

Modified Logistic Model

$$y = \frac{a}{[1 + e^{(b-cx)}]}$$

$$Y = P2 + \frac{A - P2}{[1 + e^{((\frac{\mu m}{A}) * (\lambda - t) + 2)}]}$$

Y = OD600

P2 = second measure point

A = three highest consecutive points

λ = lag phase

t = time

Determining the μ_{max}

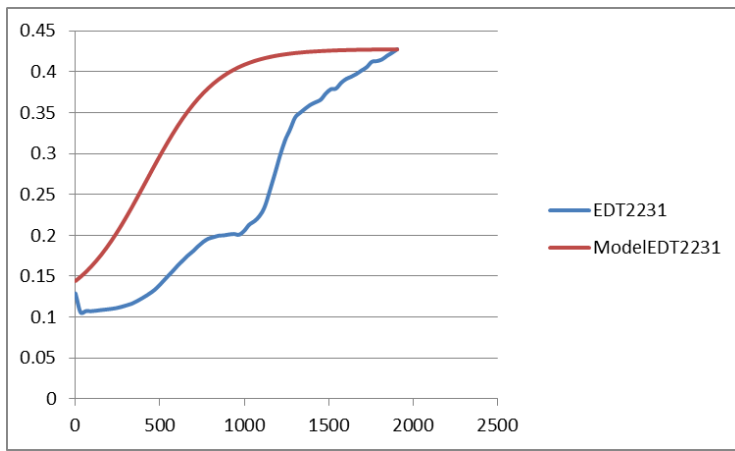
- Input data: OD600 measures in time
- Averages of replicates are divided in overlapped windows of data points
- Window sizes are adjusted to assure the same time interval in each window if different experimental time intervals are compared.
- In every overlapping window, the doubling rate of the extreme points are used to determine the rate of growth:

$$\mu = \frac{2.303(\log(OD_2) - \log(OD_1))}{(t_1 - t_2)}$$

- The highest one is sorted out

Determining The λ

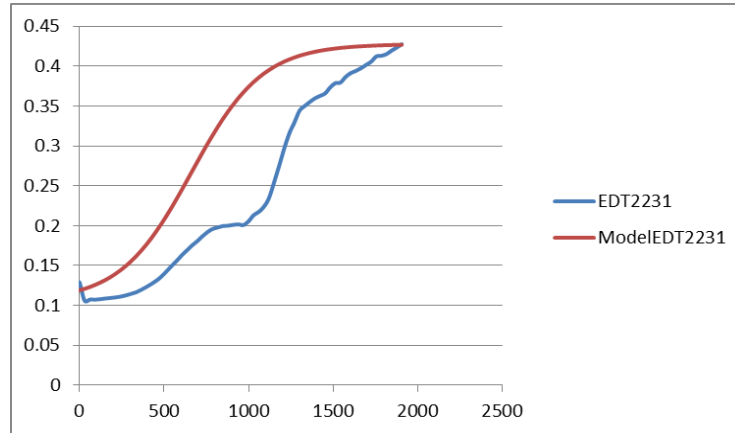
- The lag phase is an issue of controversy in the microbial kinetics literature.
- Many concepts are available, but there are only a few objective ways to actually calculate it from the data.
- To overcome this we plotted our data to the logistic model with $\lambda = 0$.
- Iterated over all time points for the lowest standard deviation between the model vs. the actual data.
- This standard deviation is also stored for a further T-test comparison (still under preperation...)



$$\lambda = 0$$

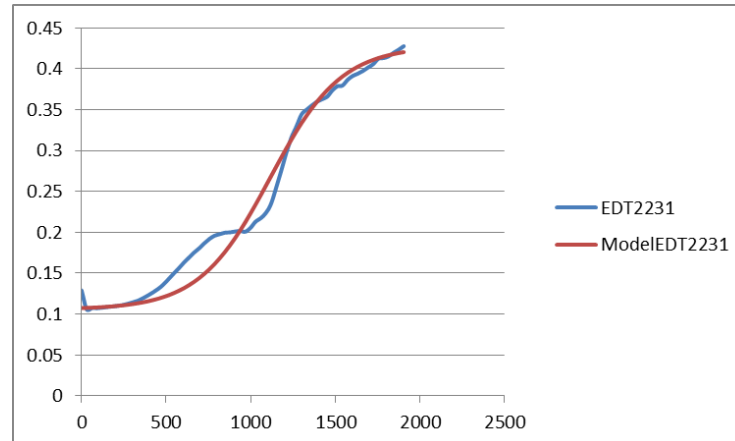
Stdev = 0.05324

$$\frac{\sum_{i=j=1}^N \sqrt{(xi - (\overline{xij}))^2}}{N}$$



$$\lambda = 242$$

Stdev = 0.035192



$$\lambda = 696$$

Stdev = 0.0062439

Scoring

- $$\frac{\sum_{i=j=1}^N \left(\frac{x_i}{x_j}\right)}{N}$$

Comparison of
control x clones

Proportions	Scores	Predicted Performance
$x \geq 2$	-4	Worst
$2 > x > 1.60$	-3	Worst
$1.60 \leq x < 1.33$	- 2	Worst
$1.33 \leq x < 1.14$	- 1	Same
$1.14 \leq x < 0.8$	0	Same
$0.8 \leq x < 0.65$	1	Same
$0.66 \leq x < 0.57$	2	Better
$0.57 \leq x < 0.5$	3	Better
$x \leq 0.5$	4	Better

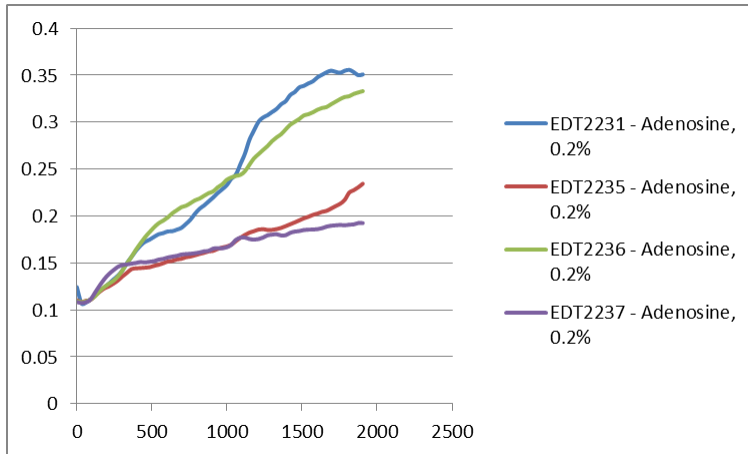
Example Output

Clone	Growth Condition	μ_{max}	λ	Asymptote	Score	Predicted performance
EDT2231	Adenosine, 0,2%	0.00131	363	0.355761		
EDT2235	Adenosine, 0,2%	0.00096	514	0.234526	- 2	worst
EDT2236	Adenosine, 0,2%	0.00144	424	0.333124	0	same
EDT2237	Adenosine, 0,2%	0.00141	242	0.192575	-2	worst

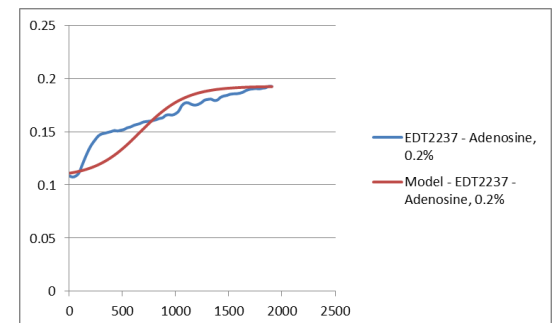
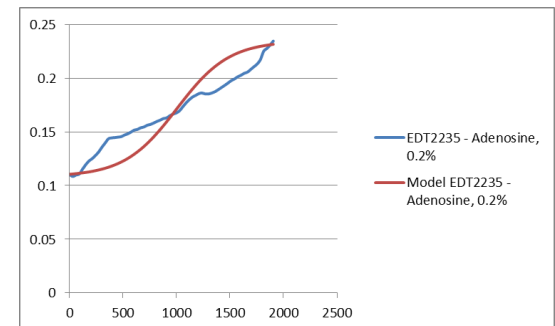
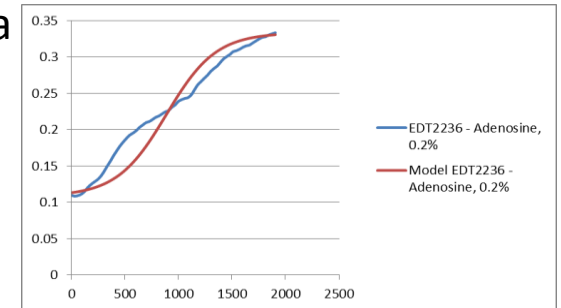
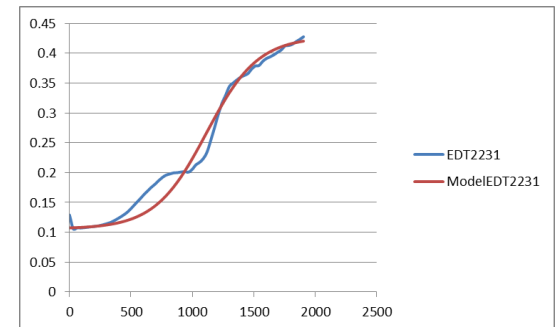
Other data are also stored in separate sheets of the same table, such as: mean and standard deviation of replicates, points used to plot the model and standard deviation of the model.

Examples

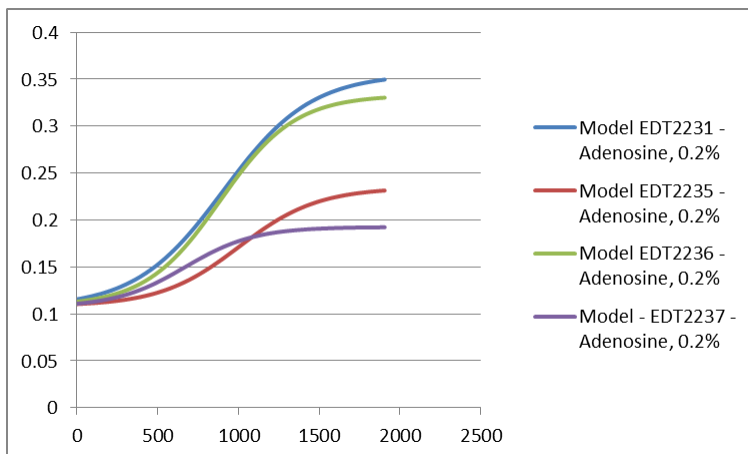
Actual Data



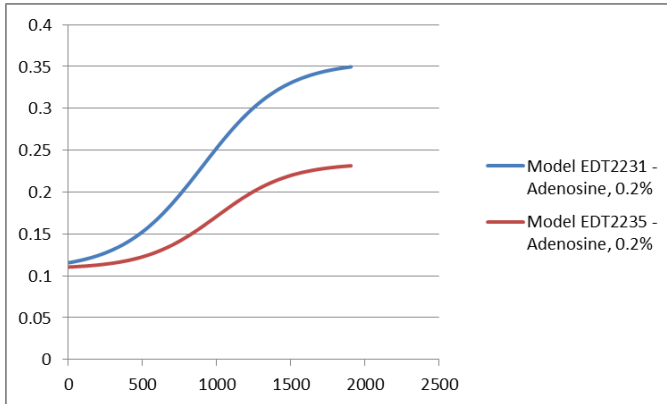
Models + Actual data



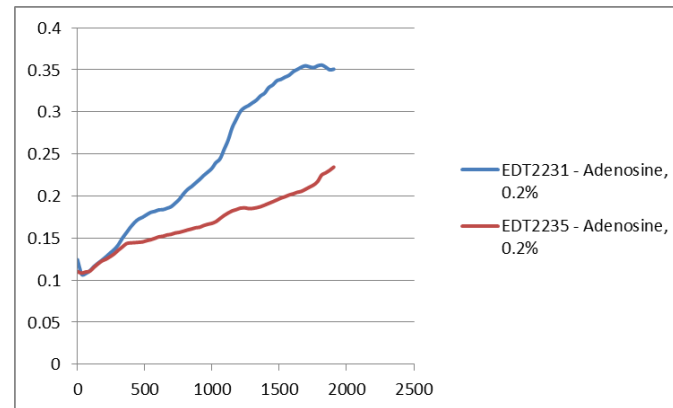
Chosen Models

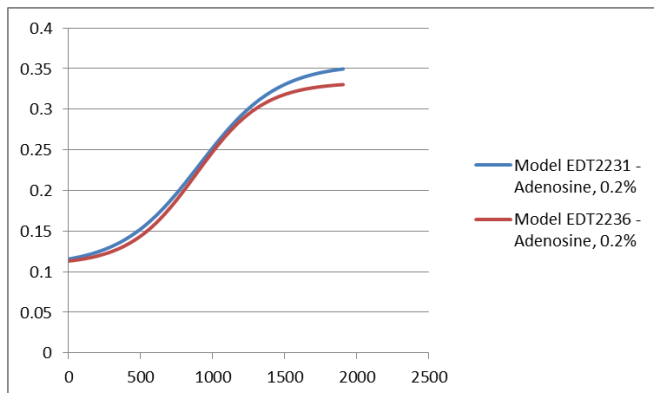


Comparisons

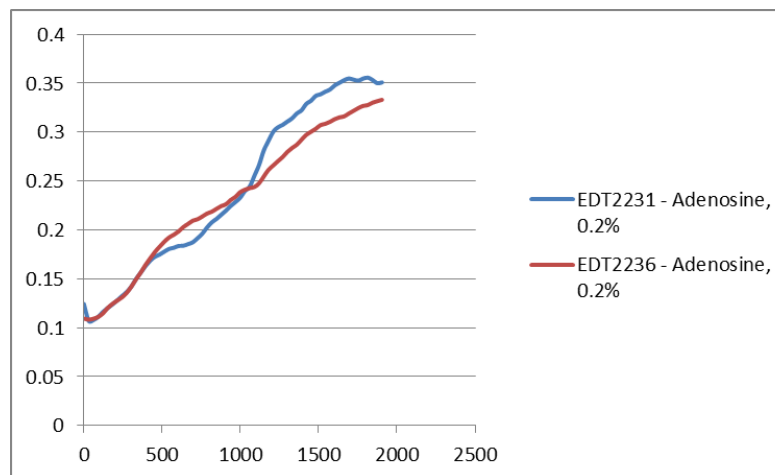


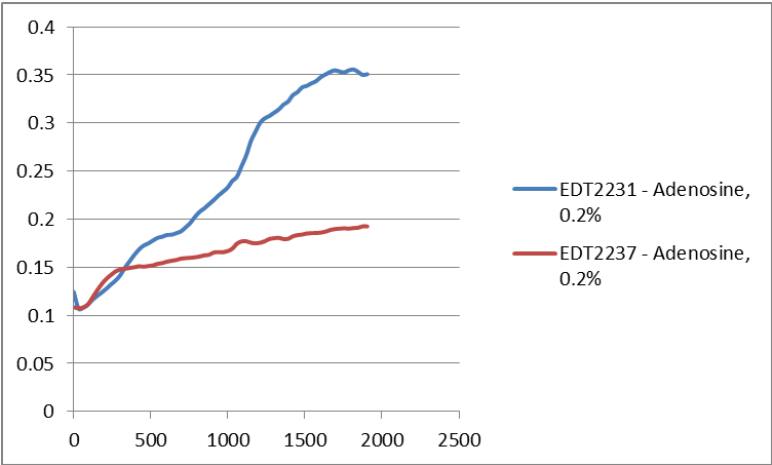
Proportion	Score	Predicted Performance
1.369	- 2	worst





Proportion	Score	Predicted Performance
1.039805	0	same





Proportion	Score	Predicted Performance
1.41373	- 2	worst

