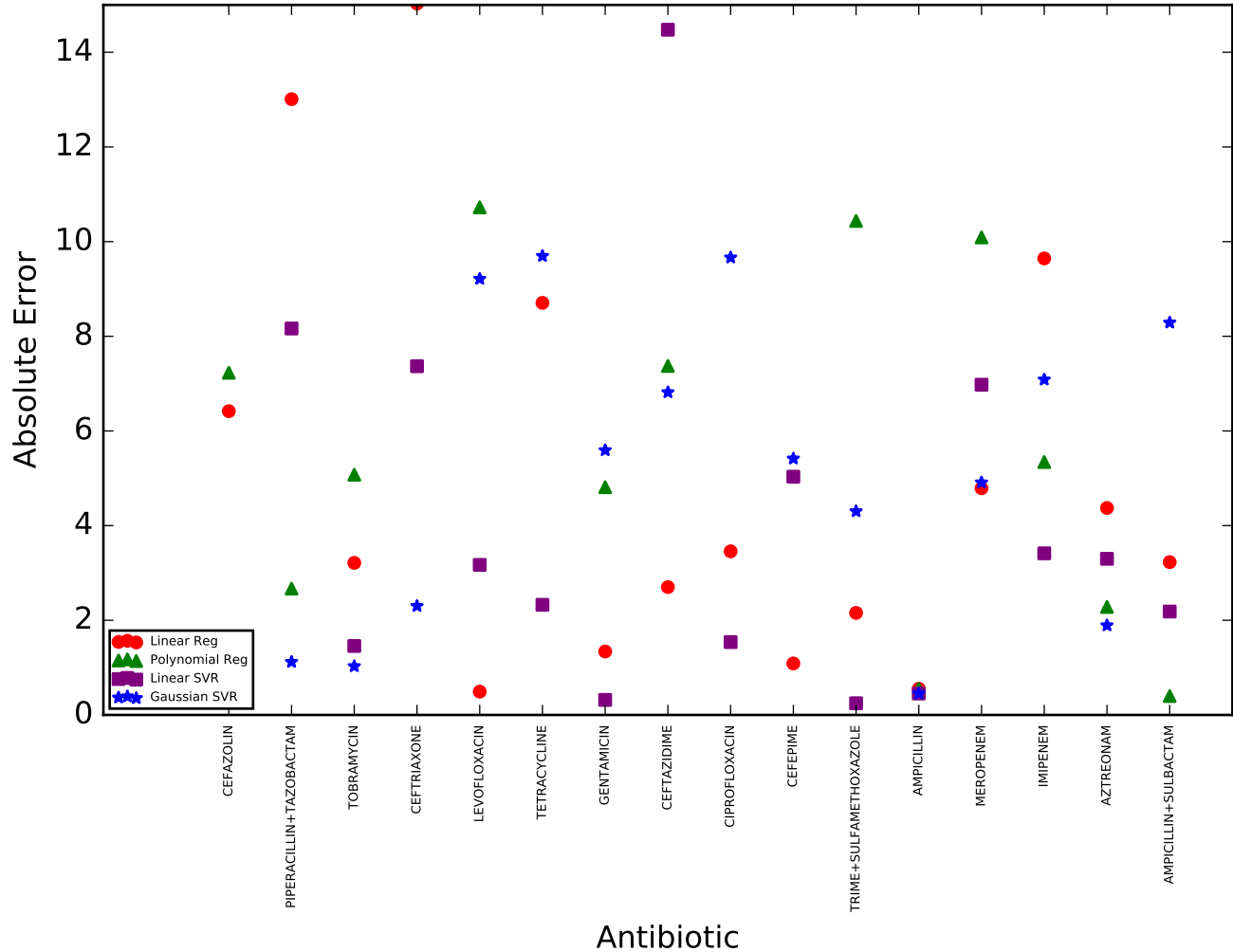


# Enterobacter cloacae

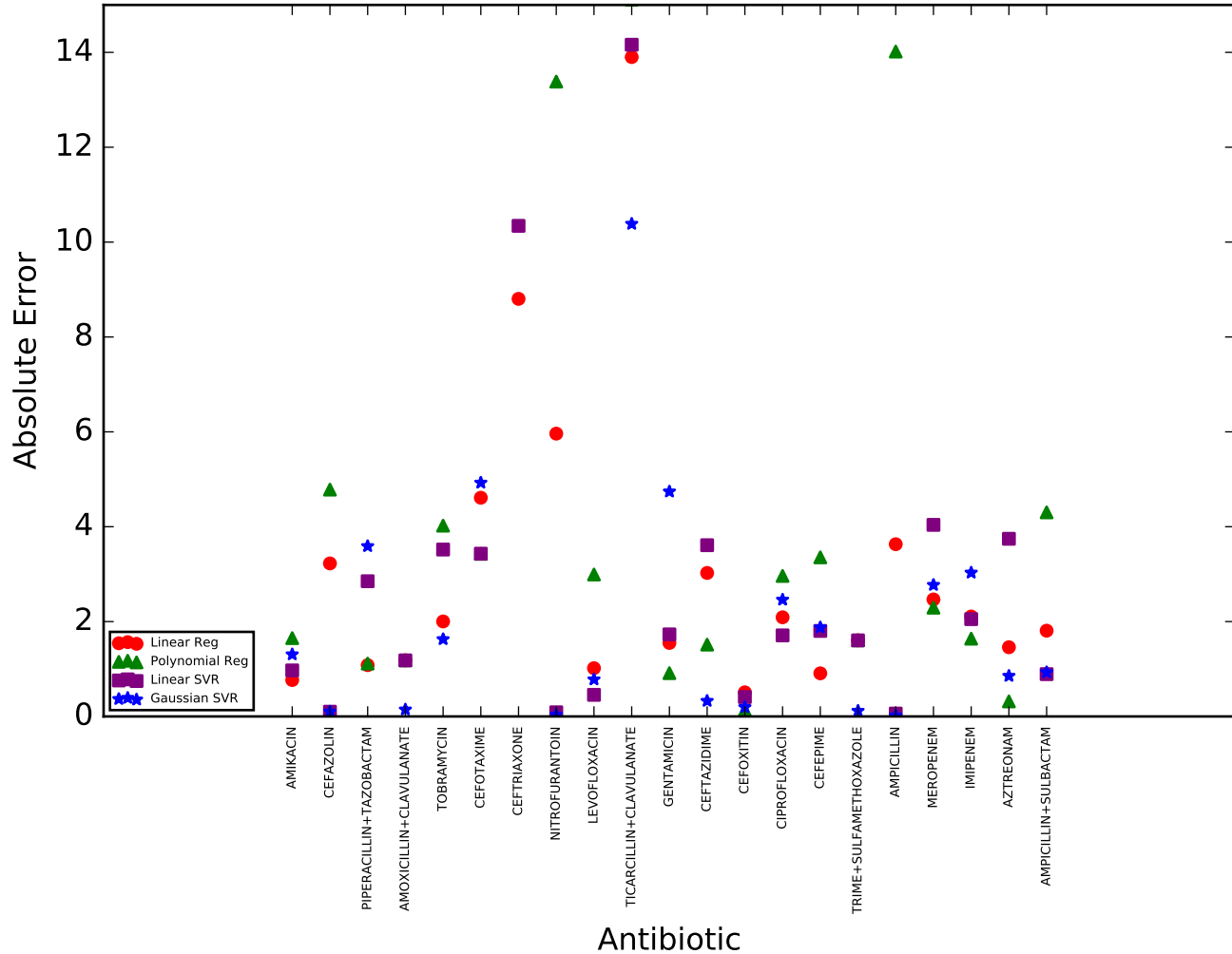
## using data from 2005-2013 to predict 2015



# Acinetobacter baumannii using data from 2005-2013 to predict 2015

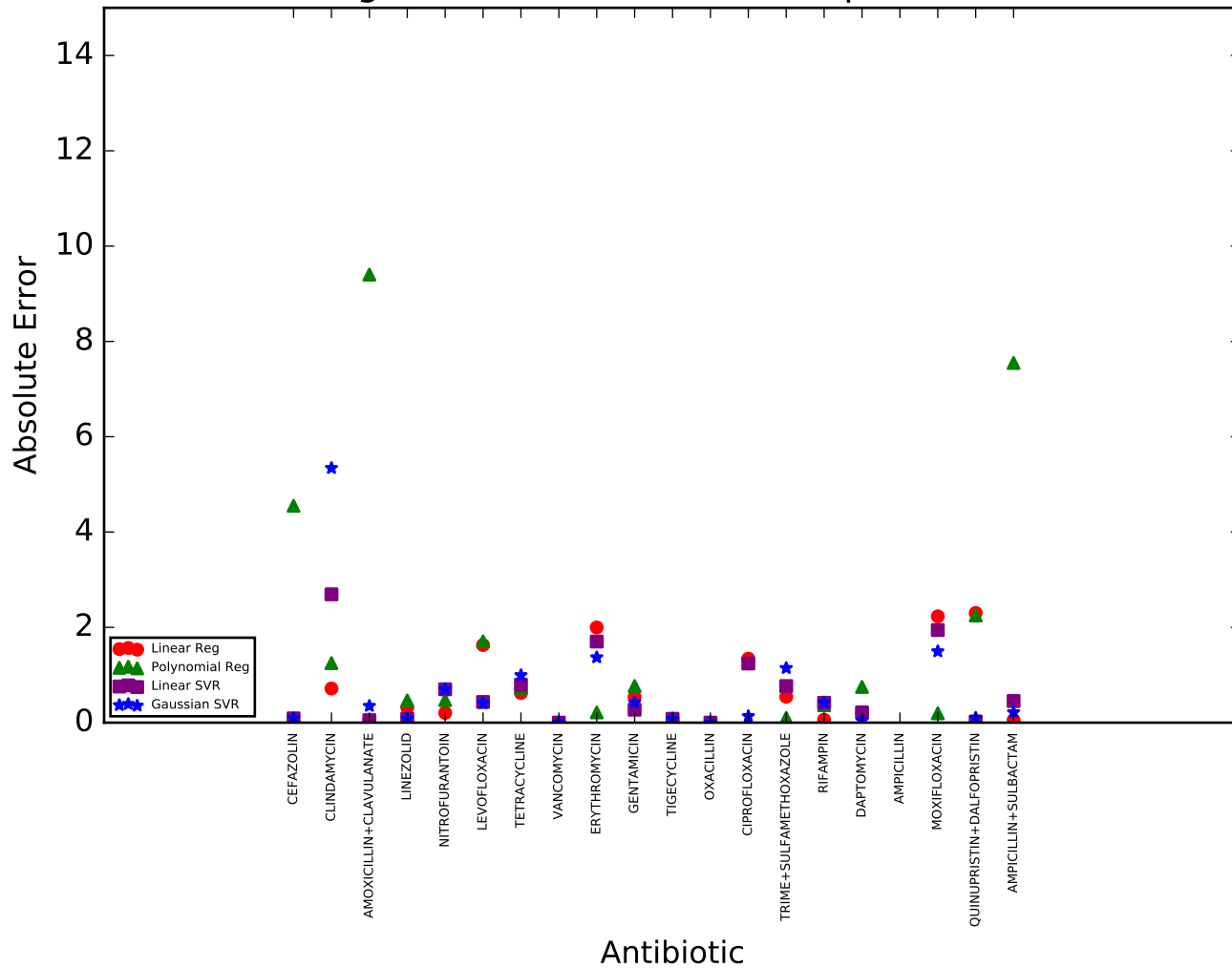


# *Pseudomonas aeruginosa* using data from 2005-2013 to predict 2015



# MSSA

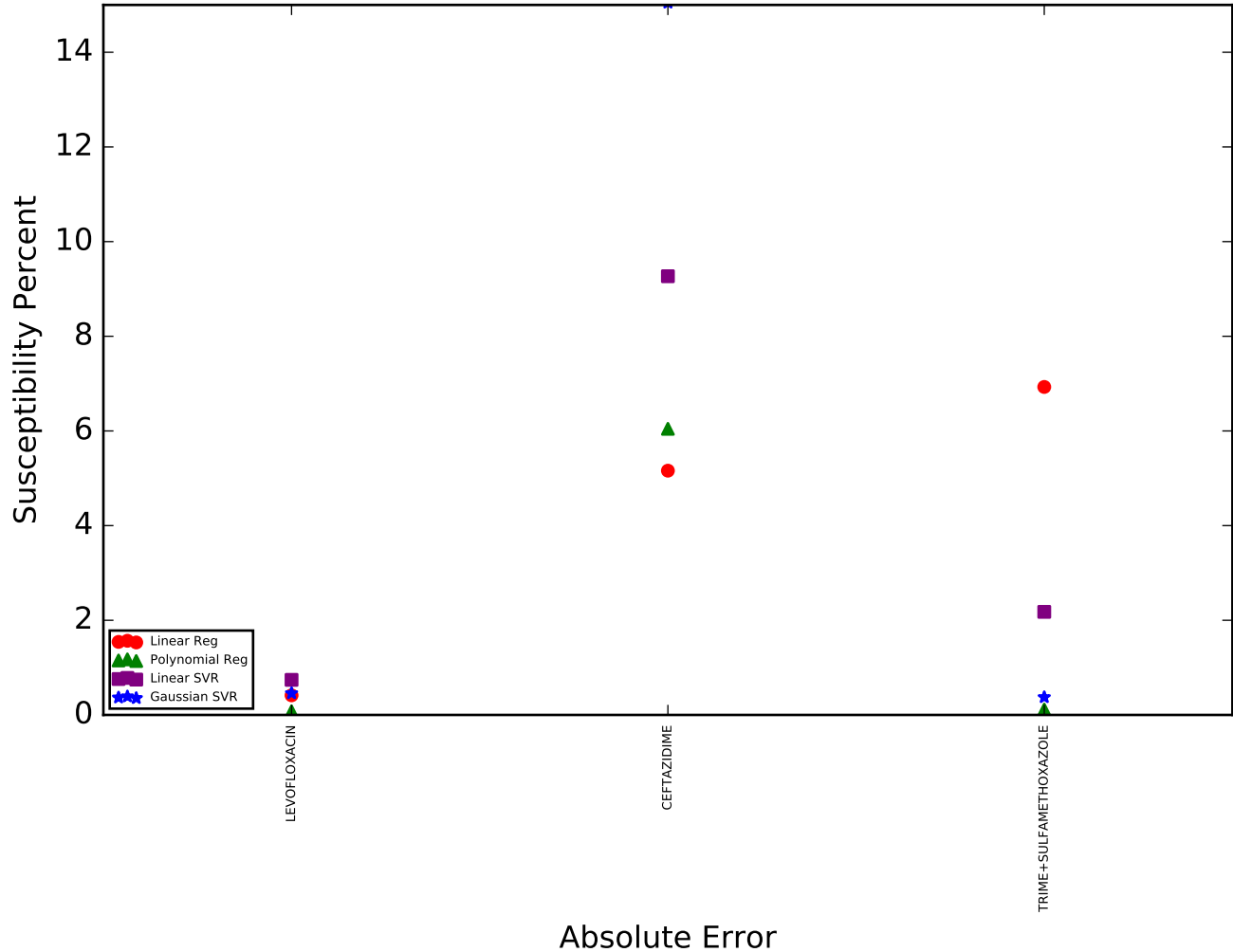
## using data from 2005-2013 to predict 2015





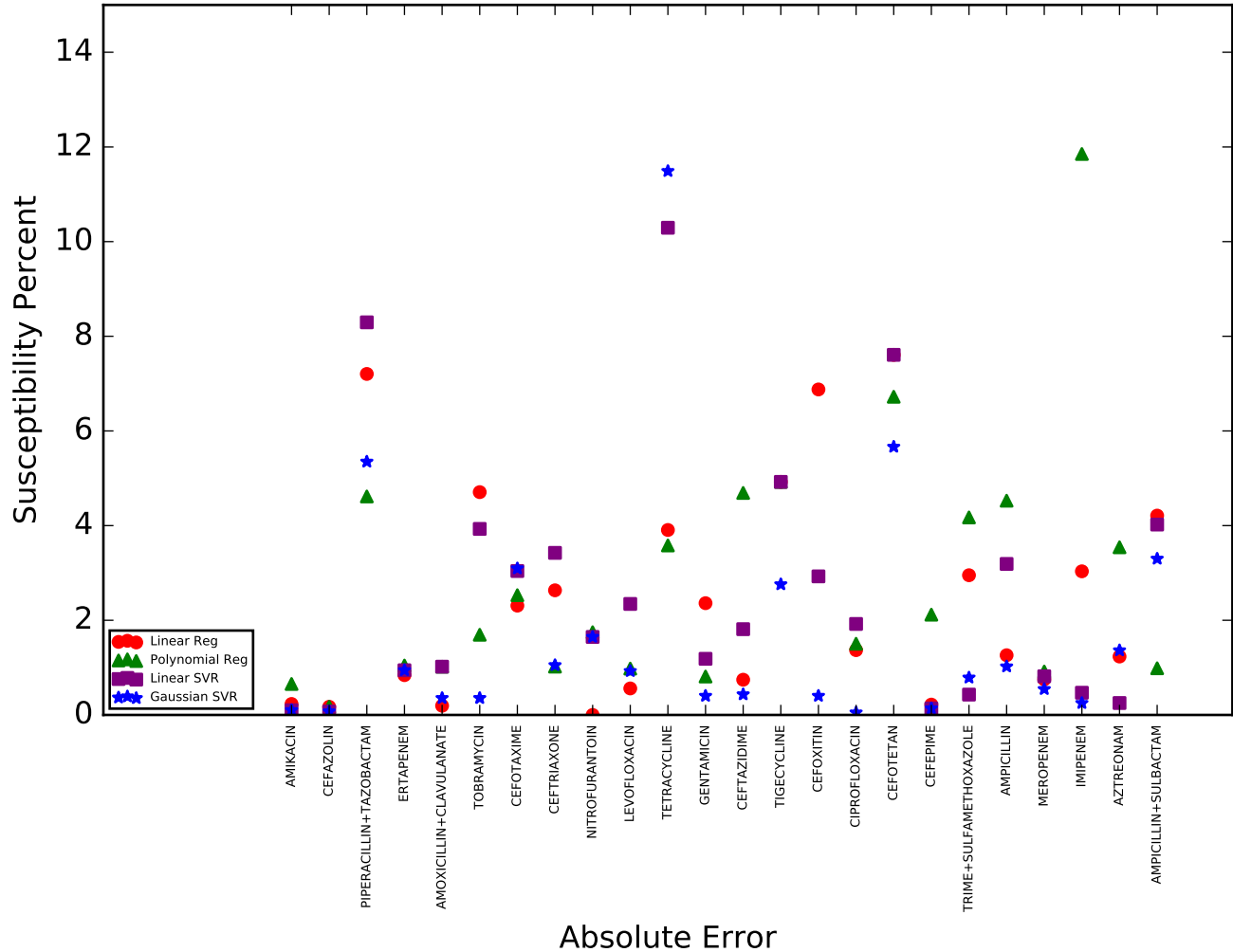
# Stenotrophomonas maltophilia

using data from 2004-2012 to predict 2015



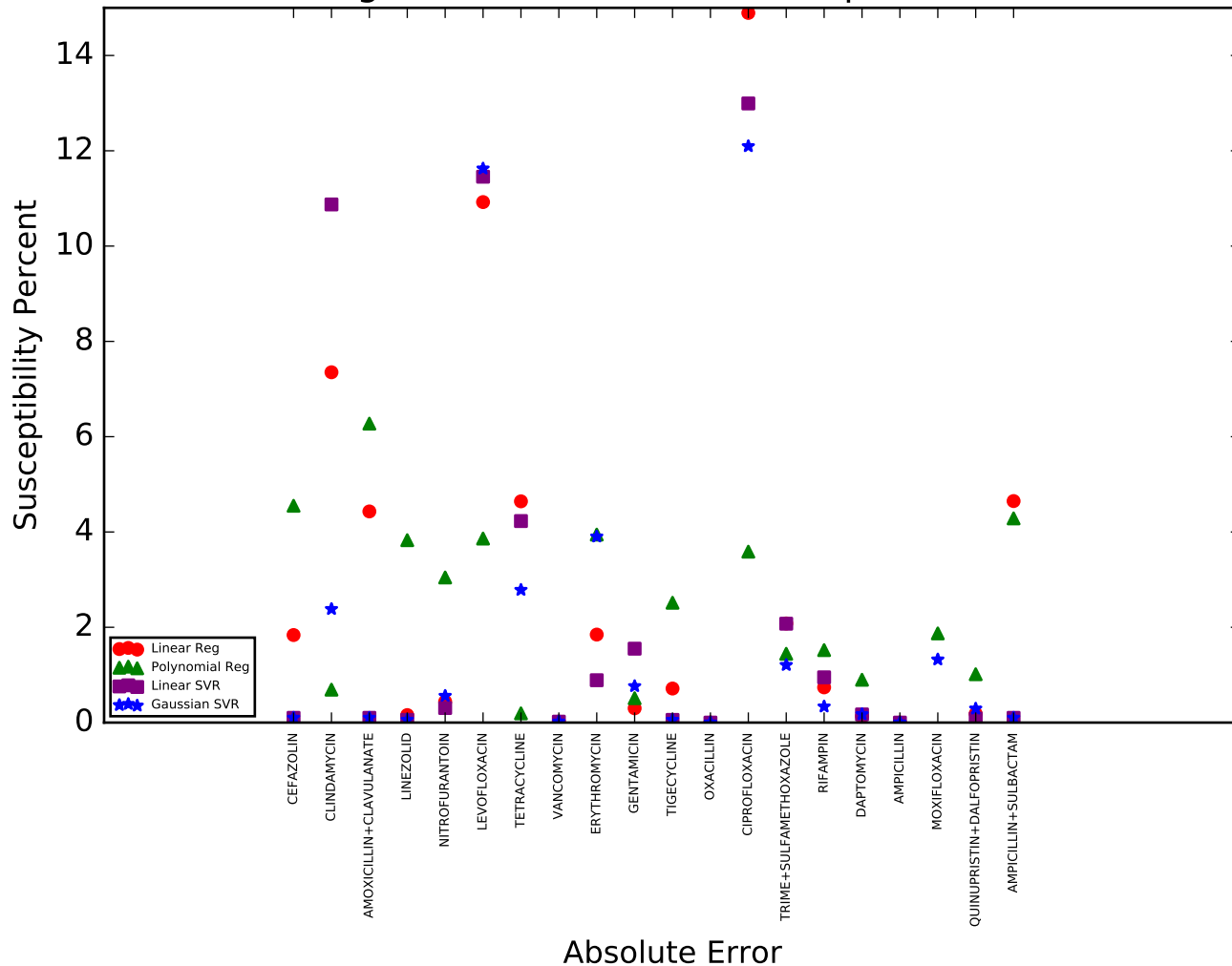
# Serratia marcescens

## using data from 2004-2012 to predict 2015

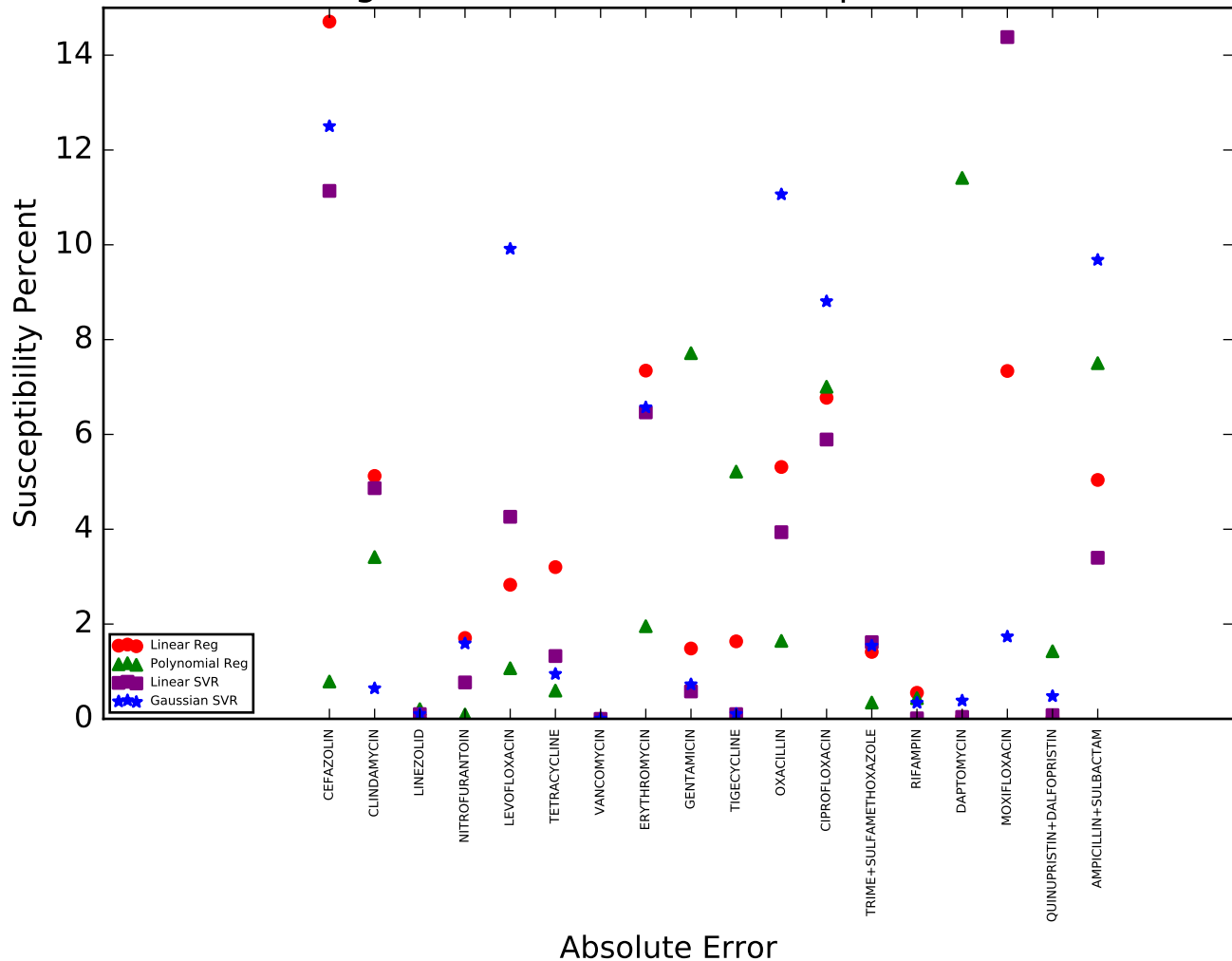


# MRSA

using data from 2004-2012 to predict 2015

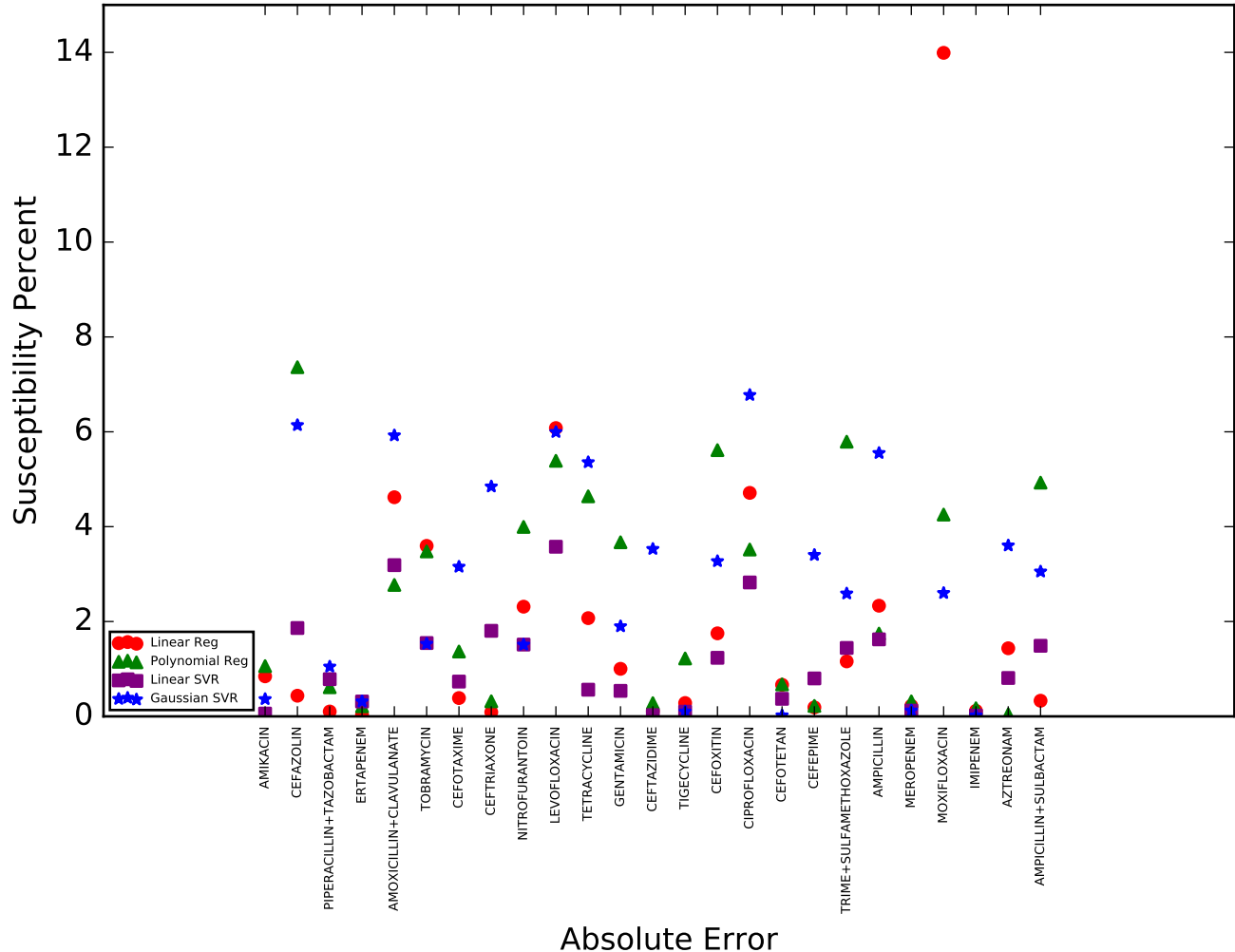


# Staphylococcus aureus using data from 2004-2012 to predict 2015



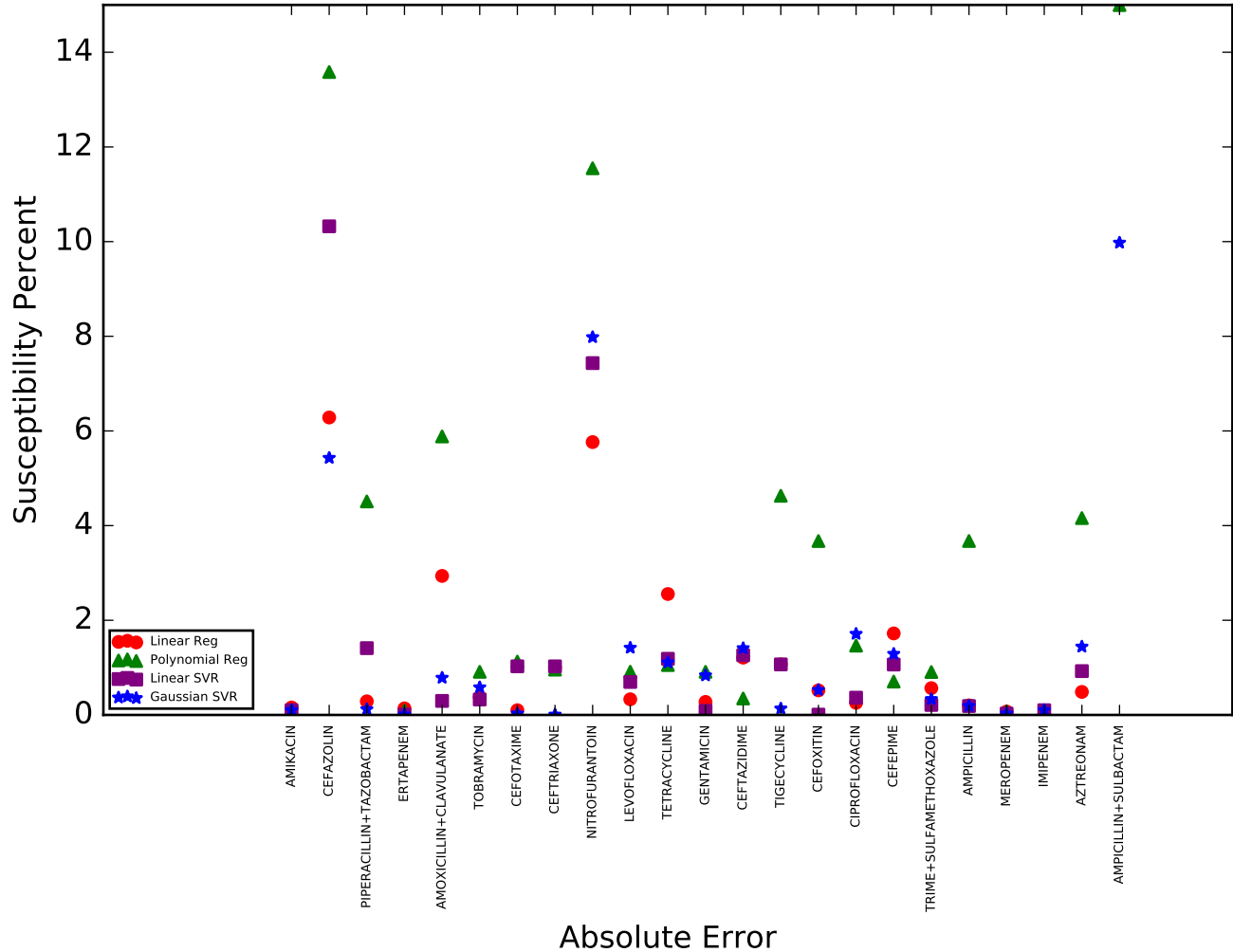
# E. coli

## using data from 2004-2012 to predict 2015



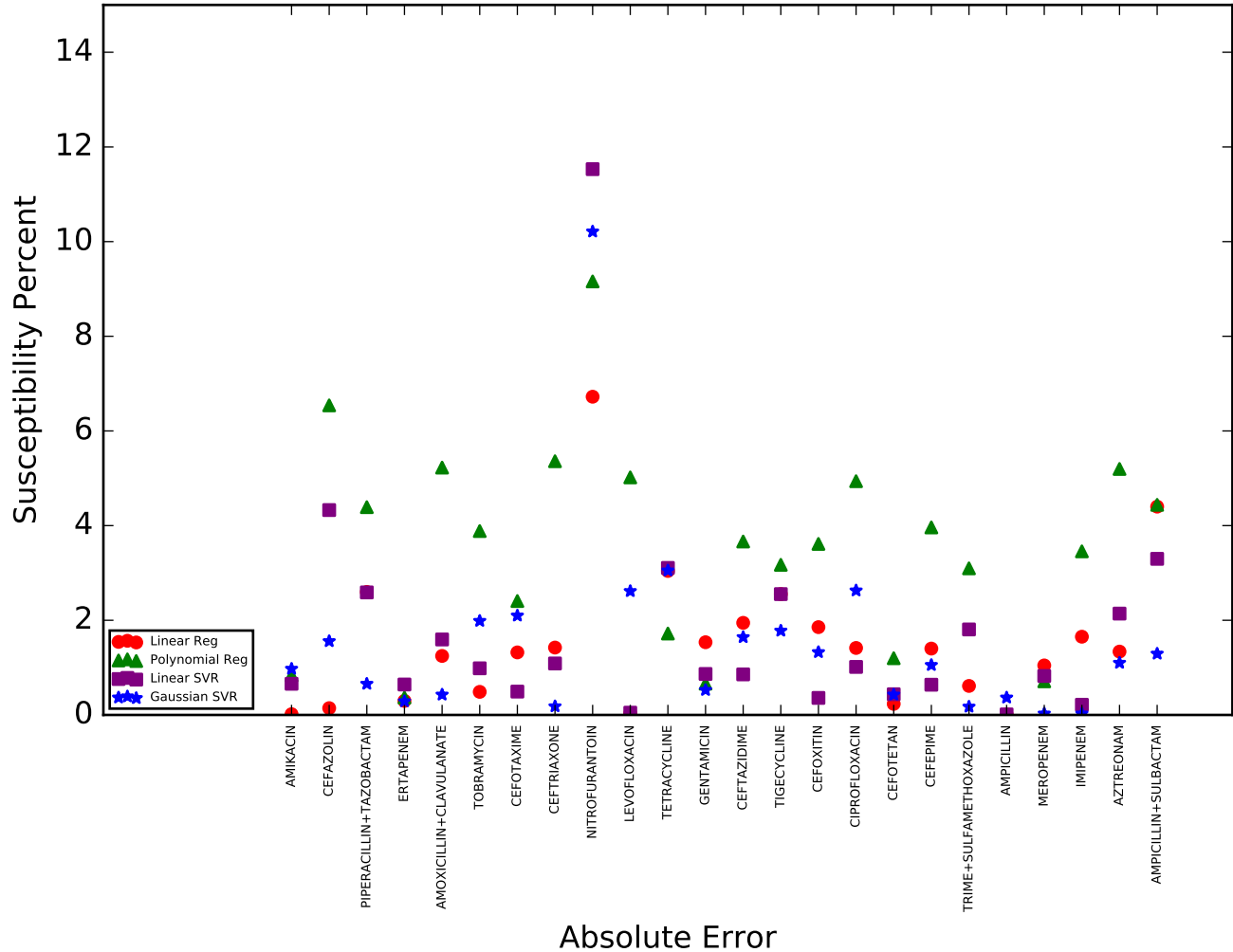
# Klebsiella oxytoca

## using data from 2004-2012 to predict 2015

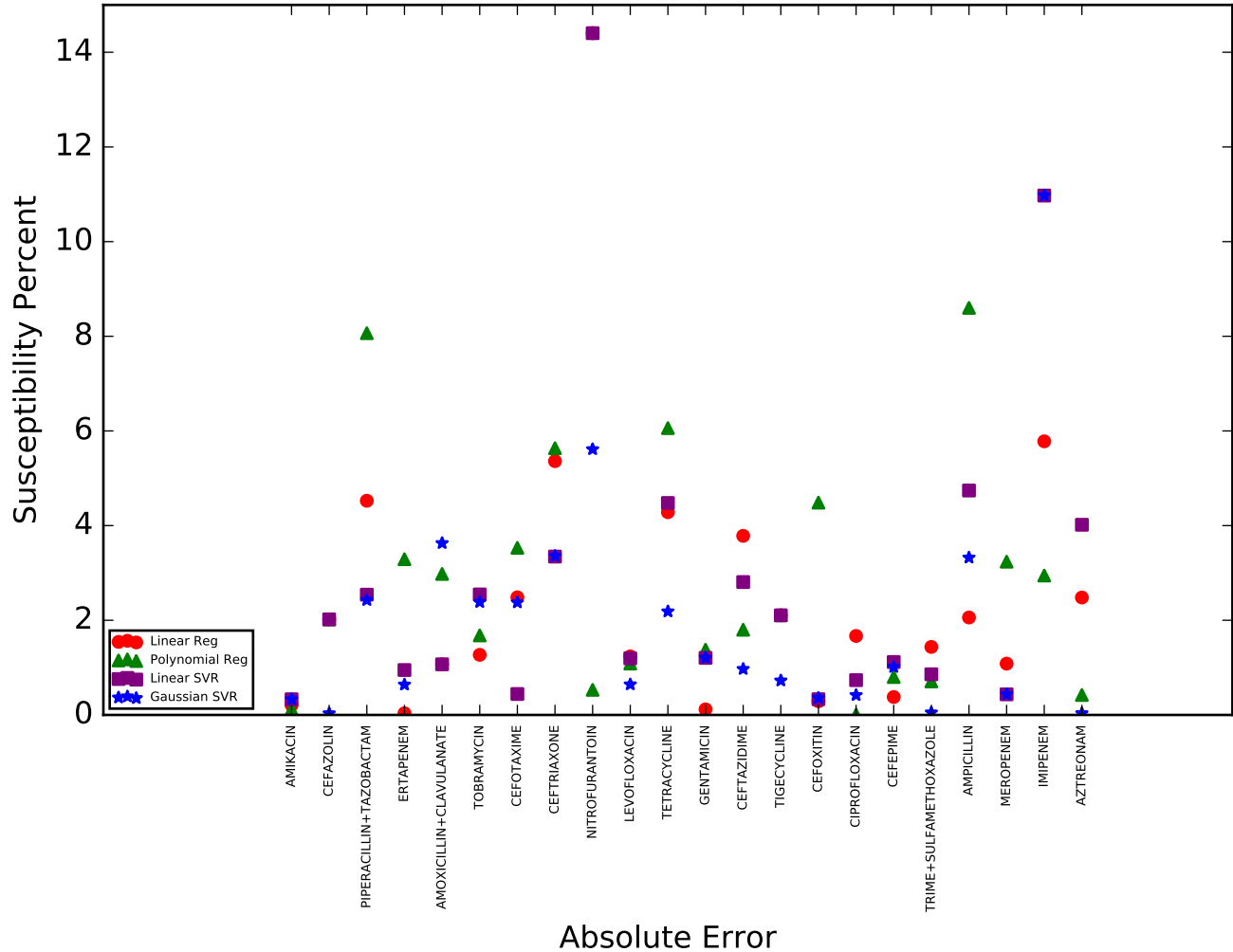


# Klebsiella pneumoniae

## using data from 2004-2012 to predict 2015

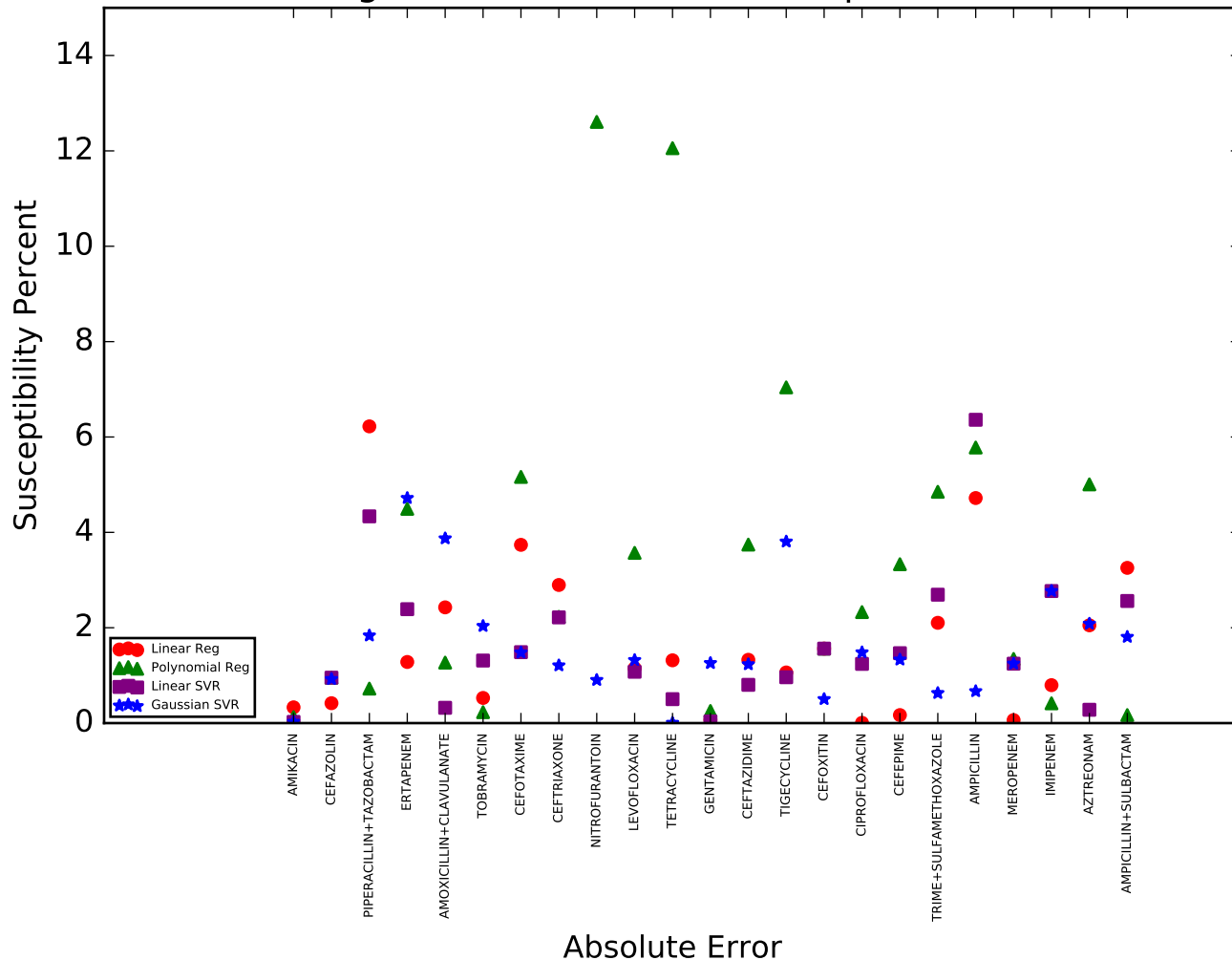


# Enterobacter aerogenes using data from 2004-2012 to predict 2015



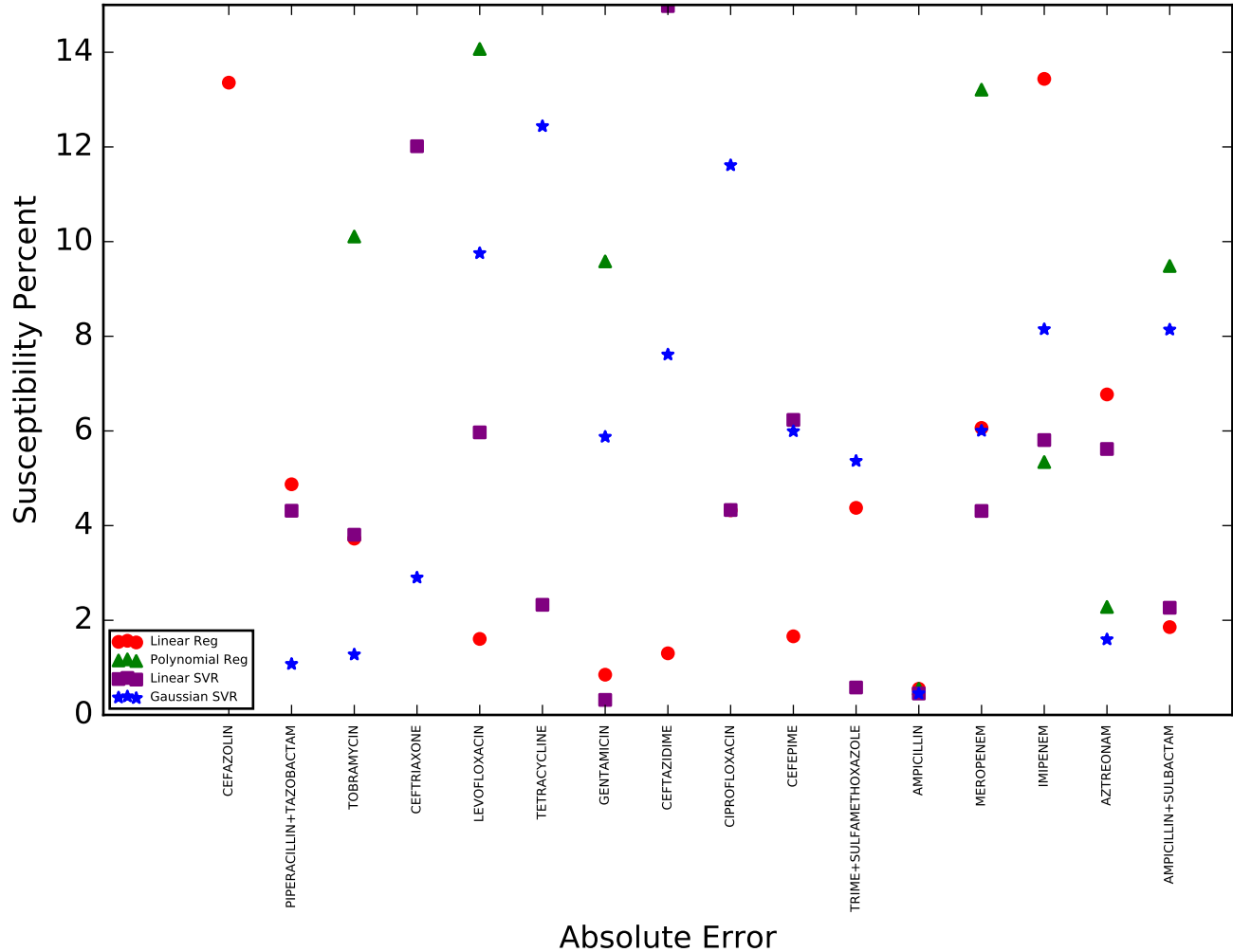


# Enterobacter cloacae using data from 2004-2012 to predict 2015

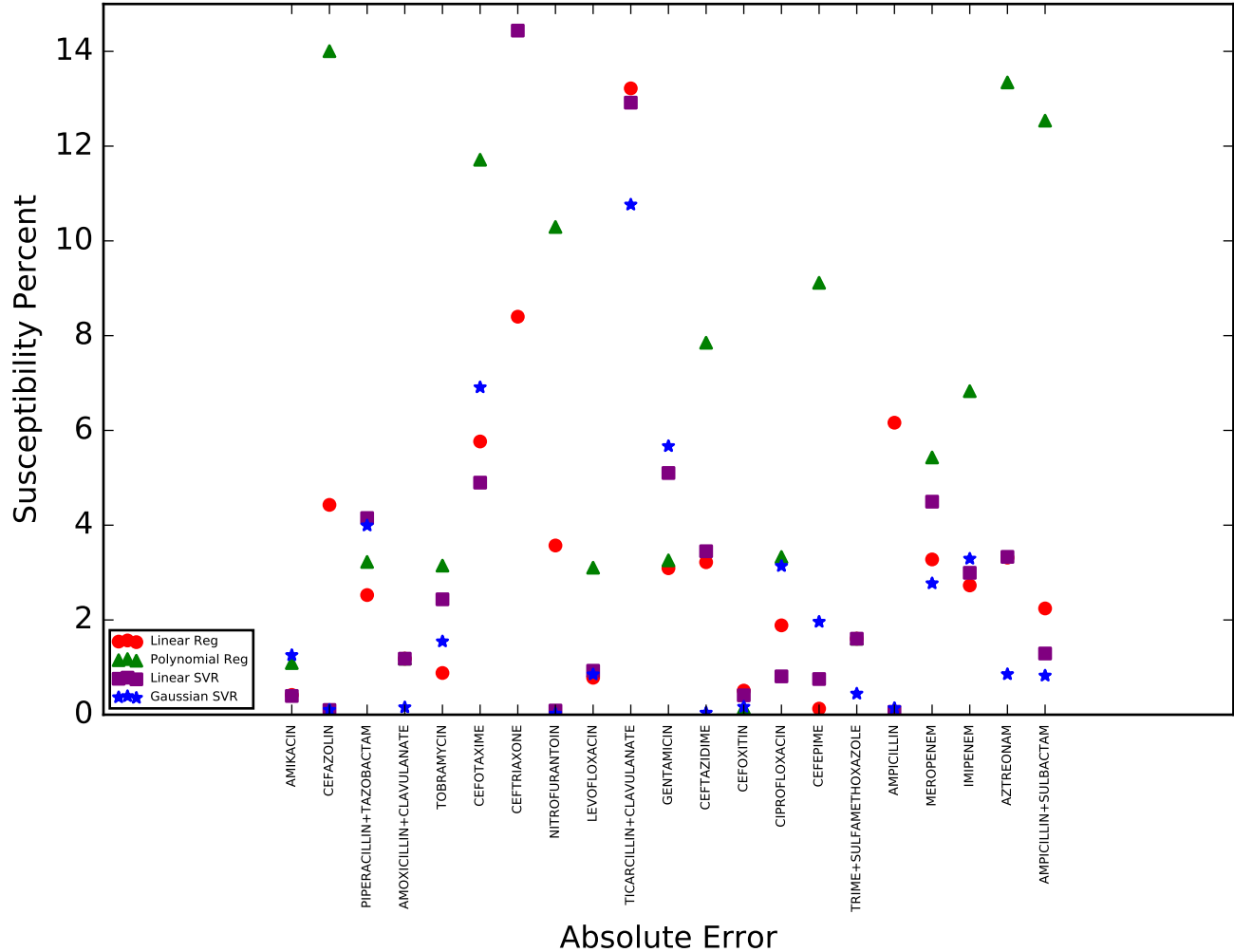


# Acinetobacter baumannii

## using data from 2004-2012 to predict 2015

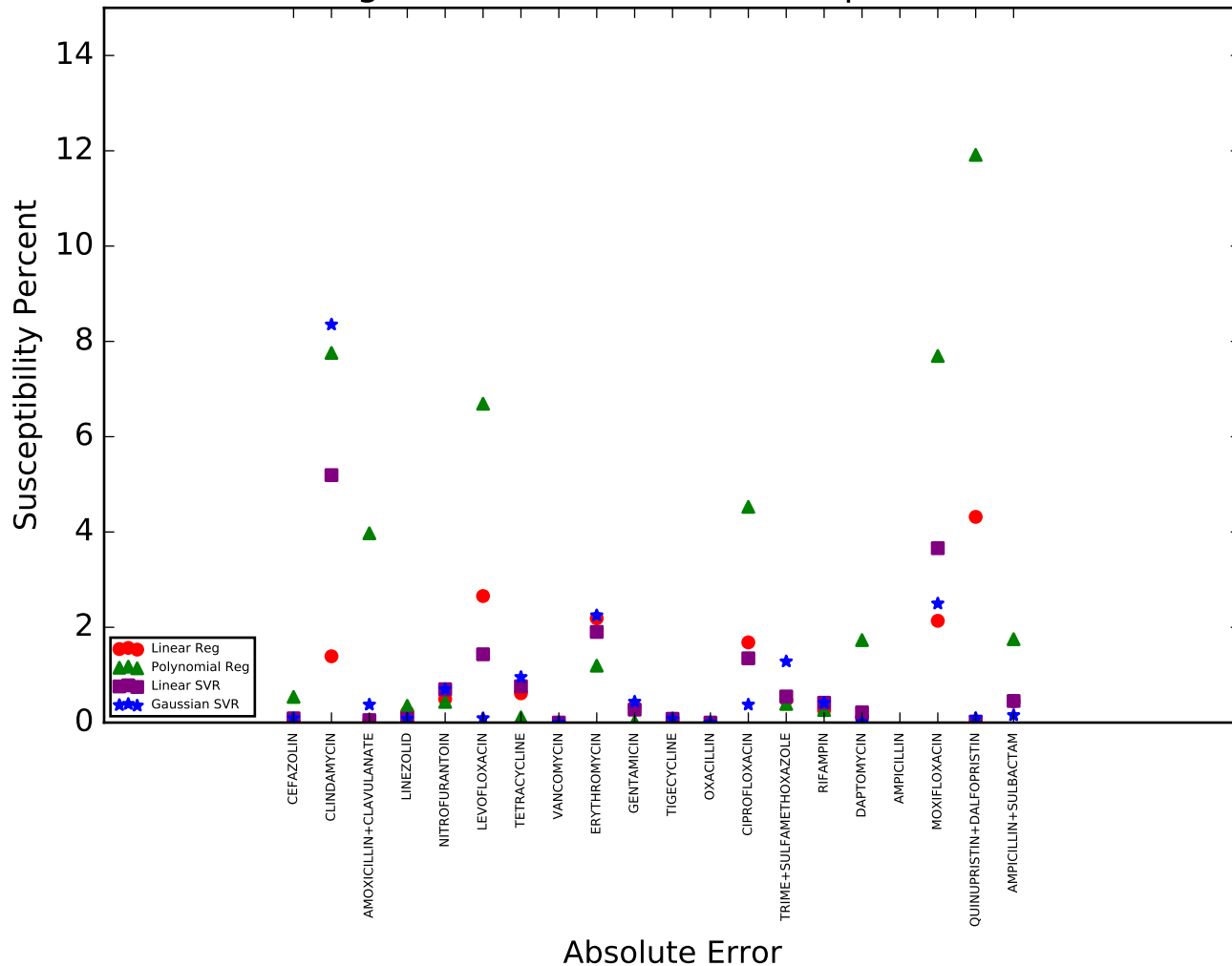


# *Pseudomonas aeruginosa* using data from 2004-2012 to predict 2015



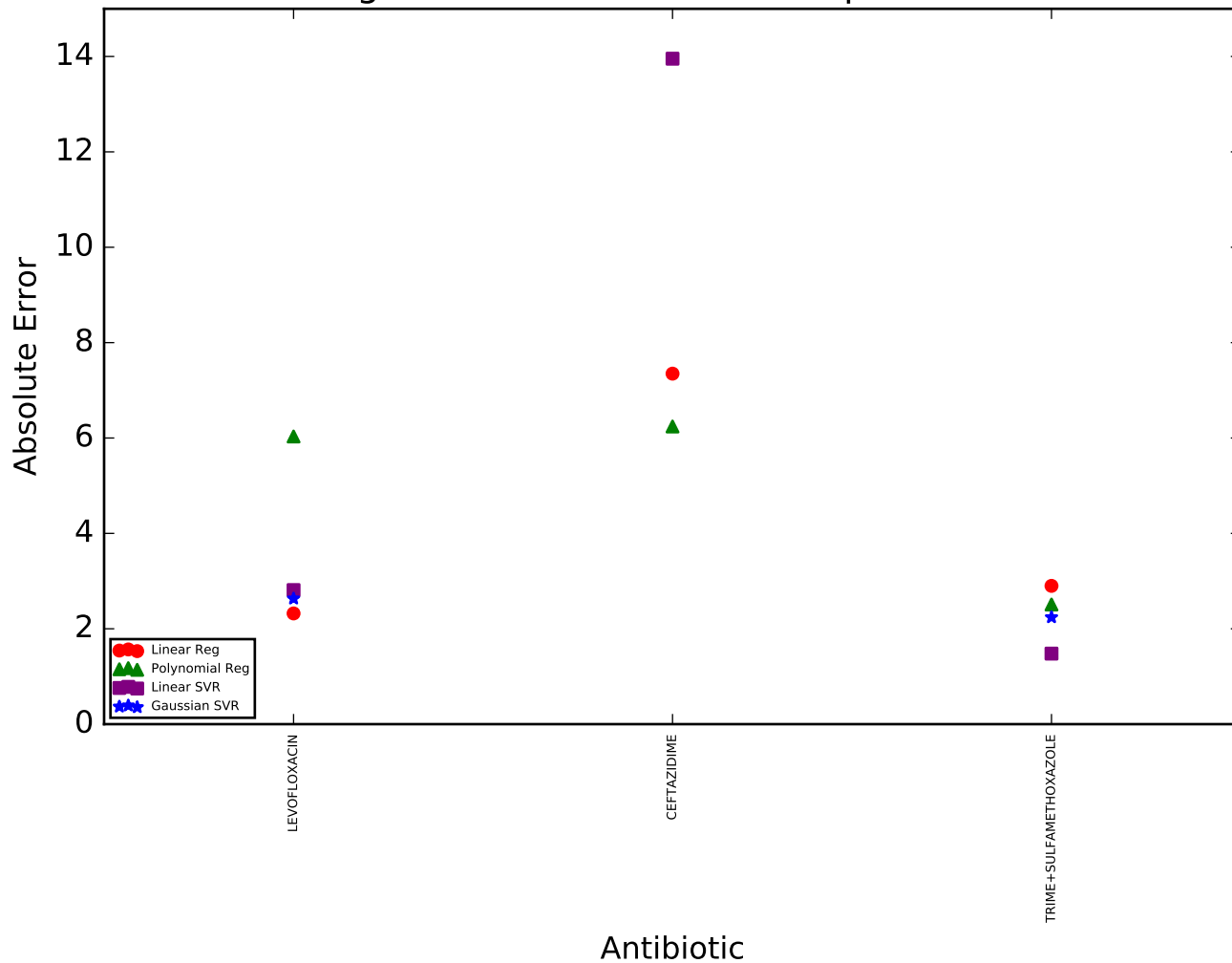
# MSSA

using data from 2004-2012 to predict 2015



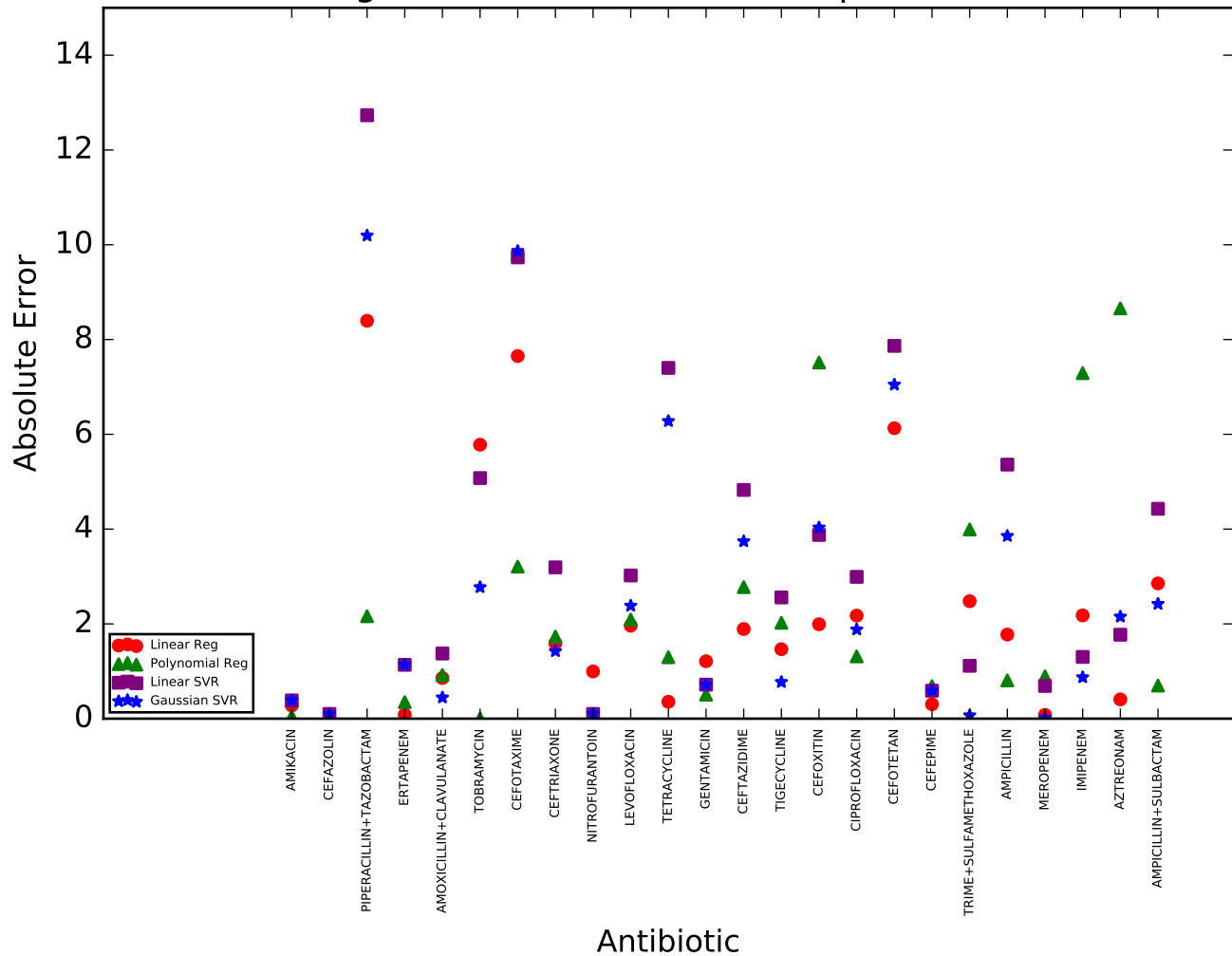
# Stenotrophomonas maltophilia

using data from 2005-2013 to predict 2014



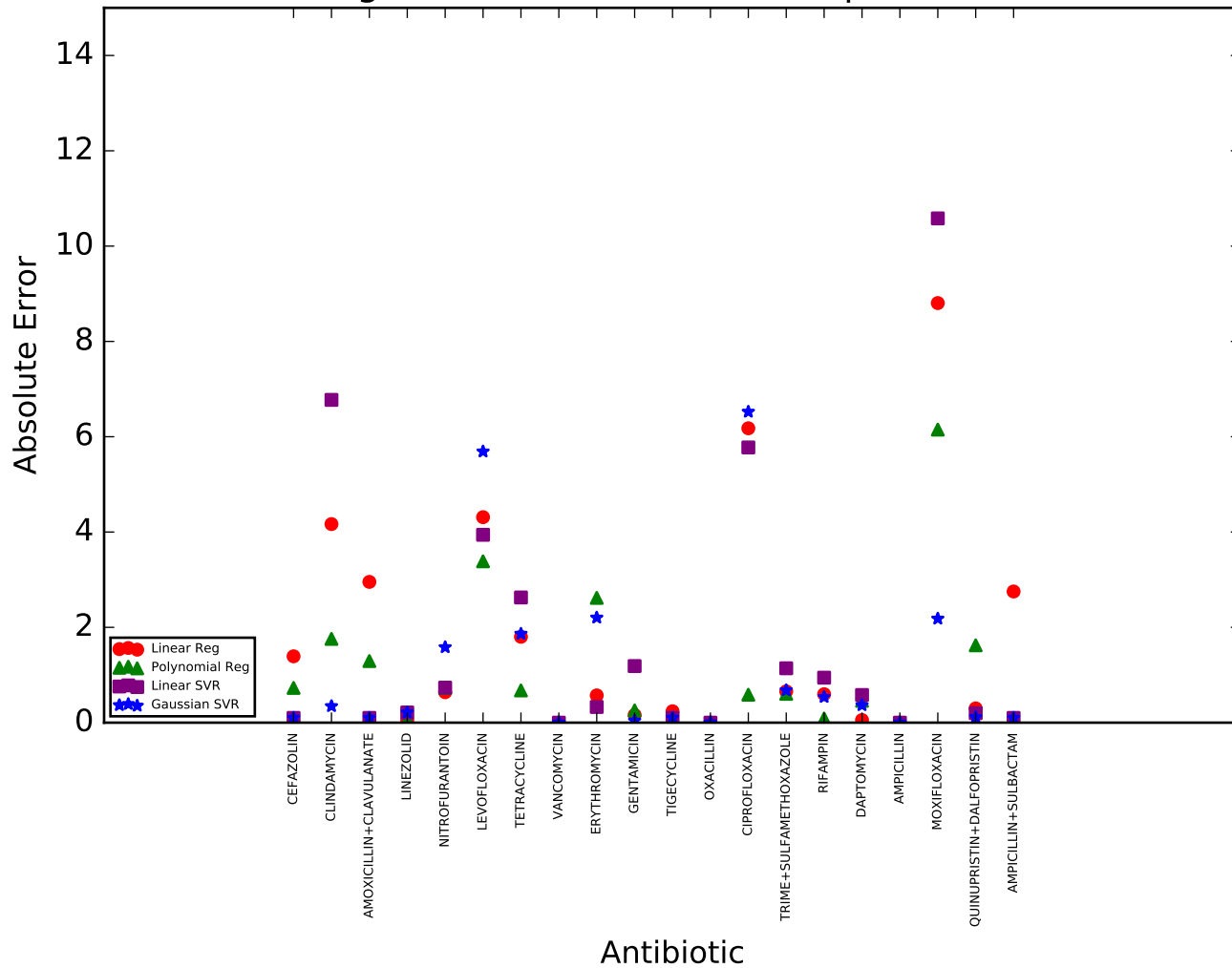
# Serratia marcescens

## using data from 2005-2013 to predict 2014



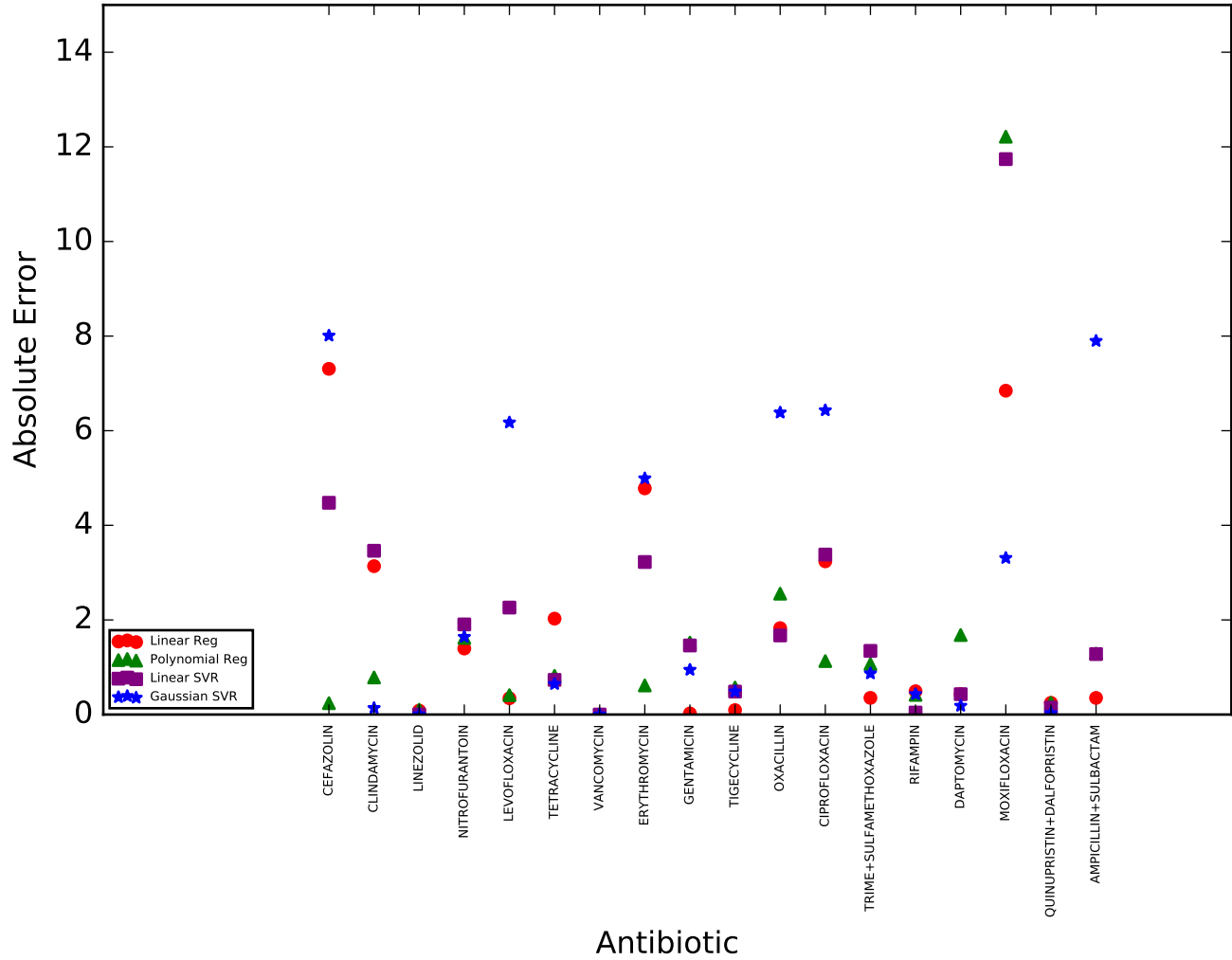
# MRSA

using data from 2005-2013 to predict 2014



# Staphylococcus aureus

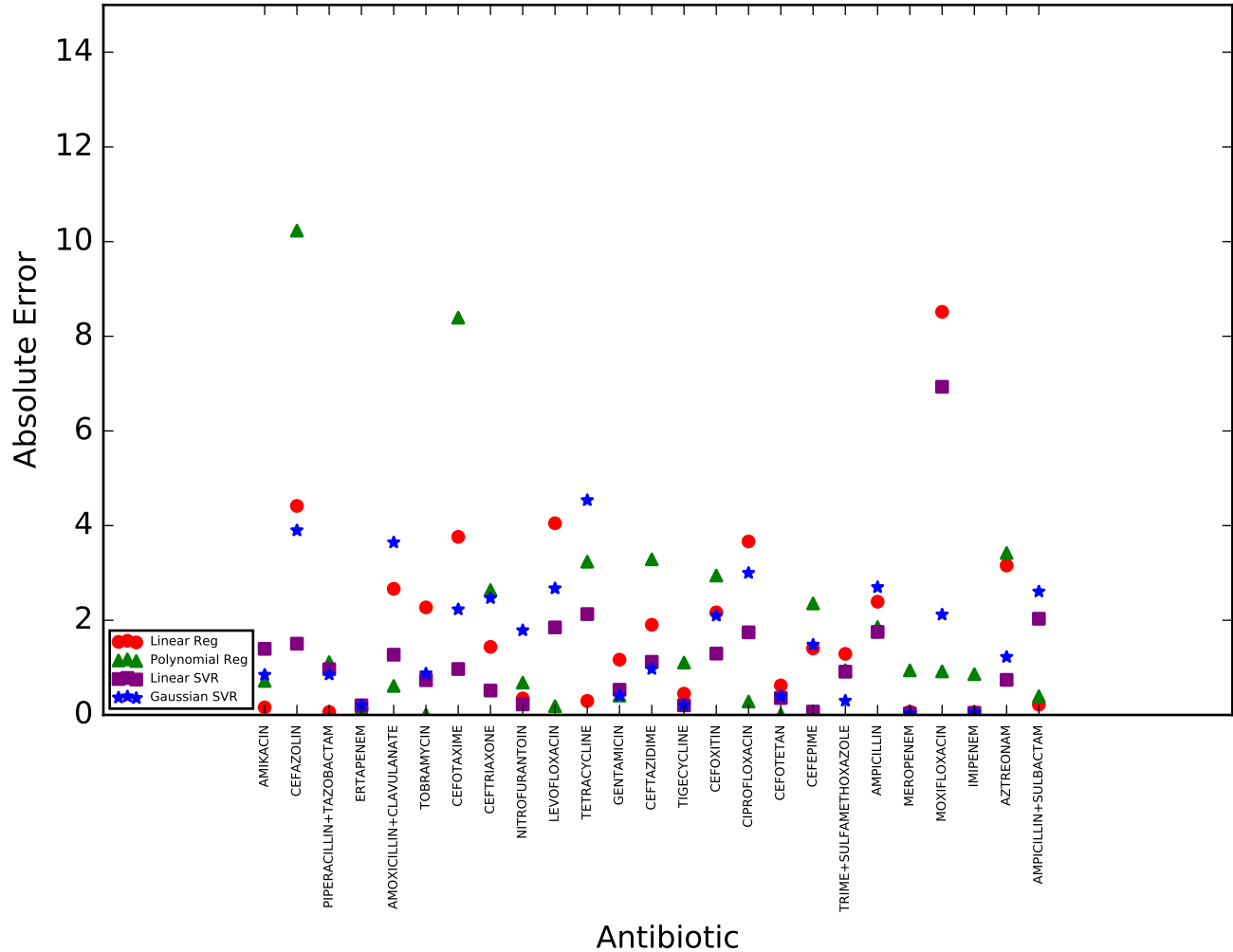
## using data from 2005-2013 to predict 2014





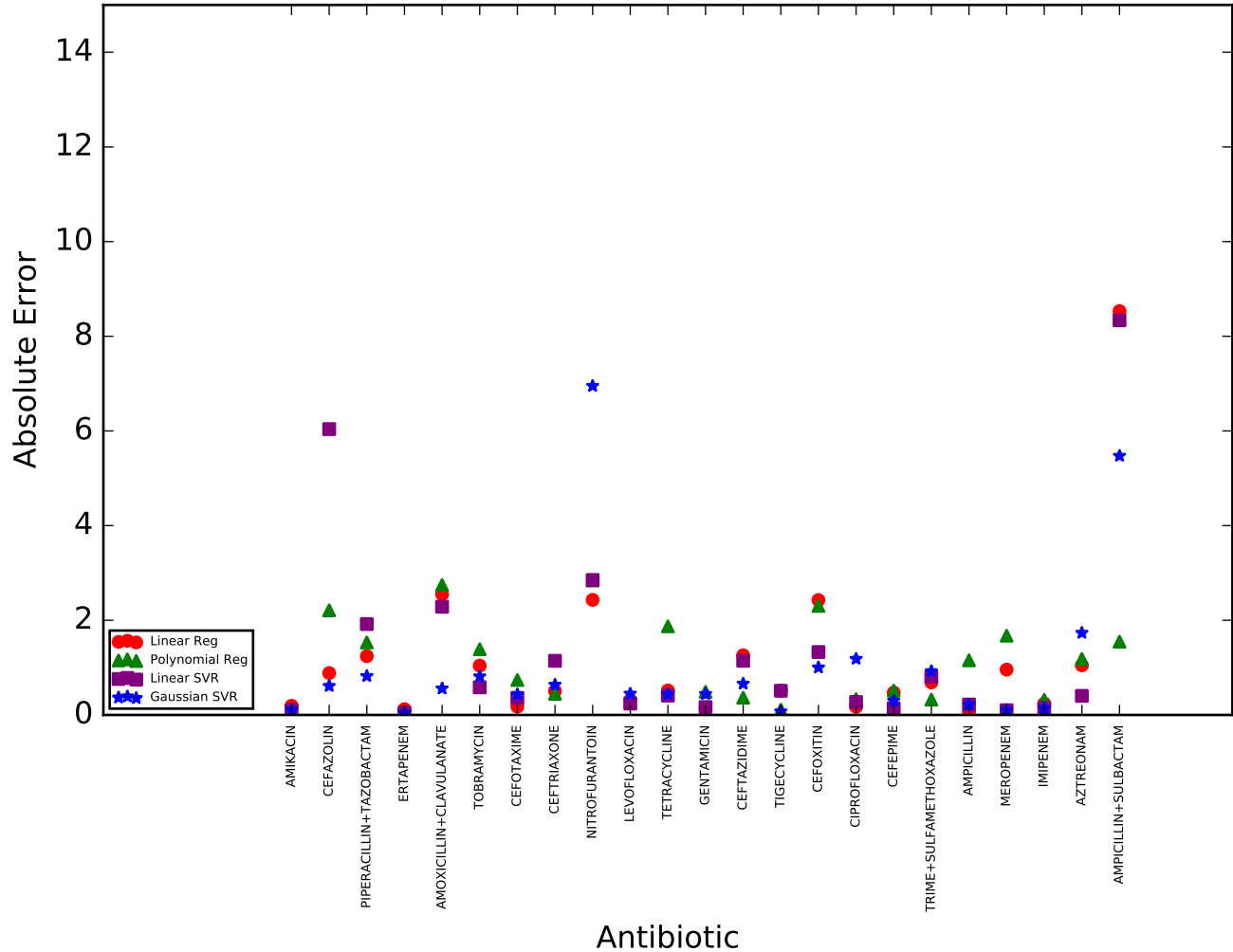
# E. coli

## using data from 2005-2013 to predict 2014



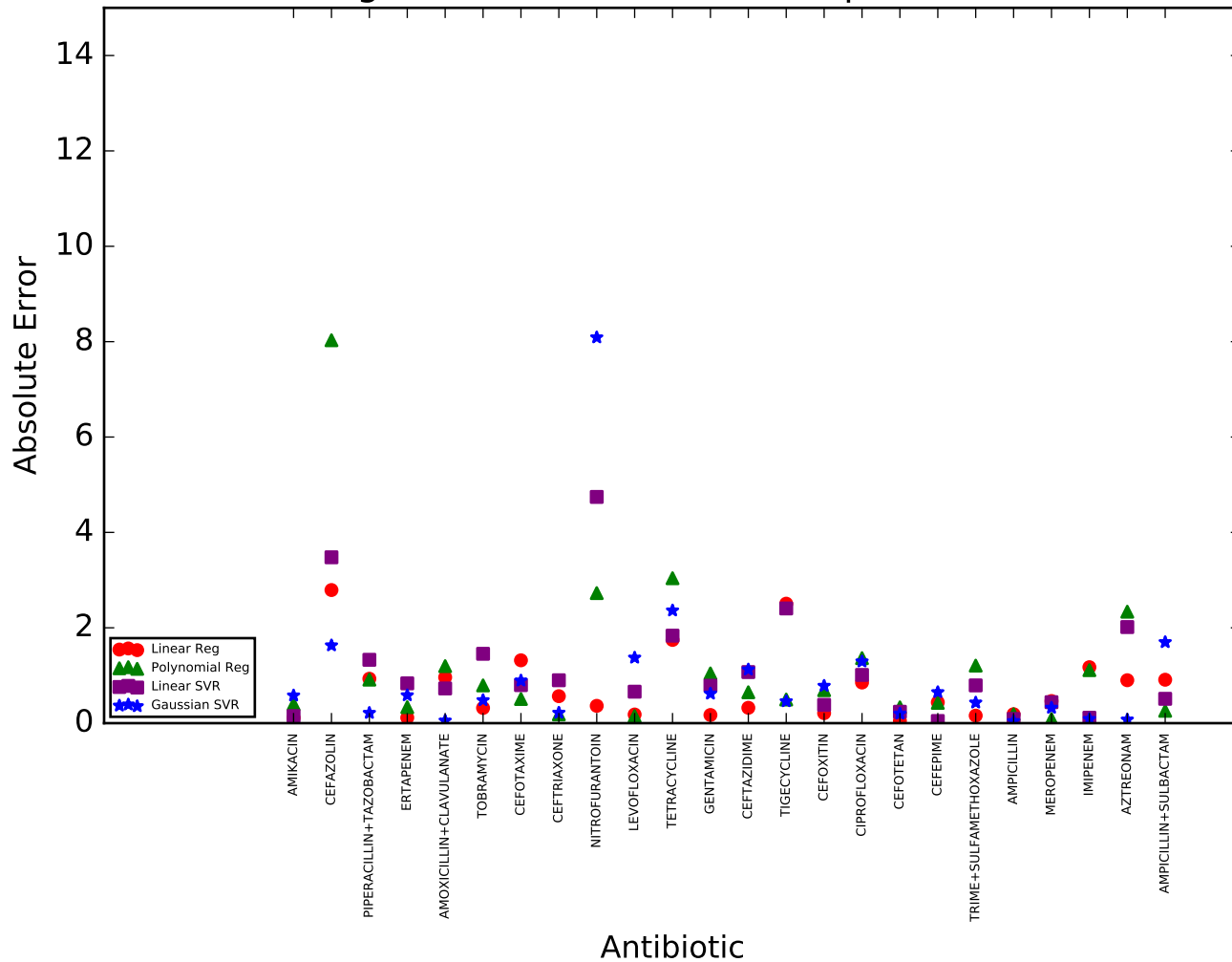
# Klebsiella oxytoca

## using data from 2005-2013 to predict 2014

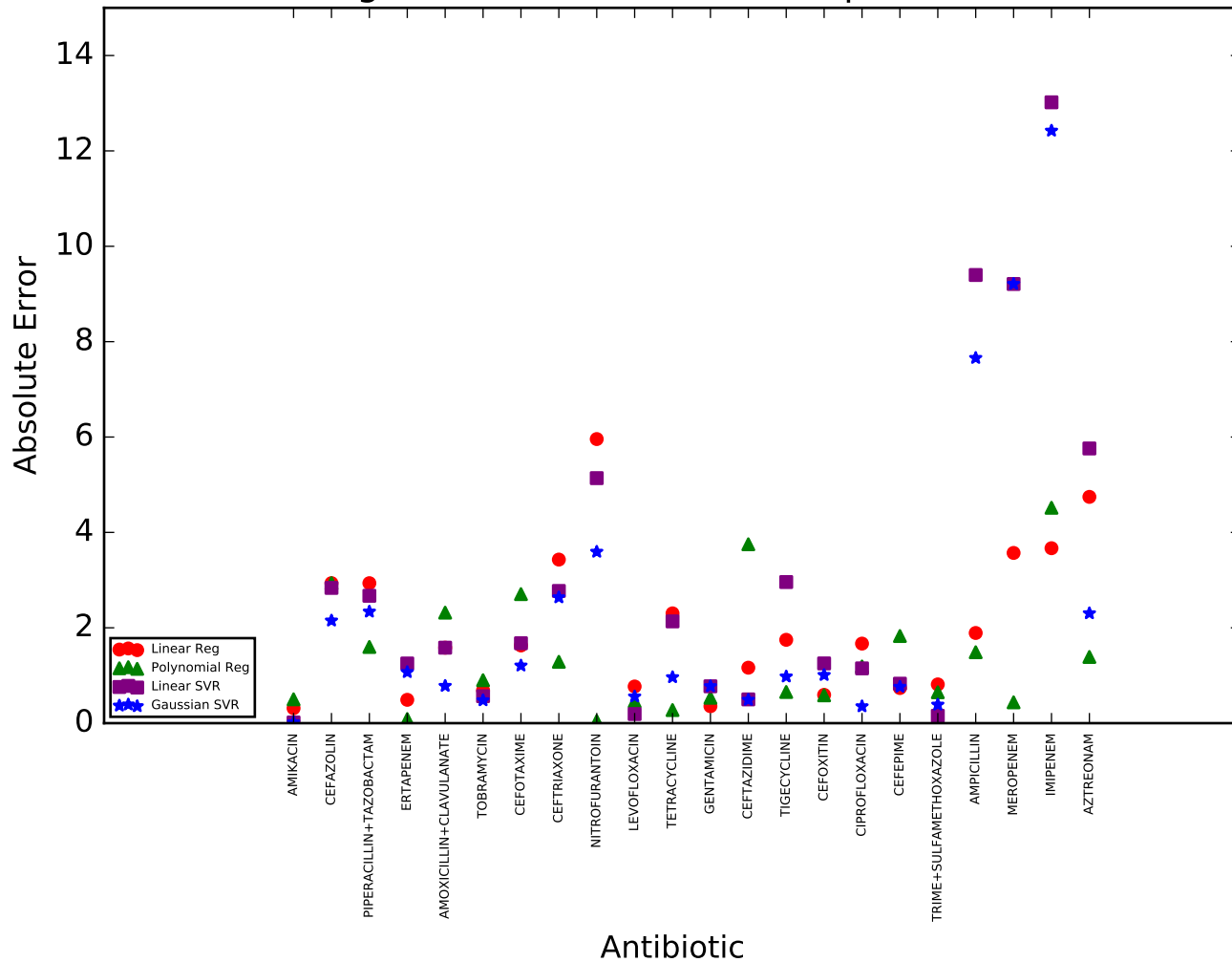


# Klebsiella pneumoniae

## using data from 2005-2013 to predict 2014

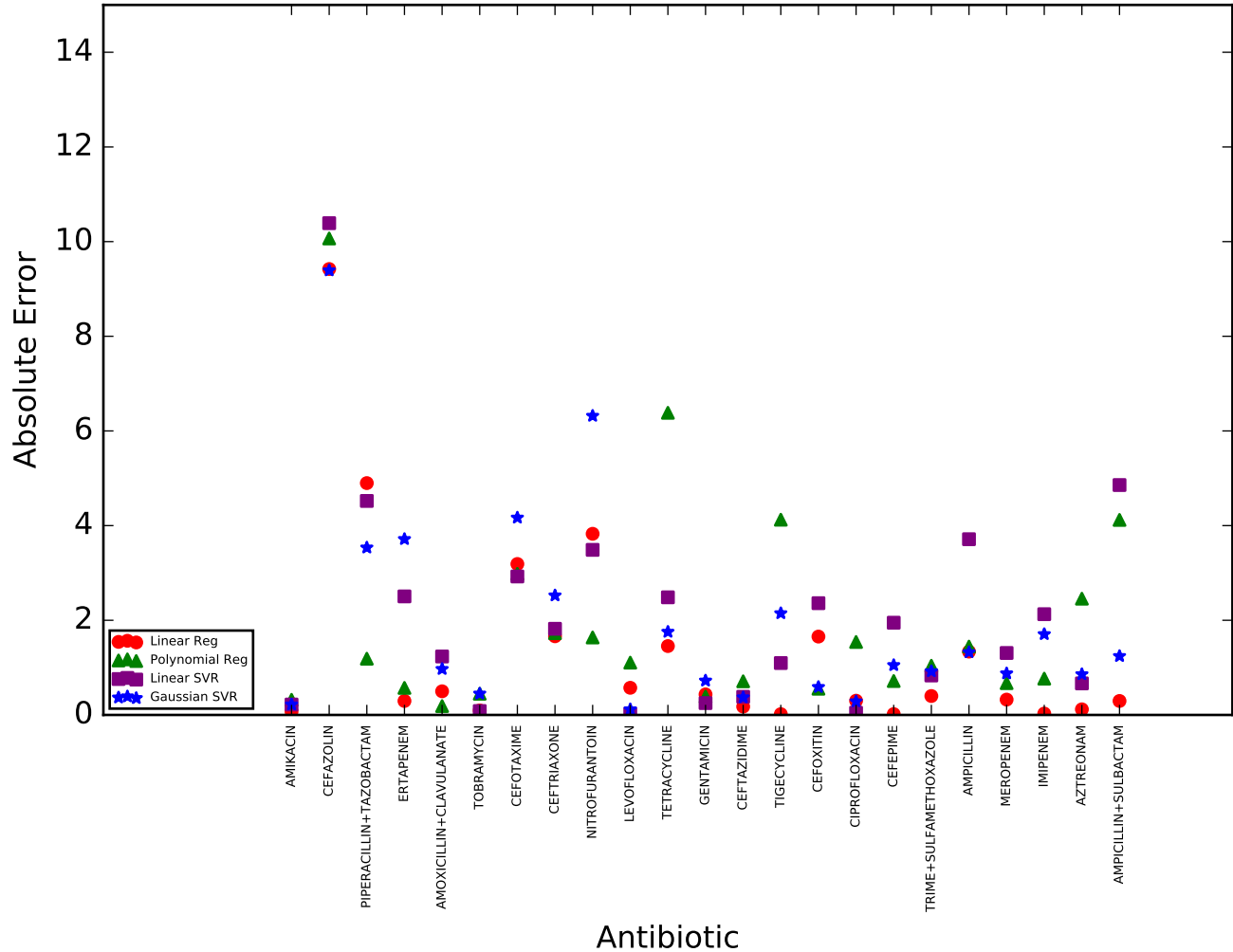


# Enterobacter aerogenes using data from 2005-2013 to predict 2014



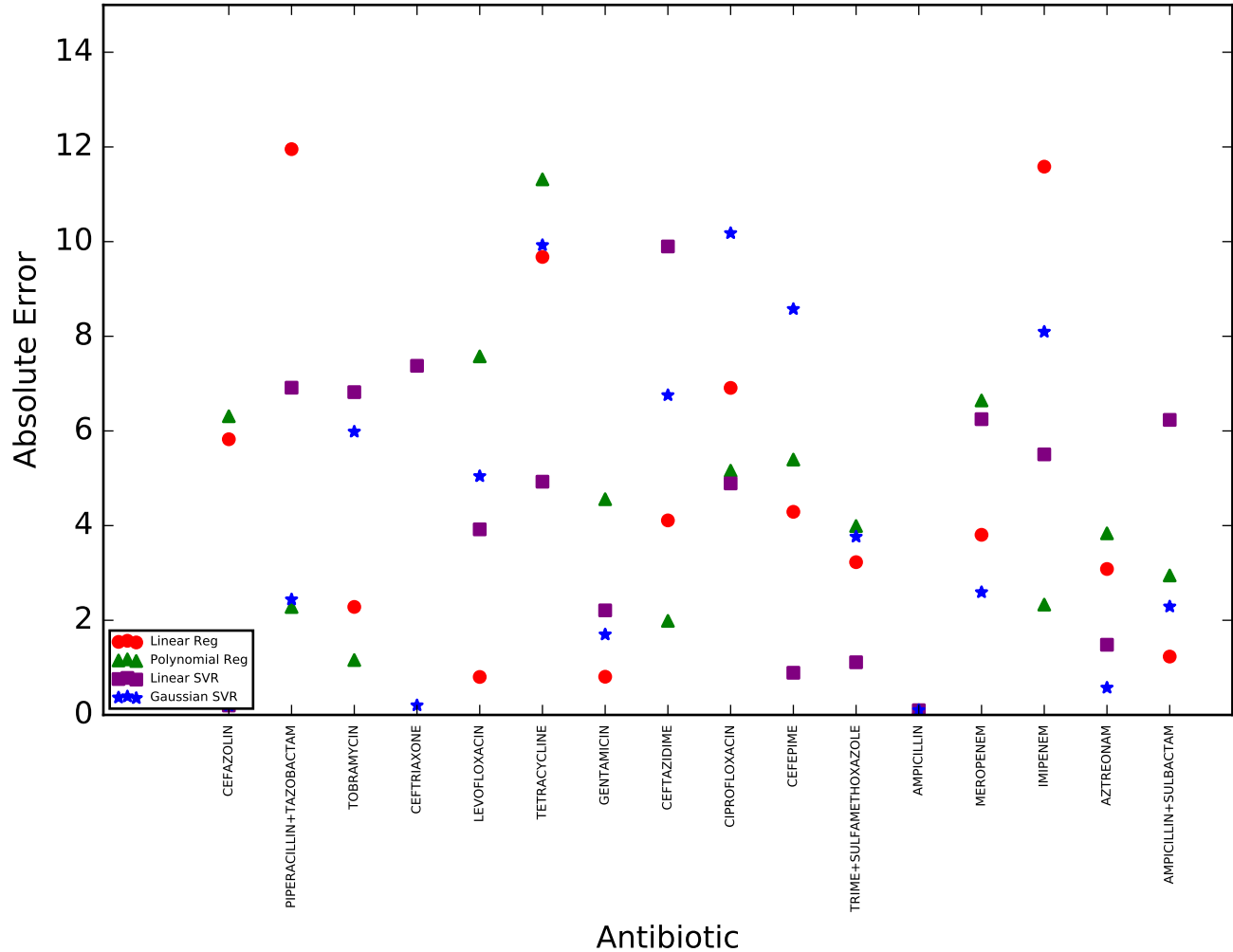
# Enterobacter cloacae

## using data from 2005-2013 to predict 2014

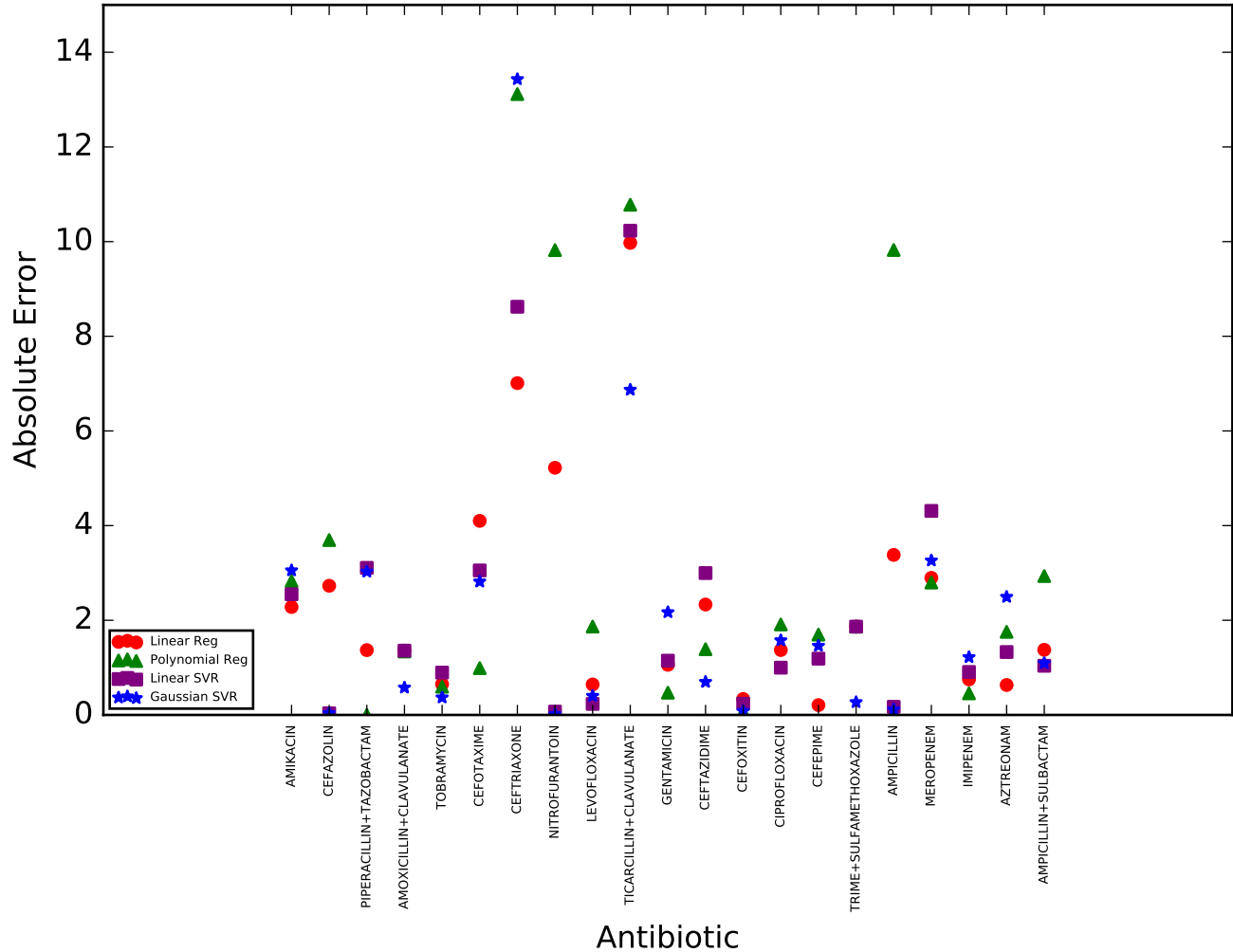


# Acinetobacter baumannii

## using data from 2005-2013 to predict 2014

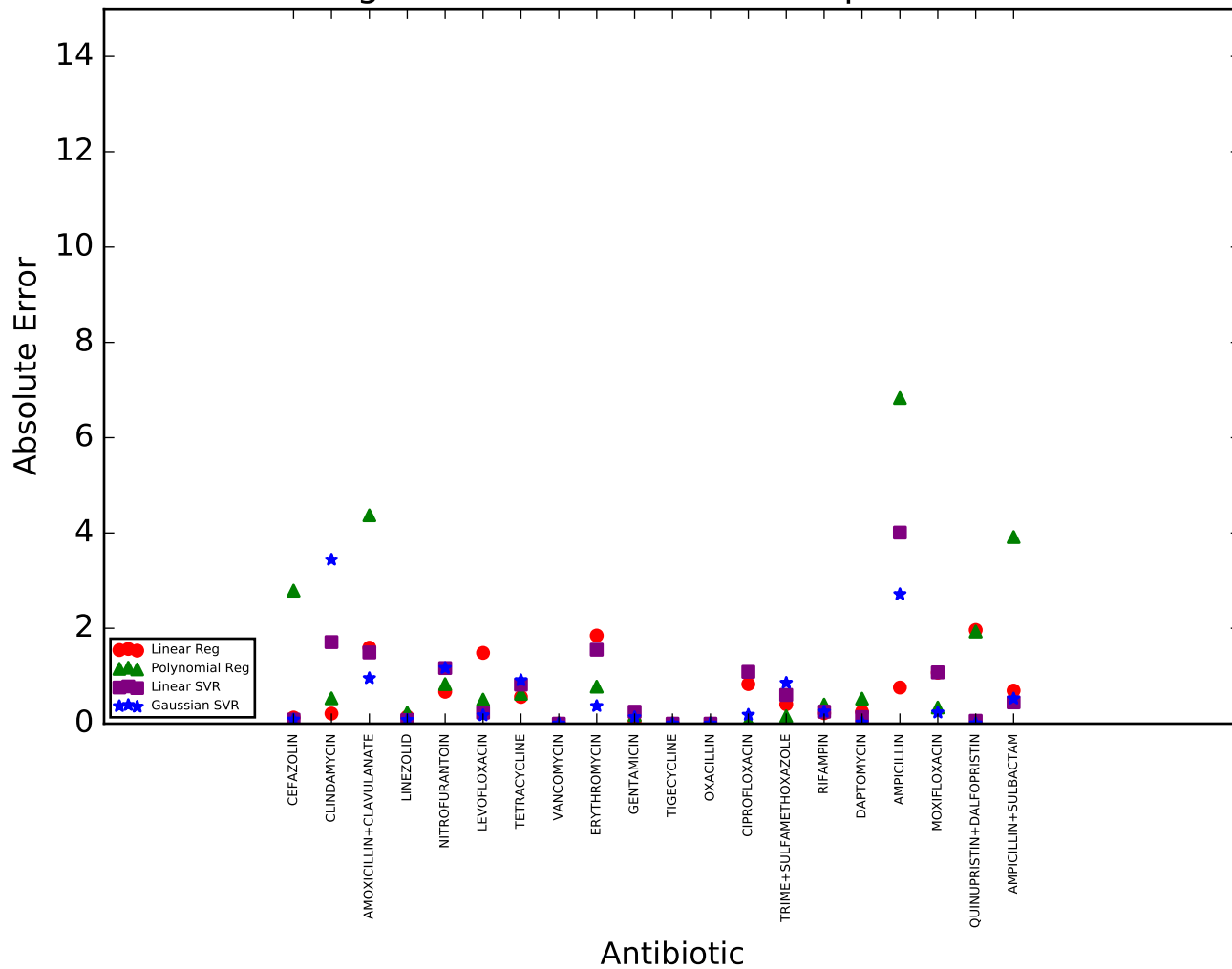


# *Pseudomonas aeruginosa* using data from 2005-2013 to predict 2014



# MSSA

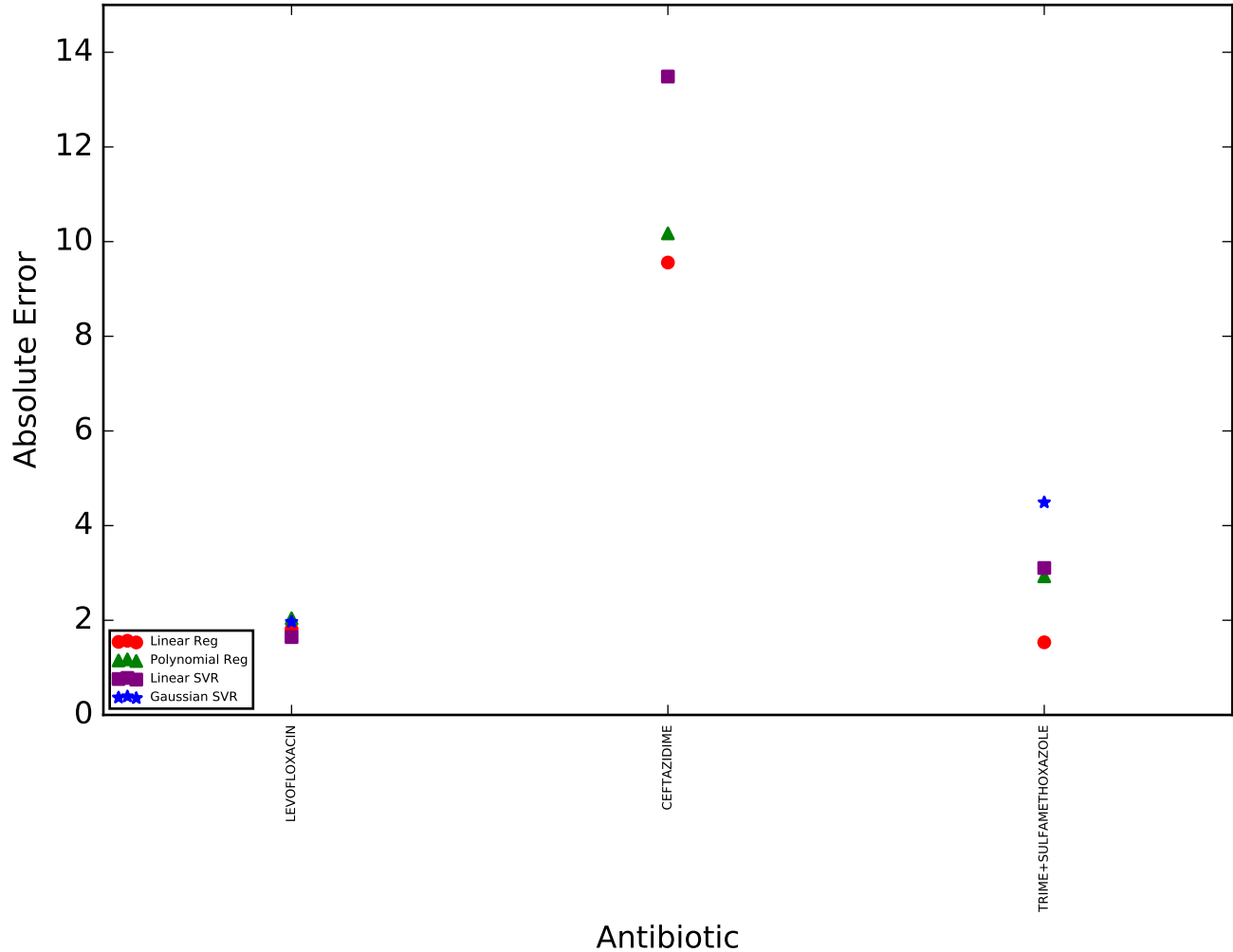
using data from 2005-2013 to predict 2014





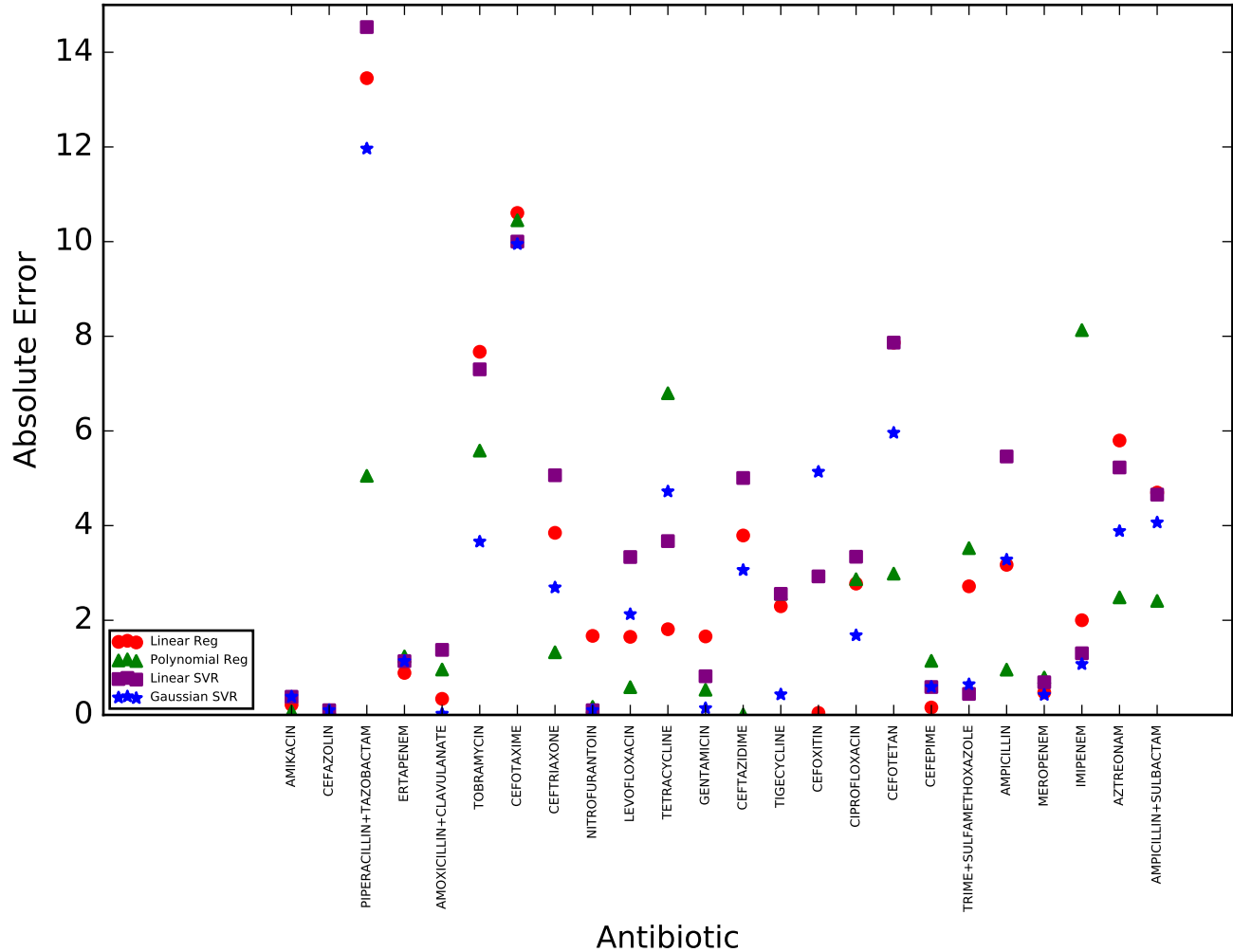
# Stenotrophomonas maltophilia

using data from 2004-2012 to predict 2014



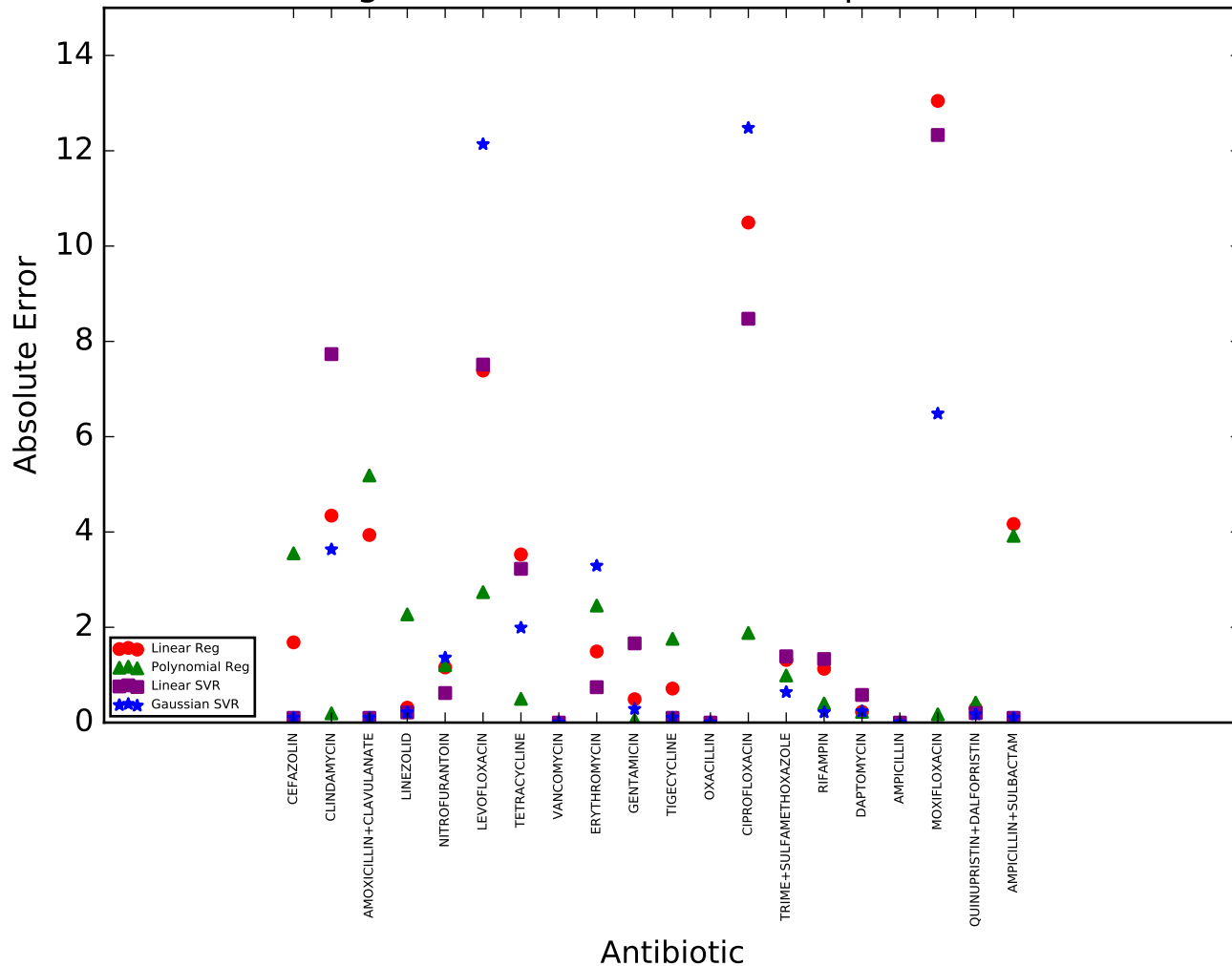
# Serratia marcescens

## using data from 2004-2012 to predict 2014



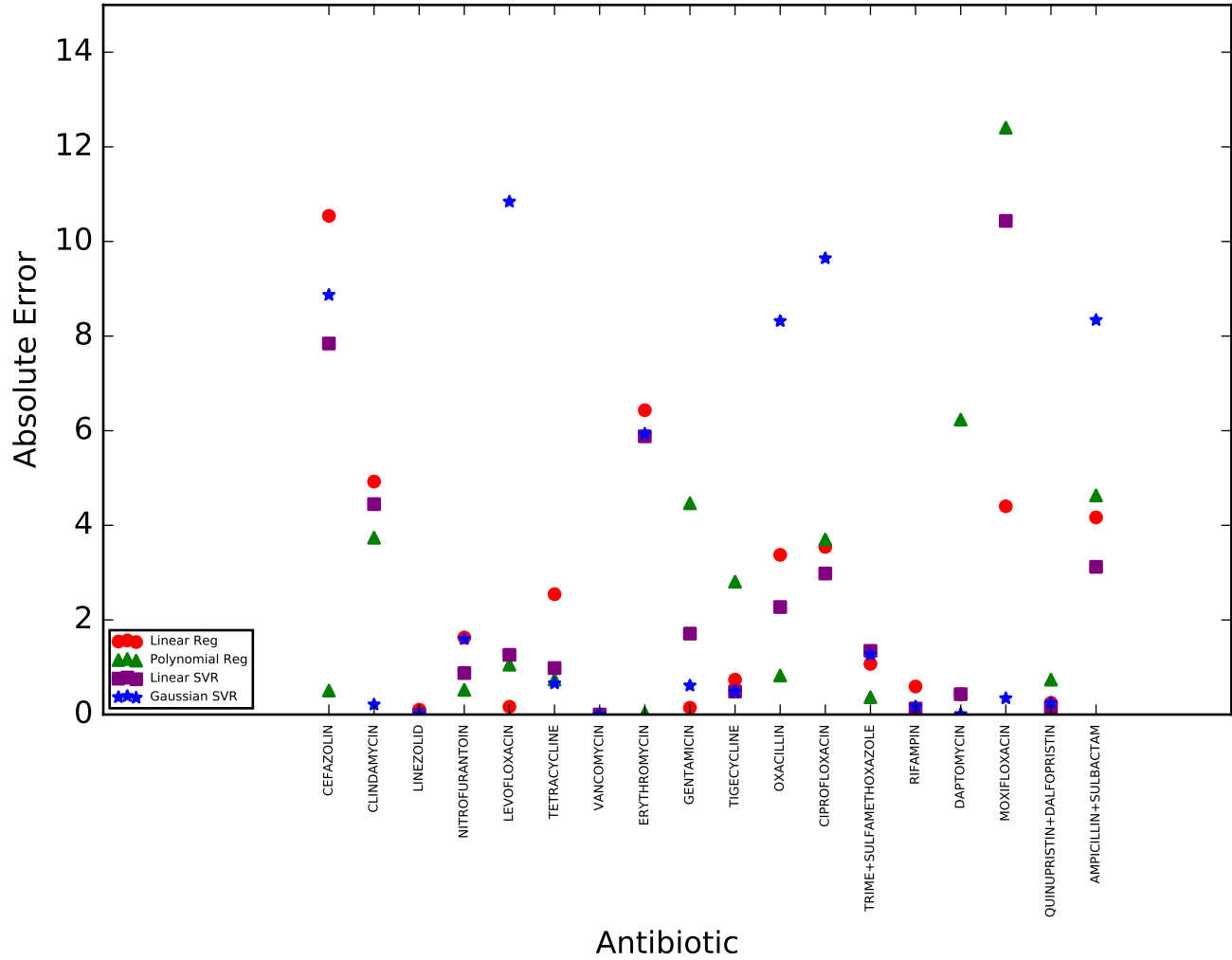
# MRSA

## using data from 2004-2012 to predict 2014



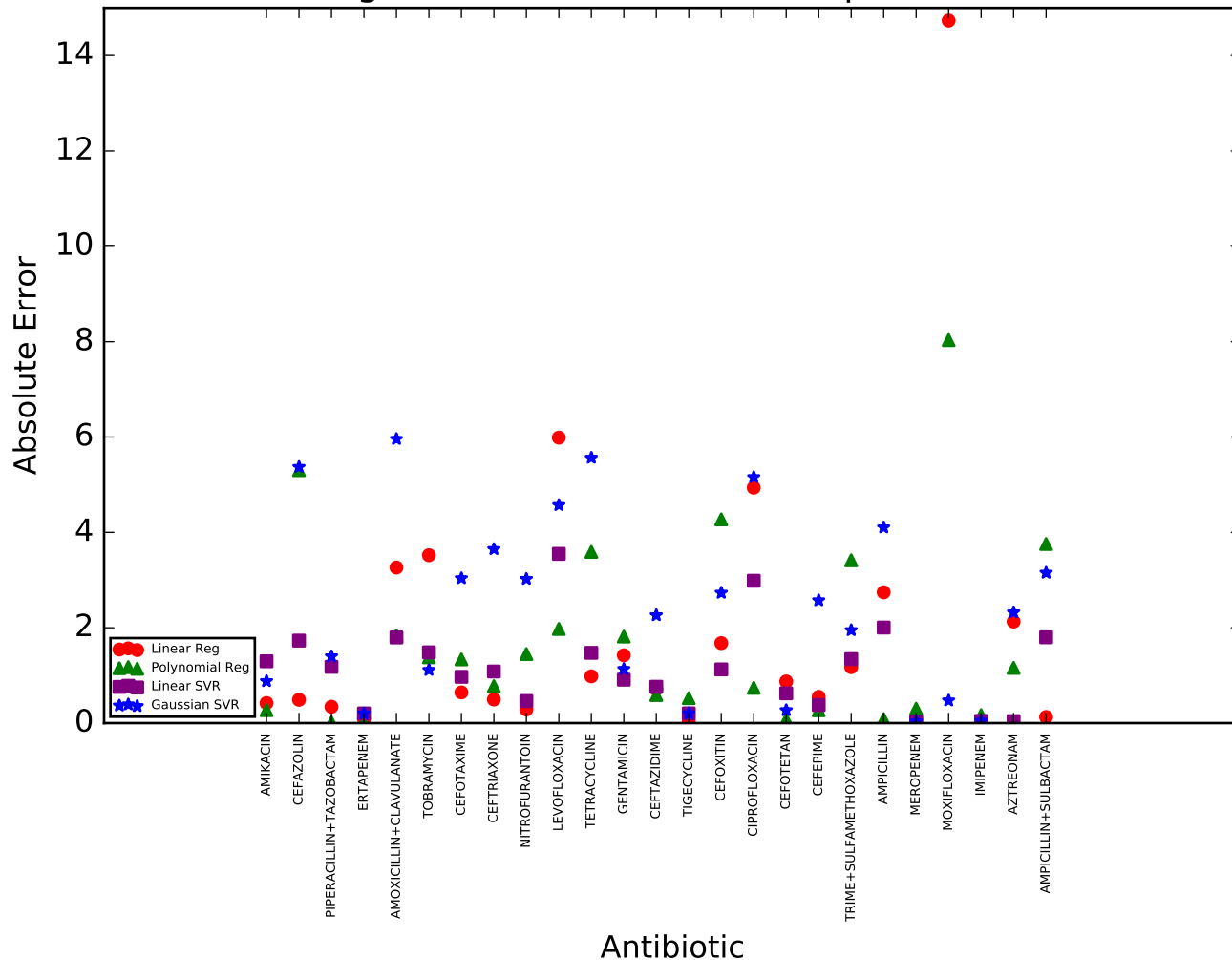
# Staphylococcus aureus

## using data from 2004-2012 to predict 2014



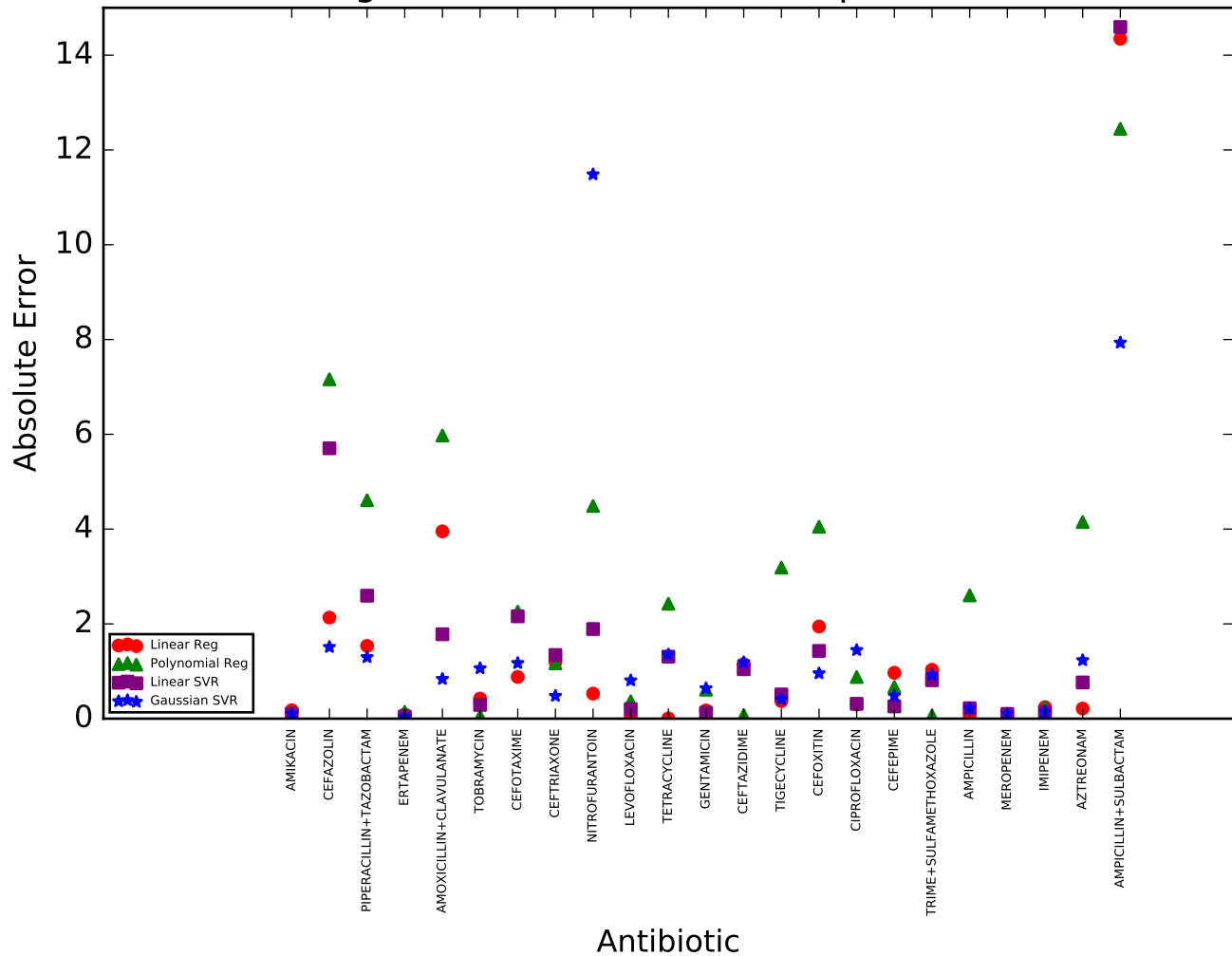
# E. coli

## using data from 2004-2012 to predict 2014



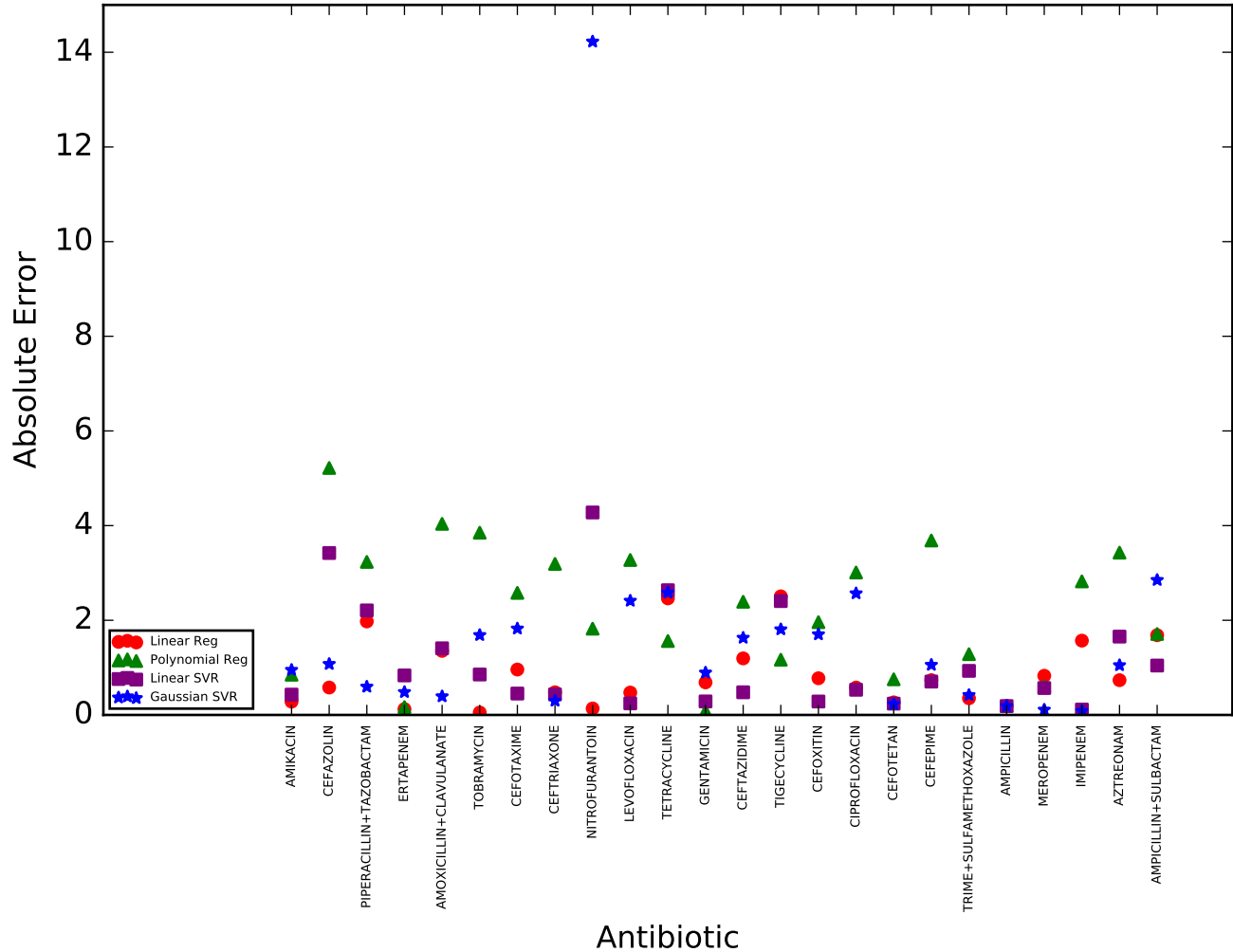
# Klebsiella oxytoca

## using data from 2004-2012 to predict 2014



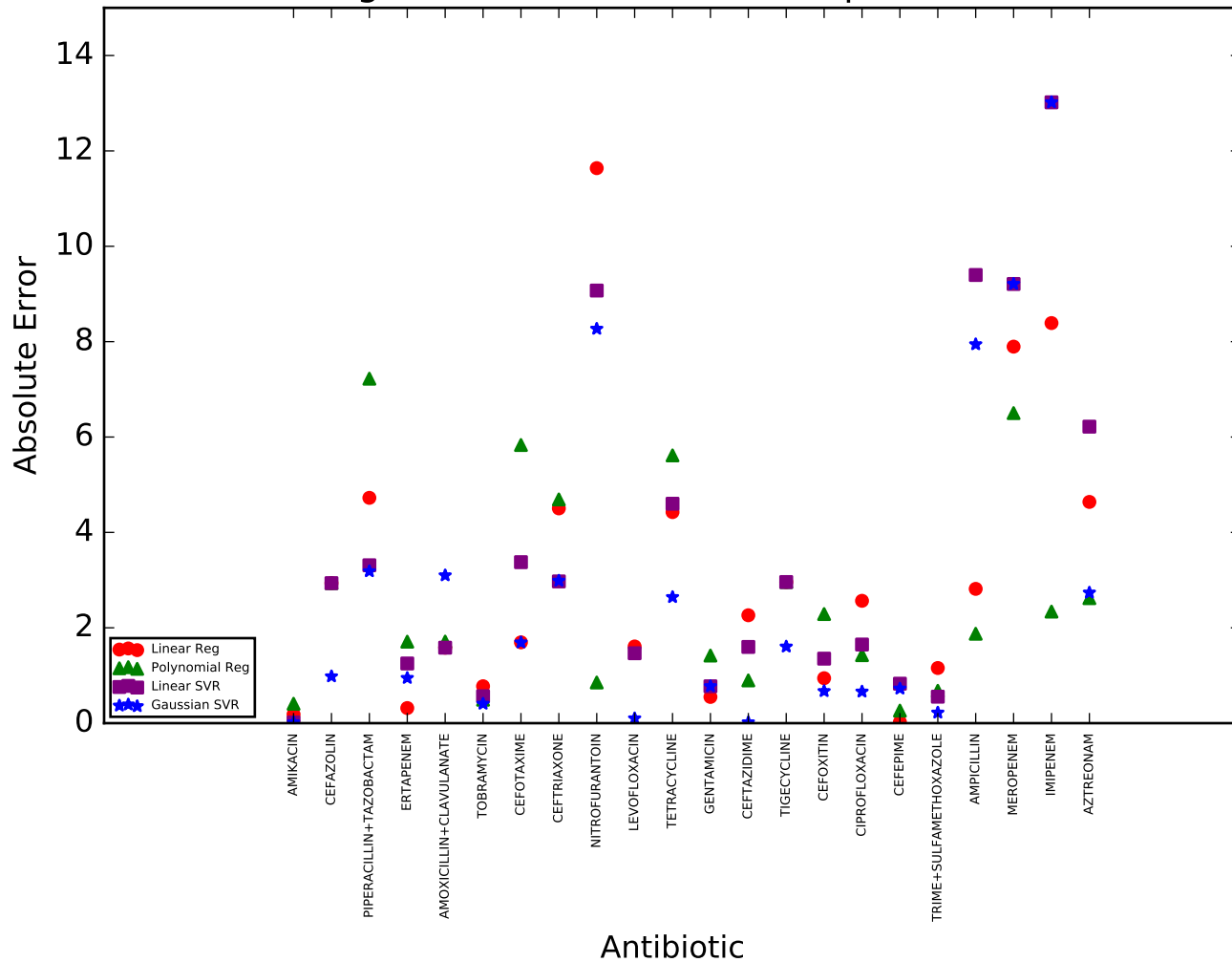
# Klebsiella pneumoniae

## using data from 2004-2012 to predict 2014



# Enterobacter aerogenes

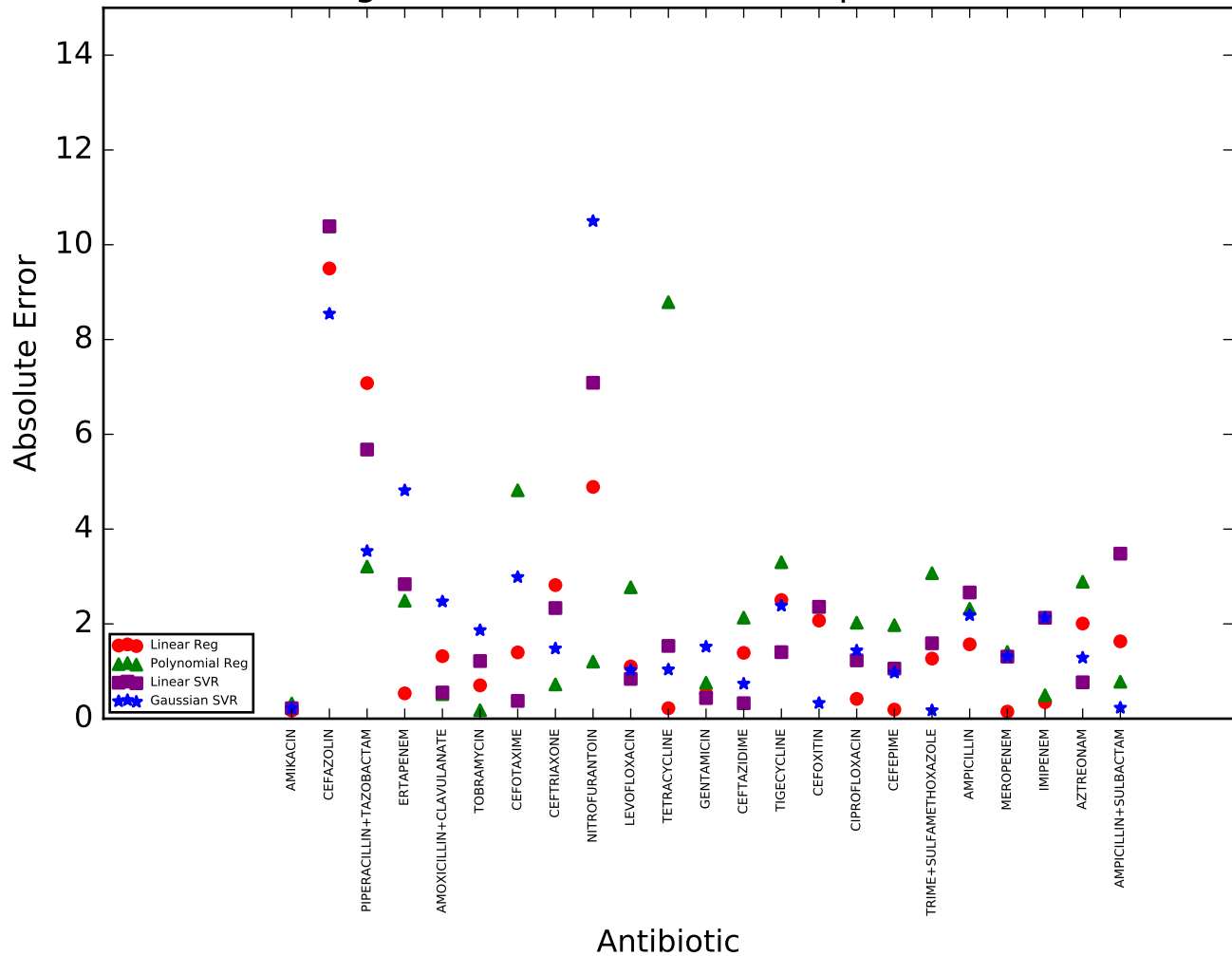
## using data from 2004-2012 to predict 2014



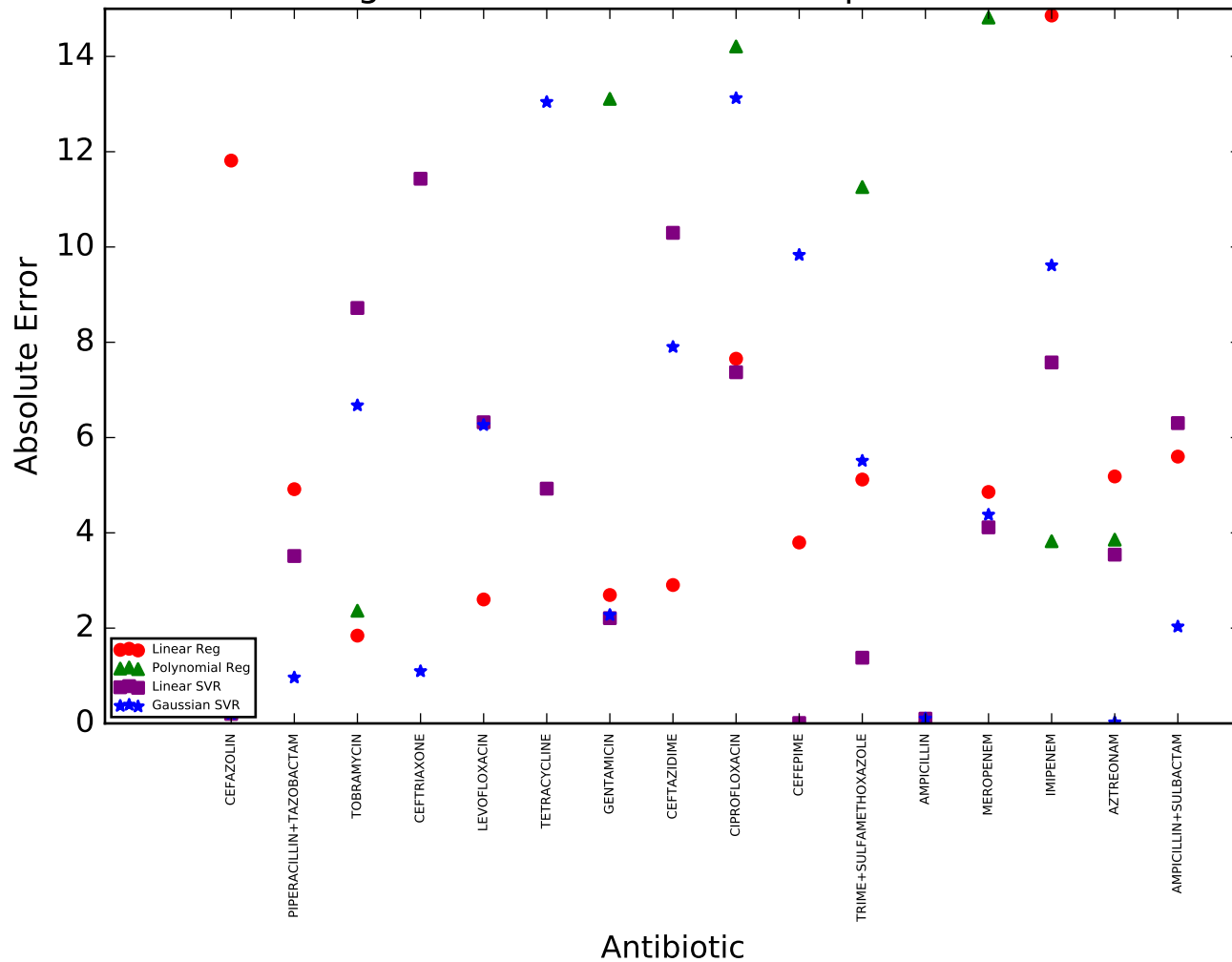


# Enterobacter cloacae

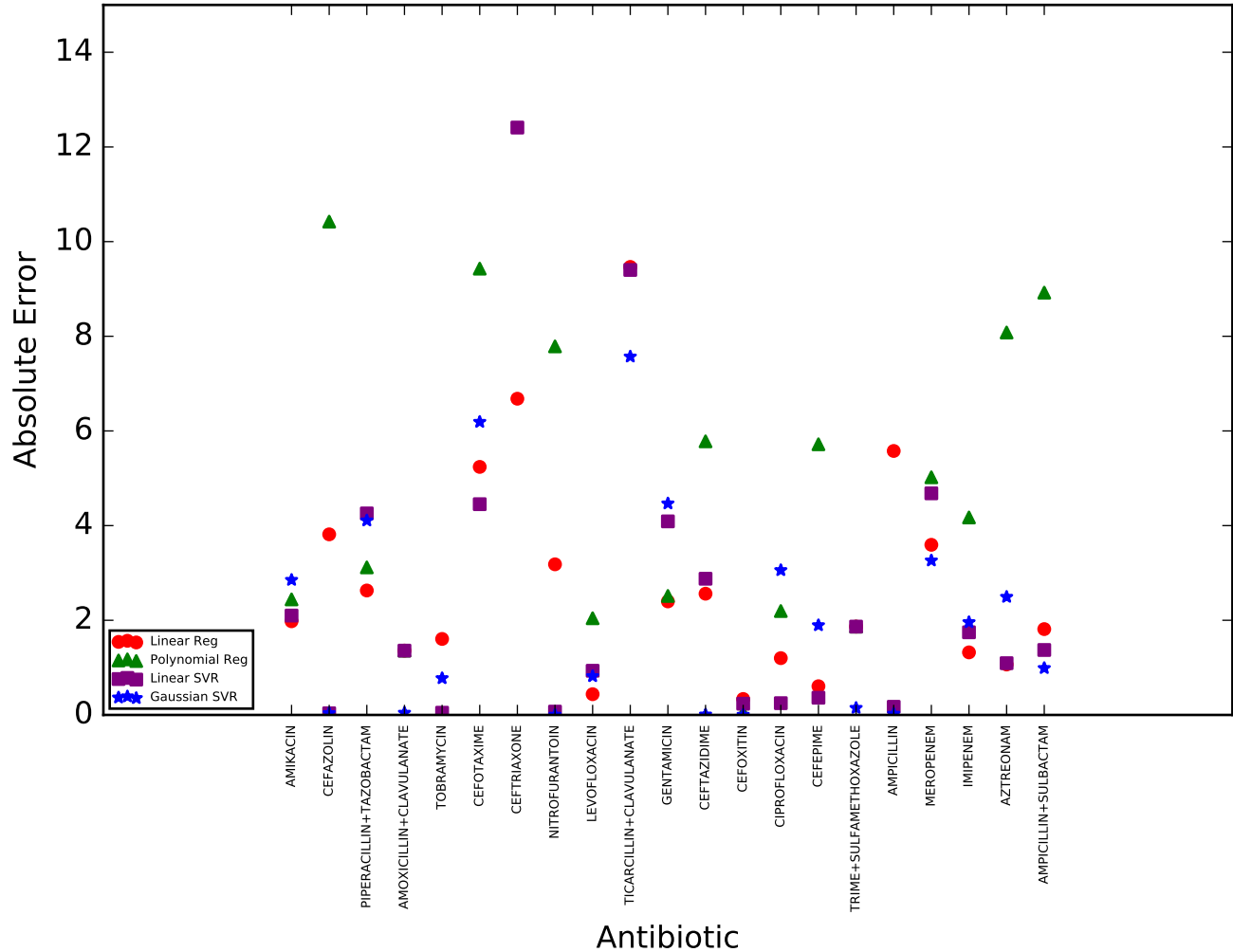
## using data from 2004-2012 to predict 2014



# Acinetobacter baumannii using data from 2004-2012 to predict 2014

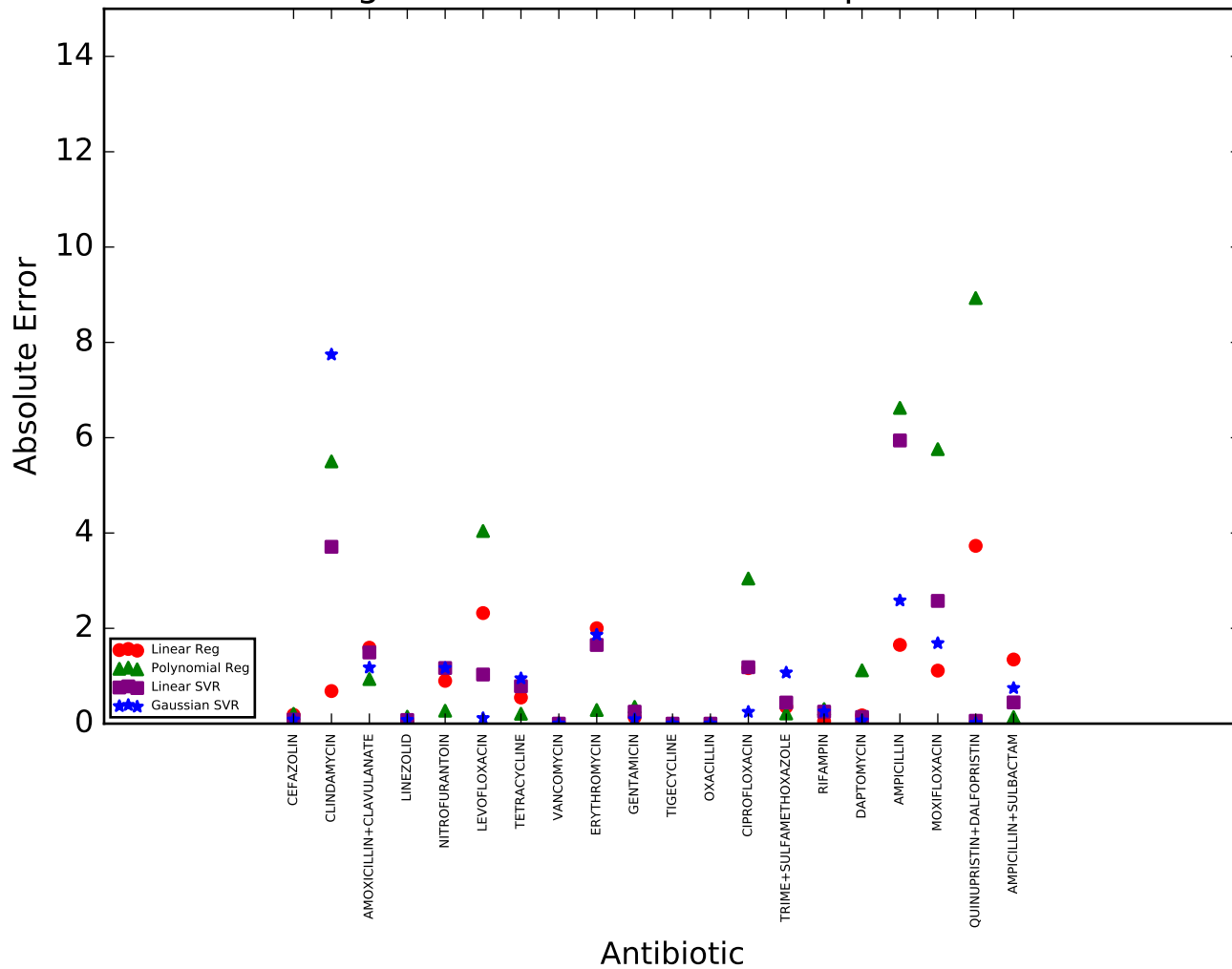


# *Pseudomonas aeruginosa* using data from 2004-2012 to predict 2014



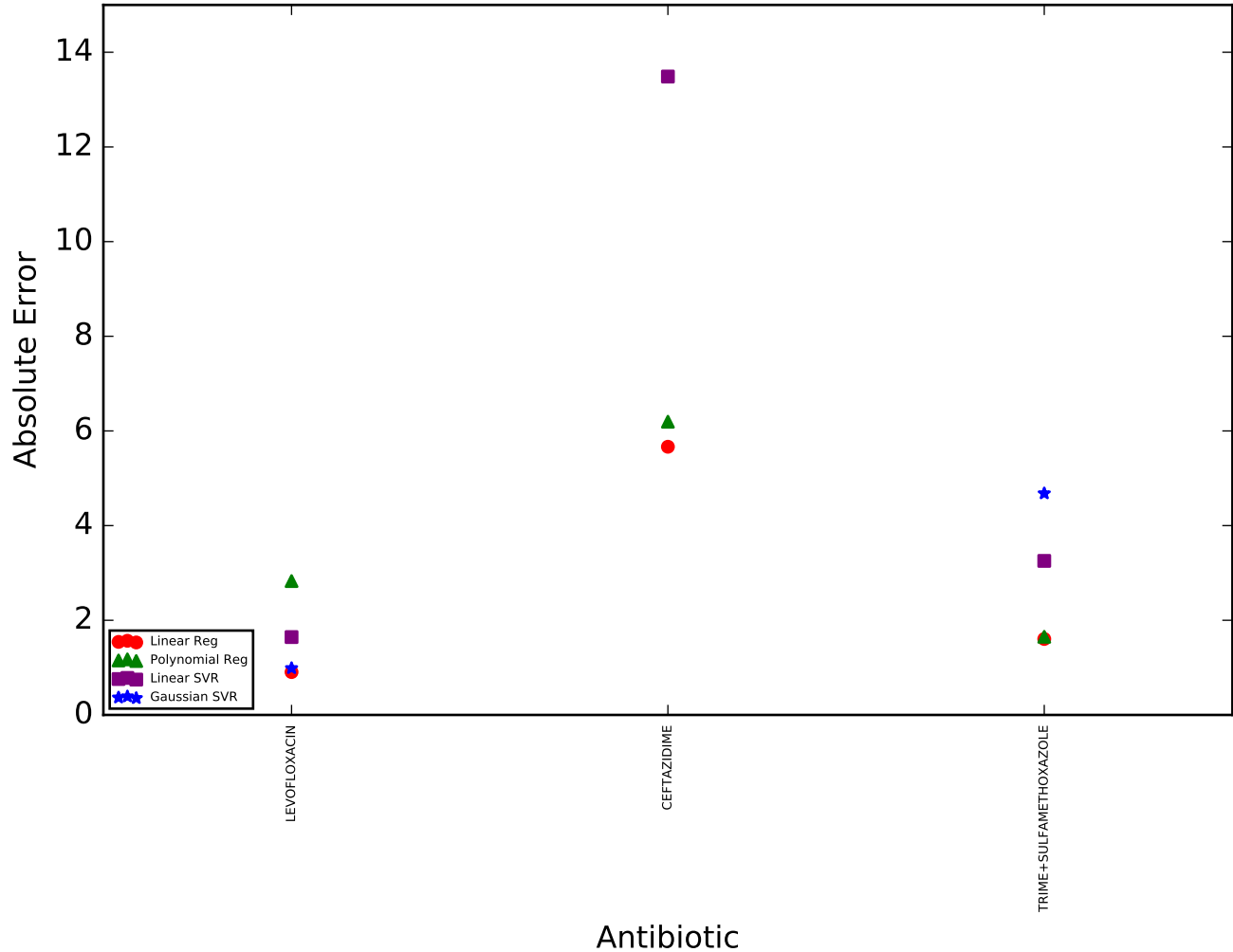
# MSSA

using data from 2004-2012 to predict 2014



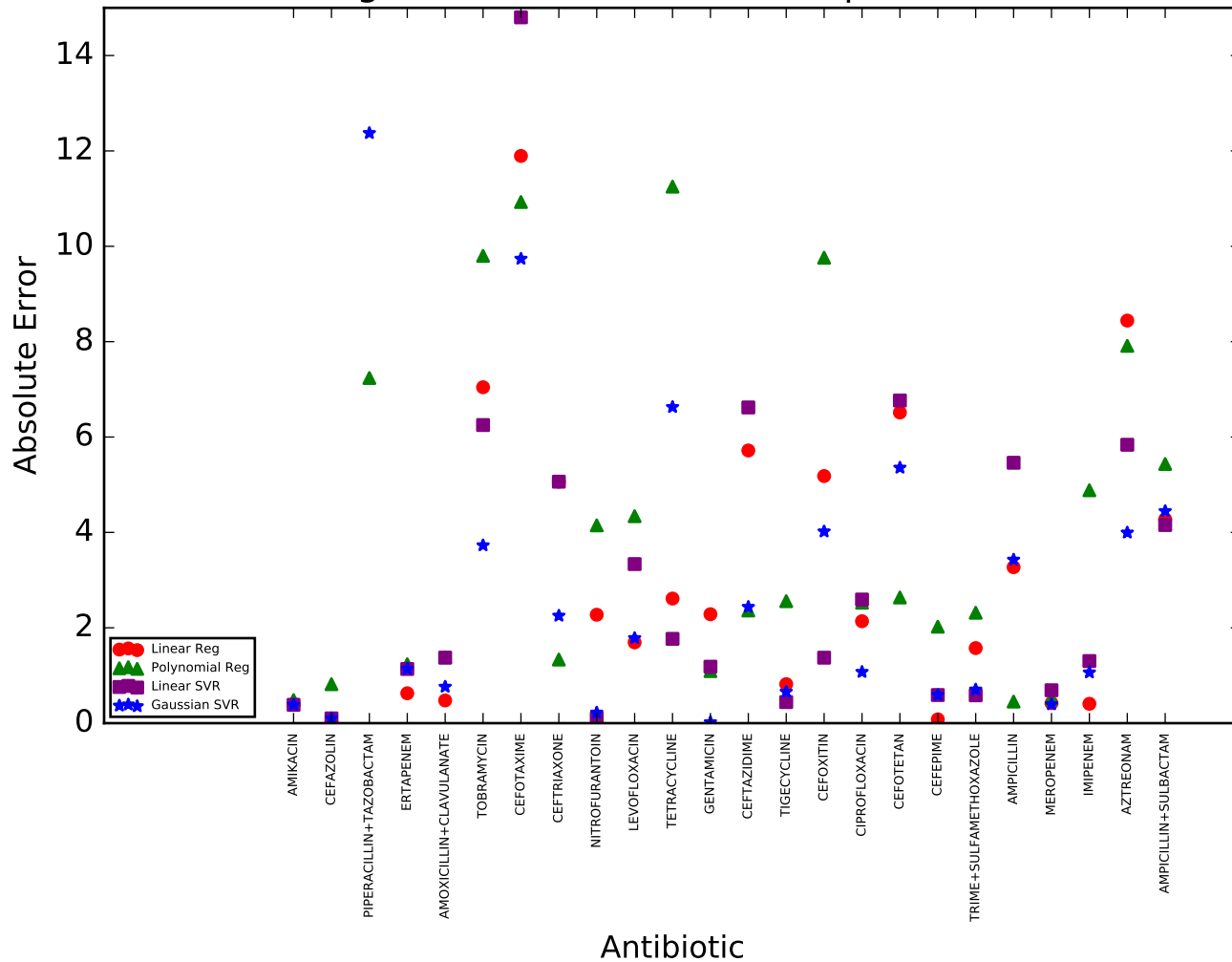
# Stenotrophomonas maltophilia

using data from 2003-2011 to predict 2014



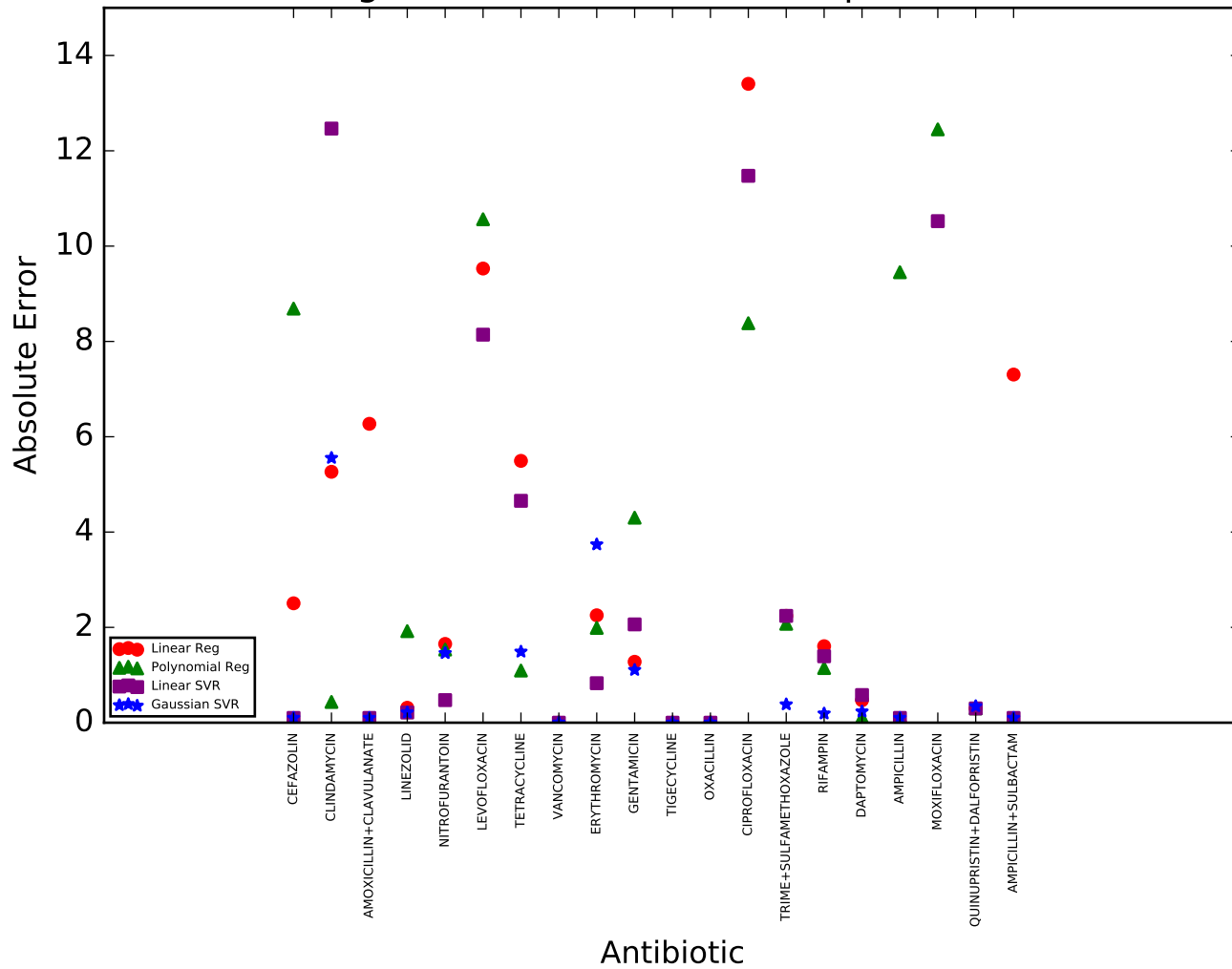
# Serratia marcescens

## using data from 2003-2011 to predict 2014



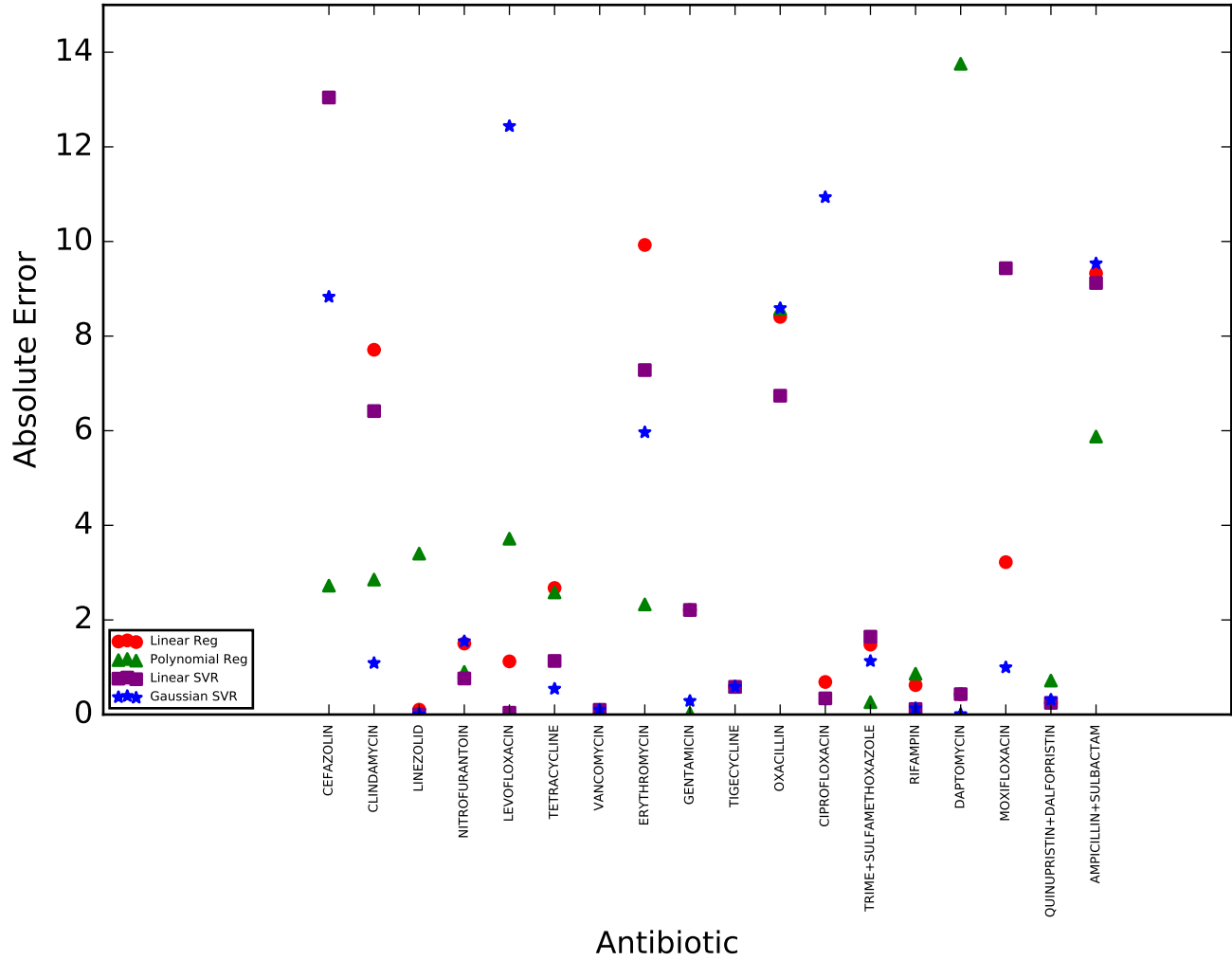
# MRSA

## using data from 2003-2011 to predict 2014



# Staphylococcus aureus

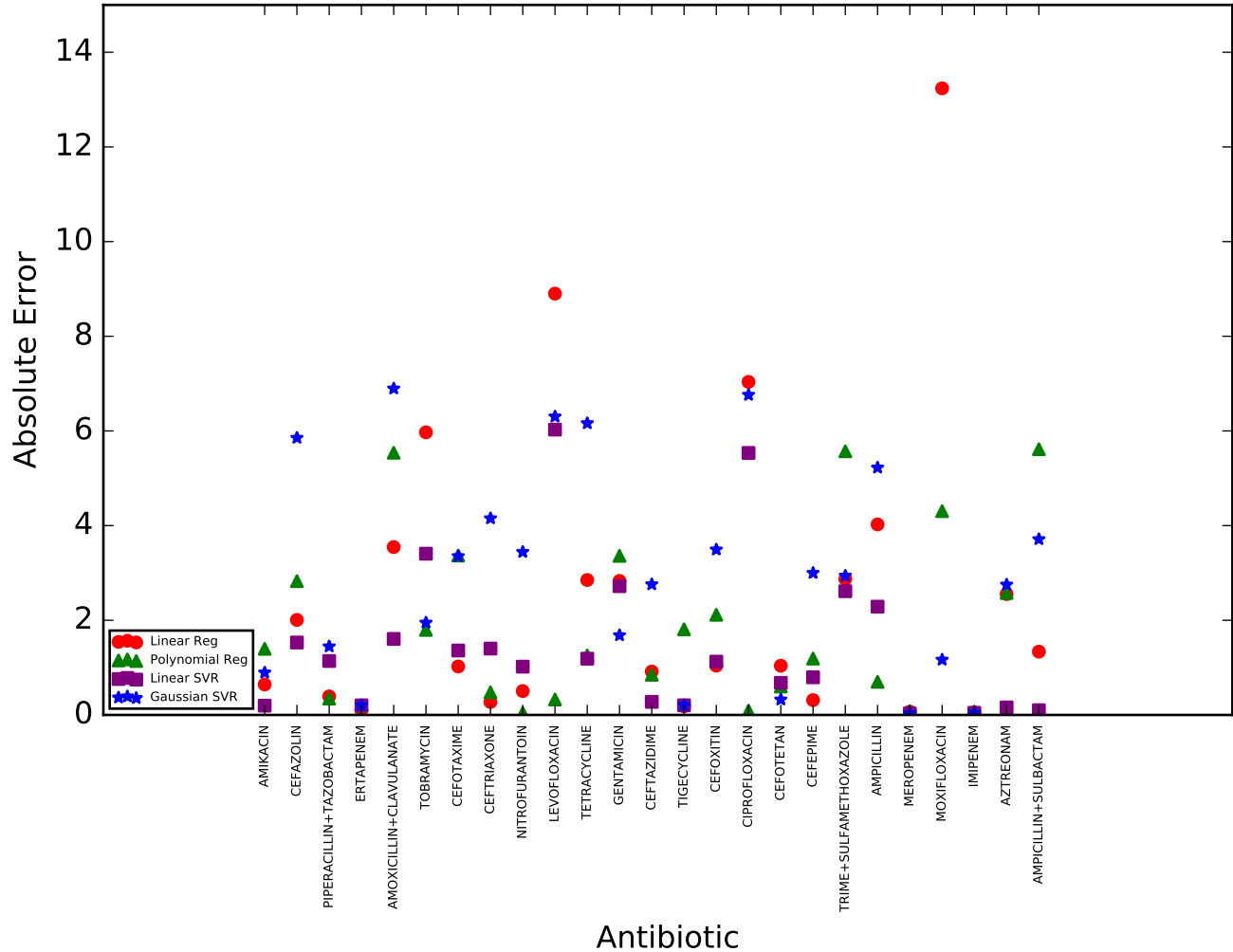
## using data from 2003-2011 to predict 2014





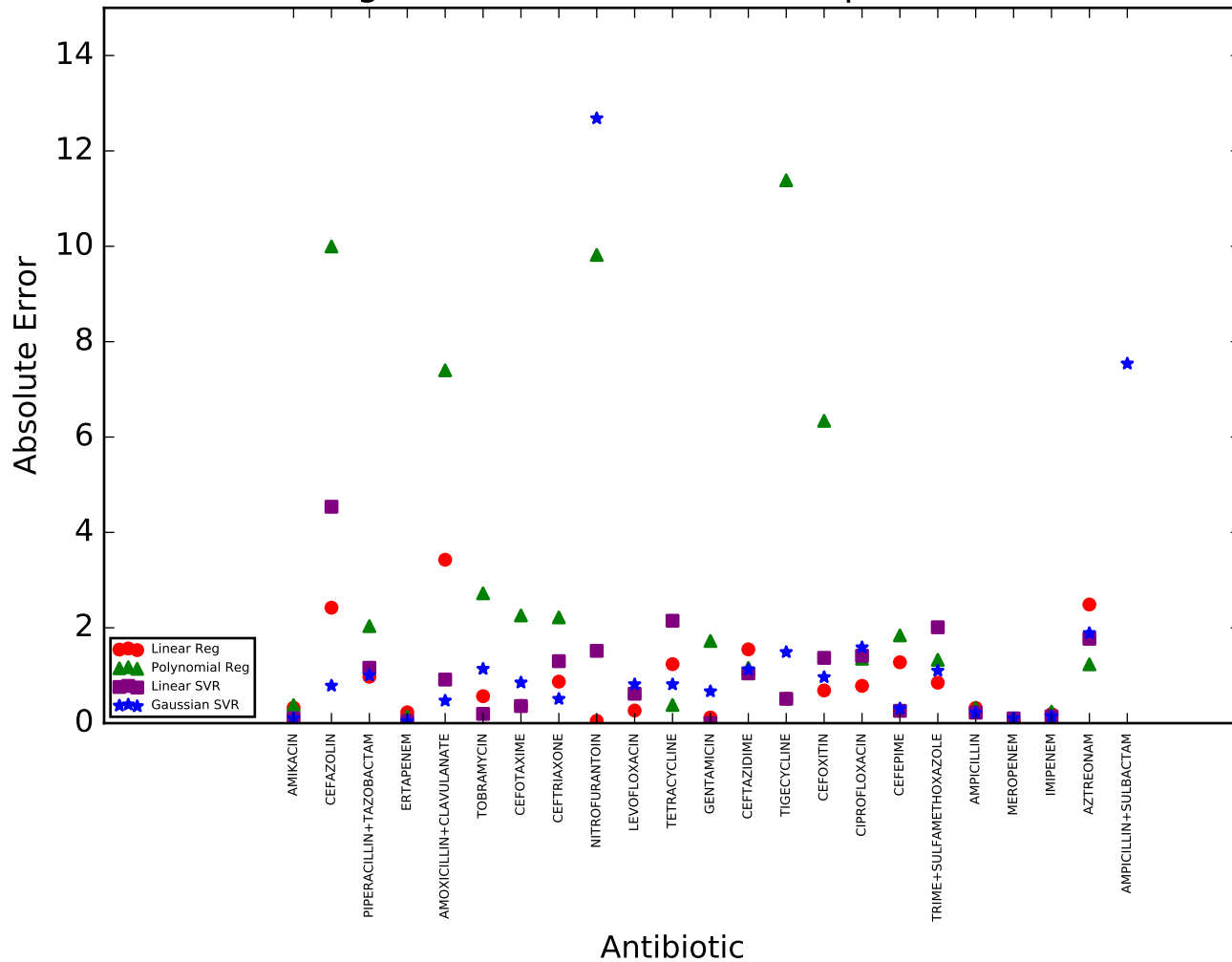
# E. coli

## using data from 2003-2011 to predict 2014



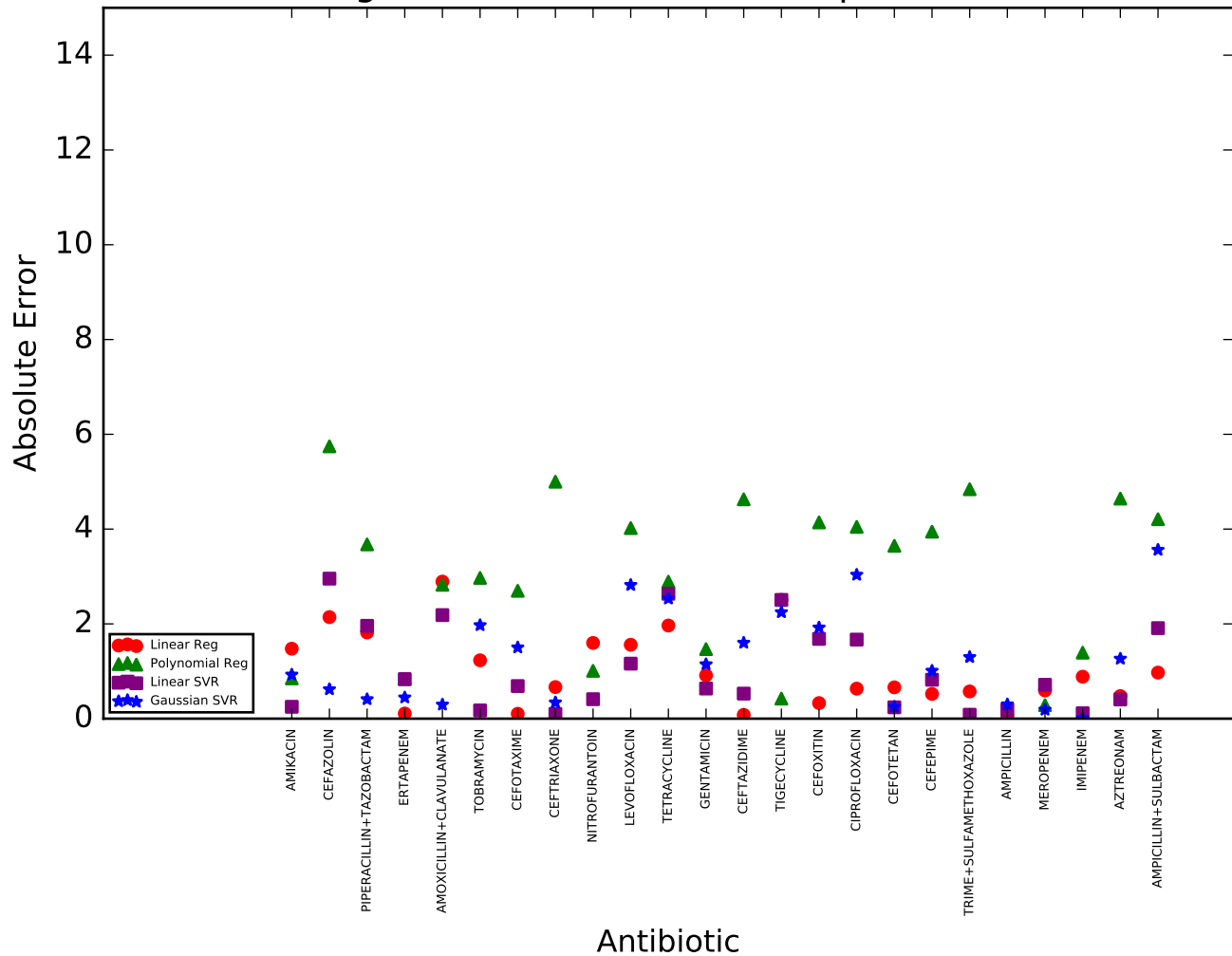
# Klebsiella oxytoca

## using data from 2003-2011 to predict 2014



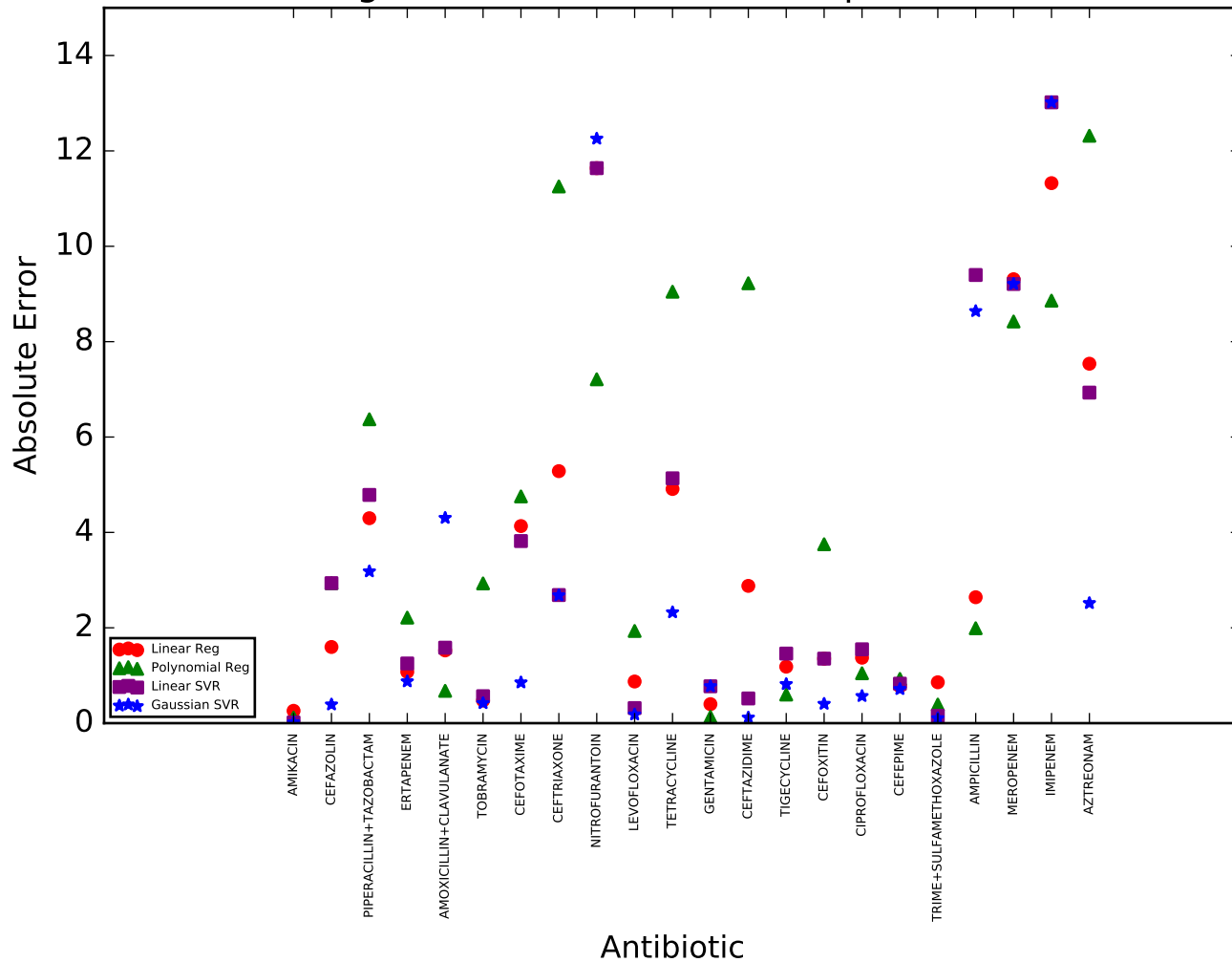
# Klebsiella pneumoniae

## using data from 2003-2011 to predict 2014



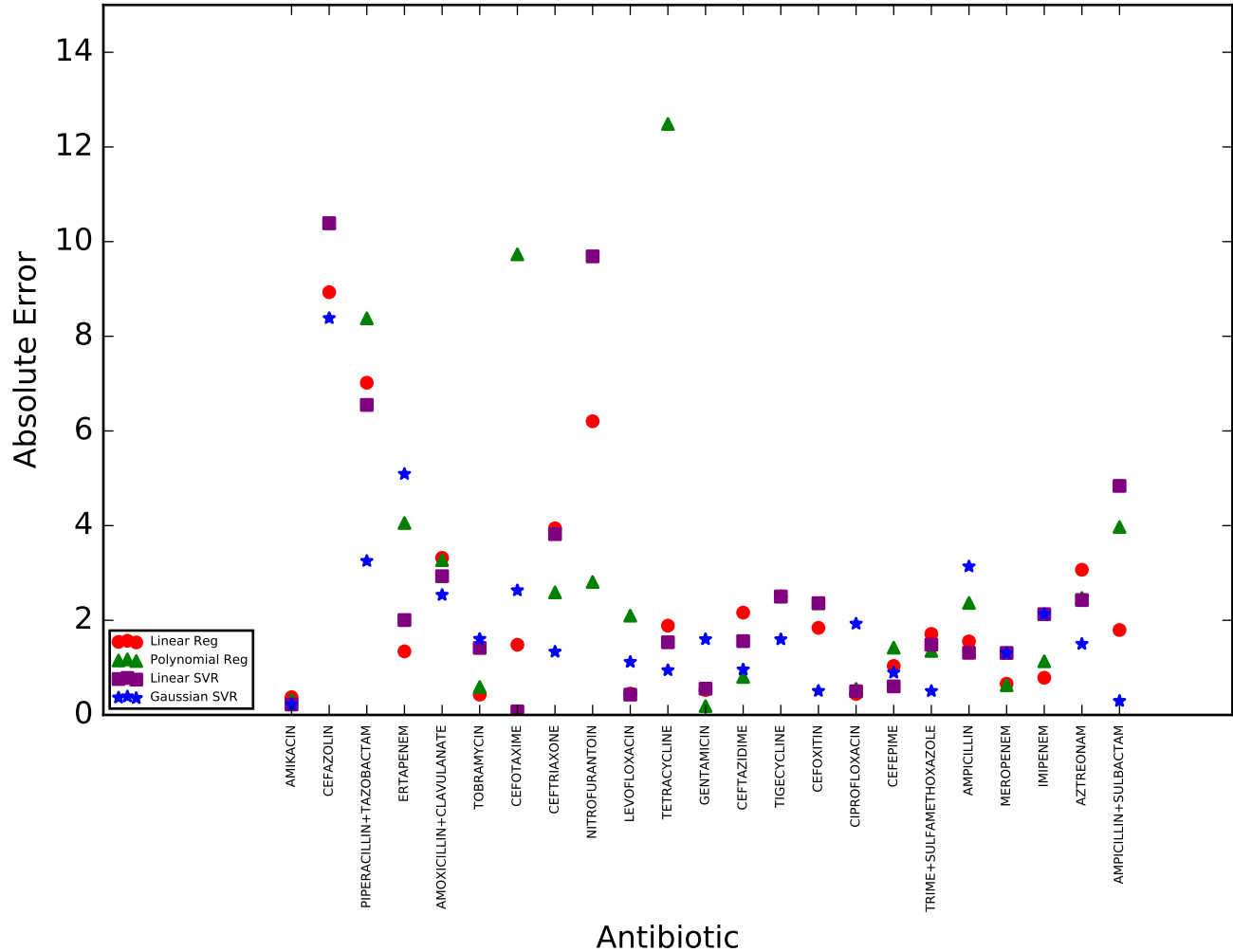
# Enterobacter aerogenes

## using data from 2003-2011 to predict 2014

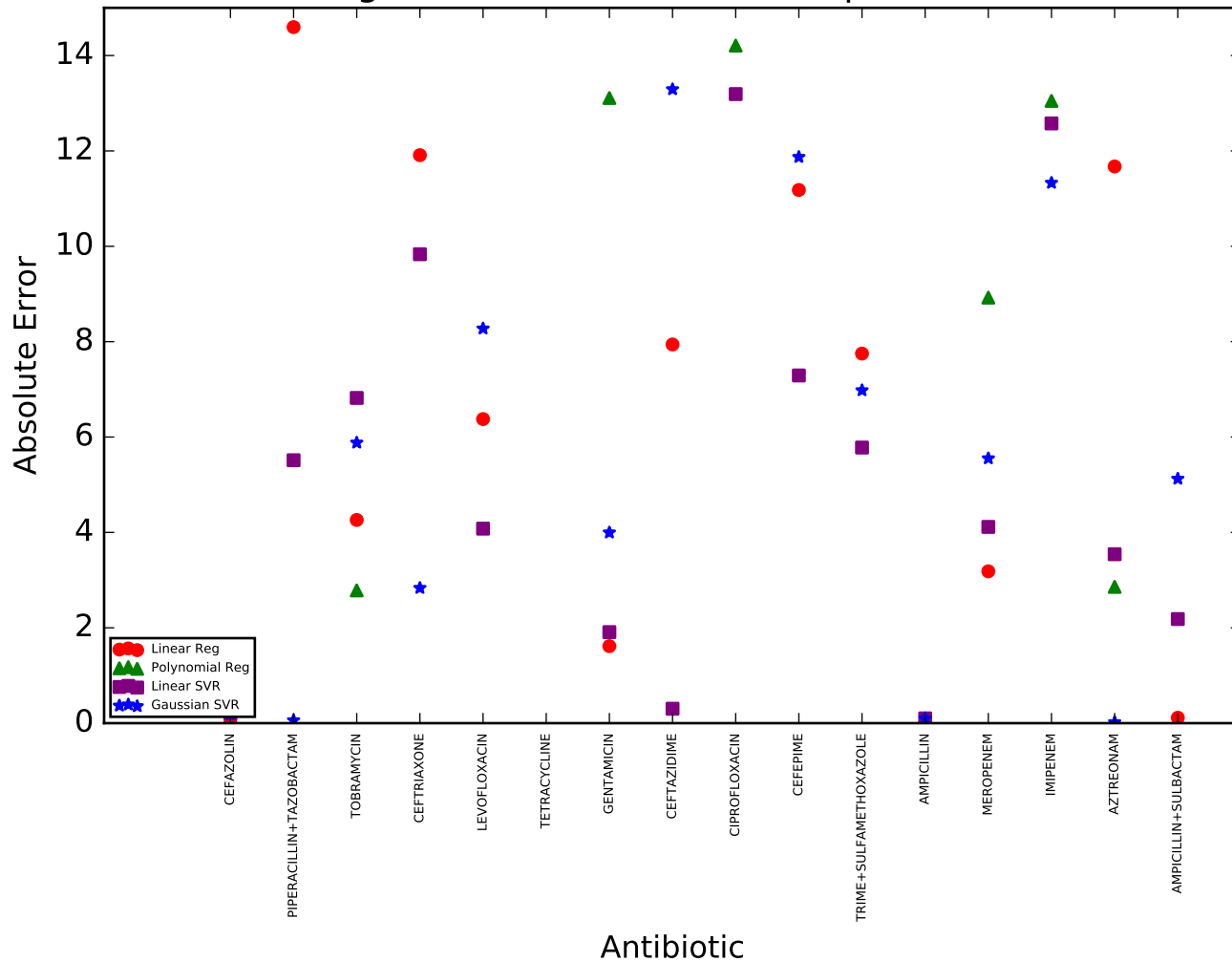


# Enterobacter cloacae

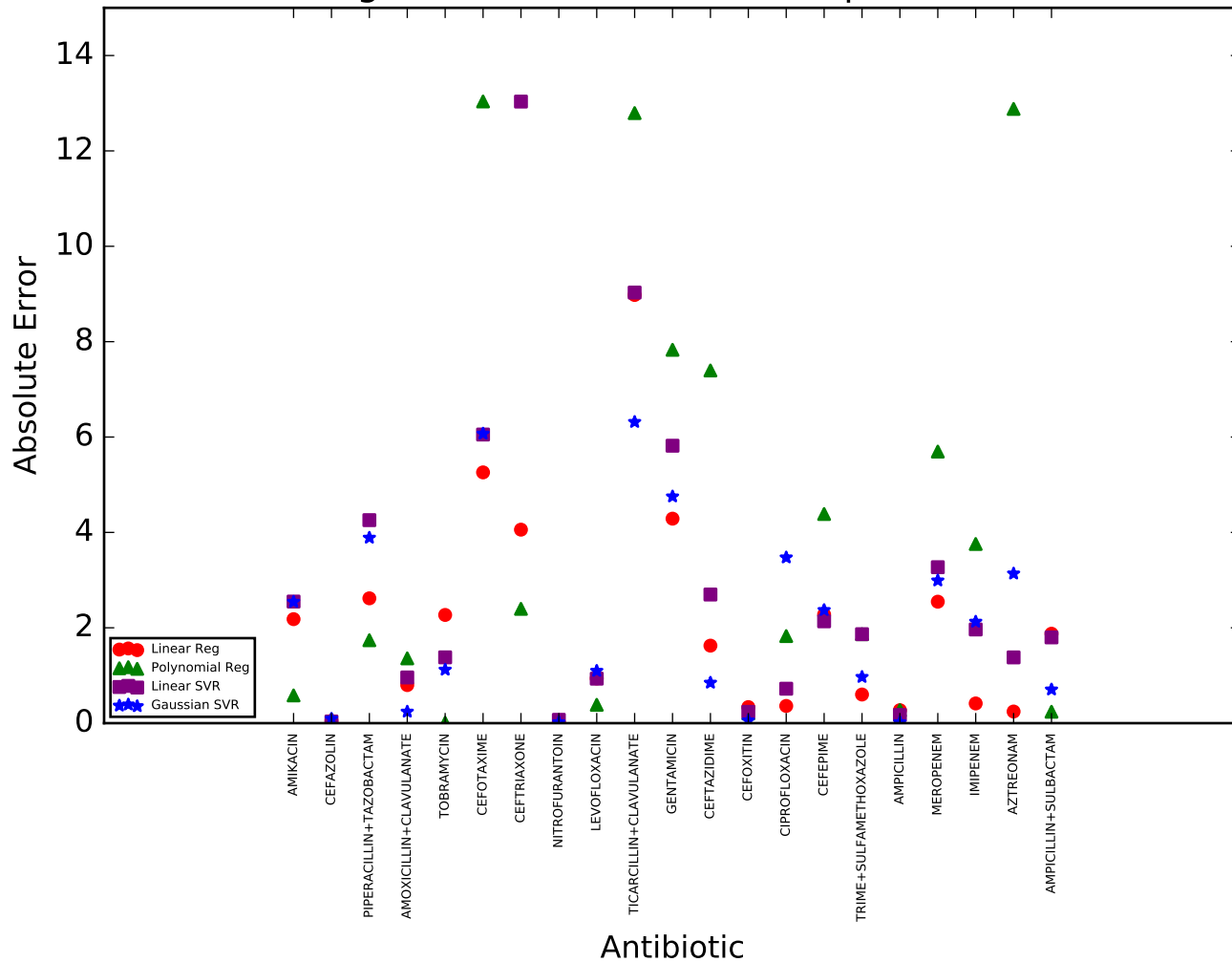
## using data from 2003-2011 to predict 2014



# Acinetobacter baumannii using data from 2003-2011 to predict 2014

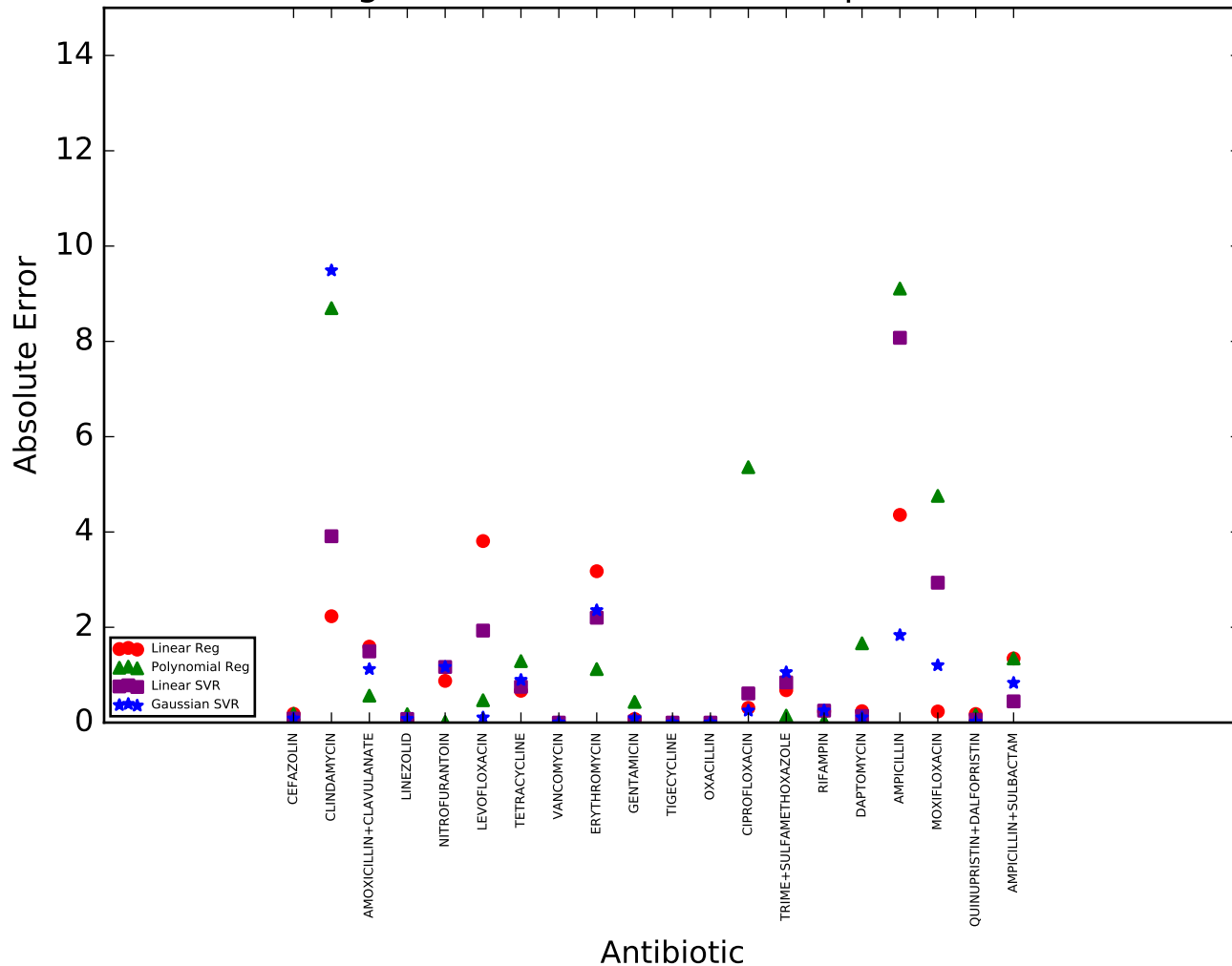


# *Pseudomonas aeruginosa* using data from 2003-2011 to predict 2014



# MSSA

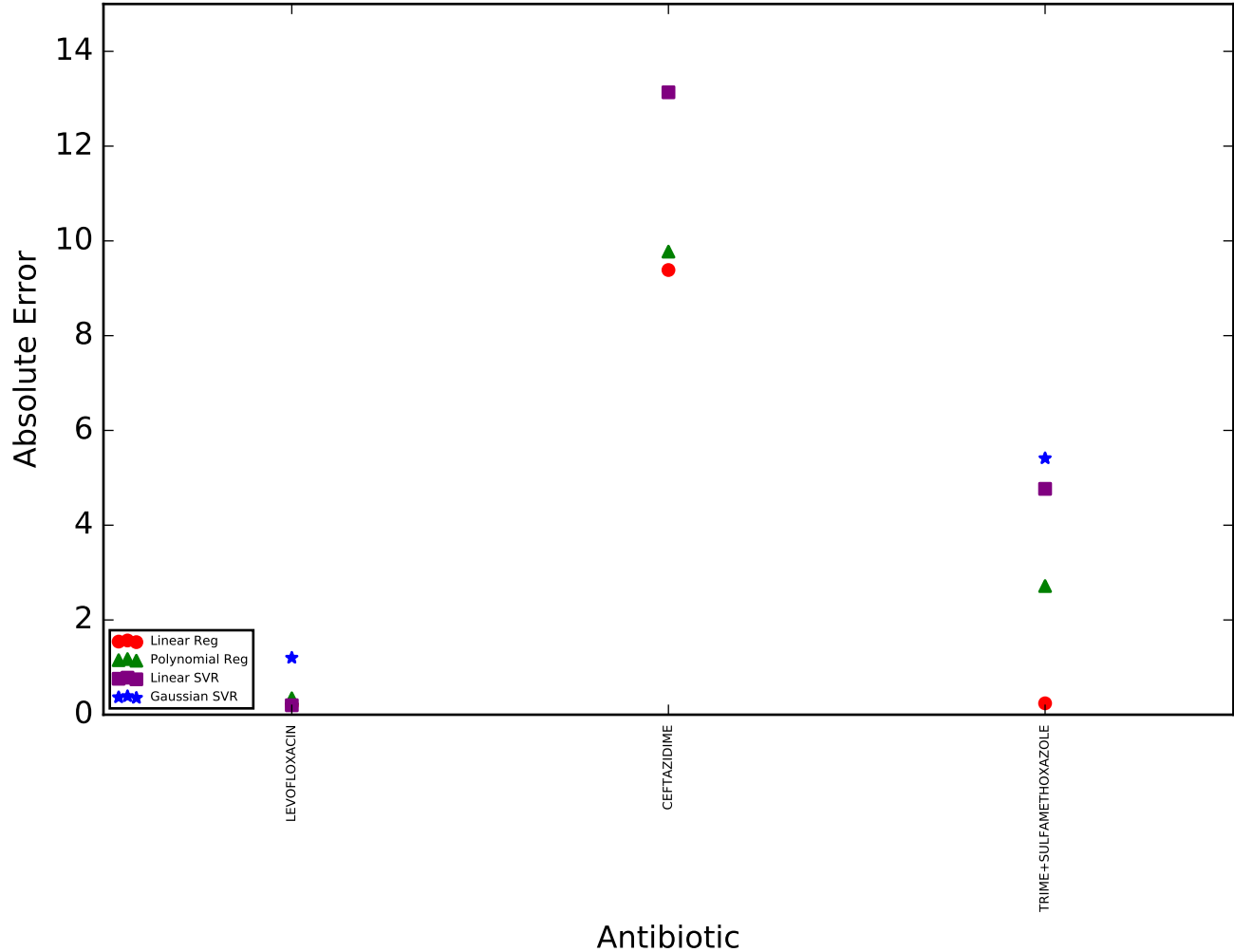
## using data from 2003-2011 to predict 2014





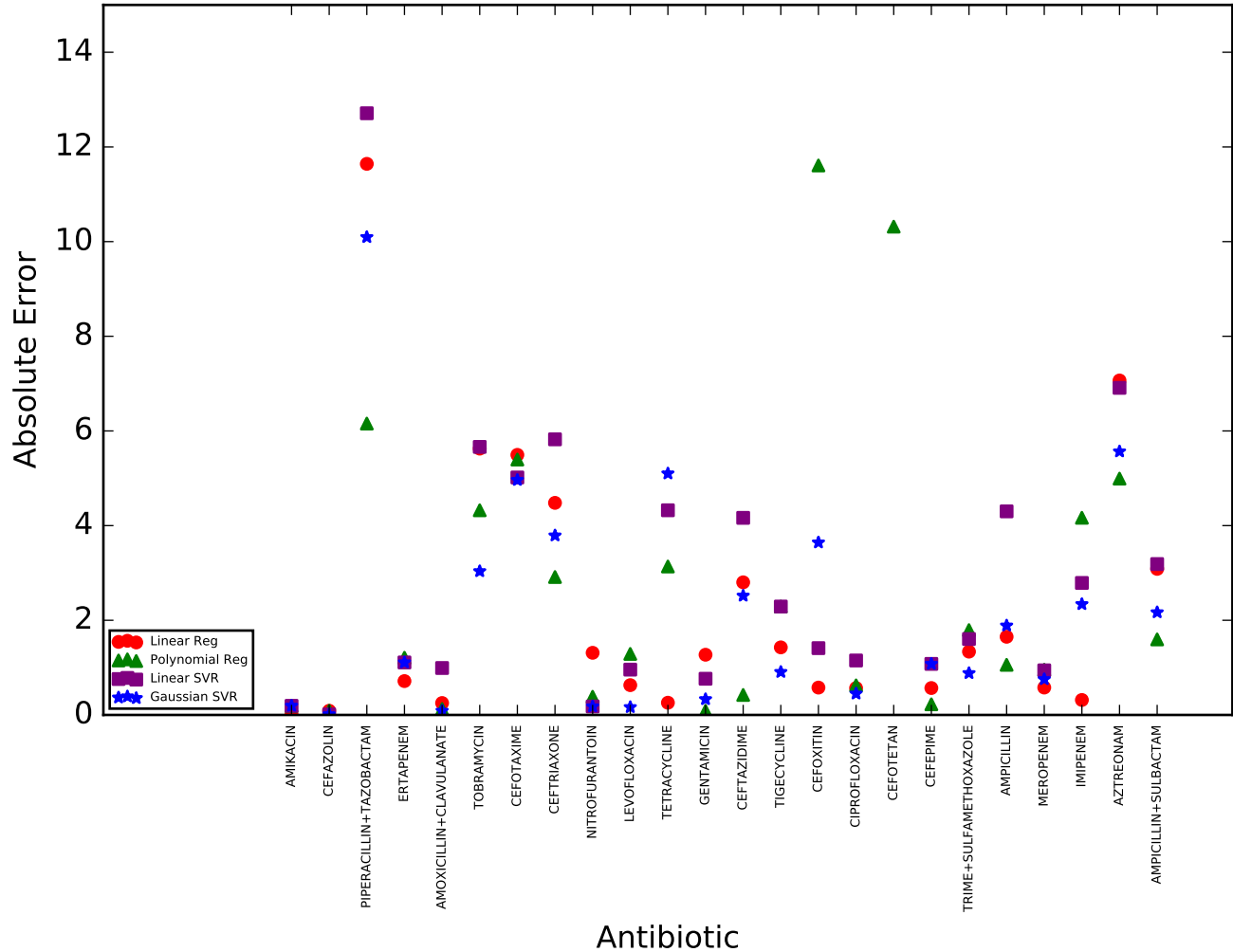
# Stenotrophomonas maltophilia

using data from 2004-2012 to predict 2013



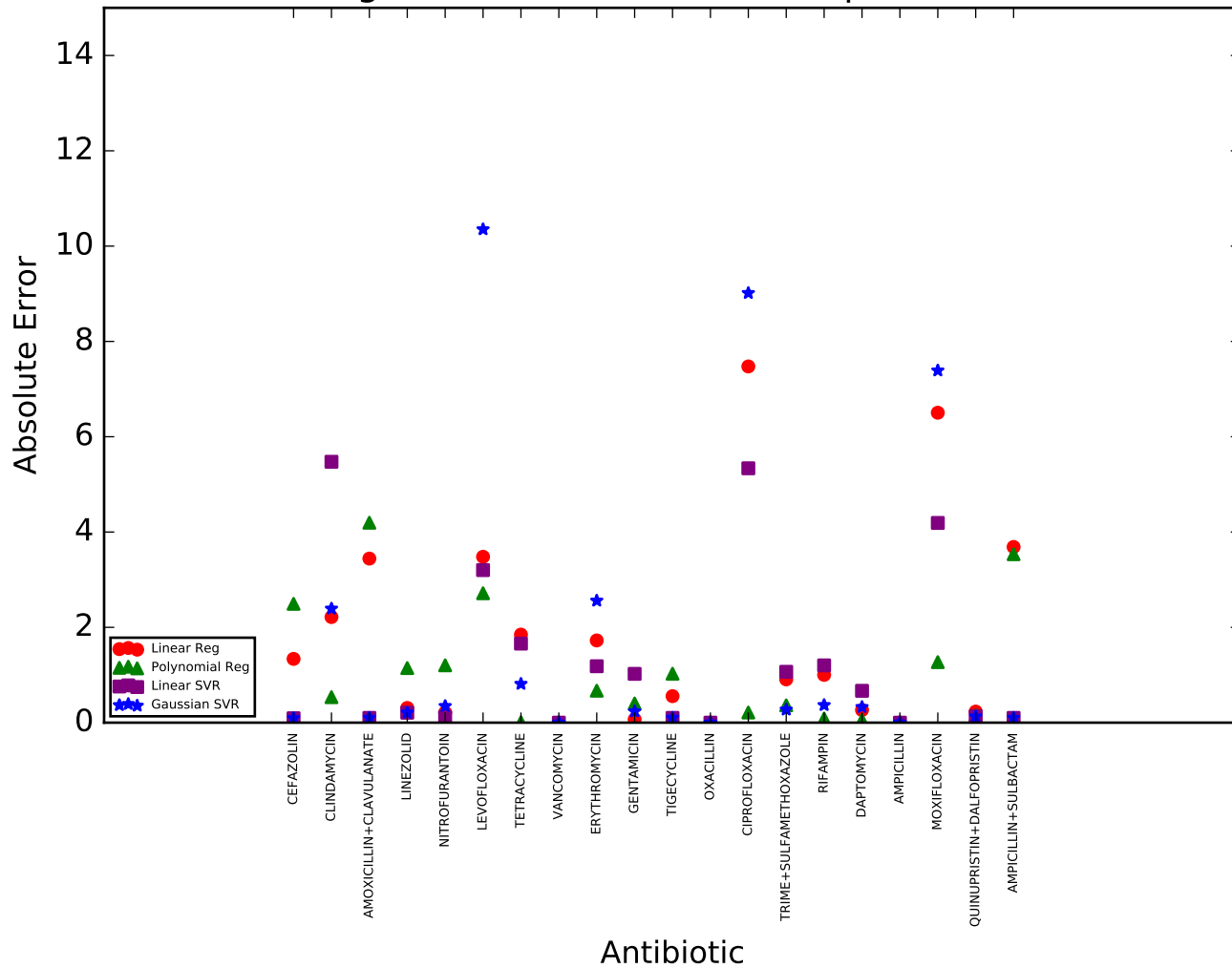
# Serratia marcescens

## using data from 2004-2012 to predict 2013



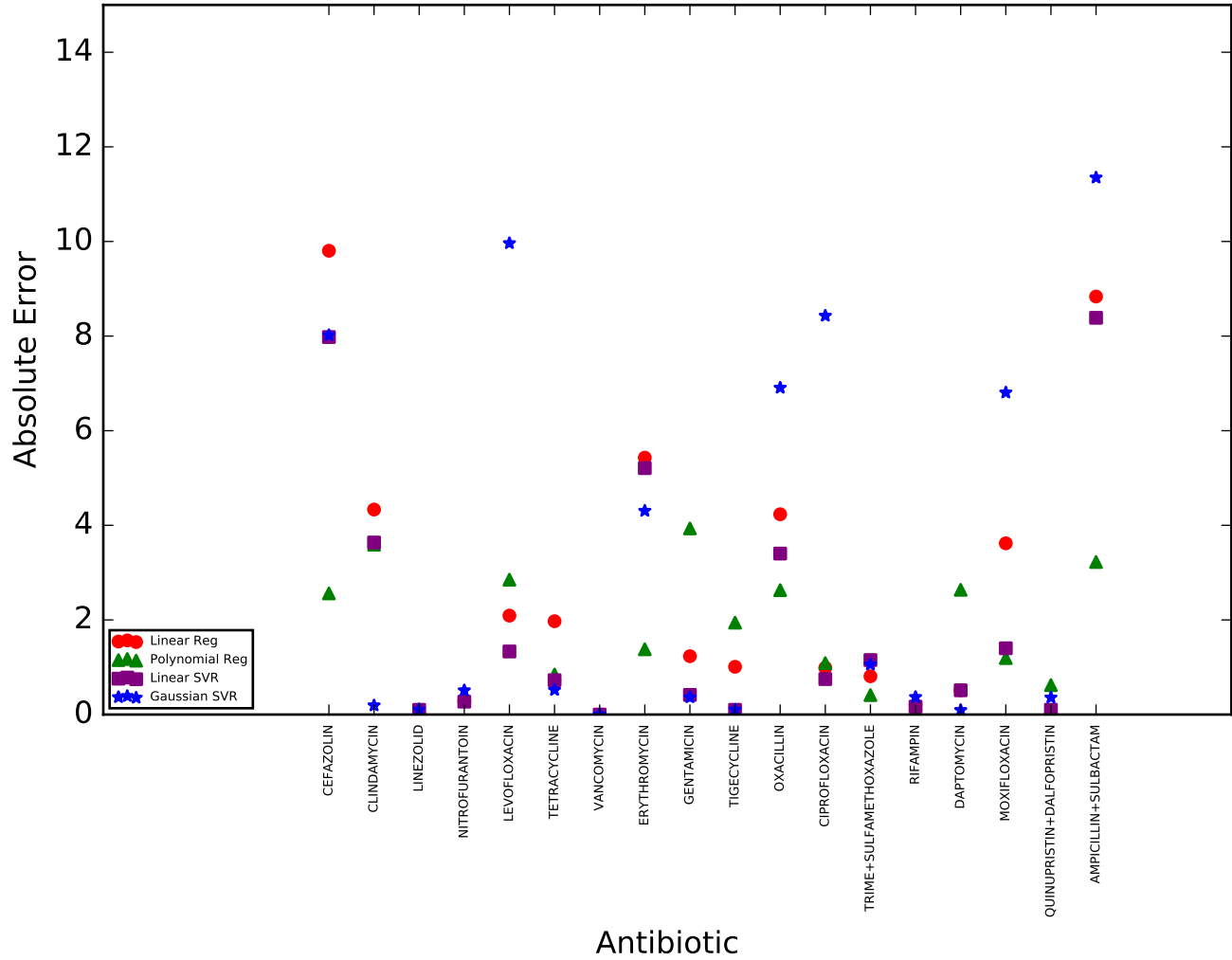
# MRSA

## using data from 2004-2012 to predict 2013



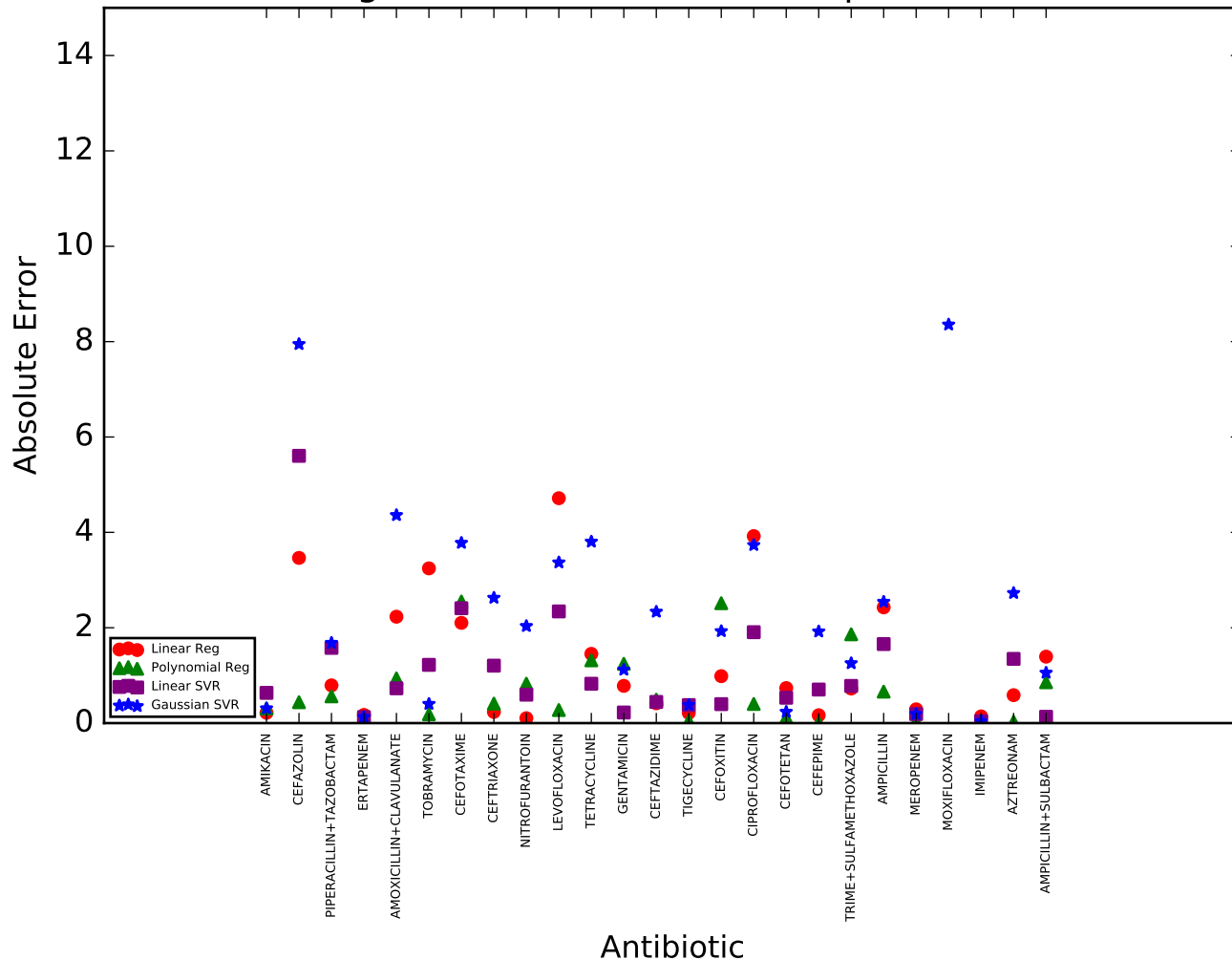
# Staphylococcus aureus

## using data from 2004-2012 to predict 2013



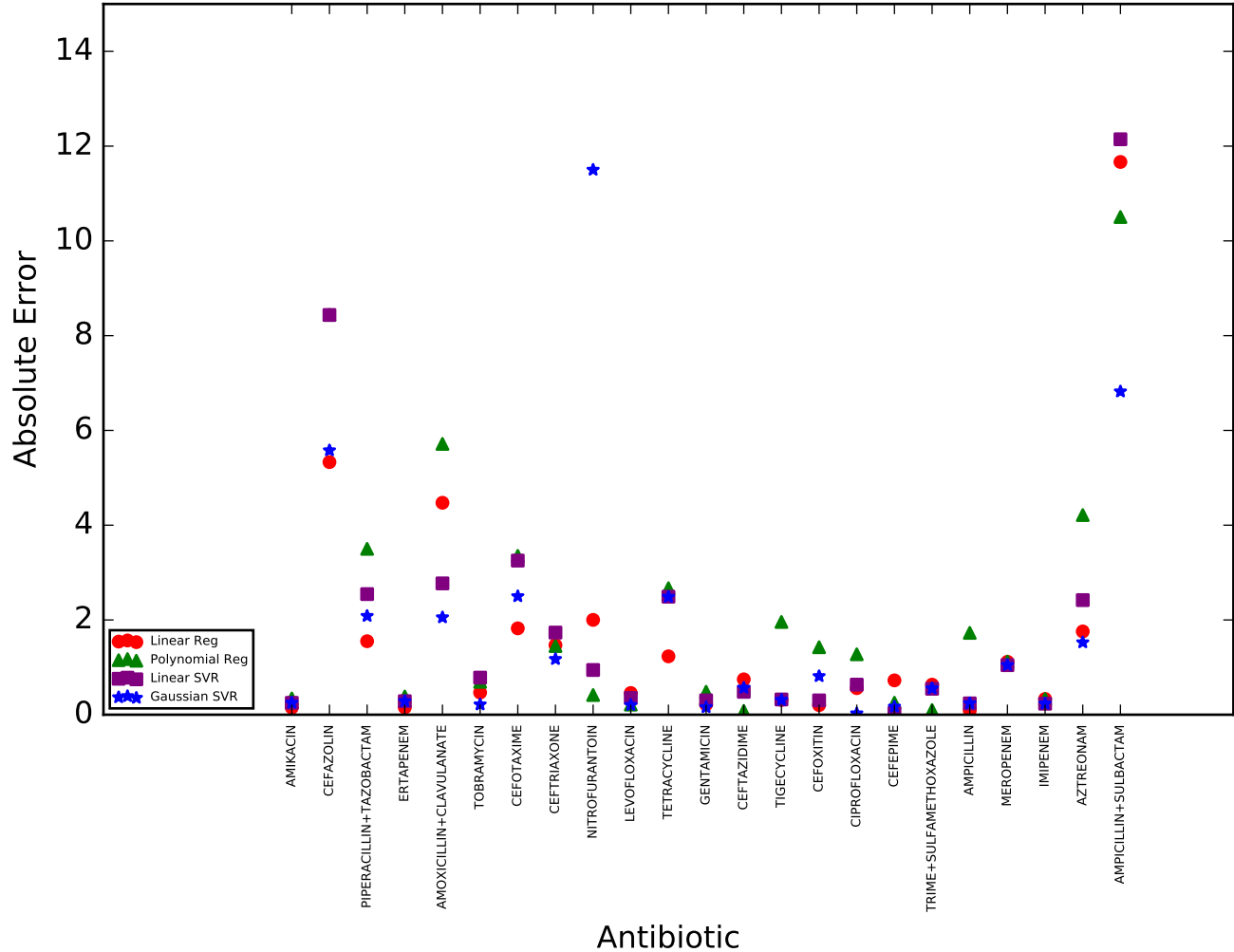
# E. coli

## using data from 2004-2012 to predict 2013



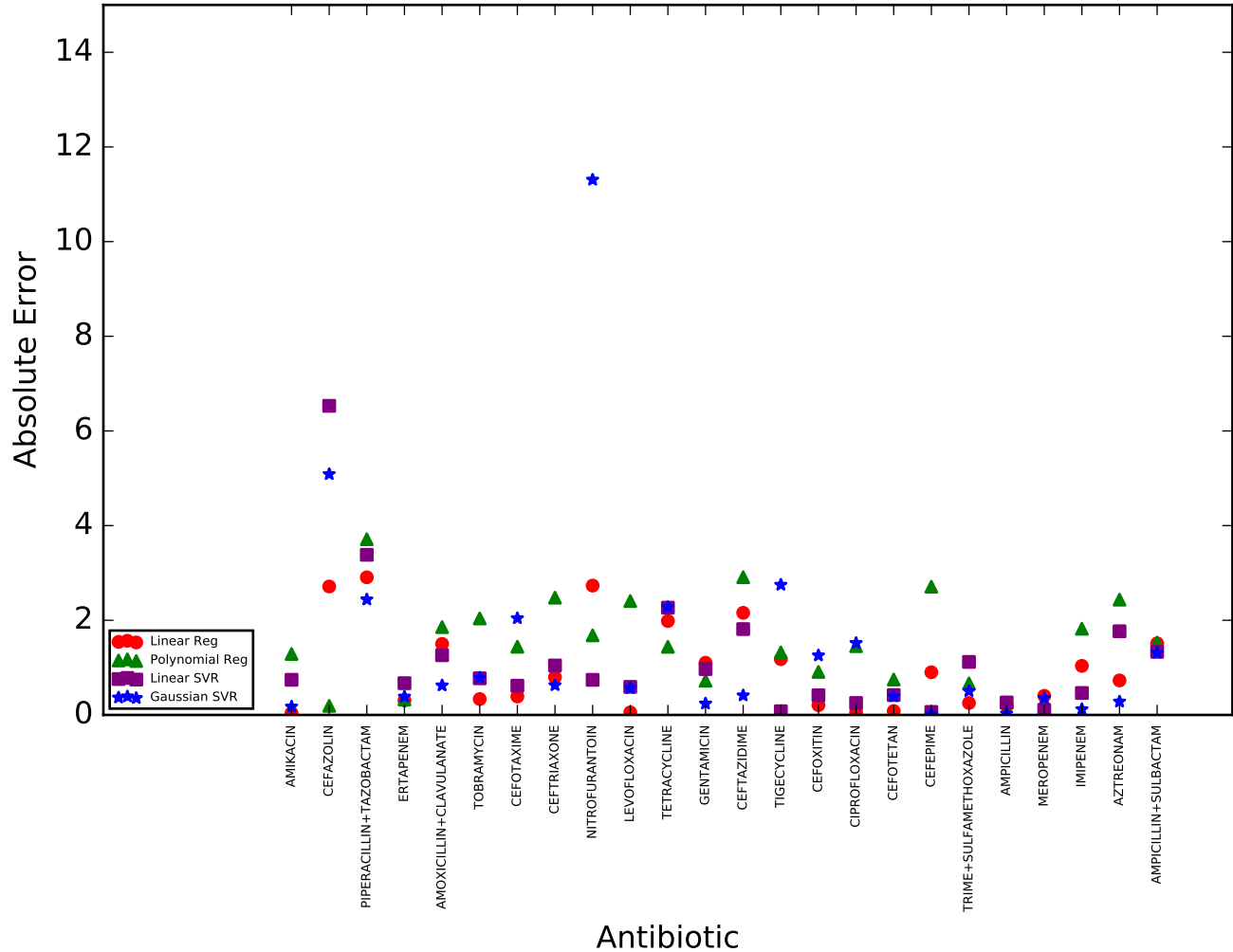
# Klebsiella oxytoca

## using data from 2004-2012 to predict 2013



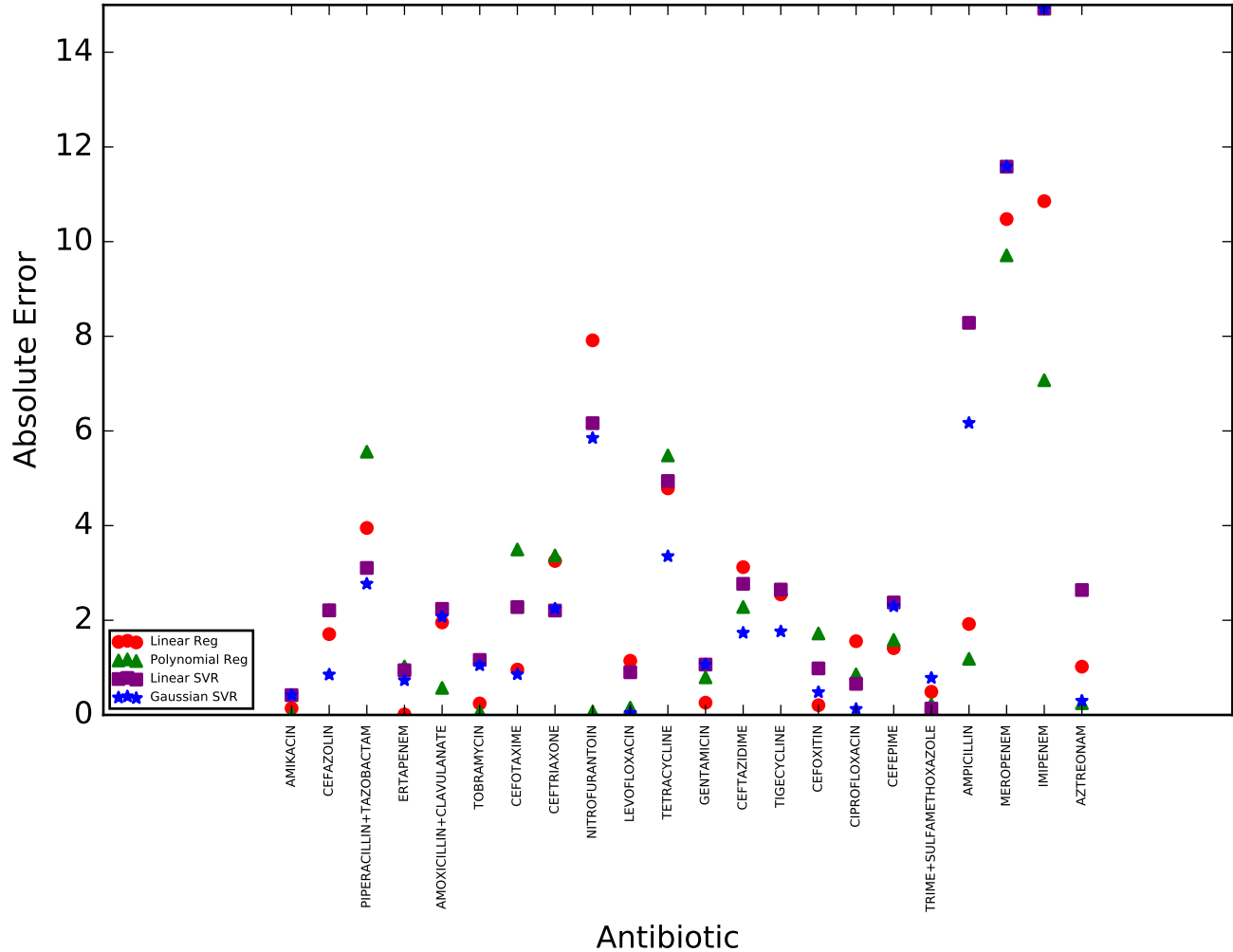
# Klebsiella pneumoniae

## using data from 2004-2012 to predict 2013



# Enterobacter aerogenes

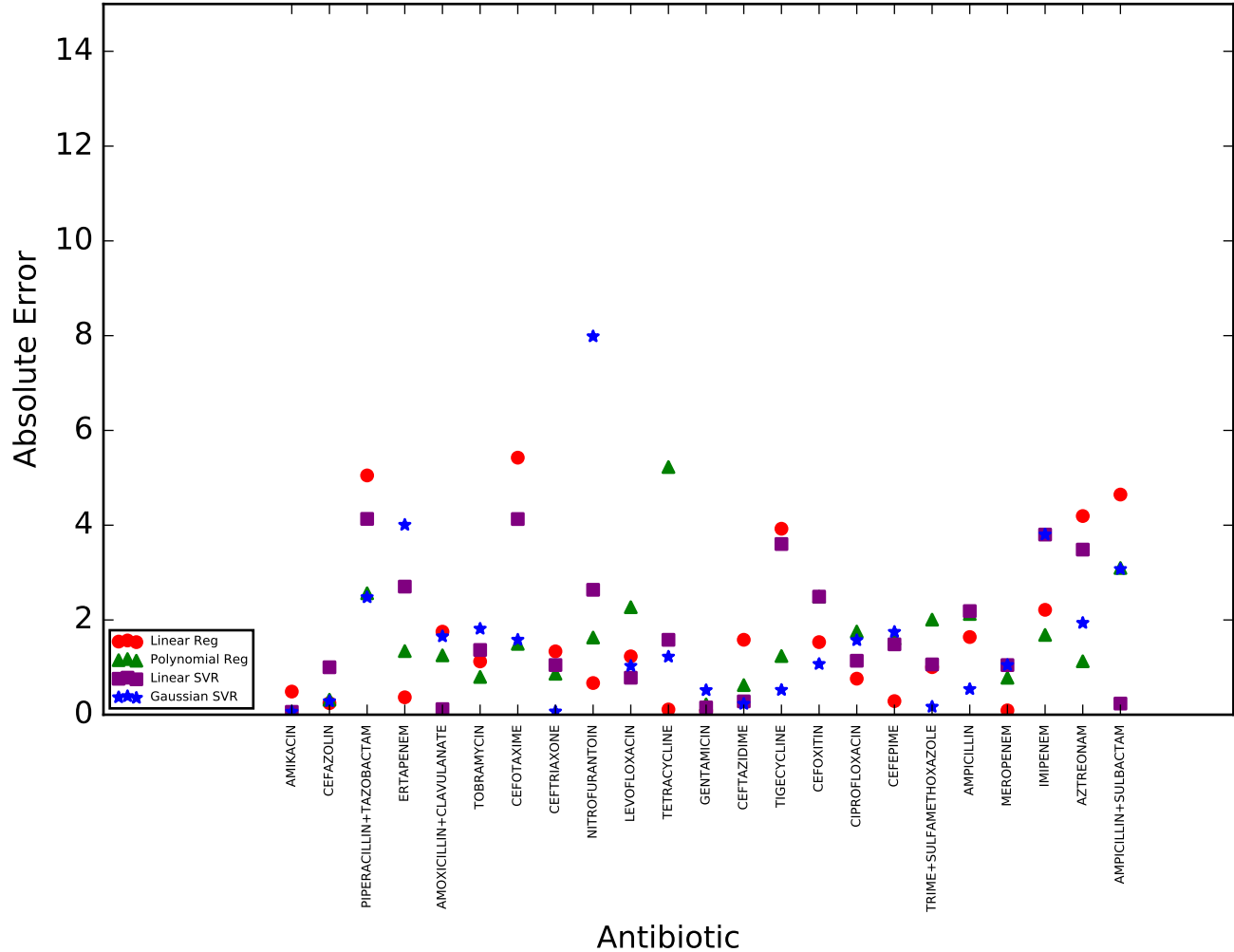
## using data from 2004-2012 to predict 2013



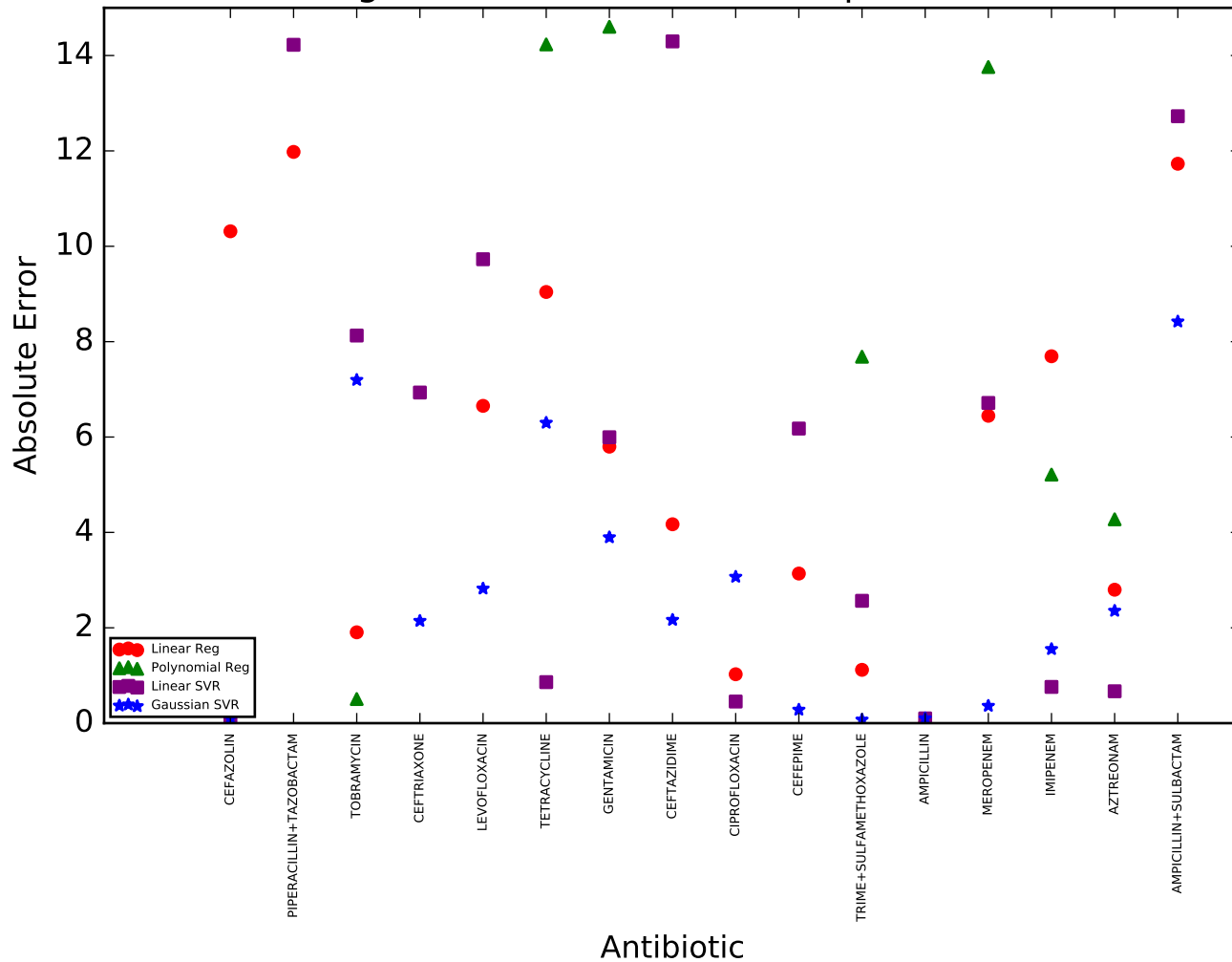


# Enterobacter cloacae

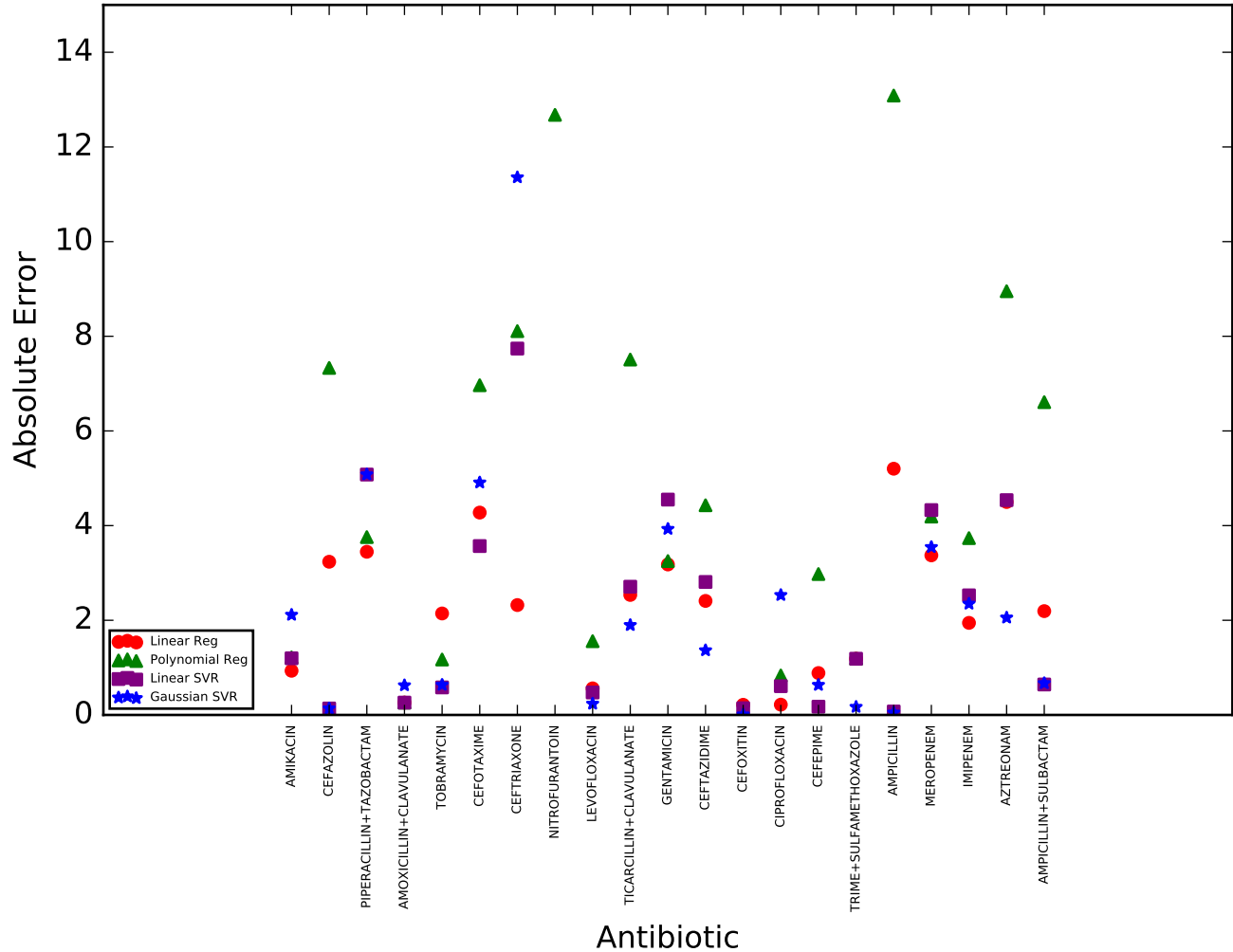
## using data from 2004-2012 to predict 2013



# Acinetobacter baumannii using data from 2004-2012 to predict 2013

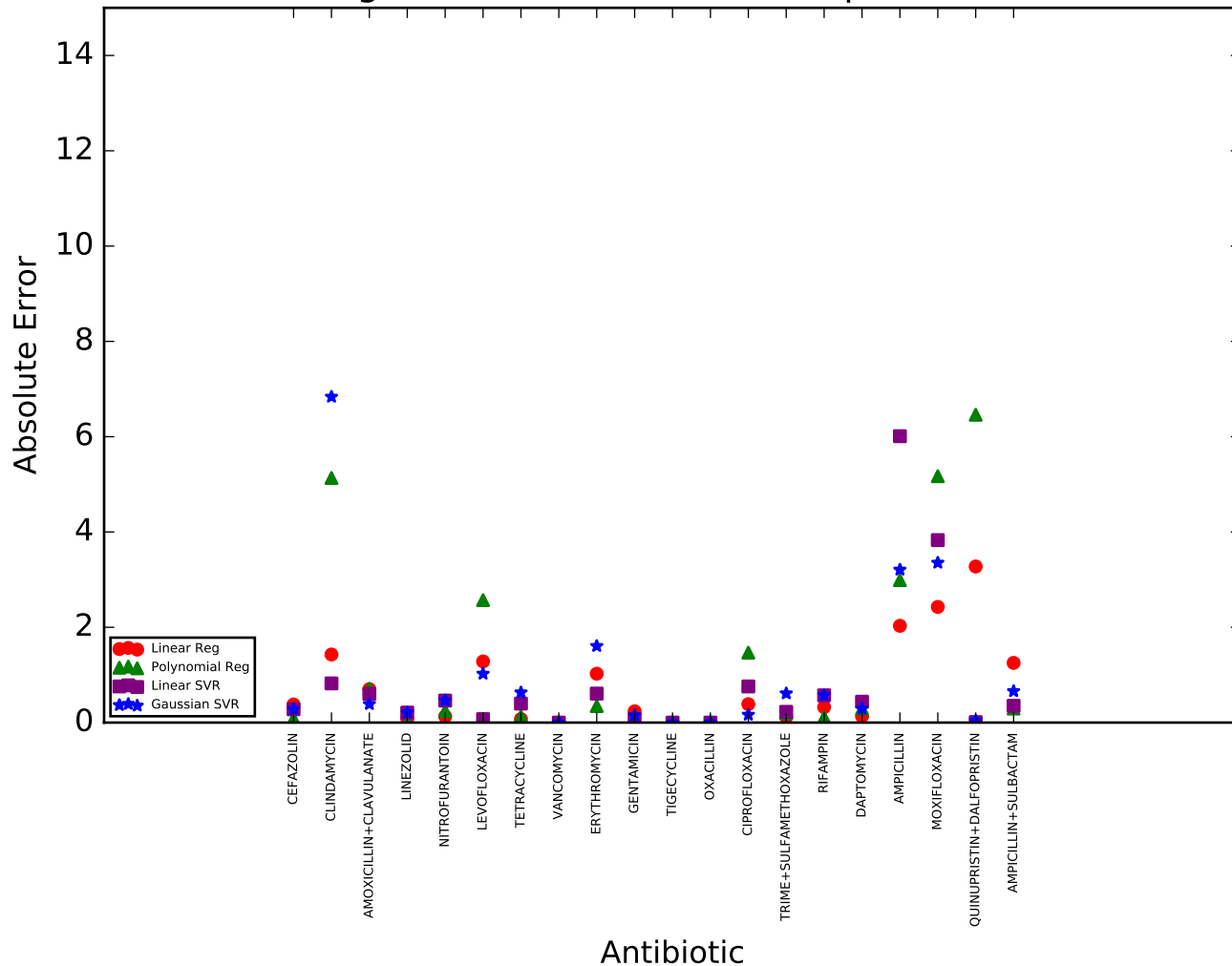


# *Pseudomonas aeruginosa* using data from 2004-2012 to predict 2013



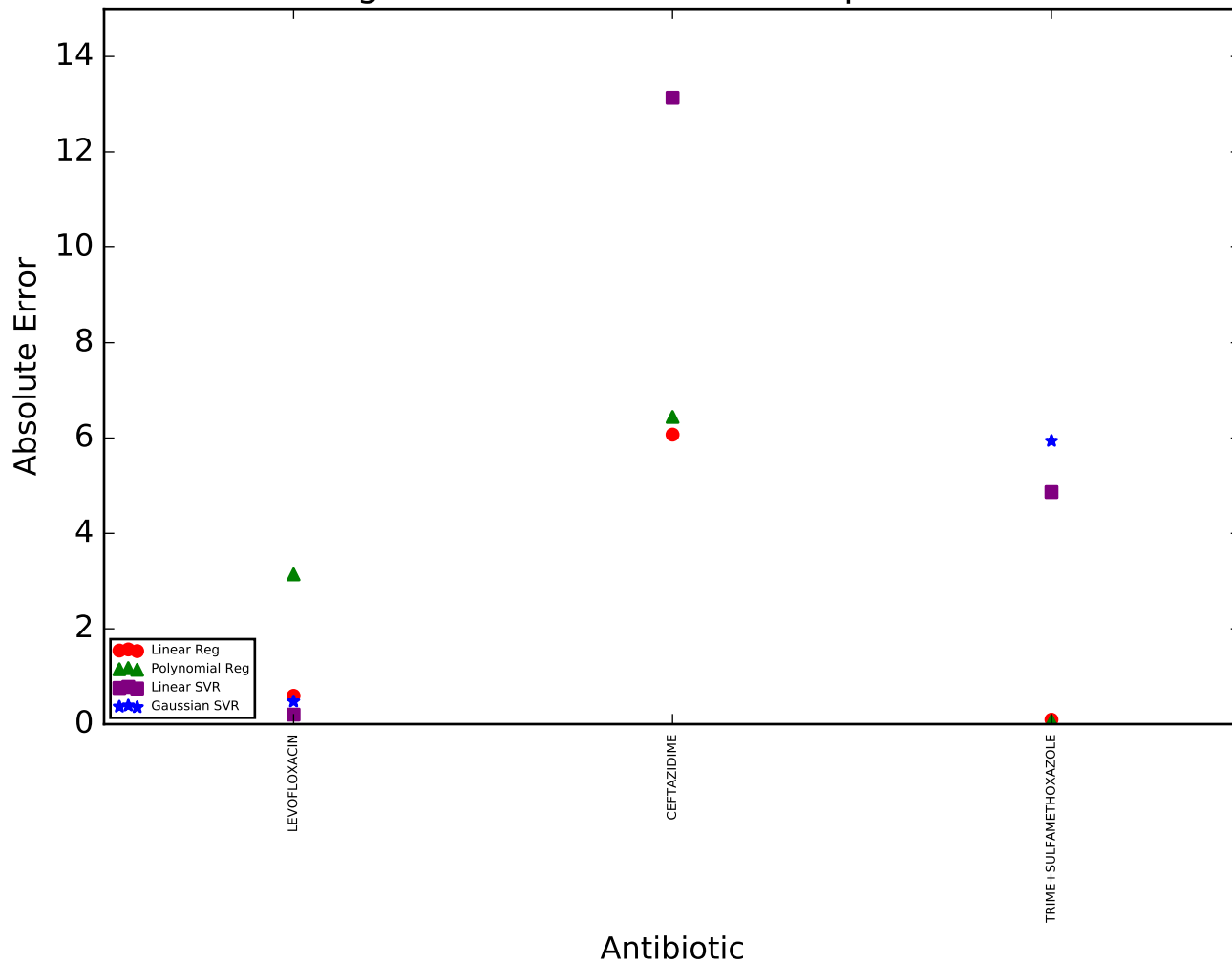
# MSSA

using data from 2004-2012 to predict 2013



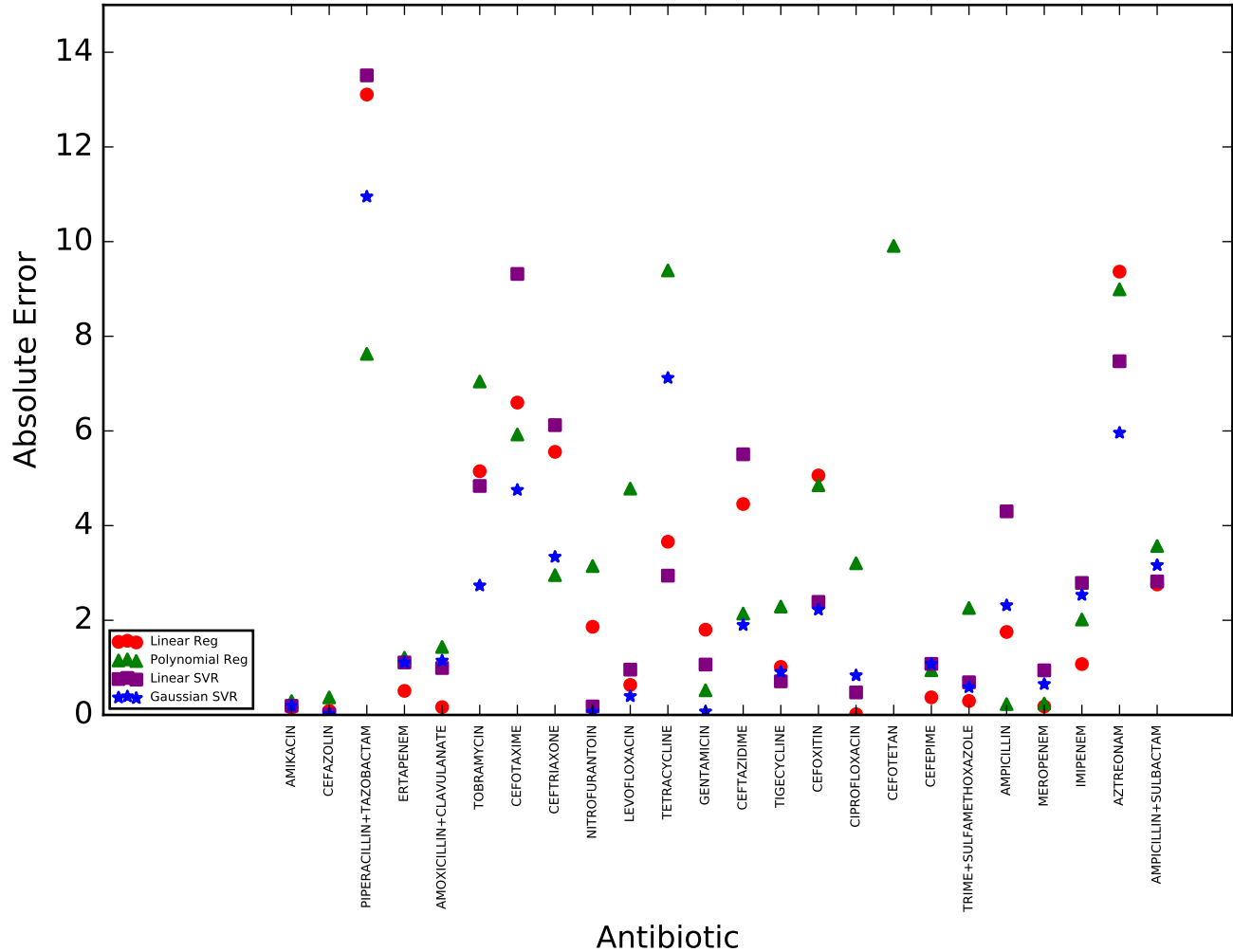
# Stenotrophomonas maltophilia

## using data from 2003-2011 to predict 2013



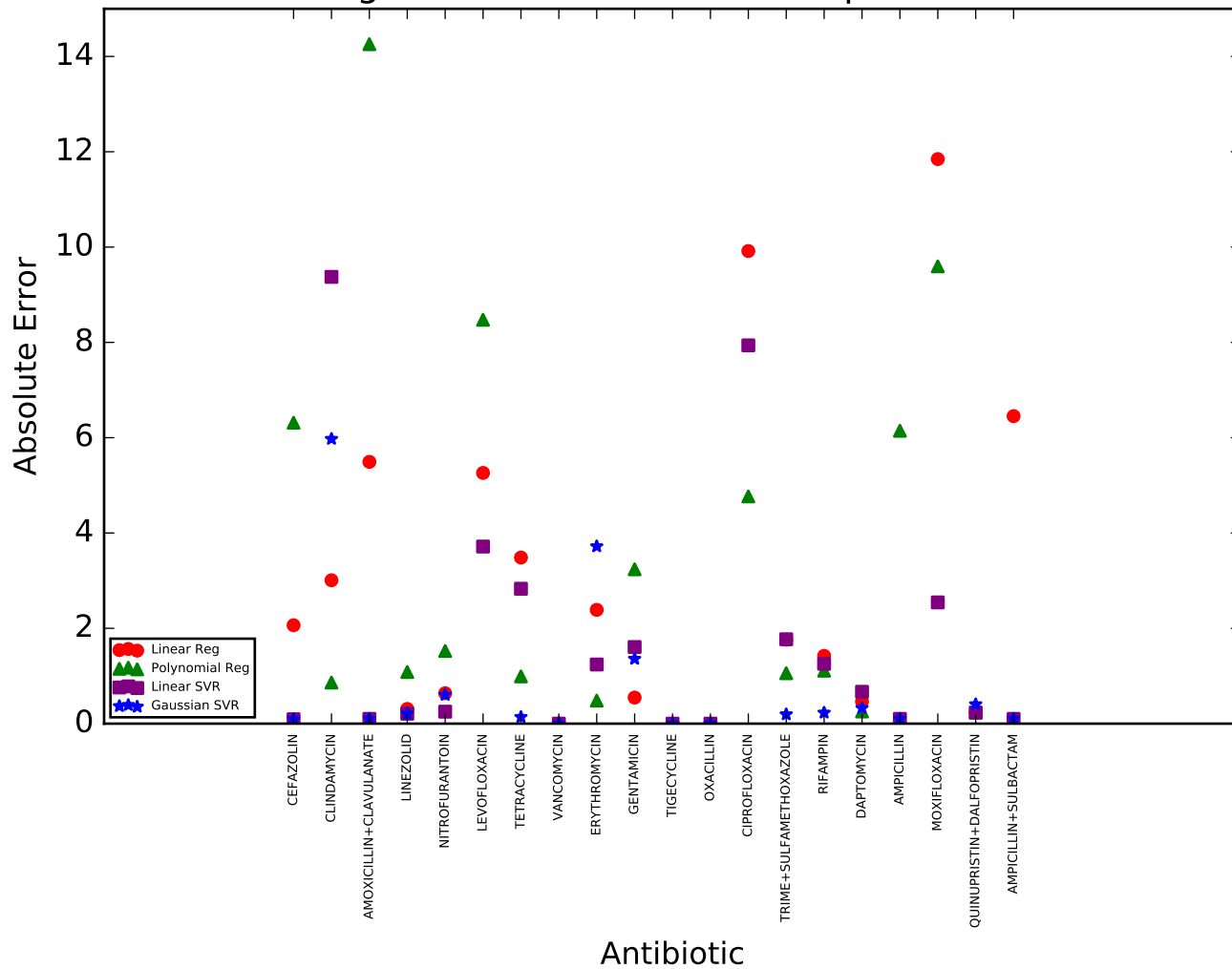
# Serratia marcescens

## using data from 2003-2011 to predict 2013



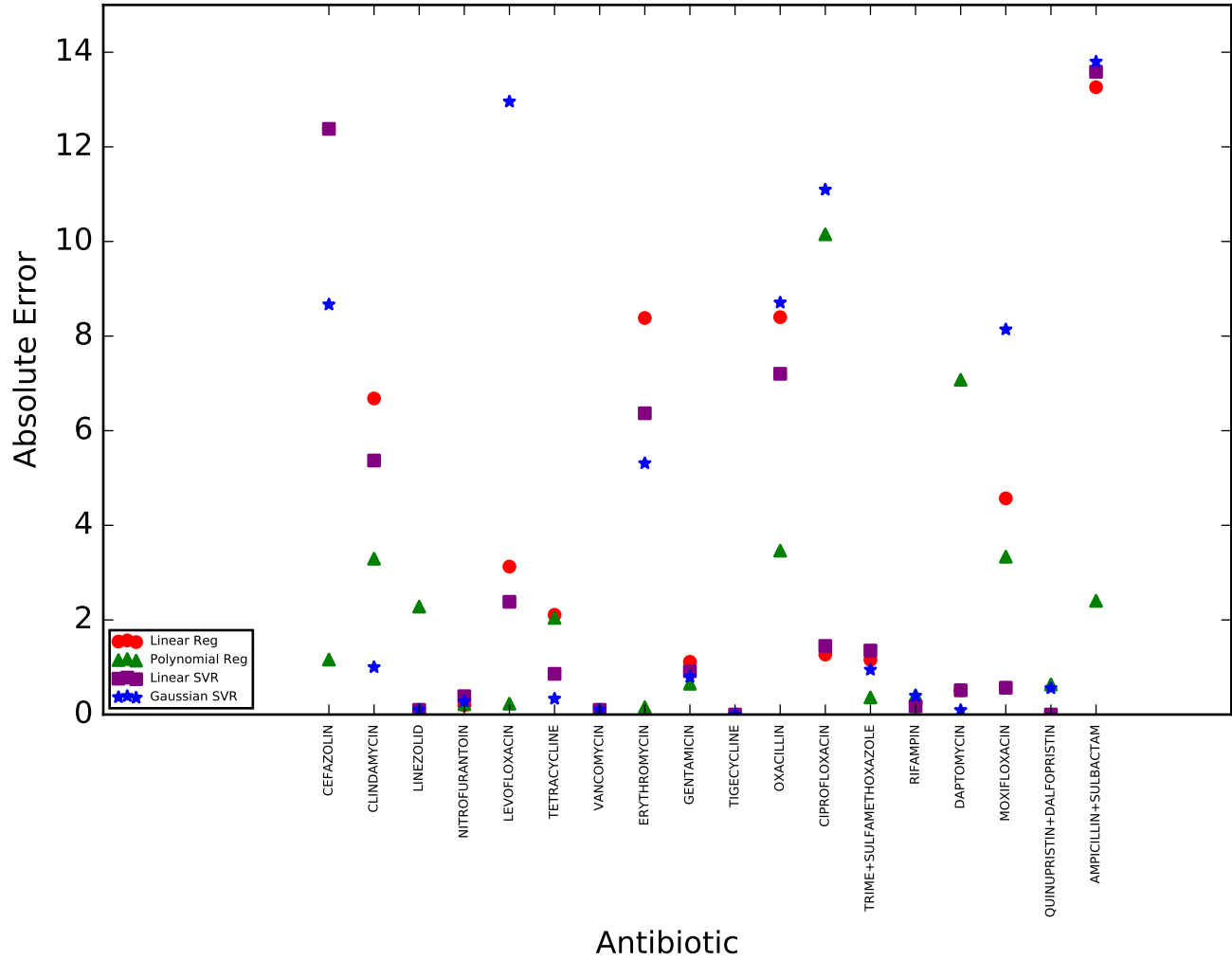
# MRSA

## using data from 2003-2011 to predict 2013



# Staphylococcus aureus

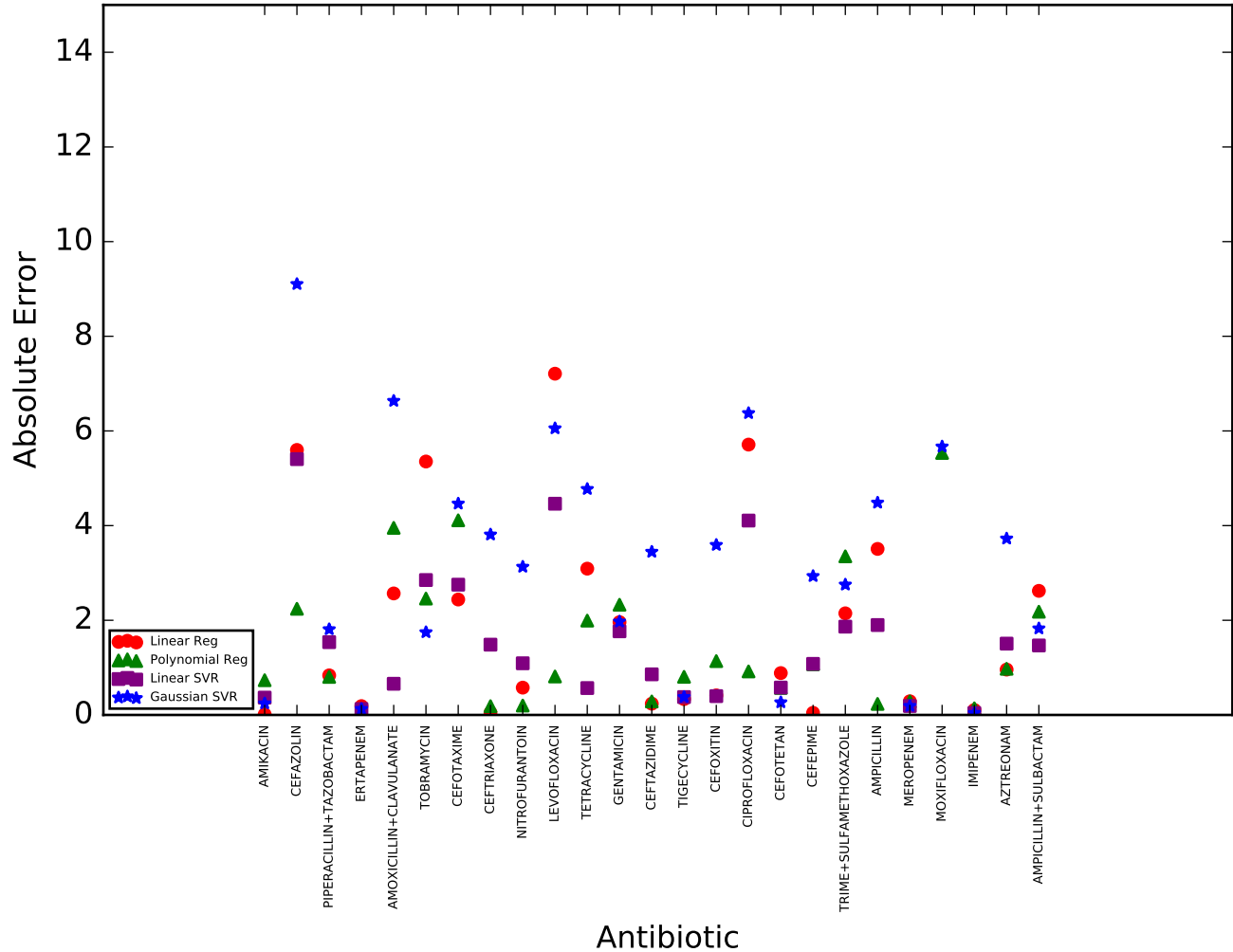
## using data from 2003-2011 to predict 2013





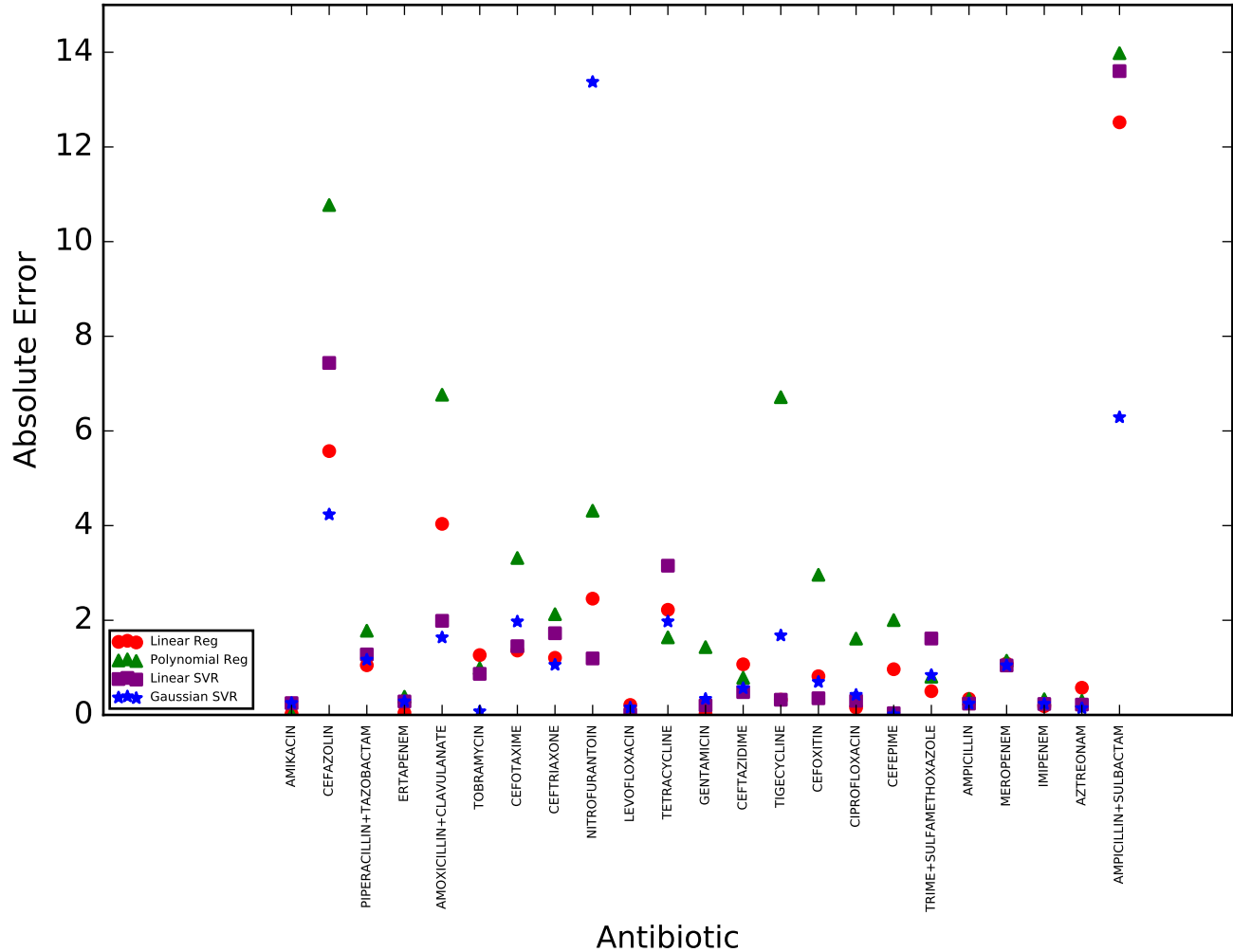
# E. coli

## using data from 2003-2011 to predict 2013



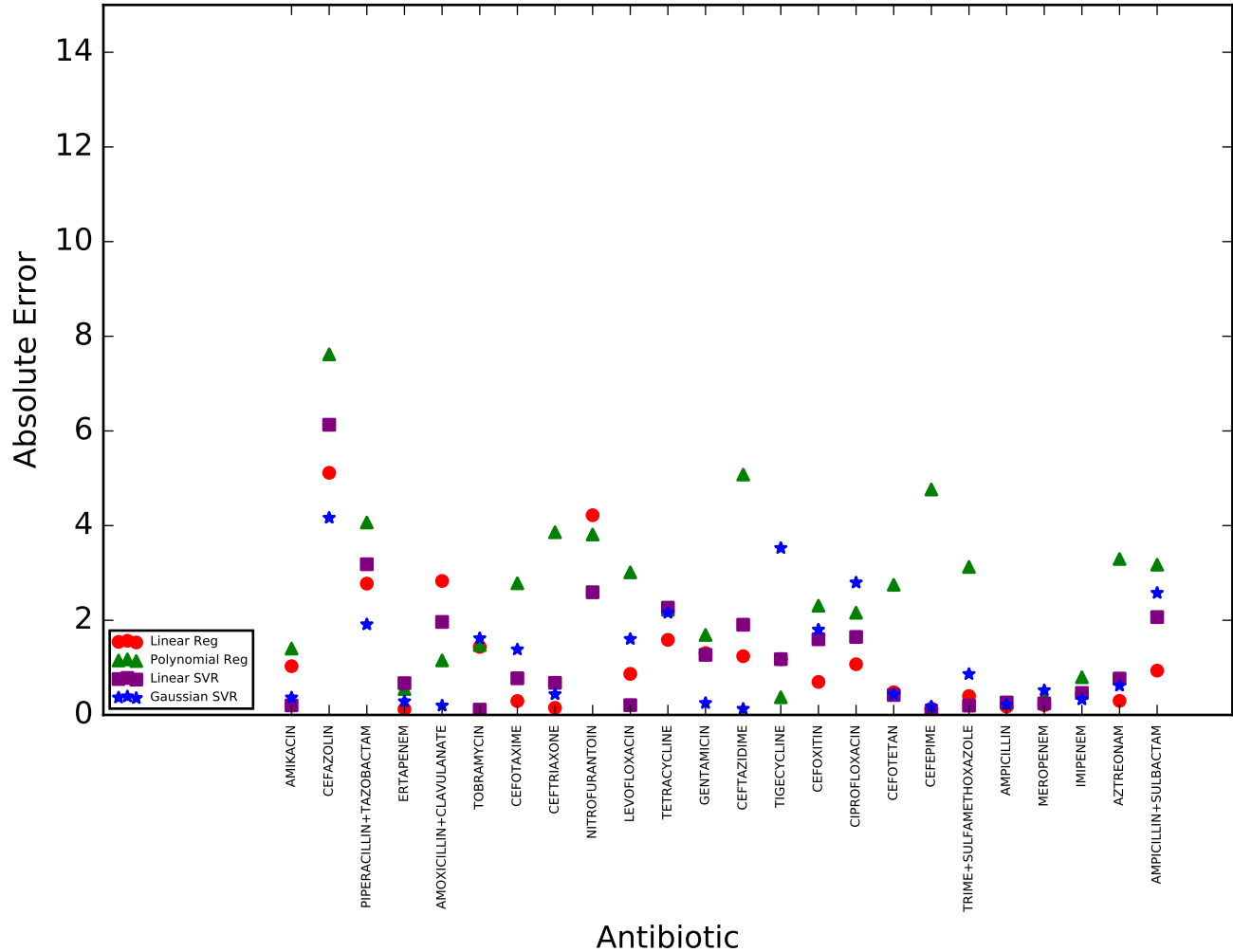
# Klebsiella oxytoca

## using data from 2003-2011 to predict 2013



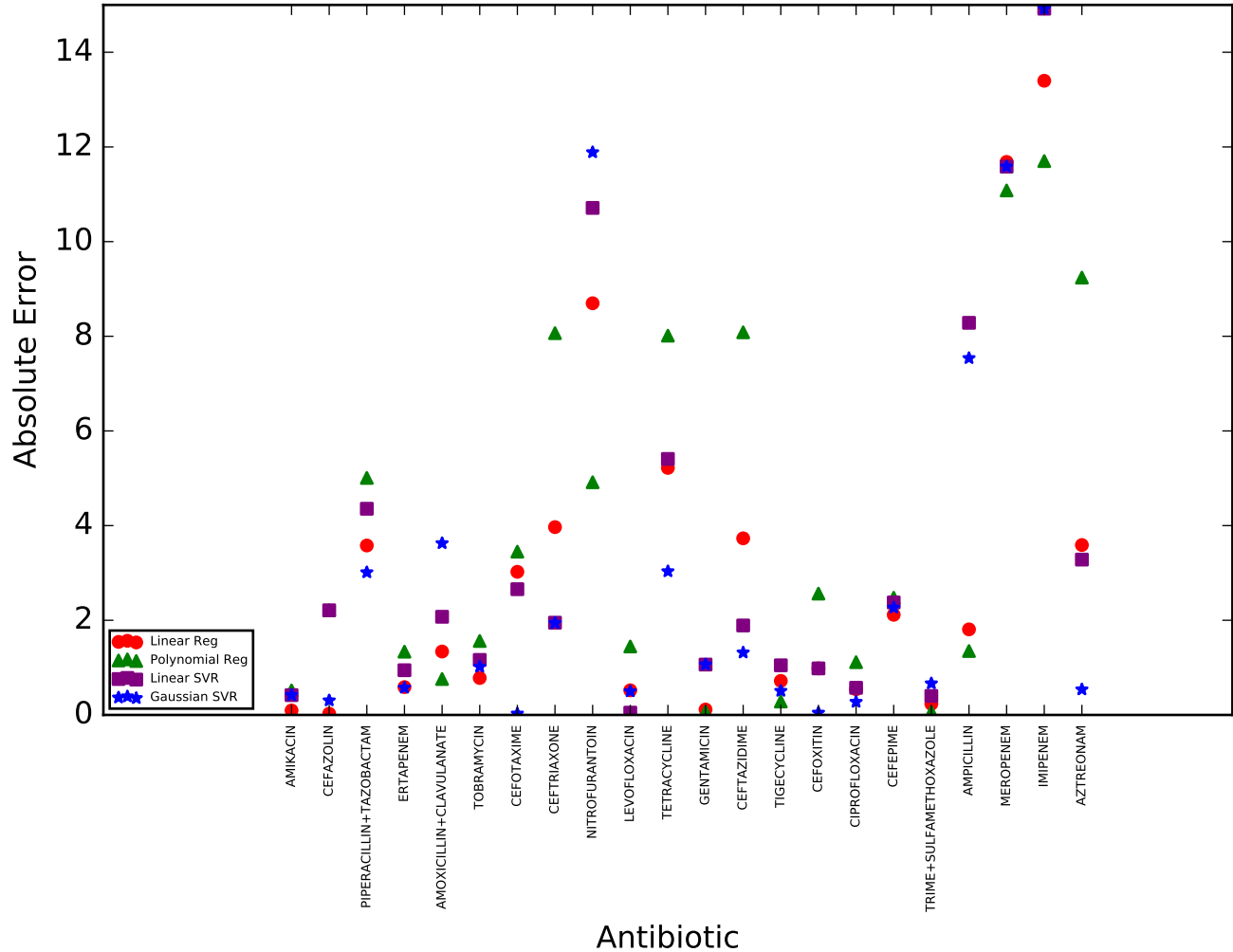
# Klebsiella pneumoniae

## using data from 2003-2011 to predict 2013



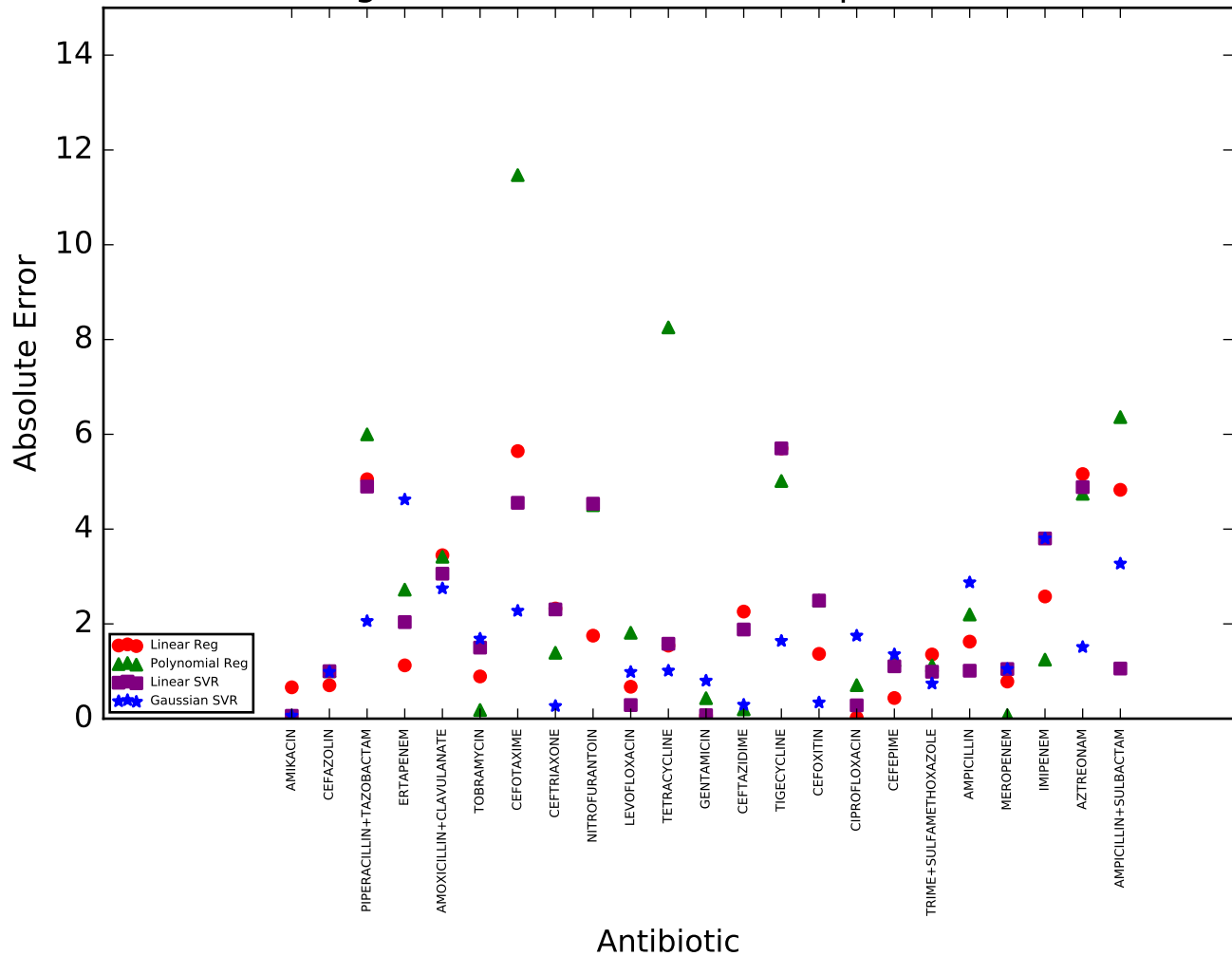
# Enterobacter aerogenes

## using data from 2003-2011 to predict 2013

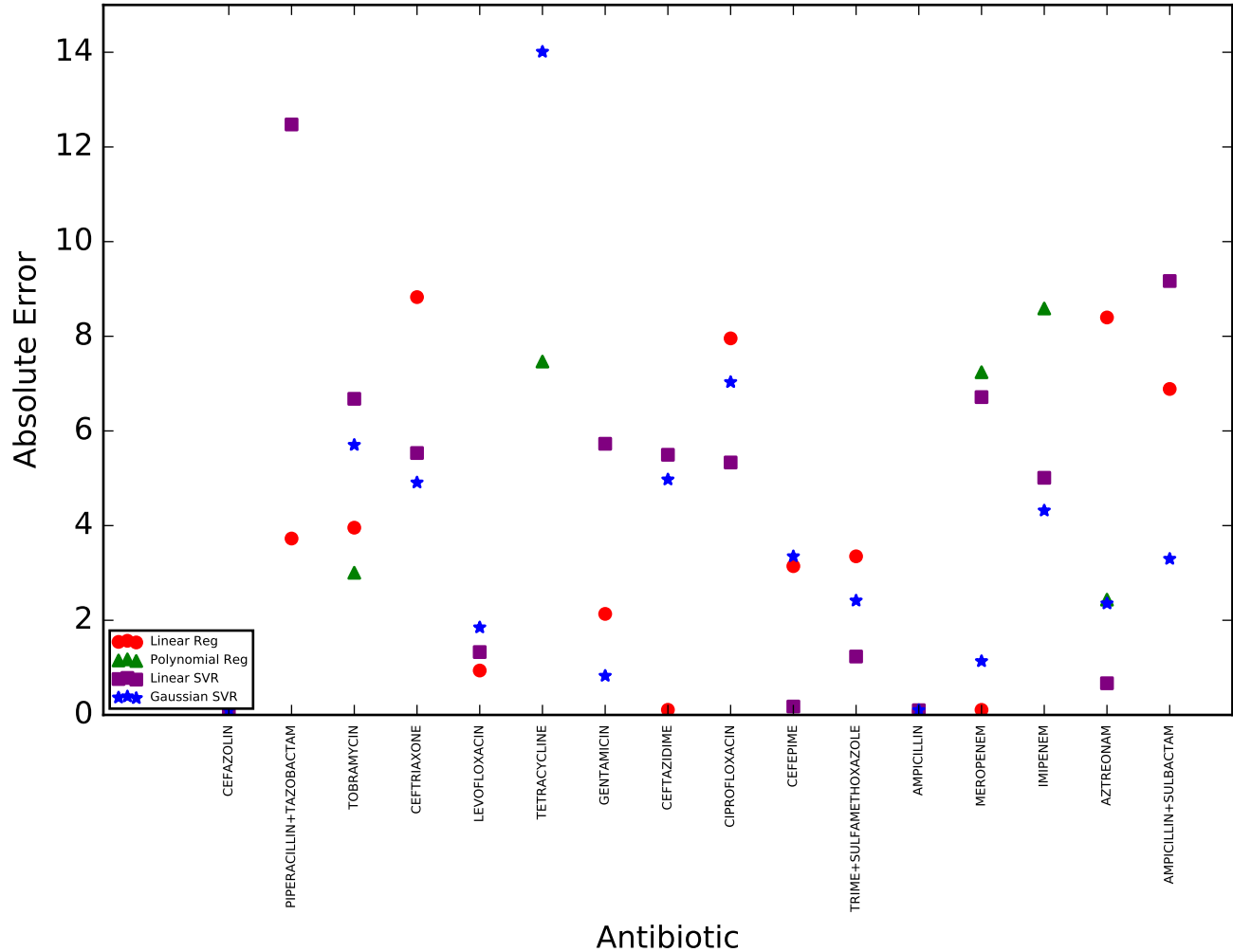


# Enterobacter cloacae

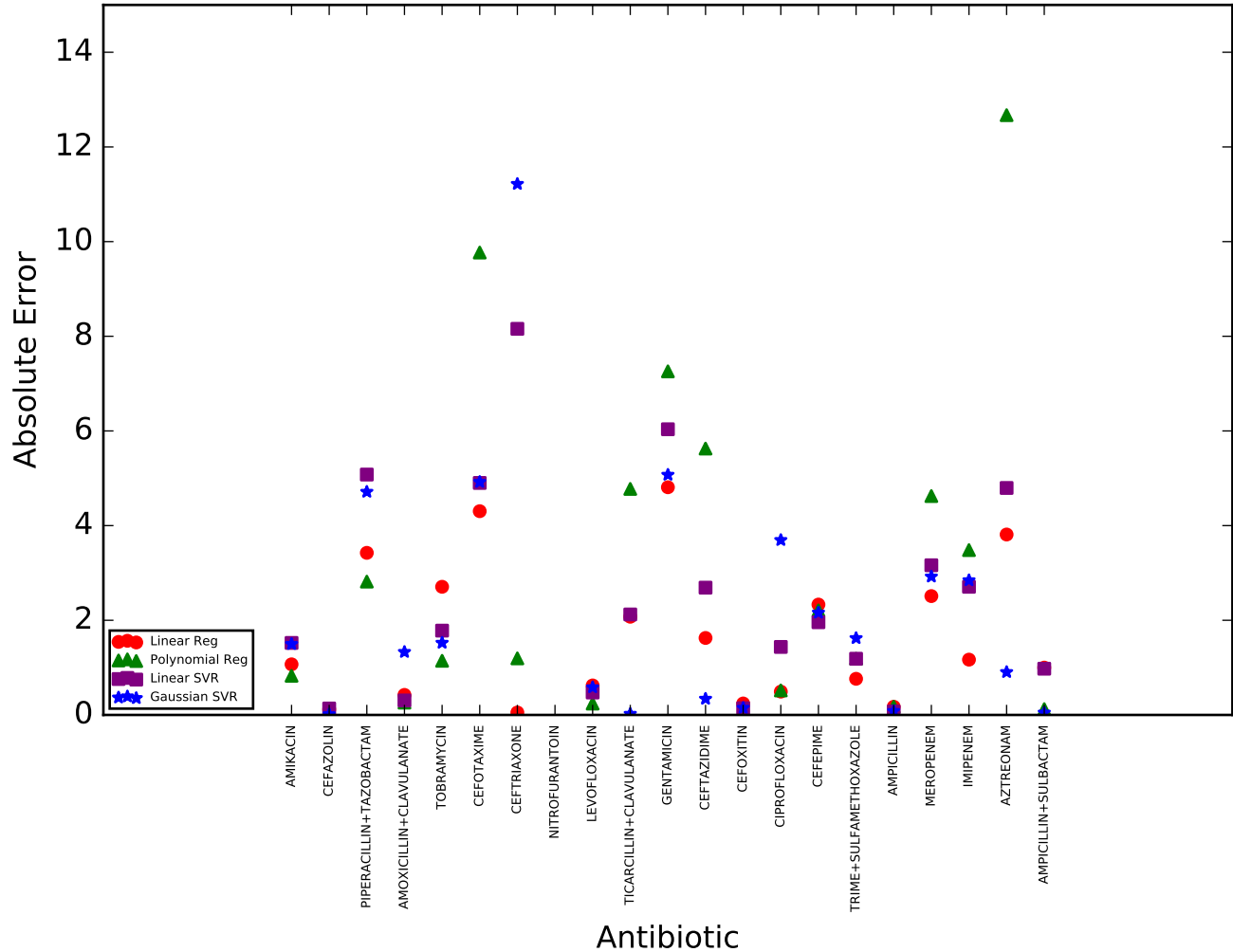
## using data from 2003-2011 to predict 2013



# Acinetobacter baumannii using data from 2003-2011 to predict 2013

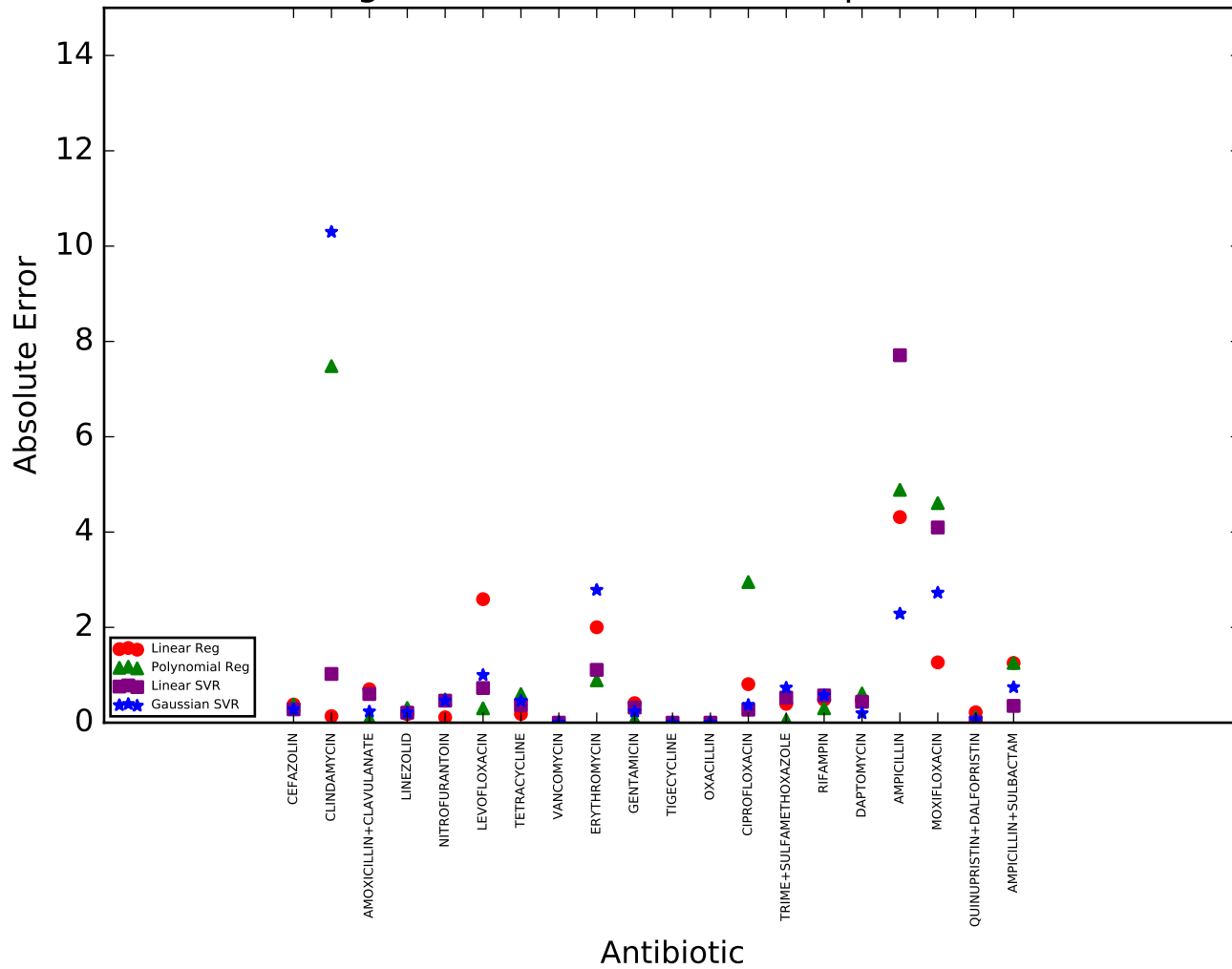


# *Pseudomonas aeruginosa* using data from 2003-2011 to predict 2013



# MSSA

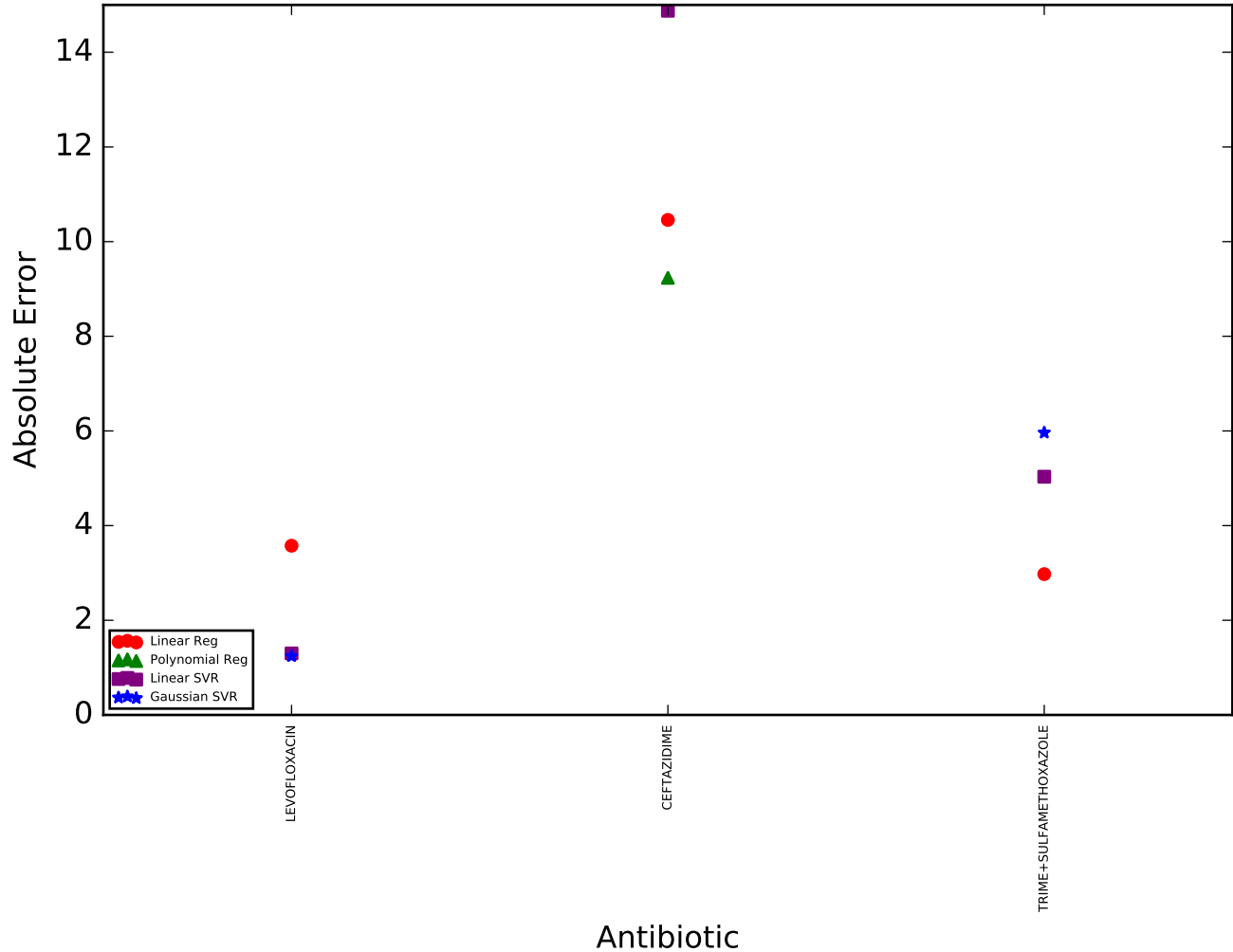
## using data from 2003-2011 to predict 2013





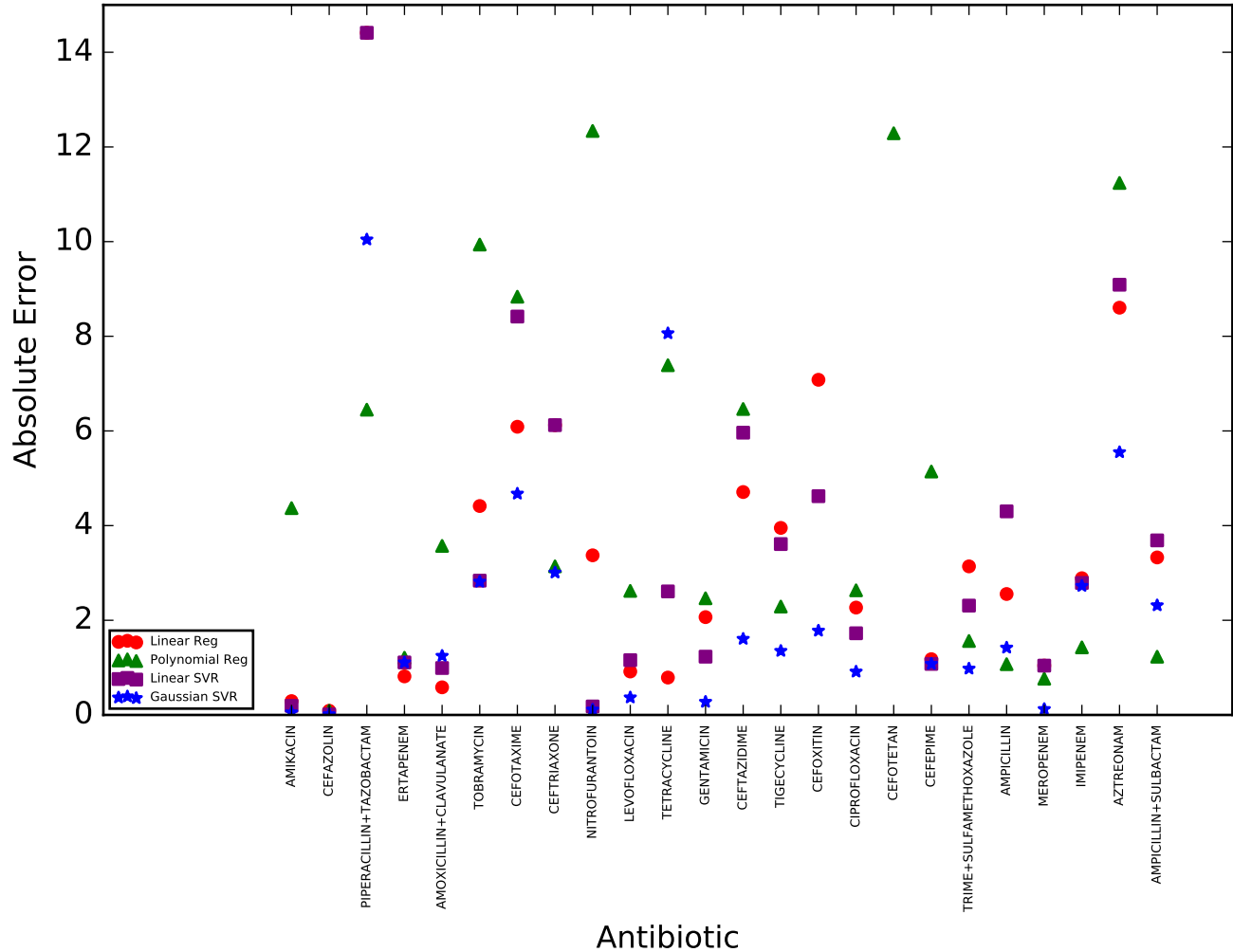
# Stenotrophomonas maltophilia

using data from 2002-2010 to predict 2013



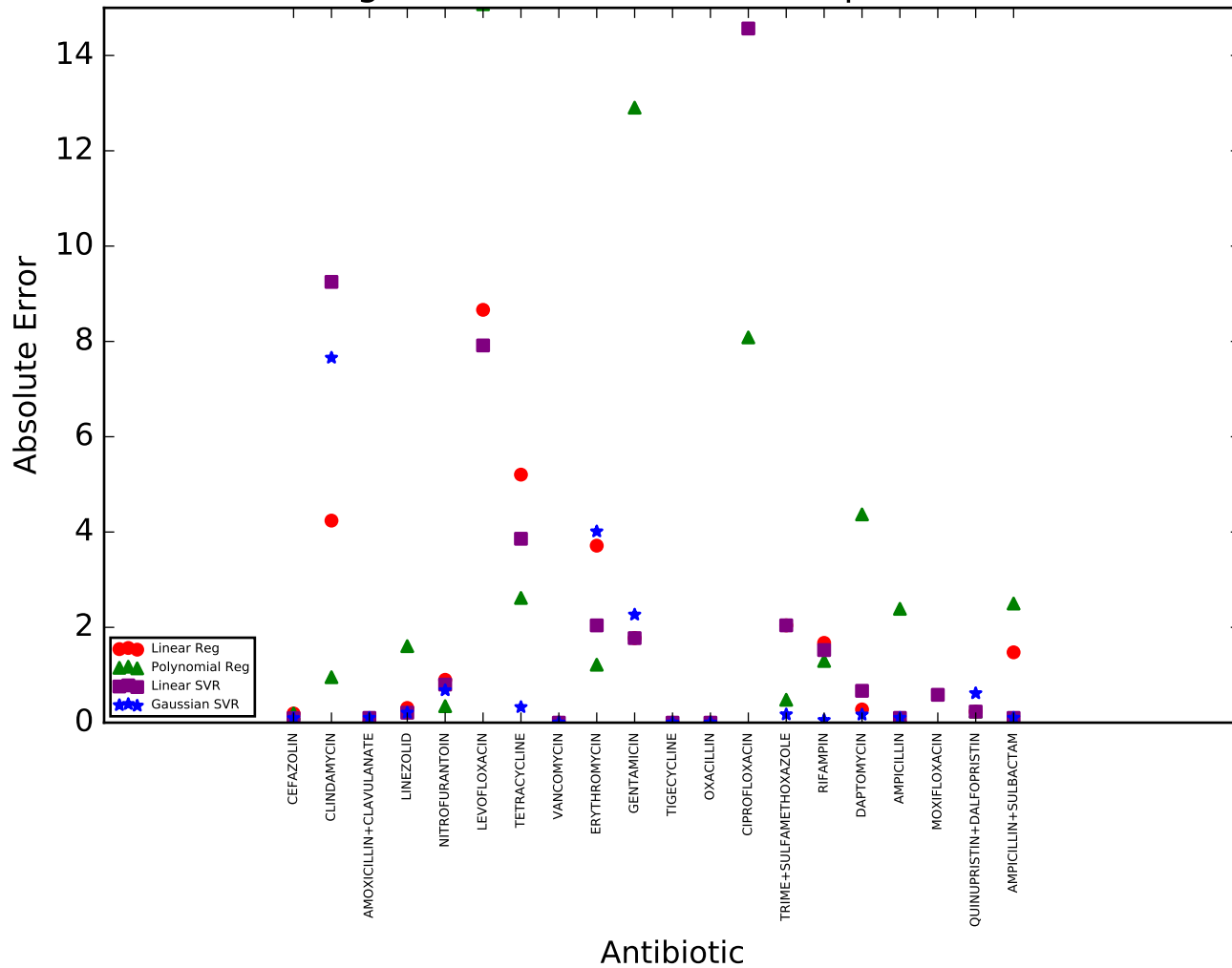
# Serratia marcescens

## using data from 2002-2010 to predict 2013



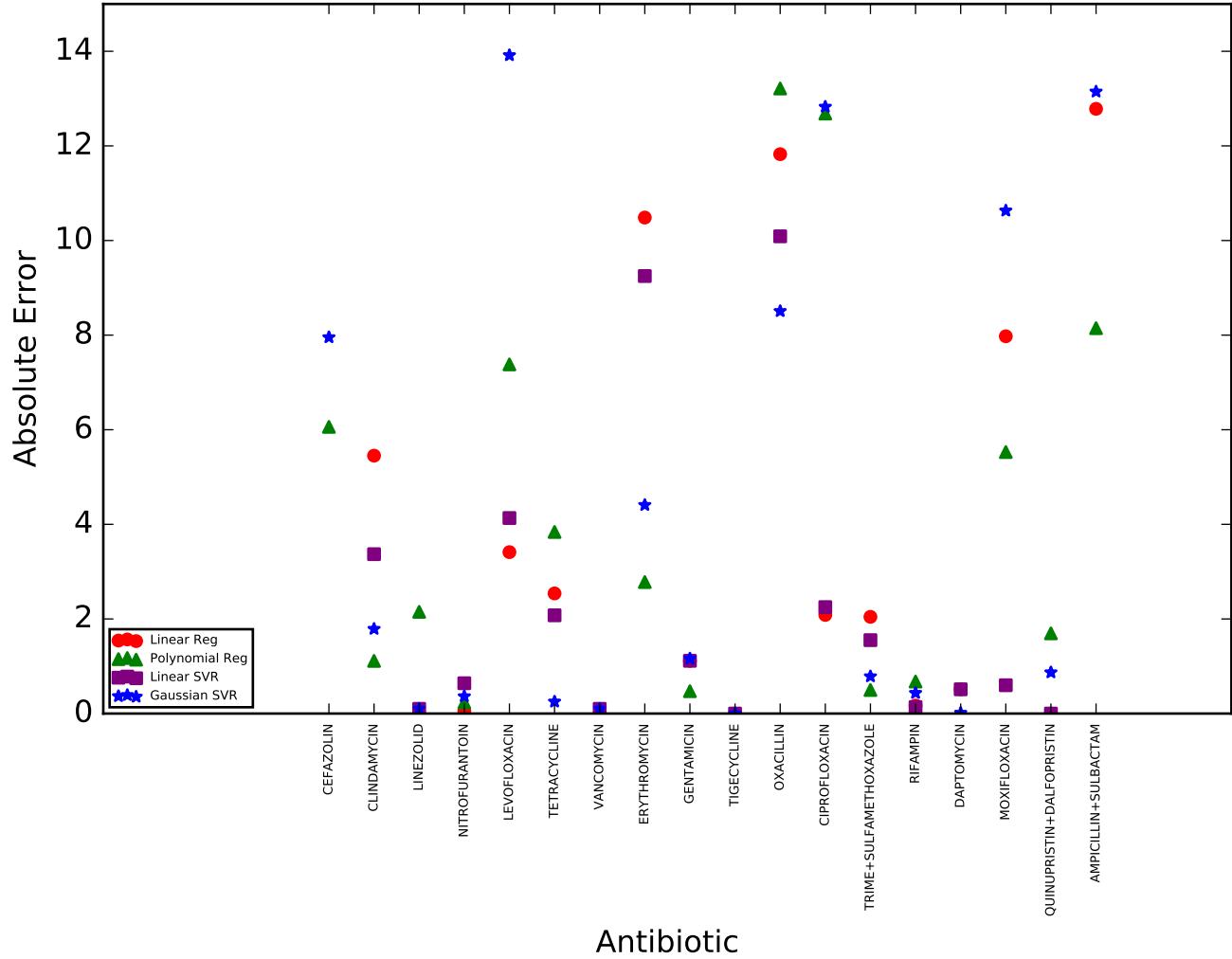
# MRSA

## using data from 2002-2010 to predict 2013



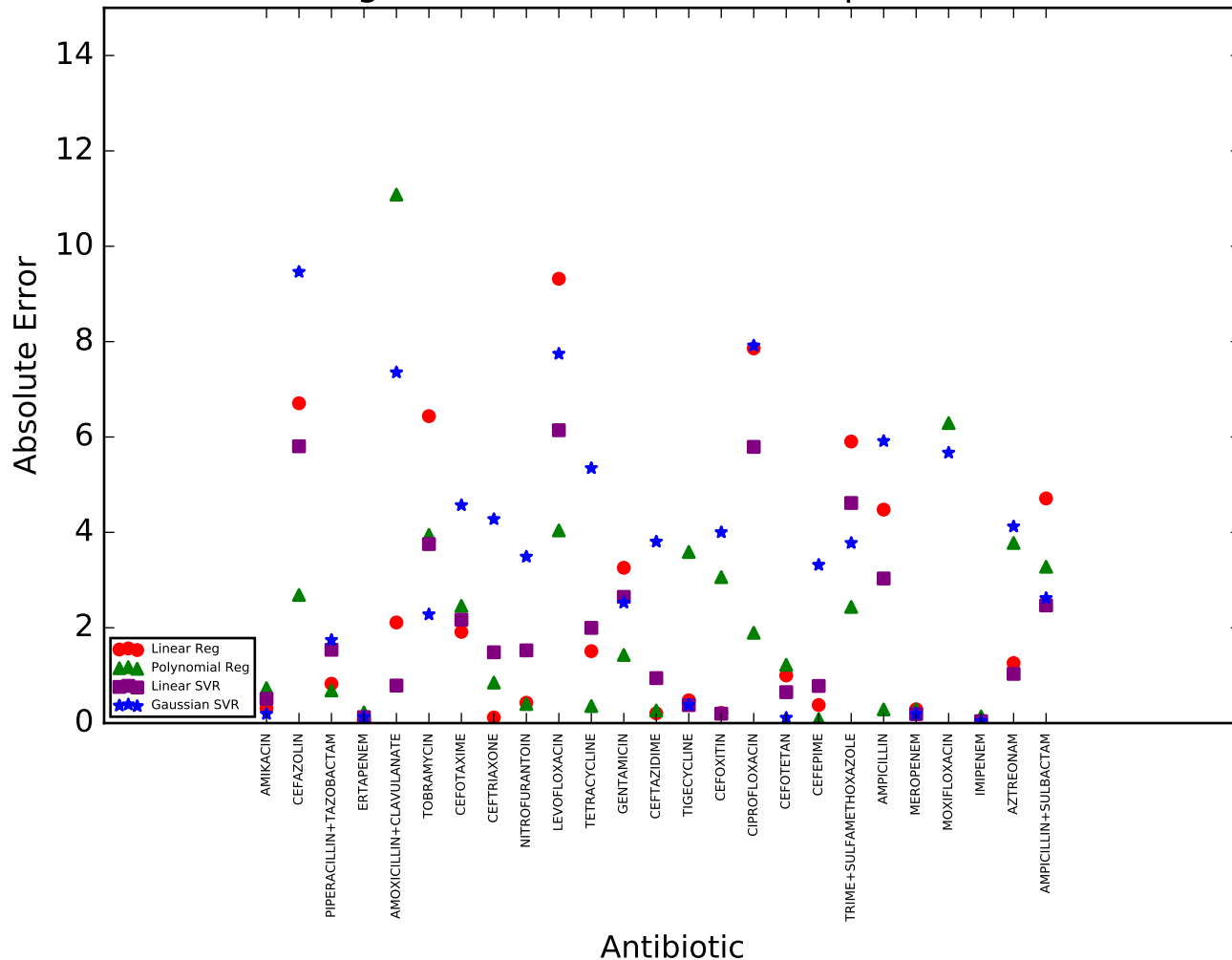
# Staphylococcus aureus

## using data from 2002-2010 to predict 2013



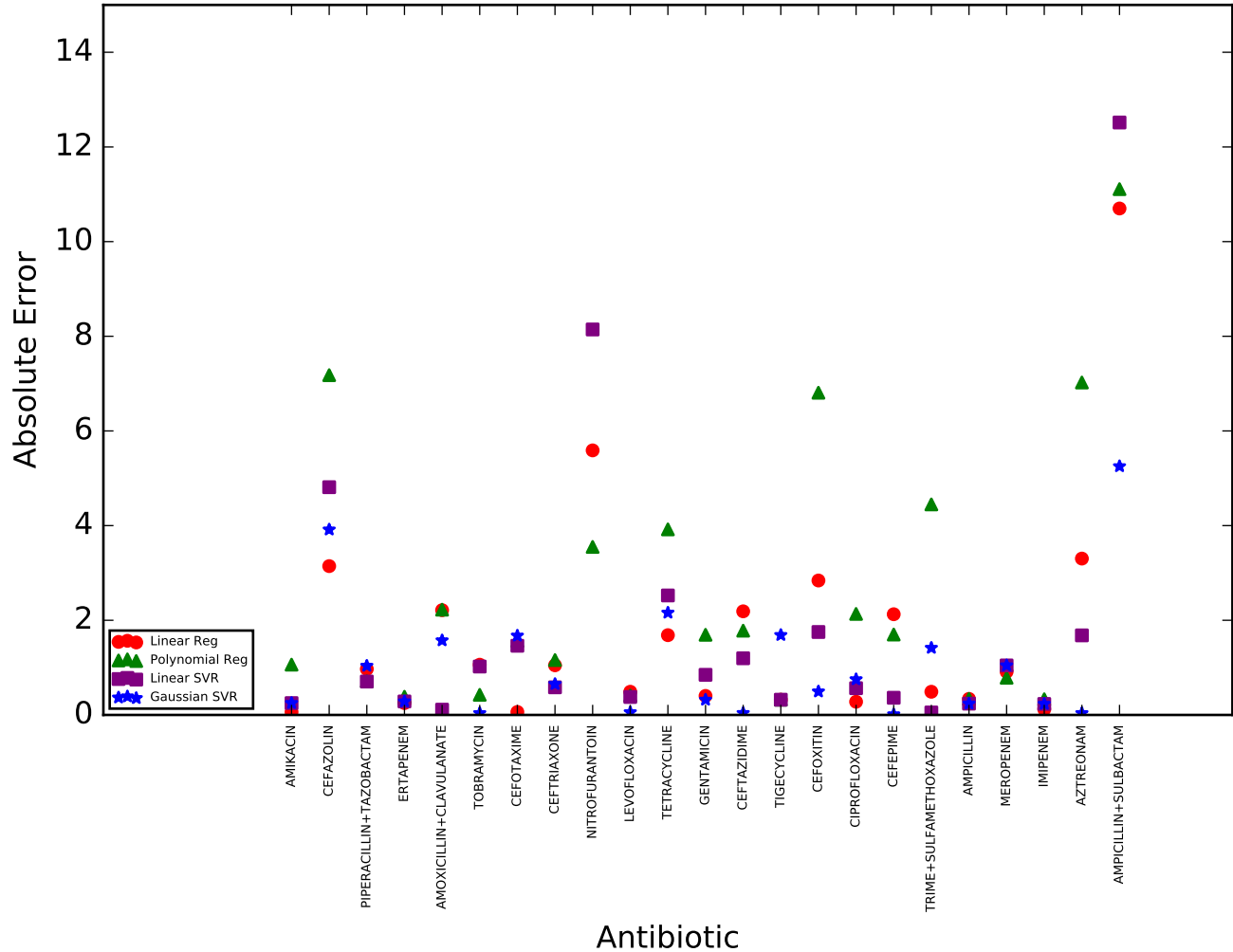
# E. coli

## using data from 2002-2010 to predict 2013



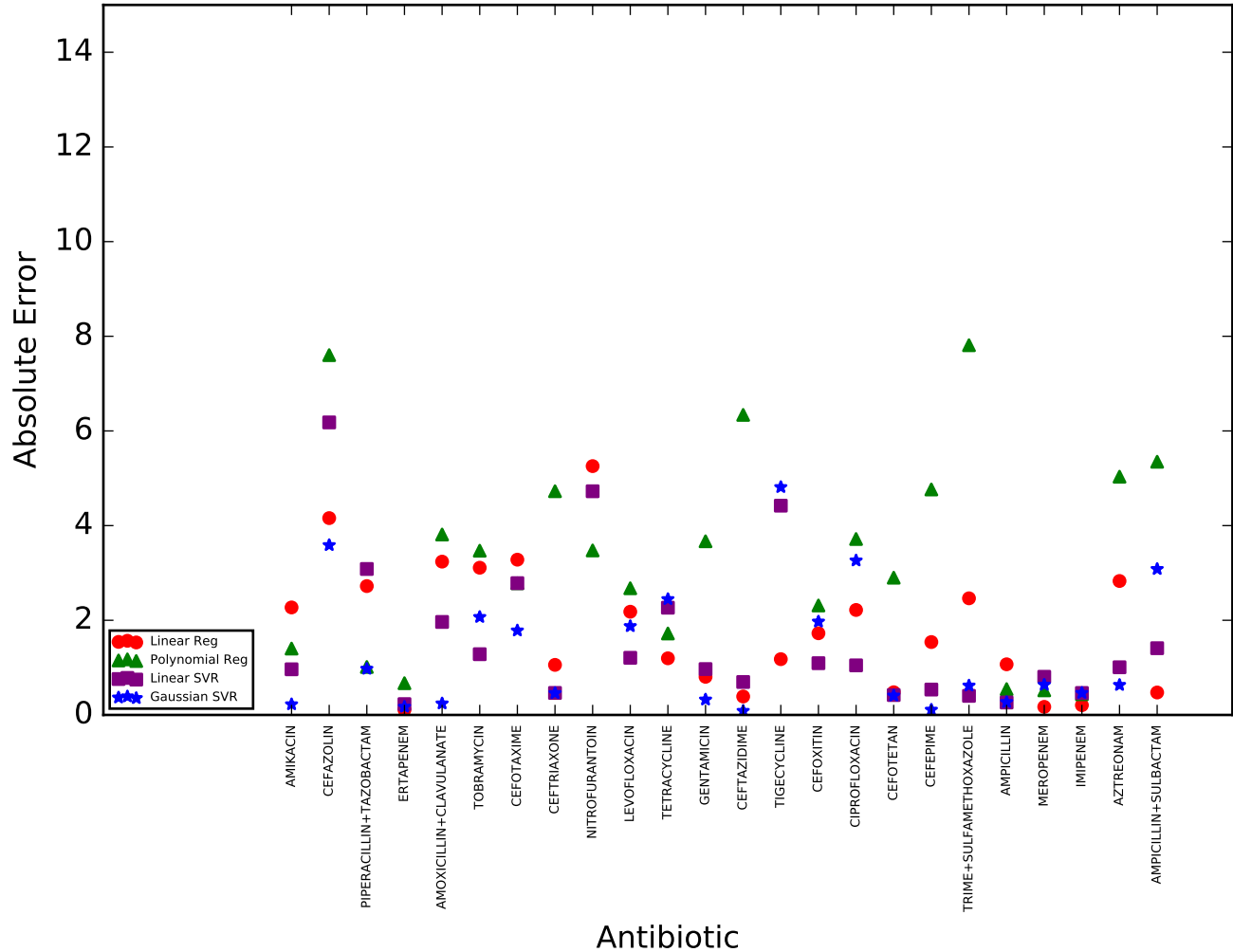
# Klebsiella oxytoca

## using data from 2002-2010 to predict 2013

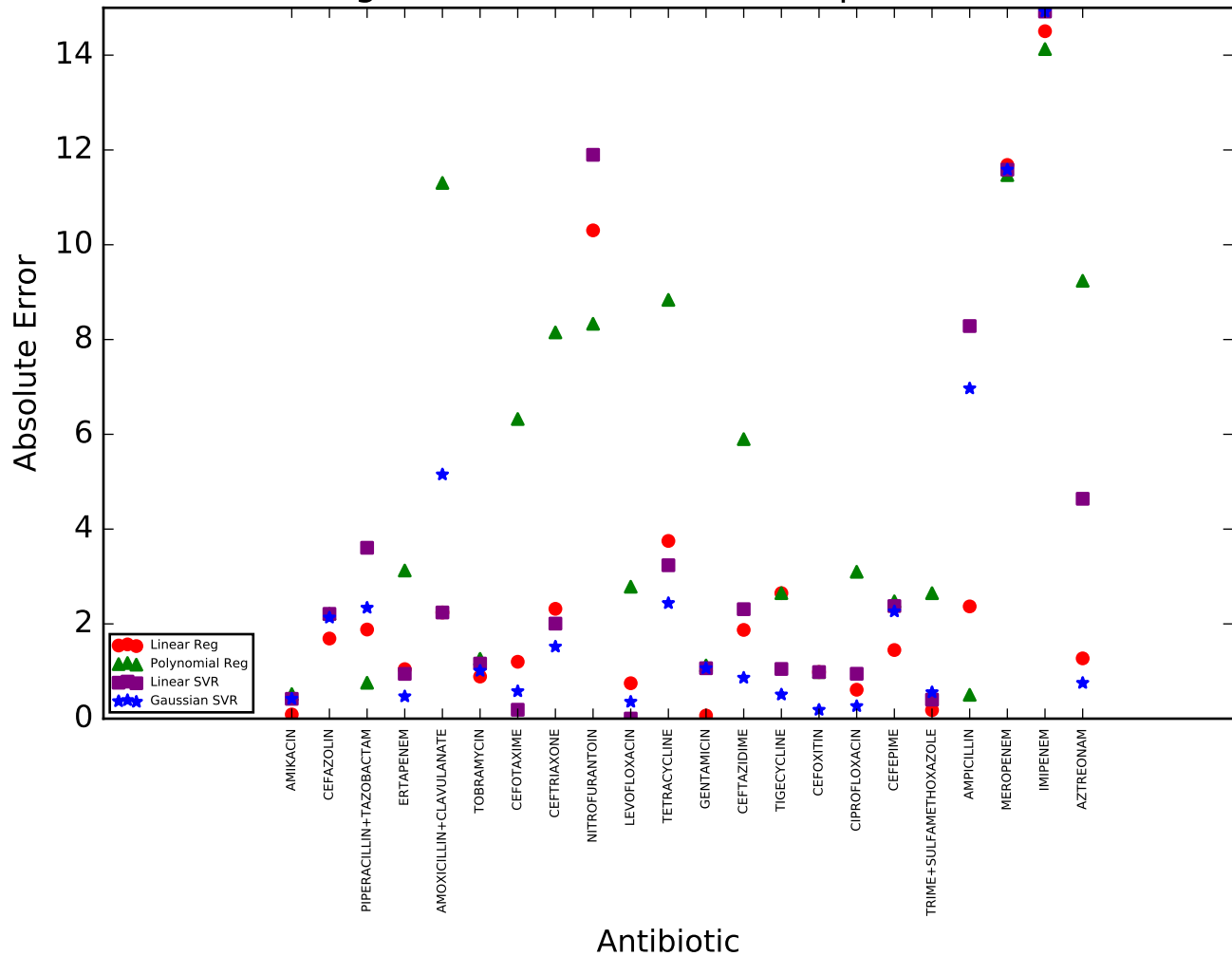


# Klebsiella pneumoniae

## using data from 2002-2010 to predict 2013



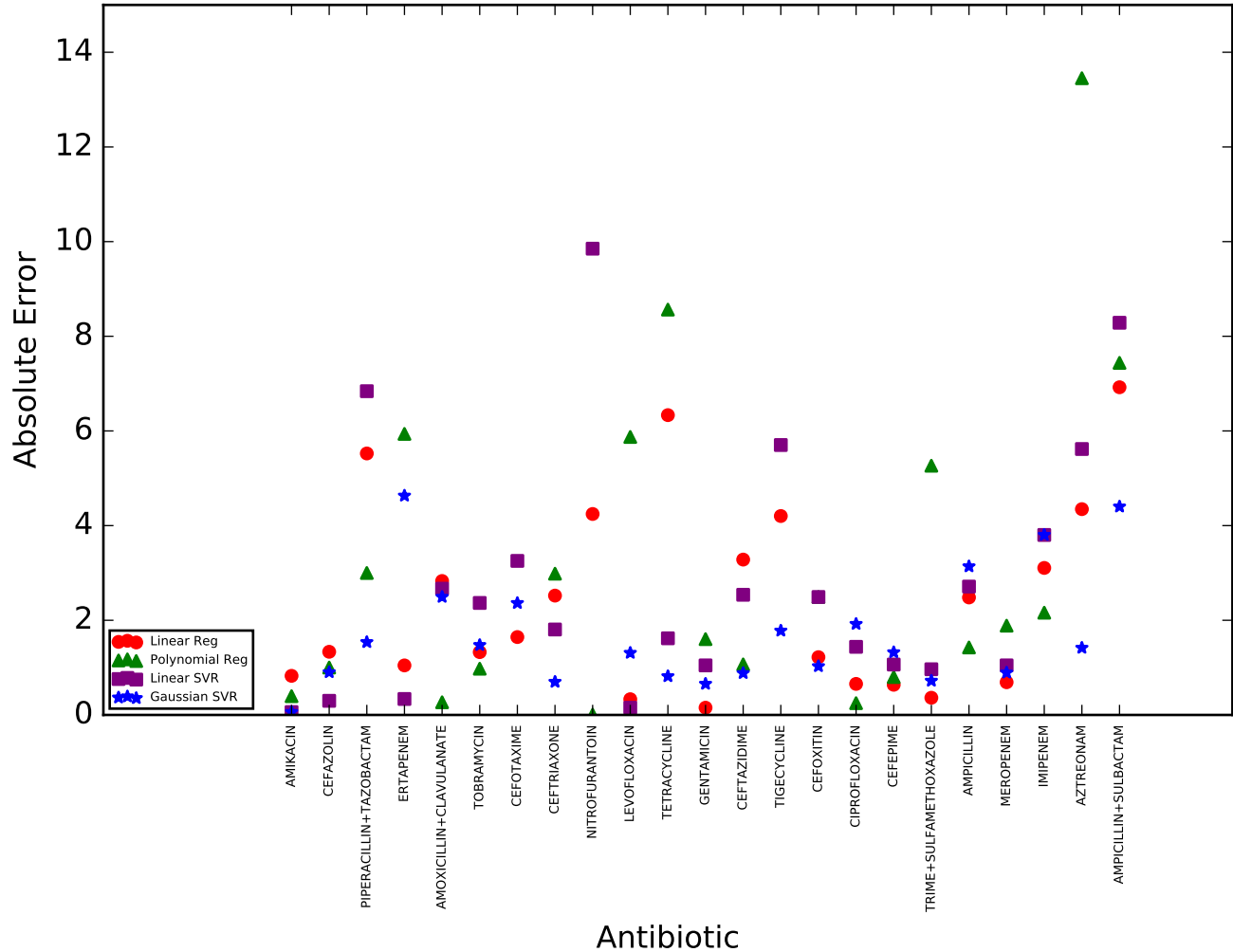
# Enterobacter aerogenes using data from 2002-2010 to predict 2013



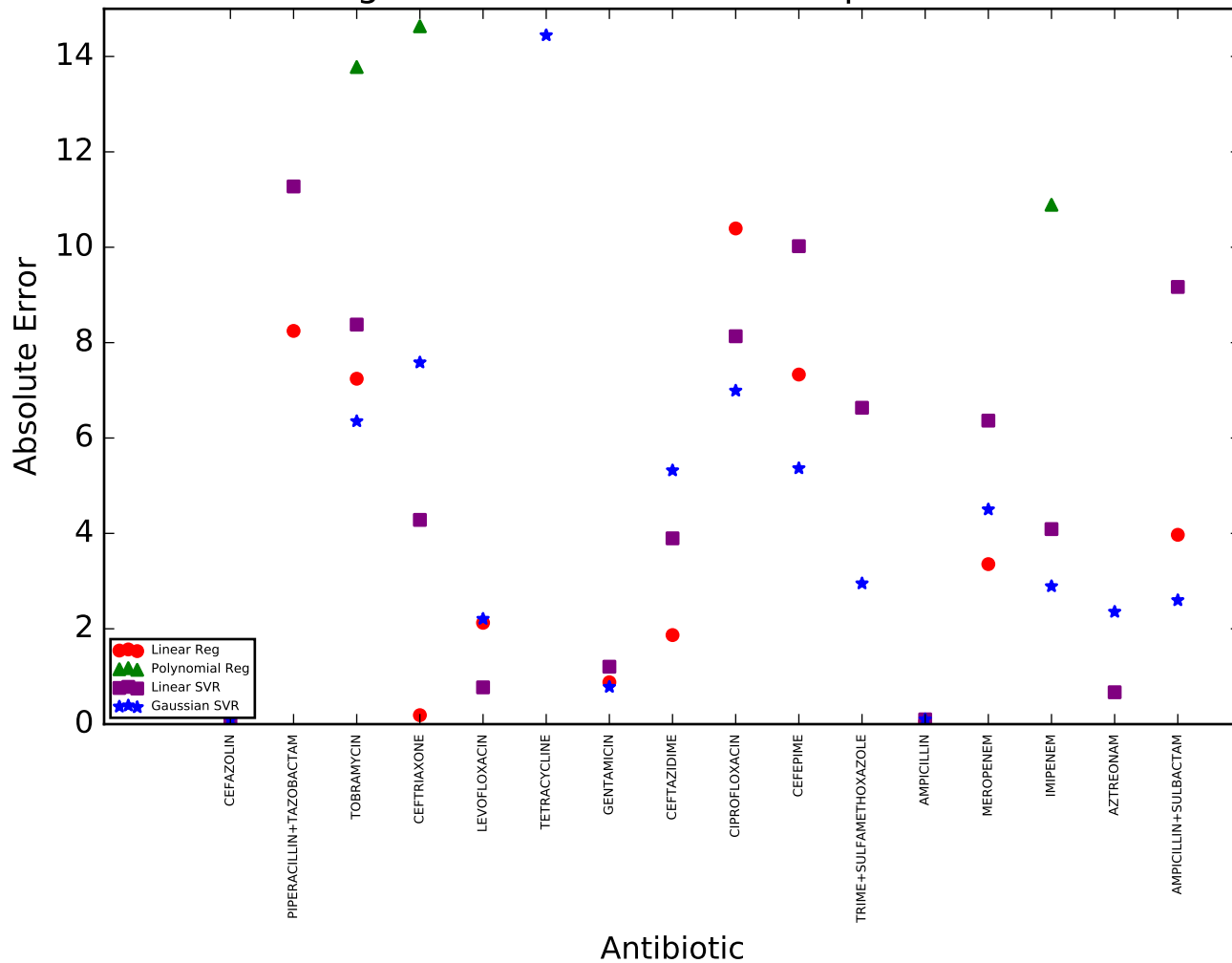


# Enterobacter cloacae

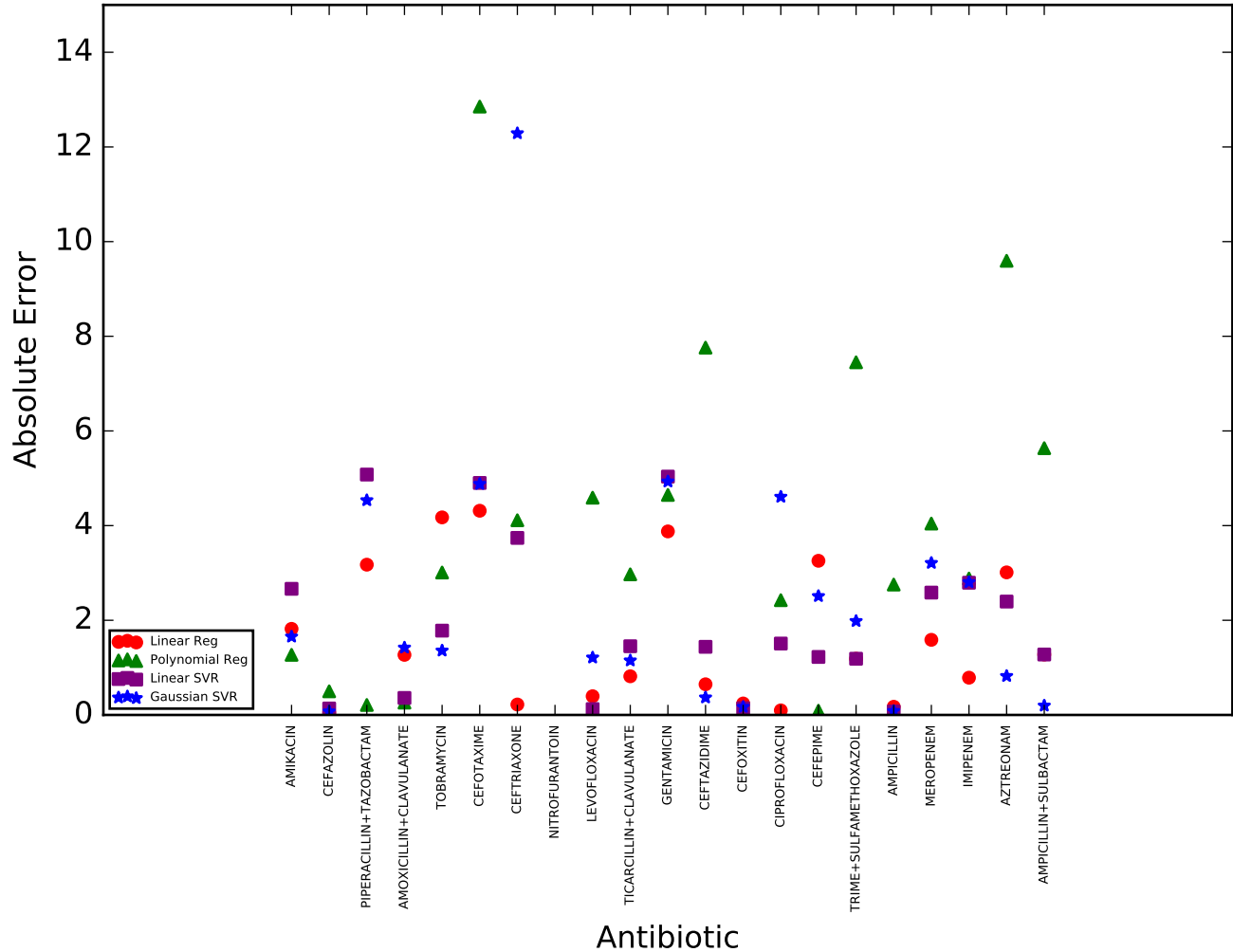
## using data from 2002-2010 to predict 2013



# Acinetobacter baumannii using data from 2002-2010 to predict 2013



# *Pseudomonas aeruginosa* using data from 2002-2010 to predict 2013



# MSSA

using data from 2002-2010 to predict 2013

