

# Auer2010 - Correlation between lag time and aggregation rate in protein aggregation



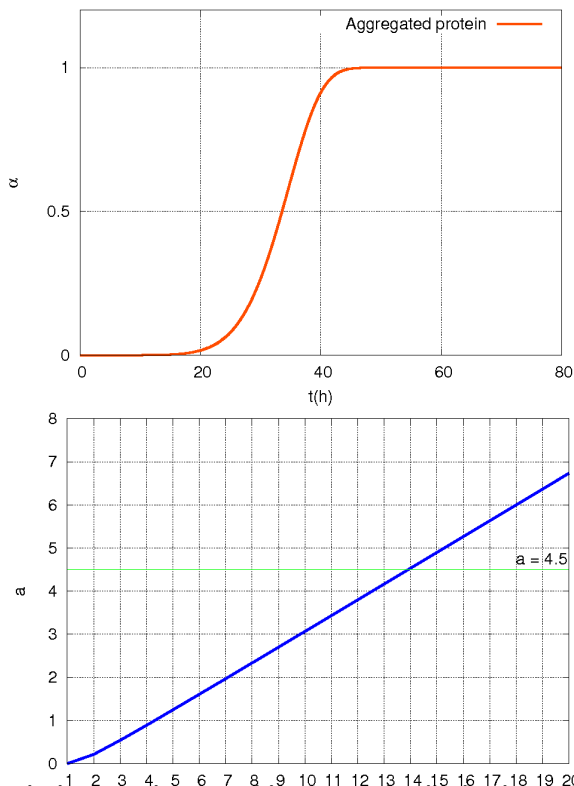
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## Curator's comment:

(added: 10 Nov 2014, 15:15:48,  
updated: 10 Nov 2014, 15:15:48)

Figure 1 (upper panel) and Figure 2 (lower panel) have been reproduced here.

Figure 1. Analysis of kinetic amyloid fibrils data for 80 hours protein aggregation.

Figure 2. Dependence of the parameter  $a$  on the kinetic index  $n$  in overall aggregation.

The simulation was done using Copasi

v4.12 (Build 81) and the plots were

generated using GnuPlot. The models

with simulation settings can be

downloaded from

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Build: [15671cb \(//bitbucket.org/biomodels/jummp-biomodels/commits/all?search=15671cb\)](https://bitbucket.org/biomodels/jummp-biomodels/commits/all?search=15671cb) | Wed, 14 Feb 2024  
21:14:41 +0000

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