### Analyzing & Testing

## **KINETICS NETZSCH** NEO Proven Excellence.

## Advantages and disadvantages of different thermokinetic approaches Unique and powerful features of Kinetics Neo Software

Dr. Elena Moukhina, Physicist, NETZSCH-Gerätebau GmbH"

NETZSCH Webinar – 29th April 2021

### Agenda



#### 1. Target of thermokinetic analysis

- 1.1 Brief overview
- 1.2 Measurements for kinetic analysis

#### 2. How to solve the problems? Different analysis approaches

#### 2.1 Degree of conversion

- 2.2 Arrhenius equation, activation energy, kinetic triplet
- 2.3 Overview of the model-free and model based approaches.
- 2.4 Why model-free is often not applicable.
- 2.5 Advantages and disadvantages
- 2.6 Limitations of solutions and prediction

#### 3. Unique Features of Kinetics Neo Software

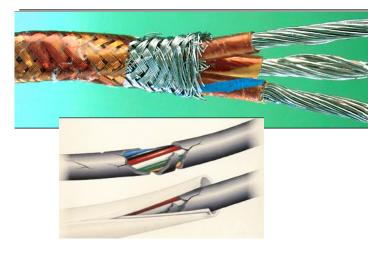
- 3.1 Fulfils all ICTAC kinetic recommendations
- 3.2 Numerical model free method
- 3.3 Special functions for curing
  - 3.3.1 Kamal-Sourour autocatalytic model
  - 3.3.2 Diffusion control, vitrification and TTT diagram
  - 3.3.3 Kinetic analysis of rheological data and prediction of dynamic viscosity
- 3.4 Crystallization kinetics
- 3.5 Multi-step model based method, including presentation of each individual reaction step
- 3.6 Small other features (Bezier baseline, export equations...)



### **1. Target of thermokinetics analysis** Brief overview of the problems solving by thermokinetics

### 1.1 Application fields



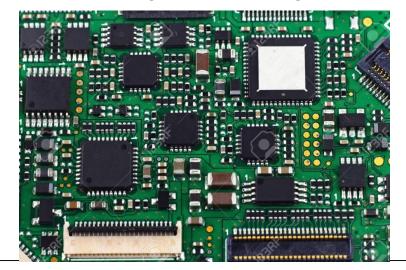


Life time predictions Recycling, pyrolysis Thermal stability



#### Curing, cross-linking







Sintering, ceramics



### 1.1 Purposes of kinetic analysis

Academic Chemical mechanism is the **subject** of study

- Find and describe the kinetic mechanism

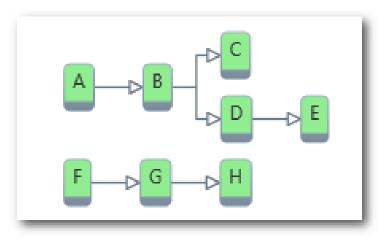
 $A \rightarrow B$ reactant product



- **Predict** conversion(Temp,time) and reaction rate for given temperature program
- **Optimize** industrial processes: decrease production time and costs and improve the quality of product

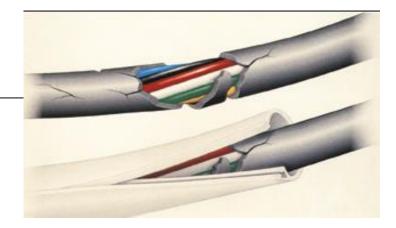


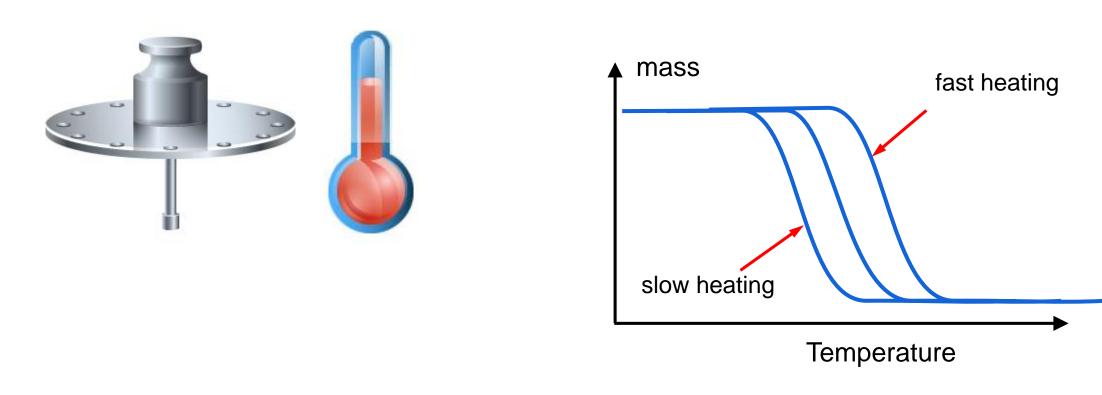






1.2 Measurements for kinetic analysis: Thermogravimetry



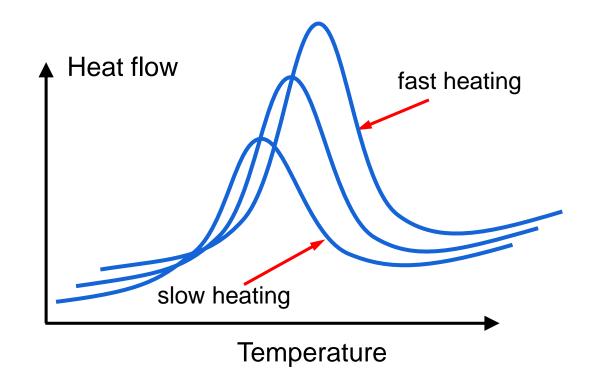


Thermogravimetry: mass change is measured during heating

1.2 Measurements for kinetic analysis: Differential scanning calorimetry



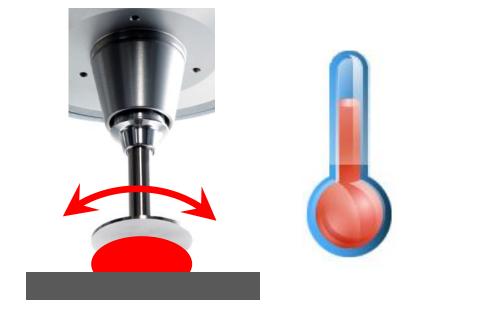


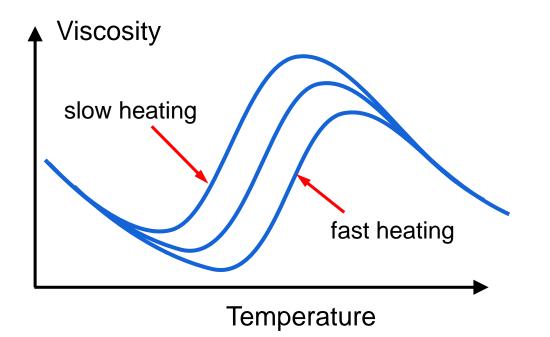


Differential scanning calorimetry: heat flow is measured during temperature





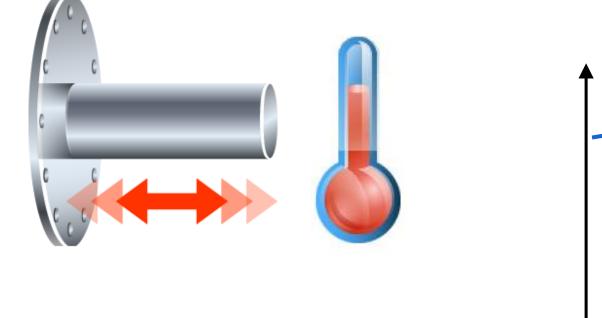


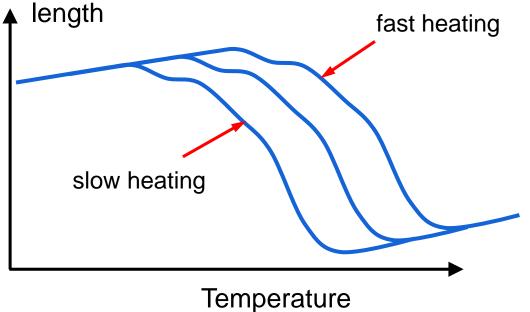


Rheometry: viscosity is measured during temperature

1.2 Measurements for kinetic analysis: Dilatometry







Dilatometry: length change is measured during heating



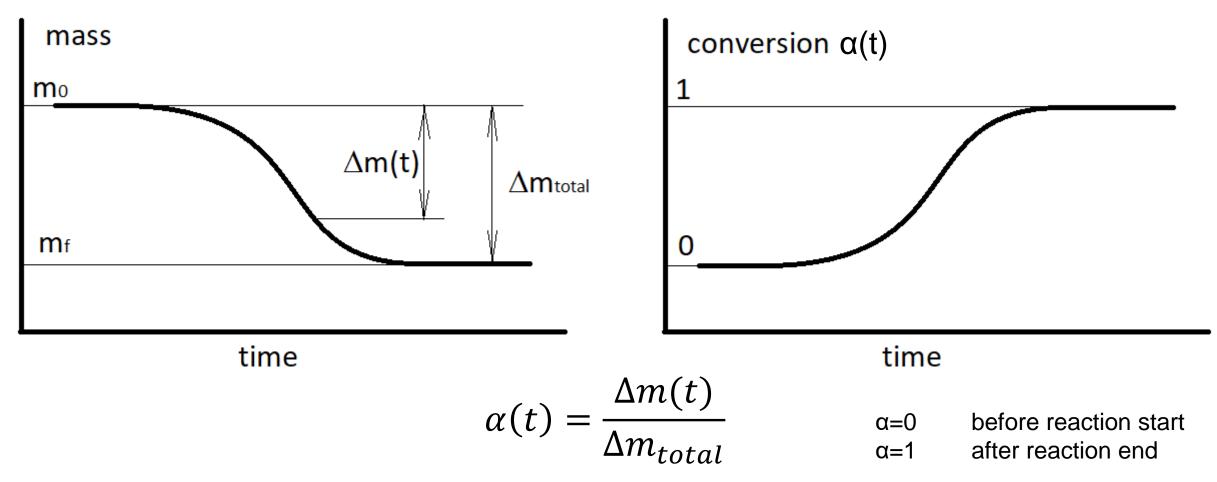
# 2. Different analysis approaches

Different approaches, advantages, disadvantages, limitations

### 2.1 Degree of conversion α(t) for TGA data (extent of conversion, conversion, extent of reaction)

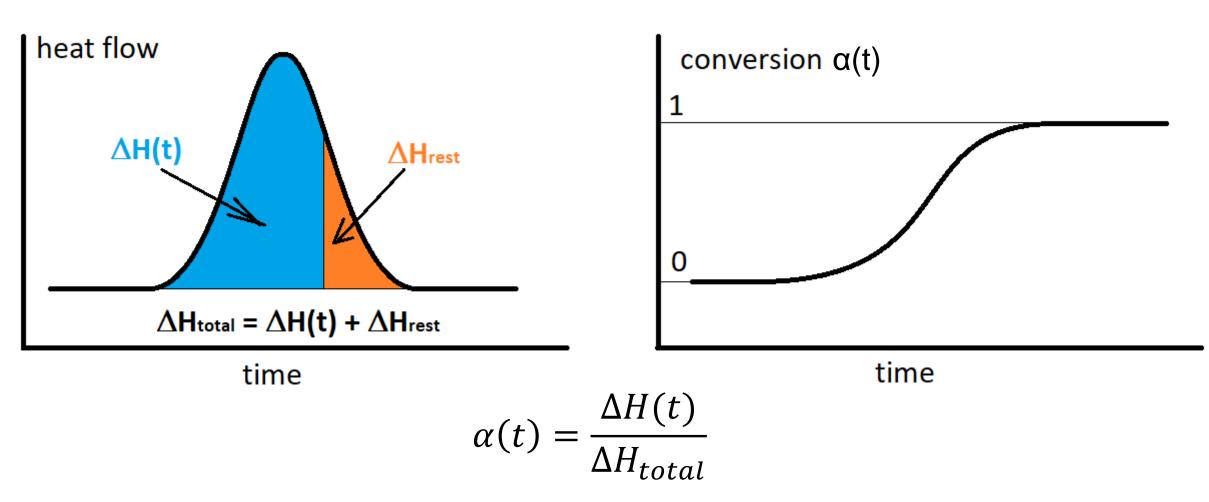


Commonly denoted by  $\alpha$  and defined as the ratio of the partial to total change of a physical property.



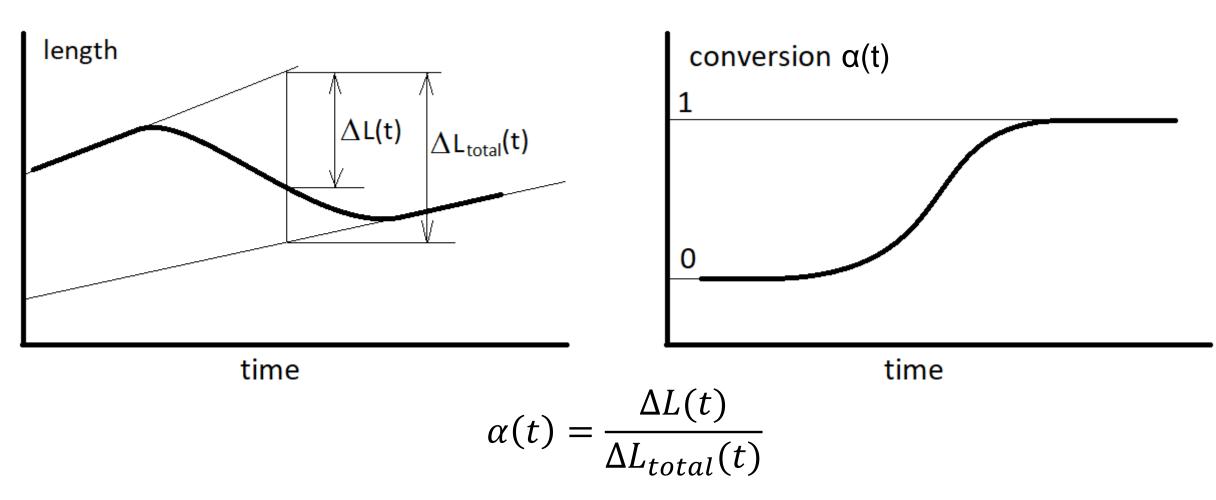
TGA: Conversion is the ratio of the partial mass loss at given time point to the total mass loss at the final time point





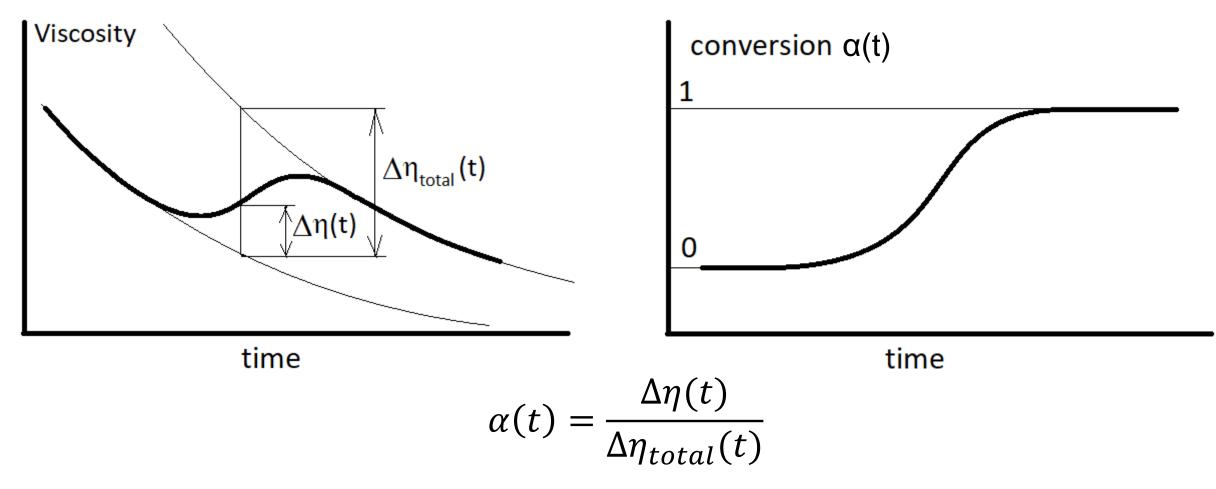
DSC: Conversion is the ratio of the partial enthalpy change at given time point to the total enthalpy change at the final time point





DIL: Conversion is the ratio of the partial length change at given time point to the total length change between extrapolated baselines at the this point

### 2.1 Conversion $\alpha(t)$ for viscosity data during heating



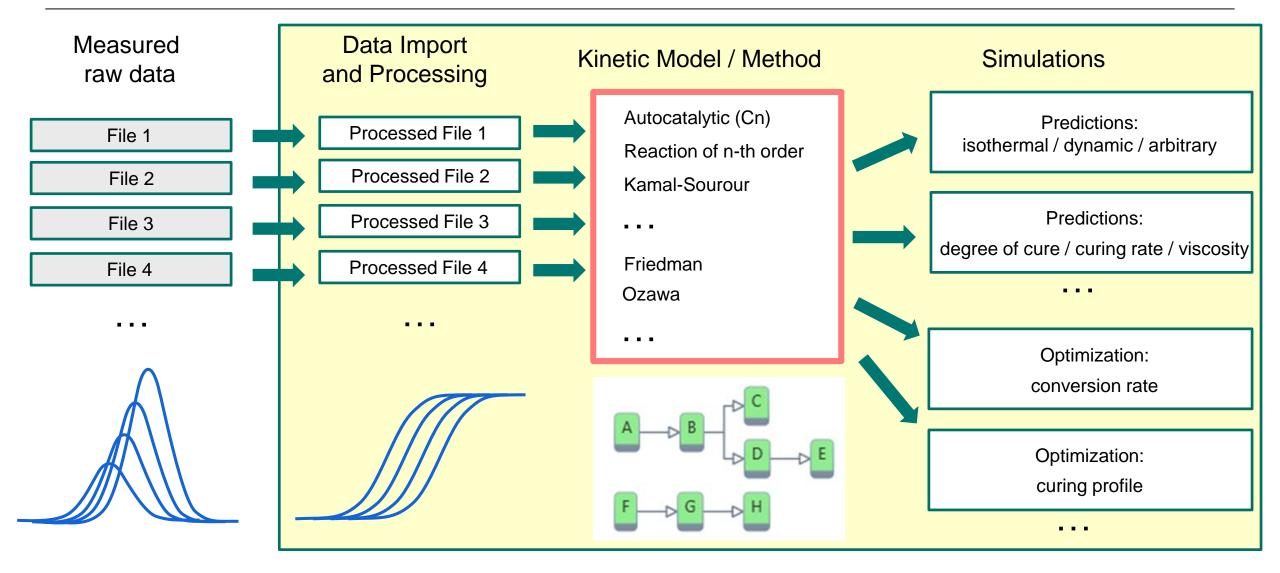
Viscosity: Conversion is the ratio of the partial viscosity change at given time point to the total viscosity change between extrapolated baselines at the this point

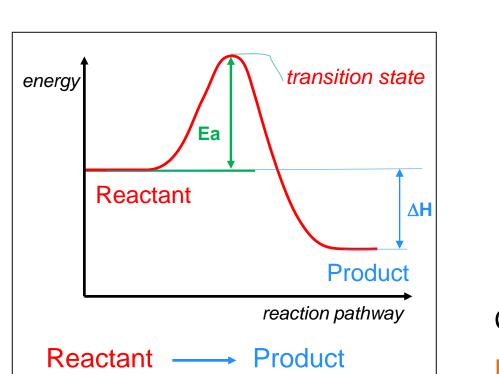
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### Steps to solve Kinetic Tasks in Kinetics Neo







Arrhenius equation (1889) for reaction rate:

$$\frac{d\alpha}{dt} = A \exp\left(\frac{-E_A}{RT}\right) f(\alpha)$$

Conversion  $\alpha$ : degree of conversion, changing from 0 to 1

Pre-exponent A: collision frequency [1/s]

Activation energy Ea [kJ/mol]

 $f(\alpha)$  Reaction type (nth order, autocatalysis, nucleation ...)

R: gas constant 8.31 [J/(gK)] T: absolute temperature [K] T[K]=T[°C]+273.15

EZ5CH

### 2.3 Approaches: model free and model based



#### Model free $A \rightarrow B$

 $\alpha$  – degree of conversion

$$\frac{d\alpha}{dt} = A(\alpha) \cdot f(\alpha) \cdot exp\left(\frac{-E_A(\alpha)}{RT}\right)$$

Unknown:  $Ea(\alpha)$  and  $A(\alpha)$ A( $\alpha$ ) can be found only with assumption of f( $\alpha$ )

#### Assumptions:

- 1. Only one kinetic equation
- 2. Ea and A depend on  $\alpha$
- Reaction rate at the same conversion is only a function of temperature
- 4. Total effect (total mass loss or total peak area) must be the same for all curves
- 5. Changes of mechanism should be at the same conversion value

### $\textbf{Model based} \qquad \textbf{A} \rightarrow \textbf{B} \rightarrow \textbf{C} \rightarrow \dots$

- a concentration of A
   b concentration of B
- c concentration of C

$$\frac{d(\mathbf{a} \to \mathbf{b})}{dt} = A_1 \cdot f_1(\mathbf{a}, \mathbf{b}) \cdot exp\left(\frac{-E_{A1}}{RT}\right)$$
$$\frac{d(\mathbf{b} \to \mathbf{c})}{dt} = A_2 \cdot f_2(\mathbf{b}, \mathbf{c}) \cdot exp\left(\frac{-E_{A2}}{RT}\right)$$

The number of unknown kinetic triplets equals the number of the steps

#### Assumptions:

- 1. Reaction consists of **several individual reaction steps** with own equations.
- 2. All kinetic parameters which are the constant values
- 3. The **total signal** is the **sum** of the signals of the single reaction steps having **own weight**

### 2.3 Model free methods in Kinetics Neo



#### Created in last century before the modern possibilities of personal computers

#### **One-point model free methods**

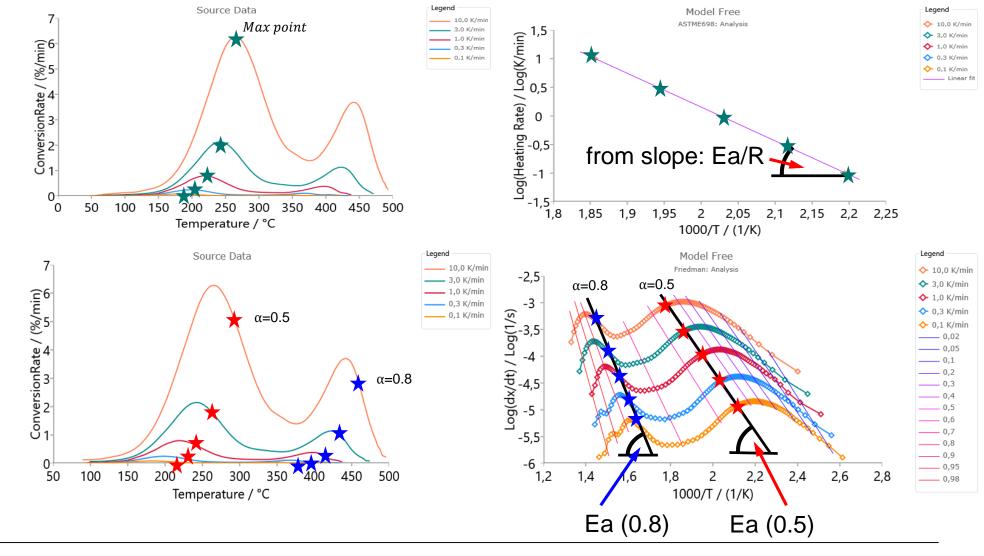
- ASTM E698
- ASTM E2890
- ASTM E1641
- ASTM E2070

Result: Value Ea

#### Multi-points model free methods

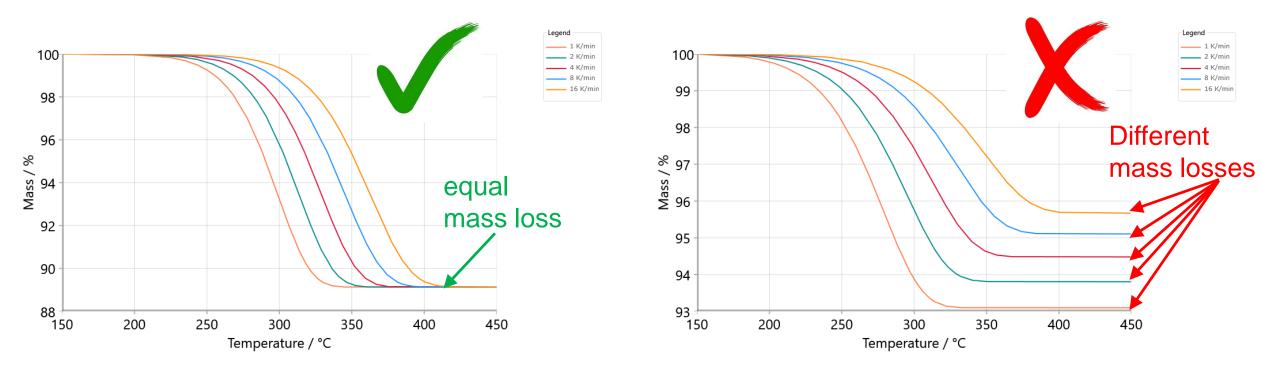
- Ozawa-Flynn-Wall (1965)
- Kissinger-Akahira-Sunose (1956)
- Friedman method (1966)
- Vyazovkin for heating (1996)
- Numerical optimization

Result: **Function Ea(α)** 





Total effect (total mass loss or total peak area) must be the same for all curves



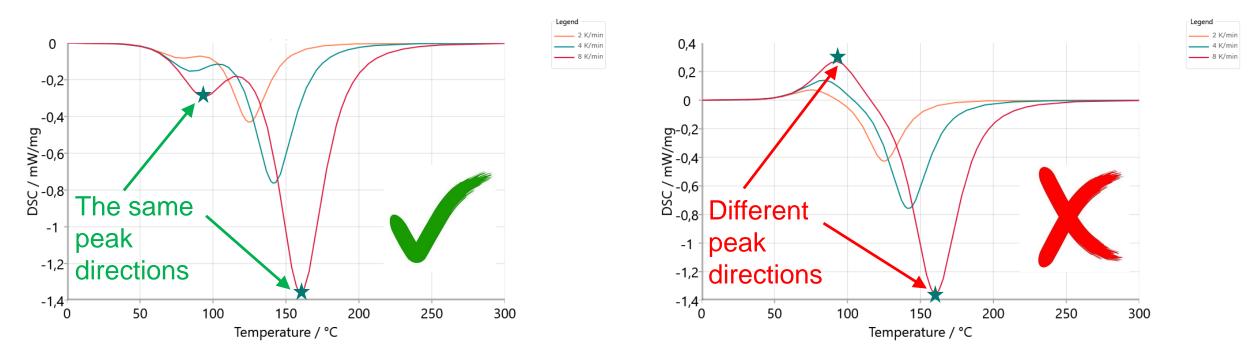
Model free is applicable

Model free is **not** applicable Solution: Model based method with competitive steps



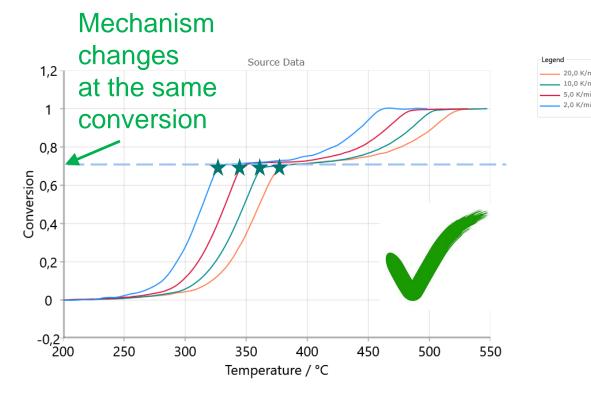


There is no reaction steps of different directions (e.g. exothermal and endothermal, mass loss and mass gain)



Model free is applicable

Model free is **not** applicable Solution: Model based method <u>with steps contributions of different signs</u>



Mechanism changes at different conversions Legend 15K/min 9 K/min 0,8 5 K/min 2 K/min ອັ **0**,4 0,2 0 -0.2 350 400 450 500 550 Temperature / °C

Model free is applicable

Model free is **not** applicable Solution: Model based method with competitive steps

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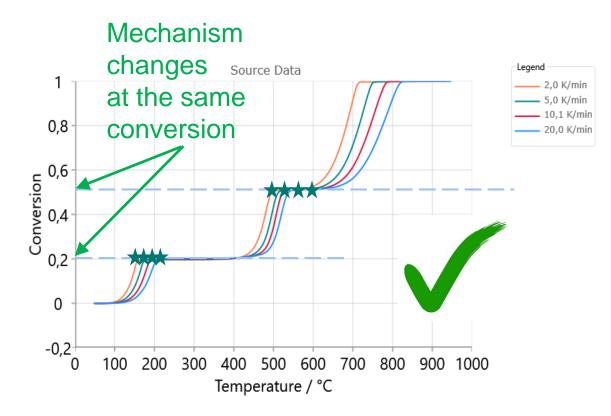
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Changes of mechanism should be at the same conversion value



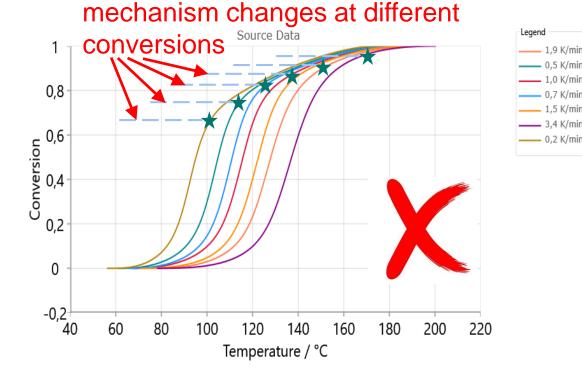


Changes of mechanism should be at the same conversion value



Model free is applicable

Reaction with diffusion control:



Model free is **not** applicable Solution: Model based method with diffusion control

### 2.5 Advantages and disadvantages







### 2.5 Advantages and disadvantages



### **Model free**

#### Advantages:

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- 1. Very fast, one click
- 2. Easy analytical methods, mostly can be done in Excel

#### Disadvantages:

- 1. Does not work for **mixtures** (independent parallel steps)
- 2. Does not work for competitive steps
- 3. Does not work for **curing with Kamal Sourour** reaction
- 4. Does not work for curing with diffusion control
- 5. Does not work for **crystallization**
- 6. Does not work for the steps of different directions
- 7. Does not work for highly overlapping steps
- 8. Does not work if reaction mechanism changes with heating rate
- 9. Has no any information about intermediate steps and reactants

### Model based

Advantages:



- 1. Works for **mixtures** (independent parallel steps)
- 2. Works for **competitive steps**
- 3. Works for curing with Kamal Sourour reaction
- 4. Works for curing with diffusion control
- 5. Works for crystallization
- 6. Works for the steps of different directions
- 7. Works for highly overlapping steps
- 8. Works if reaction mechanism changes with heating rate
- 9. Has information about intermediate steps and reactants

#### Disadvantages:

- 1. Requires elementary chemical knowledge about the process
- 2. Number of steps usually should not exceed the number of visible peaks

It is necessary to use ICTAC kinetic recommendations

https://doi.org/10.1016/j.tca.2020.178597



# 3. Unique features of Kinetics Neo software

- 3.1 Fulfils all ICTAC kinetic recommendations
- 3.2 Numerical model free method
- 3.3 Special functions for curing (Kamal-Sourour, Diffusion control, TTT diagram, viscosity prediction)
- 3.4 Crystallization kinetics
- 3.5 Multi-step model based method, including presentation of each individual reaction step
- 3.6 Small other features (Bezier baseline, export equations...)

### All Kinetics Neo features: <u>https://kinetics.netzsch.com/en/docs</u>



Documents - Kinetics Neo × +	Te <u>kin</u>
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A Docs	Ur
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Kinetics Neo: Docu	
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Brochure	
Technical Datasheet	
License Agreement	
License Agreement	Kii
Software Lifecycle Policy	
Literature about Kinetics: Theory, Metho	
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Model-Free Methods	
Definition	Model-free analysis allows to find the activation energy of the reaction without assumption of a kinetic model for it.
Methods based on a single conversion	ASTM E698     ASTM E2890     ASTM 1641     Isothermal Arrhenius for time-to-event
Conversion-dependent methods	Dynamic Arrhenius for failure temperature Friedman Ozawa-Flynn-Wall (OFW) Kissinger-Akahari-Sunose (KAS) ASTM E2070(A) for isothermal data Vyazovkin for dynamic data Numerical Optimization The numerical anodel-free method ensures fast determination of the best model-free solution to achieve best the agreement between simulated and experimental curves
Results	Analysis graph     Plot of activation energy vs degree of conversion     Plot of pre-exponential factor vs degree of conversion     Master plot f(q)     Conversion fit for signal, conversion and conversion rate
- Model-Based Methods	
Multi-step analyzing engine	Model-free methods allow for analyzing only one-step kinetic processes. However, approxi- mately 95% of all chemical reactions are multi-step reactions. This requires the multi-step analyzing engine of the Kinetics Neo software.
Unique features	The model-based kinetic analysis is based on an unlimited number of models including an unlimited number of reaction steps where the individual steps are linked as independent, parallel, competing or following.
Kinetic models	Visual arceation of a kinetic model. Visual adding, removing or editing of each reaction step. Visual adjustment of position, contribution, activation energy and peak shape of each step. Optimization of kinetic parameters for one individual step. Optimization of kinetic parameters for the complete kinetic model.
Reaction types	Each individual reaction step in each model can be one of 19 reaction types including Reaction of 1 <sup>st</sup> , 2 <sup>ad</sup> and n-th order without autocatalysis Reaction of 1 <sup>st</sup> , 2 <sup>ad</sup> and n-th order without autocatalysis Reaction of 1 <sup>st</sup> , 2 <sup>ad</sup> and n-th order without autocatalysis reactions 2-/3-dimensional phase boundary reactions 1-/2-/3-dimensional diffusion (Jander's type and Ginstling-Brounstein) Prout-Tompkings reaction 2-/3-dimensional phase boundary reactions 2-/3-dimensional autoleation according to Avrami Reactions with diffusion control Crystallization according to the Nakamura equation using the Hoffman-Lauritzen Theory Models for glass transition function for diffusion control D iB enedetto model Splines
Kinetic results	The software determines the kinetics model including  Number of reaction steps Step contribution to the total effect Standard parameters for each reaction step: Reaction type Activation energy Reaction of ruce and parameters for some reaction types Order of autocatalysis Dimension of nucleation or diffusion Parameter for crystallization Dimension of nucleation Melting temperature and glass transition temperature Nakamura parameter K <sub>G</sub> Correlation coefficient
Stastical results	Correlation coefficient Sum of the squares of deviations Mean residual t-value Durbin-Watson value Durbin-Watson test F-test for fit quality F-test for the number of steps
	2 Prescior the number of steps

#### Technical Datasheet: **Kinetics Neo 2.5** <u>kinetics.netzsch.com</u>



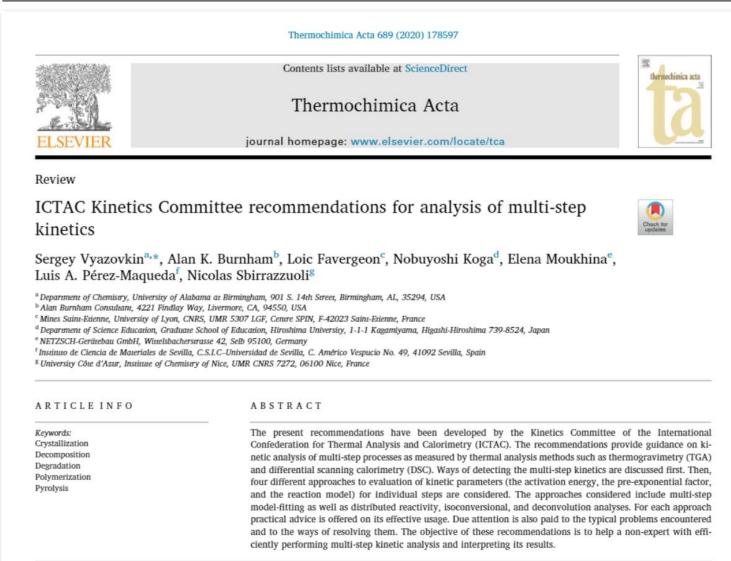
Predictions for several isothermal temperatures				
Predictions for given conversion at several isothermal temperatures				
Predictions for several heating rates for reactions or for several cooling rates for crystallization				
Prediction for a user-defined sequence of dynamic and isothermal segments with the possibility of export/import of multi-step program to/from text file				
Prediction of a step-iso temperature program, representing a stepwise temperature increase				
Prediction of a modulated temperature program which is the sum of an underlying constant temperature or constant heating and a sinus-shaped temperature oscillation				
Calculation of the adiabatic temperature increase for various initial temperatures.				
Find the start temperature for adiabatic process at maximum heating rate in 24 hours				
Prediction for the real atmospheric temperatures for a selected point on the Earth, from the selected day of the year, for the selected durations, which is usually several months or years.				
Prediction for temperature program loaded from external text file (ASCII of CSV) which includes time and temperature columns				
This method uses standard fire presets loaded into external temperature profile prediciton				
Time-Temperature-Transition diagram for reactions with diffusion control.				
After input of the temperature program by the user, the software will make a simulation of the system behavior. The following values can be simulated: Measurement output (signal) Conversion Conversion rate Concentration of the reactants for model-based method only Presentation of the reactants for model-based method only Presentation of conversion rate as the sum of individual reaction steps Predictions of raw viscosity for viscosity project type The simulated values can be presented: Curves as a function of time Curves as a function of temperature Table with simulated values, time and temperature				
Finding a temperature program for a given system behavior (optimization). It is the typcial ques- tion arising during a production process. The temperature program for optimal time and quality must be found. Without the Kinetics Neo software, it is necessary to make an adjustment of the temperature program by hand and to measure several times while hoping to achieve the expected signal curve. The software – in contrast – saves time and finds such a temperature program • For a given reaction rate • For a given output signal • For a given rate of the final product production • For user-defined function of conversion versus time				
Temperature program     Measurement output (signal)     Conversion     Conversion     Conversion rate     Concentration of the reactants, for model-based method only     Reaction rate for individual reaction steps, for model-based method only     Presentation of conversion rate as the sum of individual reaction steps     Prediction of raw viscosity for viscosity project type     Sum of peaks presents the total conversion-rate curve as the sum of the individual reaction     peaks				

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### 3.1 Unique: Kinetics Neo fulfils ICTAC kinetics recommendations



International Confederation for Thermal Analysis and Calorimetry

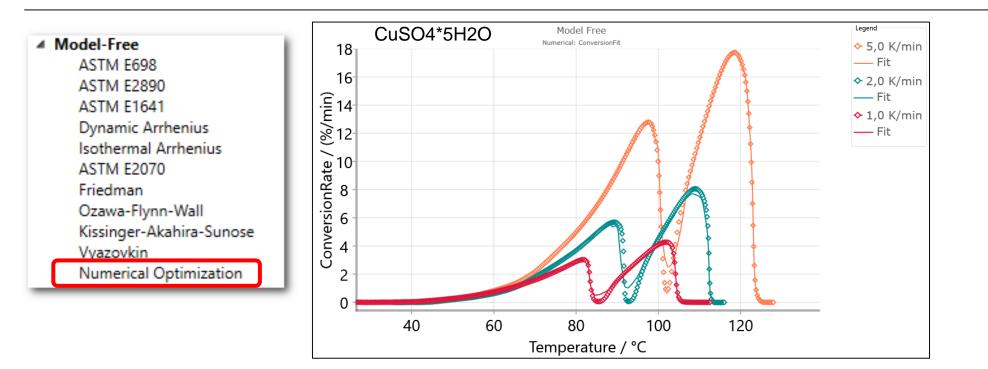




- Model free analysis
- Multi-step model-fitting (model based)
- Diffusion control for curing
- Crystallization kinetics
- Kamal model for curing
- Deconvolution analysis (sum of peaks)
- Distributed reactivity analysis

### 3.2 Unique: Model-free methods, Numerical method





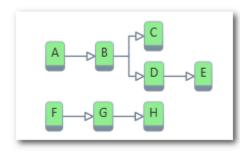
#### Please always check the fit visually

Method/Model	Fit To	R <sup>2</sup>	Sum of dev. squares	Mean Residual
Numerical	Signal	0,99655	2,092	0,023
Friedman	Signal	0,99237	4,612	0,030
Vyazovkin	Signal	0,97562	14,611	0,054
KissingerAkahiraSunose	Signal	0,87774	69,648	0,138
OzawaFlynnWall	Signal	0,85547	81,358	0,165
ASTME1641	Signal	0,71256	149,337	0,205
ASTME2890	Signal	0,70314	153,383	0,264
ASTME698	Signal	0,70182	153,947	0,264

### 3.3 Unique: Model based analysis

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#### F1, 1st order

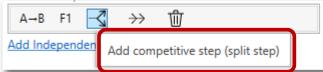
- F2, 2nd order
- Fn, n-th order
- R2, 2D phase bound.
- R3, 3D phase bound.
- D1, 1D diffusion
- D2, 2D diffusion
- D3, 3D diff. Jander
- D4, 3D diff.Ginstling-Brounstein
- **B1**, Prout-Tompkins
- Bna, expanded Prout-Tompkins
- C1, 1st order with autocat.
- Cn, n-th order with autocat.
- Cnm, n-th order, m-Power with autocat.
- KS, Kamal-Sourour
- A2, 2D Avrami
- A3, 3D Avrami

An, n-Dim. Avrami

Nk, Nakamura crystallization

#### Modelling

- unlimited number of models
- unlimited number of reaction steps
- individual steps are linked as independently, parallel, competing or following
- Visual creation of a kinetic model
- Visual adding, removing or editing of each reaction step.
- Optimization of kinetic parameters for one individual step.
- Optimization of kinetic parameters for the complete kinetic model.
- Reaction types
  - Reaction of 1st, 2nd and n-th order without autocatalysis
  - Reaction of 1st, 2nd and n-th order with autocatalysis including Prout-Tompkins and Kamal-Sourour reactions
  - 2-/3-dim. phase boundary reactions
  - 1-/2-/3-dim. diffusion (Jander's type and Ginstling-Brounstein)
  - 2-/3-/n-dim. nucleation according to Avrami
  - Curing Reactions with diffusion control (using DiBenedetto model or splines for dependence Tg vs Alpha)
  - Crystallization according to the Nakamura equation using the Hoffman-Lauritzen theory



### 3.3.1 Unique: Kinetic Modelling for Curing

Chemical process is **generally** described by Arrhenius equation:

$$\frac{d\alpha}{dt} = A \cdot f(\alpha) \cdot exp\left(\frac{-Ea}{RT}\right)$$

Curing can be described by the equation Kamal-Sourour for autocatalytic reaction:

$$\frac{d\alpha}{dt} = \underbrace{A \cdot (1 - \alpha)^n \cdot \exp\left(\frac{-E_{a1}}{RT}\right)}_{\text{n-th order}} + \underbrace{A \cdot K \cdot (1 - \alpha)^n \cdot \alpha^m \cdot \exp\left(\frac{-E_{a2}}{RT}\right)}_{\text{autocatalysis}}$$

This equation with its parameters A,  $E_{a1}$ ,  $E_{a2}$ , K, m, n, is the kinetic model.

Kinetic parameters are found from the best fit for all experimental data

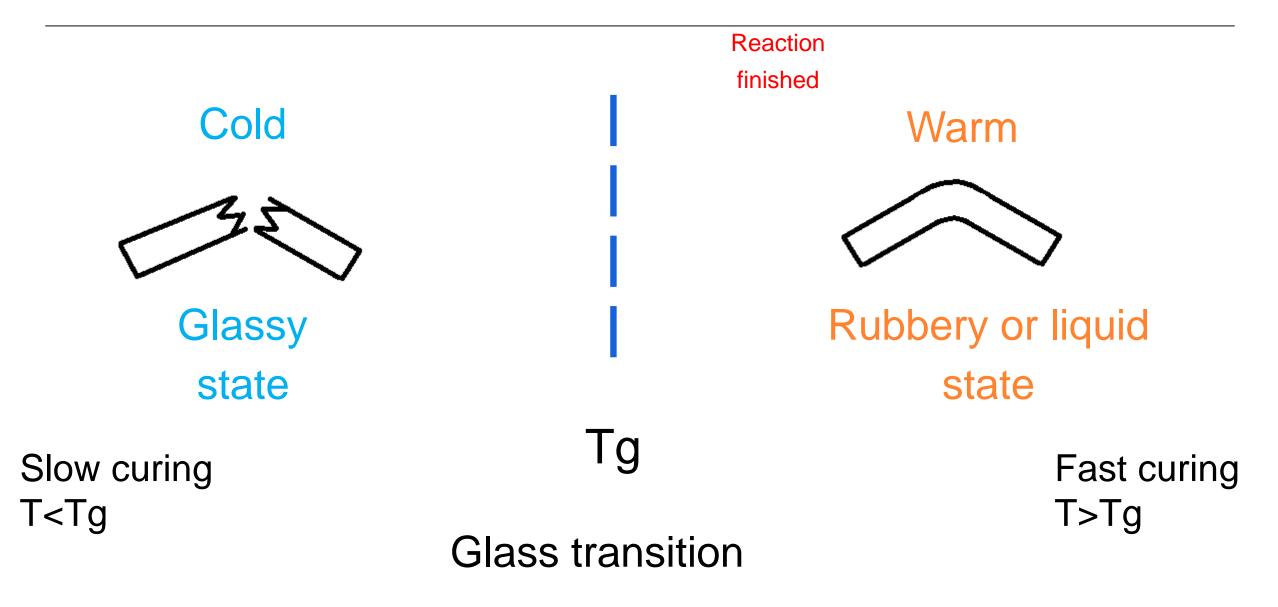




n-th order

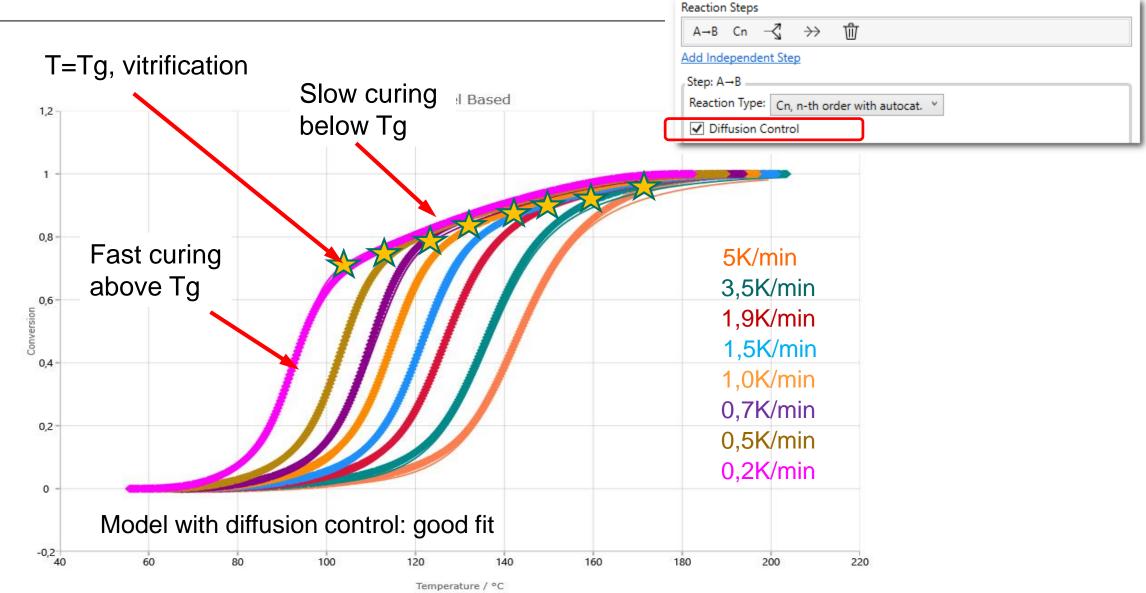
autocatalysis





### 3.3.2 Unique: Experiment and Model Fit for diffusion control

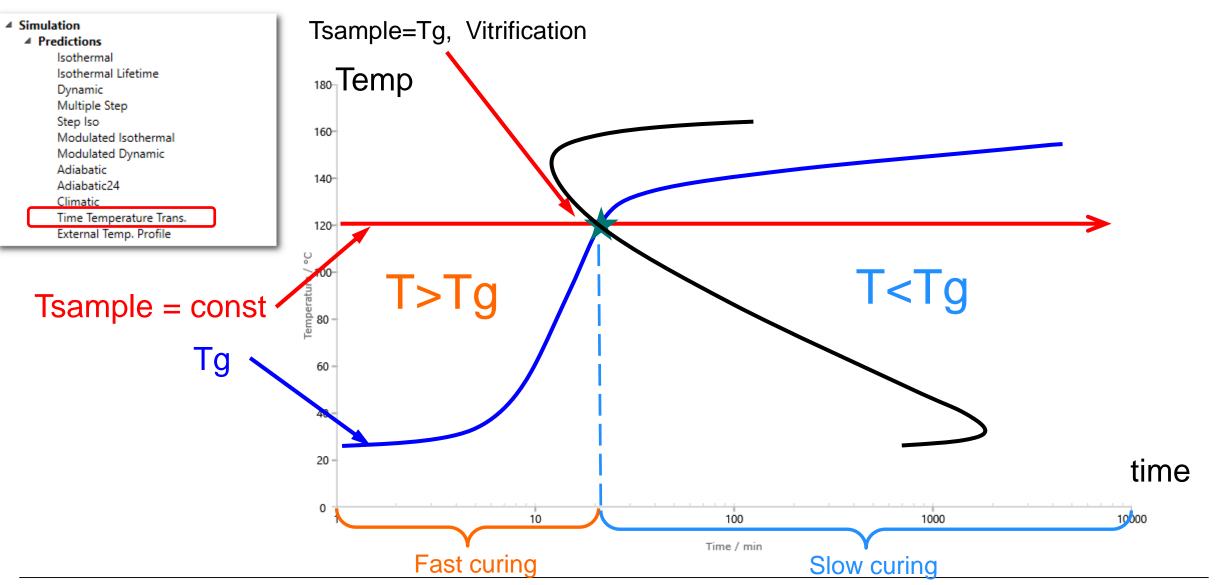




#### See Theory of diffusion control in

S. Vyazovkin, A.K. Burnham, L.Favergeon, N.Koga, E.Moukhina, L.A.Perez-Maqueda, N. Sbirrazuoli. Thermochimica Acta 689 (2020) 1785977, ICTAC Kinetics Committee recommendations for analysis of multi-step kinetics

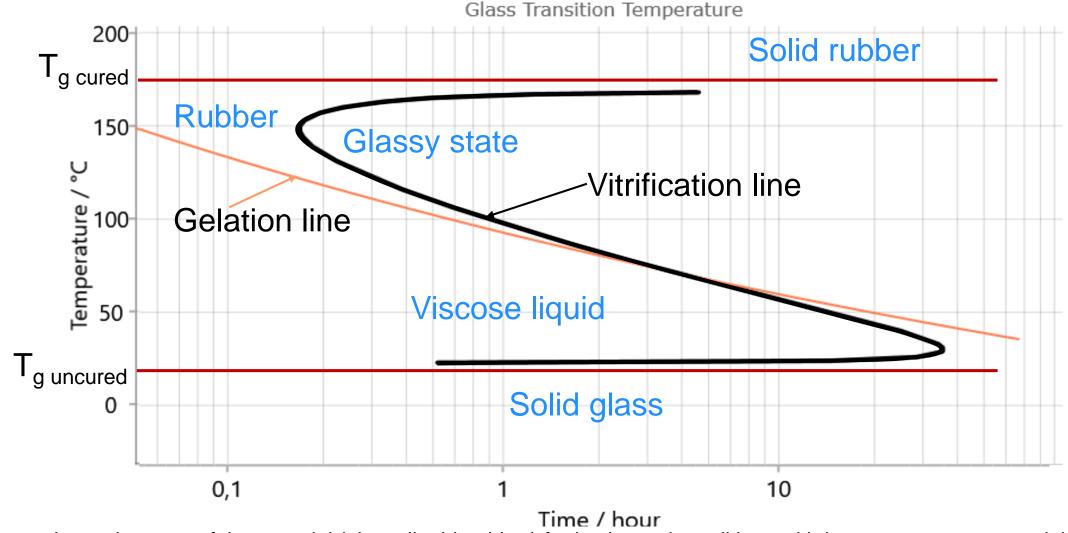
### 3.3.2 Unique: Isothermal Predictions of glass transition for Time-Temperature-Transition diagram



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### 3.3.2 Unique: Time-Temperature-Transition diagram for curing

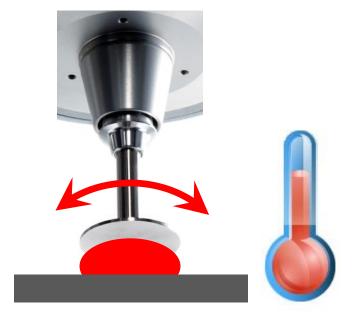




TTT Diagram shows the state of the material (glass, liquid, rubber) for isothermal conditions with known temperature and time

# 3.3.3 Unique: Kinetic analysis of rheological data and prediction of dynamic viscosity



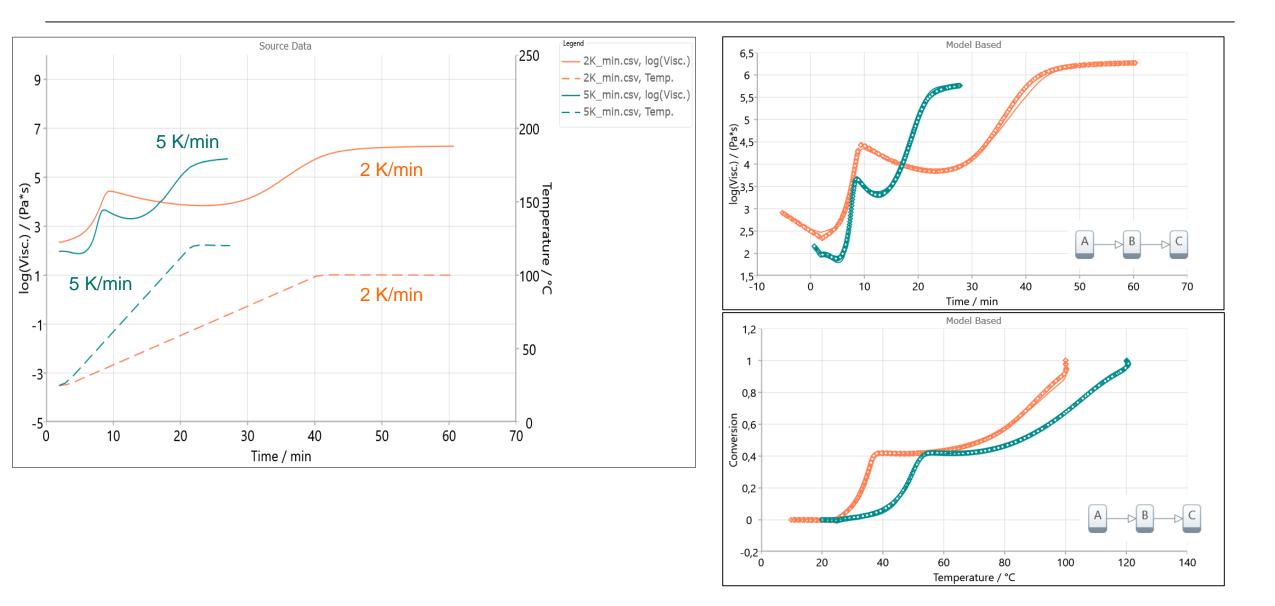




2-component adhesive"Very fast setting": 5 minutes...Complete curing 24 hours

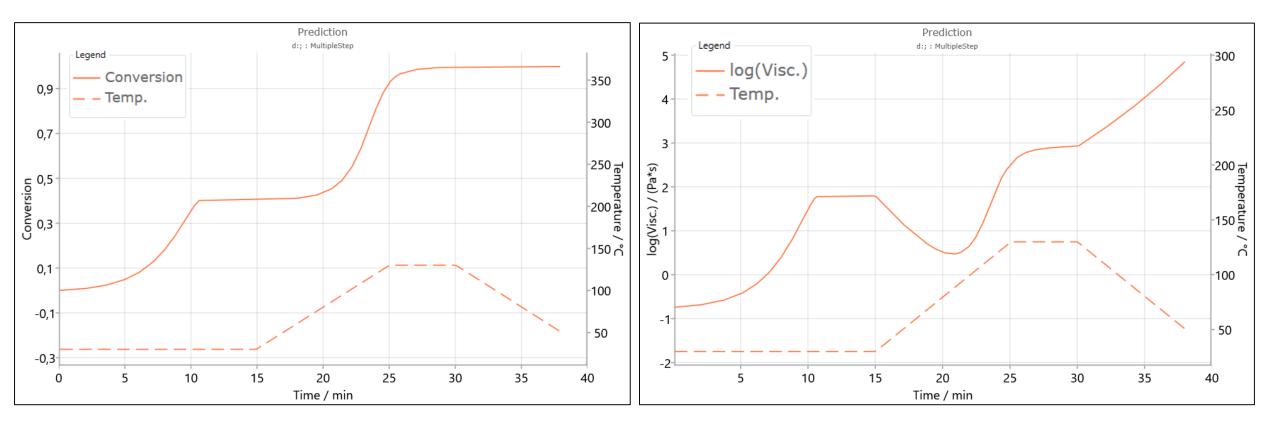
### 3.3.3 ARALDITE, Rheometry measurements for kinetic analysis





### 3.3.3 Prediction of conversion and viscosity for complex temperature program

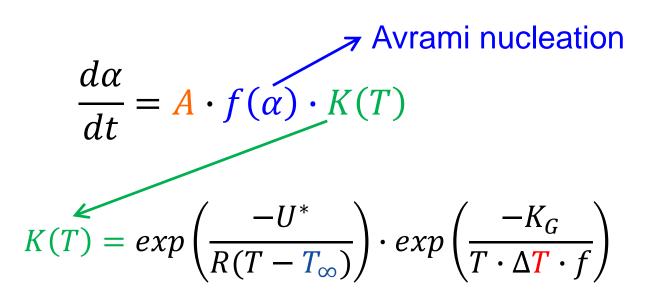




### 3.4 Unique: Crystallization kinetics:



non-Arrhenius approach: Nakamura und Hoffman-Lauritzen



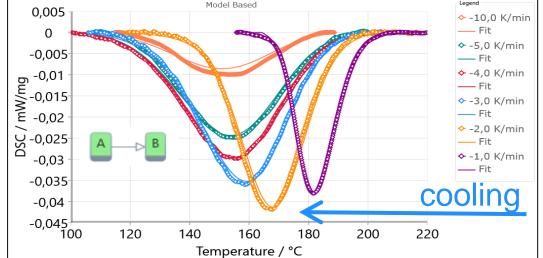
activation energy of segmental jump in polymers, this parameter has universal value 6.3kJ/mol

- K<sub>G</sub> kinetic parameter for nucleation
- $\Delta T=Tm-T$  undercooling from the equilibrium *melting point Tm*
- T<sub>w</sub>=Tg-30 temperature at which crystallization transport is finished, this temperature is 30K below the *glass transition temperature Tg*.

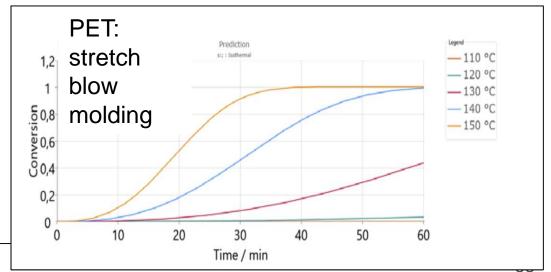
f=2T/(T+Tm) correction factor

U\*

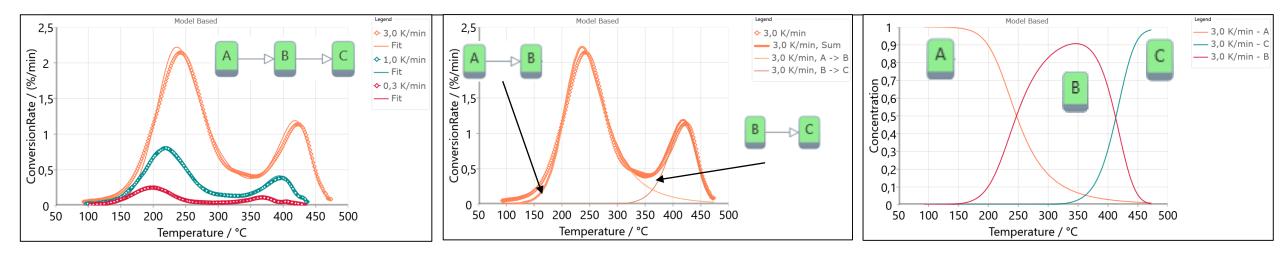
Measurement and Model for PET: Model Based



#### **Predictions: Conversion**



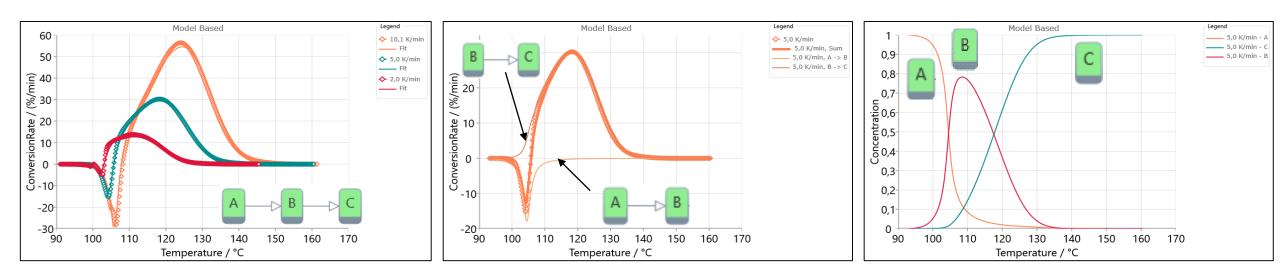
### 3.5 Unique: Model based method with reaction steps and concentrations **NETZSCH**



Kinetic model

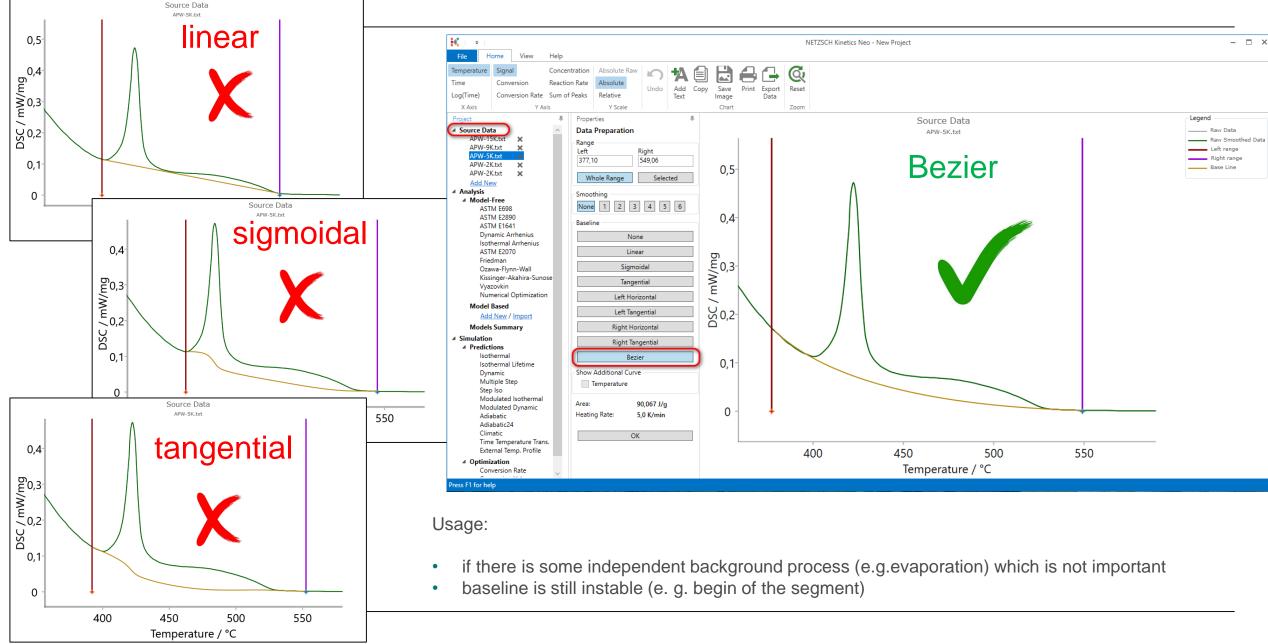
Individual reaction steps

#### Concentration of reactants



### 3.6 Unique: Bezier baseline for DSC with background process





### 3.6 Unique: Export equations



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	- File Edit Format View Help	
File     Home     View     Help	Project: Glucose_Analysis.kinx Model: t:	
Temperature     Signal     Concentration     Absolute Raw       Time     Conversion     Reaction Rate       Log(Time)     Conversion Rate Sum of Peaks     Mobilitie       X Avis     Y Avis	Model Scheme: A-BTC D	
Arady 1FrozenicModel to the second se	<pre>Model Reaction Steps: A → B B → C B → D Concentration Equations: da/dt=-d(a-&gt;b)/dt db/dt=d(a-&gt;b)/dt-d(b-&gt;c)/dt-d(b-&gt;d)/dt dc/dt=d(b-&gt;c)/dt dd/dt=d(b-&gt;c)/dt dd/dt=d(b-&gt;d)/dt Balance Equation: Mass=InitialMass-TotalMassChange*[Contribution(a-&gt;b)*Integral[d(a-&gt;b)/dt]dt + Contribution(b-&gt;c)*Integral Step: A → B</pre>	
Add New / Import Add New / Import Models Summary 4 Simulation 4 Predictioner Press F1 for help Reaction Type: Cn, n-th order with autocat. * Equation d(a->b)/dt=PreExp*a^n*(1+AutocatPreExp*b)*Exp[- ActivationEnergy/(RT)] Press F1 for help	<pre>Reaction Type: Cn Equation: d(a-&gt;b)/dt=PreExp*a^n*(1+AutocatPreExp*b)*Exp[-ActivationEnergy/(RT)] ActivationEnergy: 98,633 kJ/mol Log(PreExp): 7,708 Log(1/s) ReactOrder n: 1,754 Log(AutocatPreExp): 0,686 Contribution: 0,268</pre>	
These equations easy in FEM (Finite Element Method) software to simulate	Step: B → C Reaction Type: Fn Equation: d(b->c)/dt=PreExp*b^n*Exp[-ActivationEnergy/(RT)] ActivationEnergy: 1,000 kJ/mol	
<ul><li>Temperature distributions</li><li>Concentration distributions</li></ul>	ActivationEnergy:     1,000 kJ/mol       Log(PreExp):     -15,713 Log(1/s)       ReactOrder n:     8,067       Contribution:     0,726	
for the volume with any complex geometry	Step: B → D Reaction Type: Fn Equation: d(b->d)/dt=PreExp*b^n*Exp[-ActivationEnergy/(RT)]	
kinetics.netzsch.com	Ln 1, Col 1         100%         Windows (CRLF)         UTF-8	

### **NETZSCH Kinetics Neo Web Site including User Guides**

### https://kinetics.netzsch.com

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Trial Version 30 days

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# **KINETICS** NEO is used to analyze kinetics of temperature dependent chemical processes.

#### Analysis

- one mathematical kinetic description for several measurements
- different approaches: model free (11 methods) and model based (unlimited number of models)
- model based: individual reaction steps, concentrations, kinetic triplet for each step, reaction mechanism
- standard reaction types (n-th order, Autocatalysis, diffusion, Avrami nucleation)
- unique reaction types (Kamal-Sourour, diffusion control, Nakamura nucleation, Hoffman-Lauritzen theory)
- Standard data types like TG or DSC
- Unique data types (Dilatometry, Rheometry, DEA)

#### **Predictions and optimizations**

- optimization of industrial chemical processes like debinding, curing, sintering
- standard predictions (isothermal, heating, multi-step, user-defined, TD24, climatic for 100 weather stations)
- unique predictions of individual peaks, concentrations, Tg, TTT diagram, temperature presets like fire presets)
- prediction of viscosity or ion viscosity for new temperature profile
- Optimization of temperature profile for constant or predefined conversion rate

### You can rely on NETZSCH.



Thank you for your attention. Any questions? webinar\_ngb@netzsch.com

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