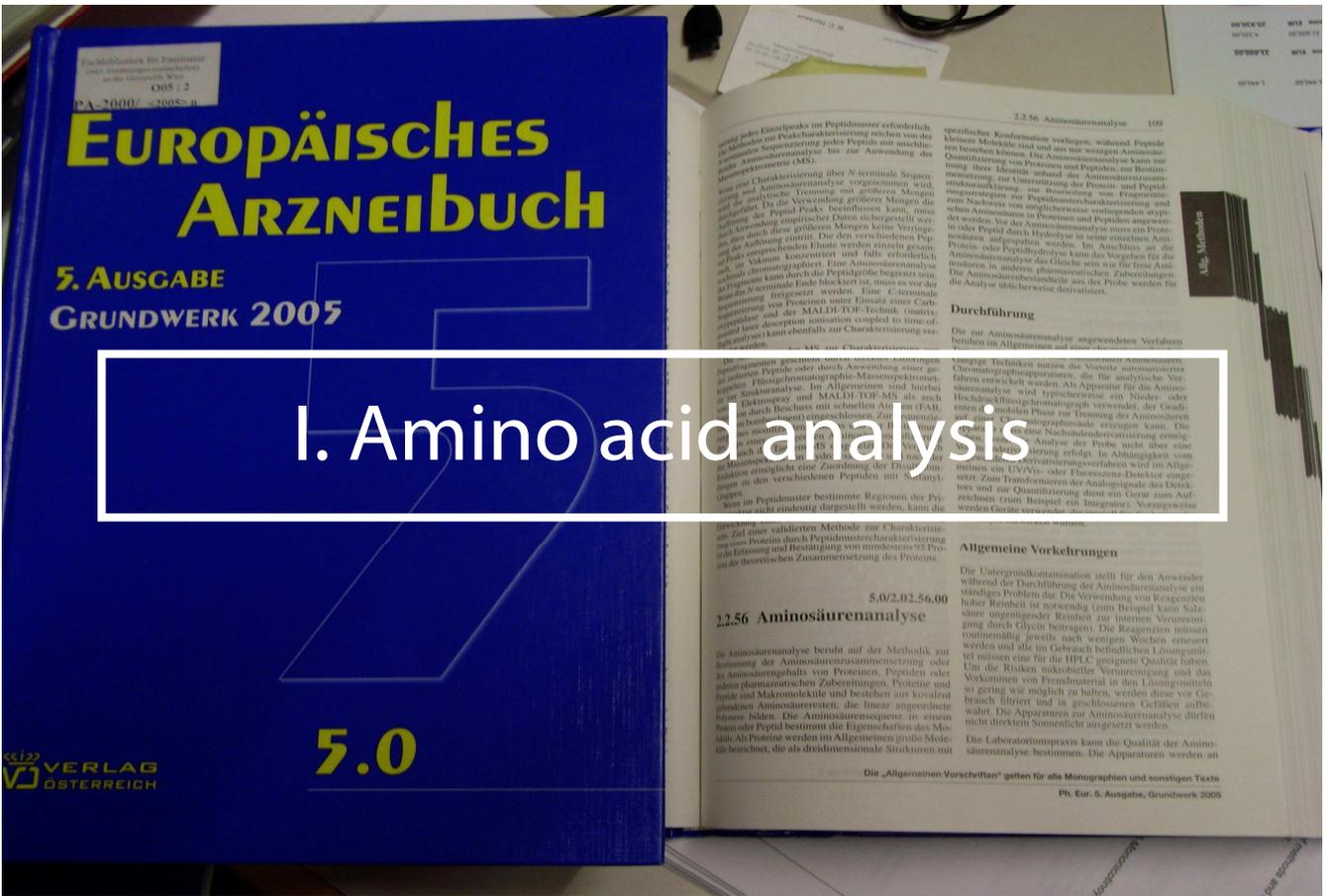


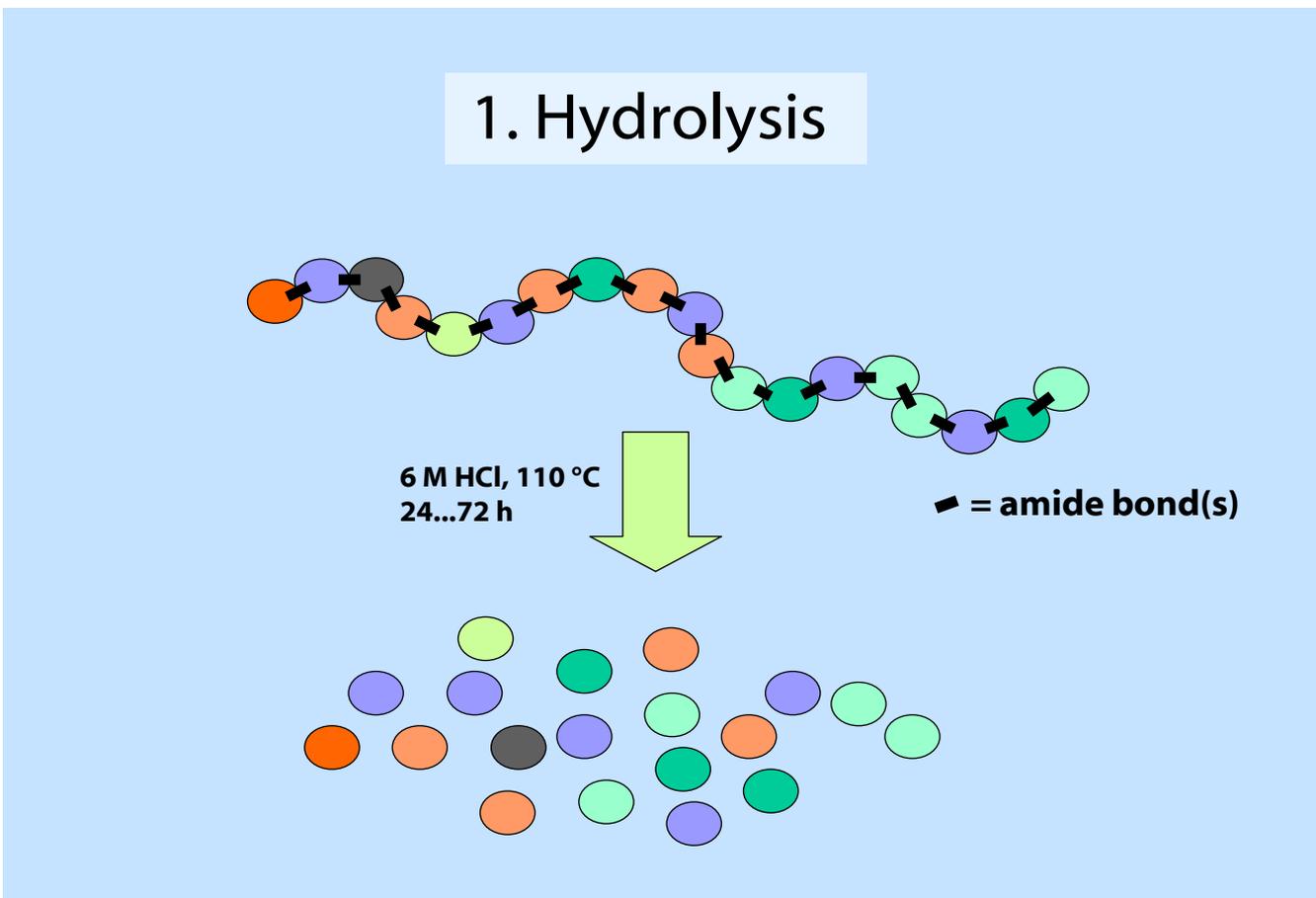
M. Kratzel

# Two Examples of interactive eLearning

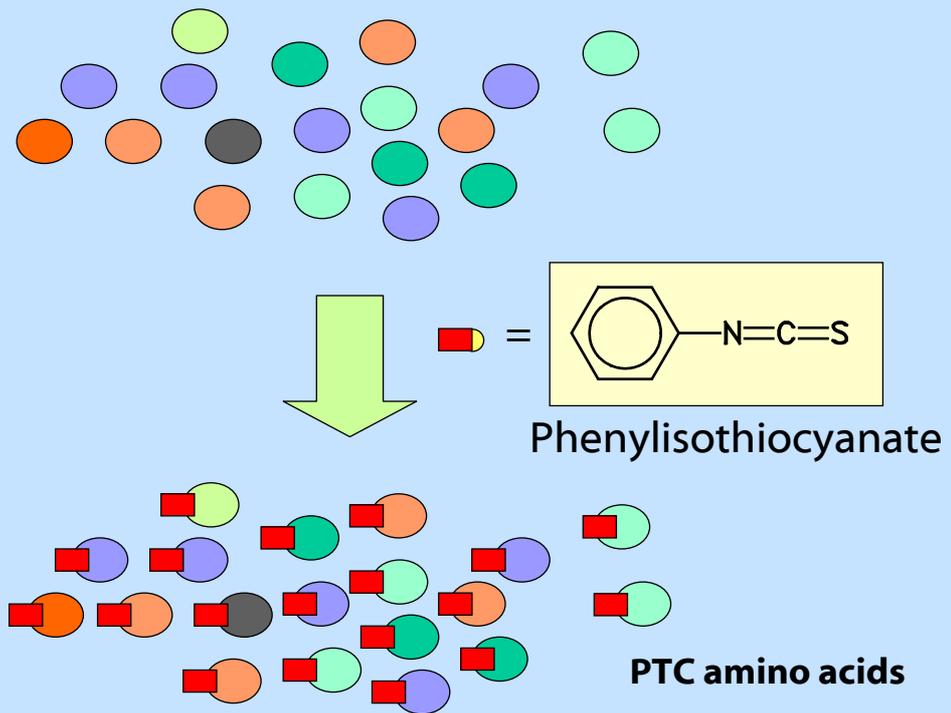
## I. Amino acid analysis



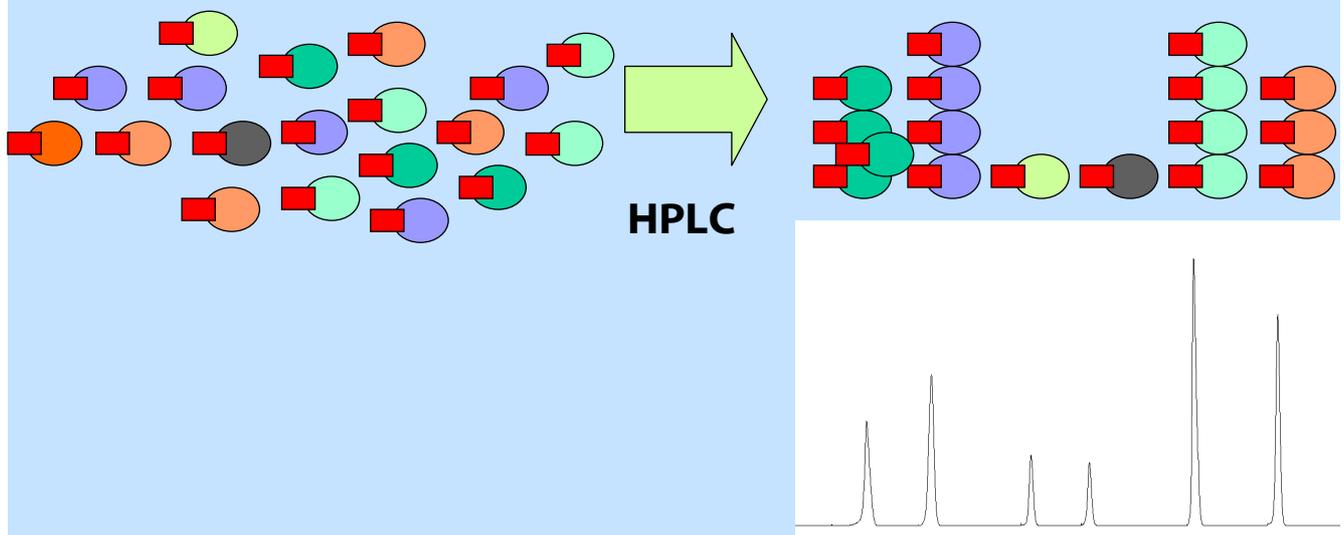
# I. Amino acid analysis



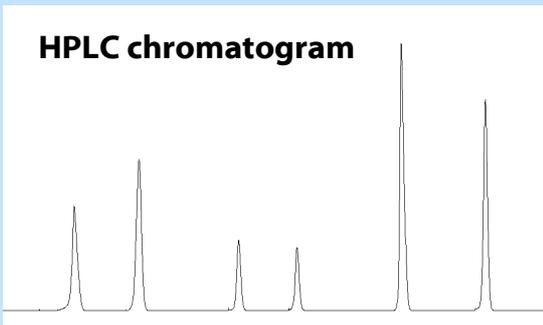
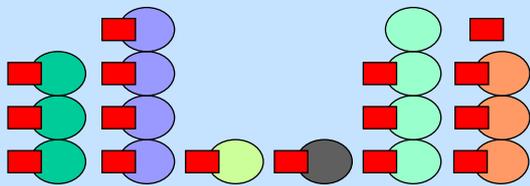
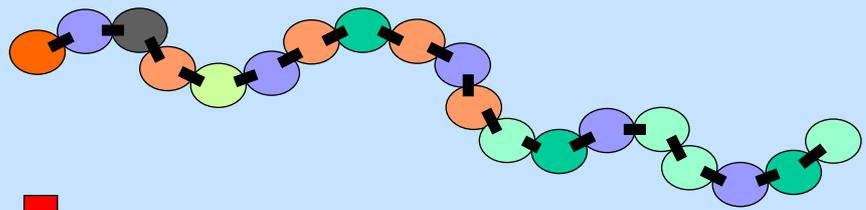
## 2. Derivatization



## 3. Separation



# 4. Calculation



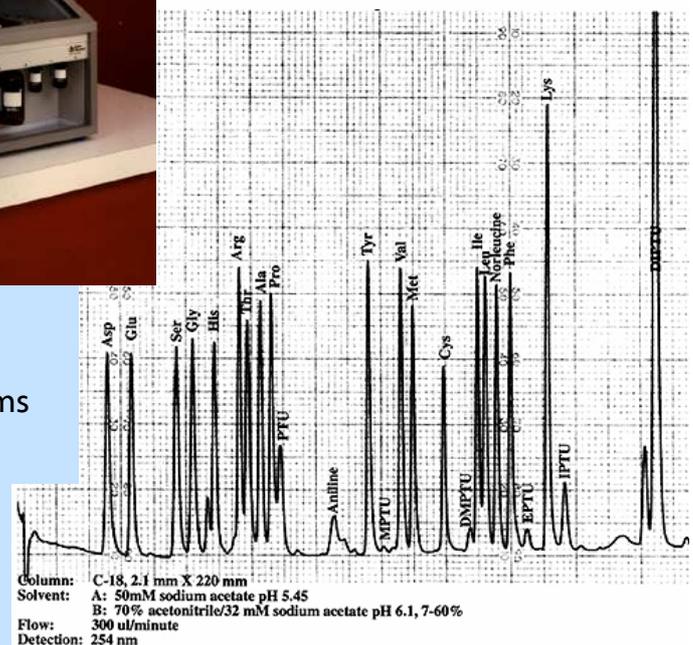
Sequence confirmation

Protein identification using databases (unique amino acid composition required)

Exact quantitation



Perkin Elmer Applied Biosystems  
**Model 420A PTC Derivatizer** with  
 an on-line Perkin Elmer Applied Biosystems  
**Model 130A PTC Amino Acid Analyzer**

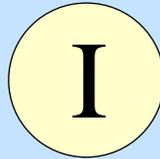


## Two didactic aims...

1. Hydrolysis

2. Derivatization

3. Separation



The analytical procedure

4. Calculation

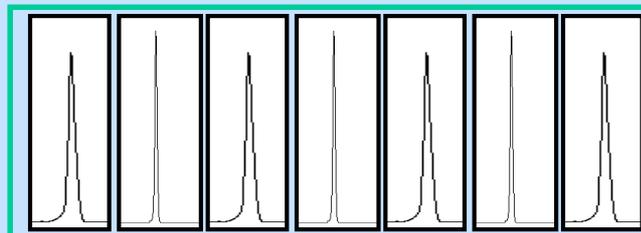
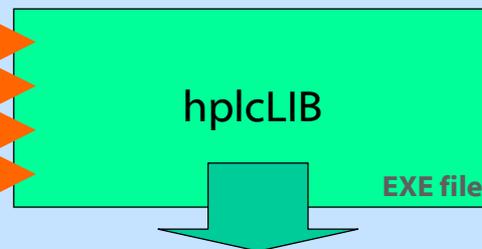
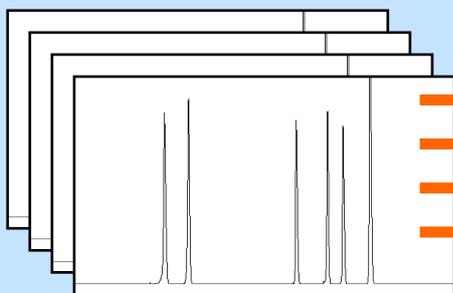


Evaluation,  
Calculation,  
Sequence confirmation



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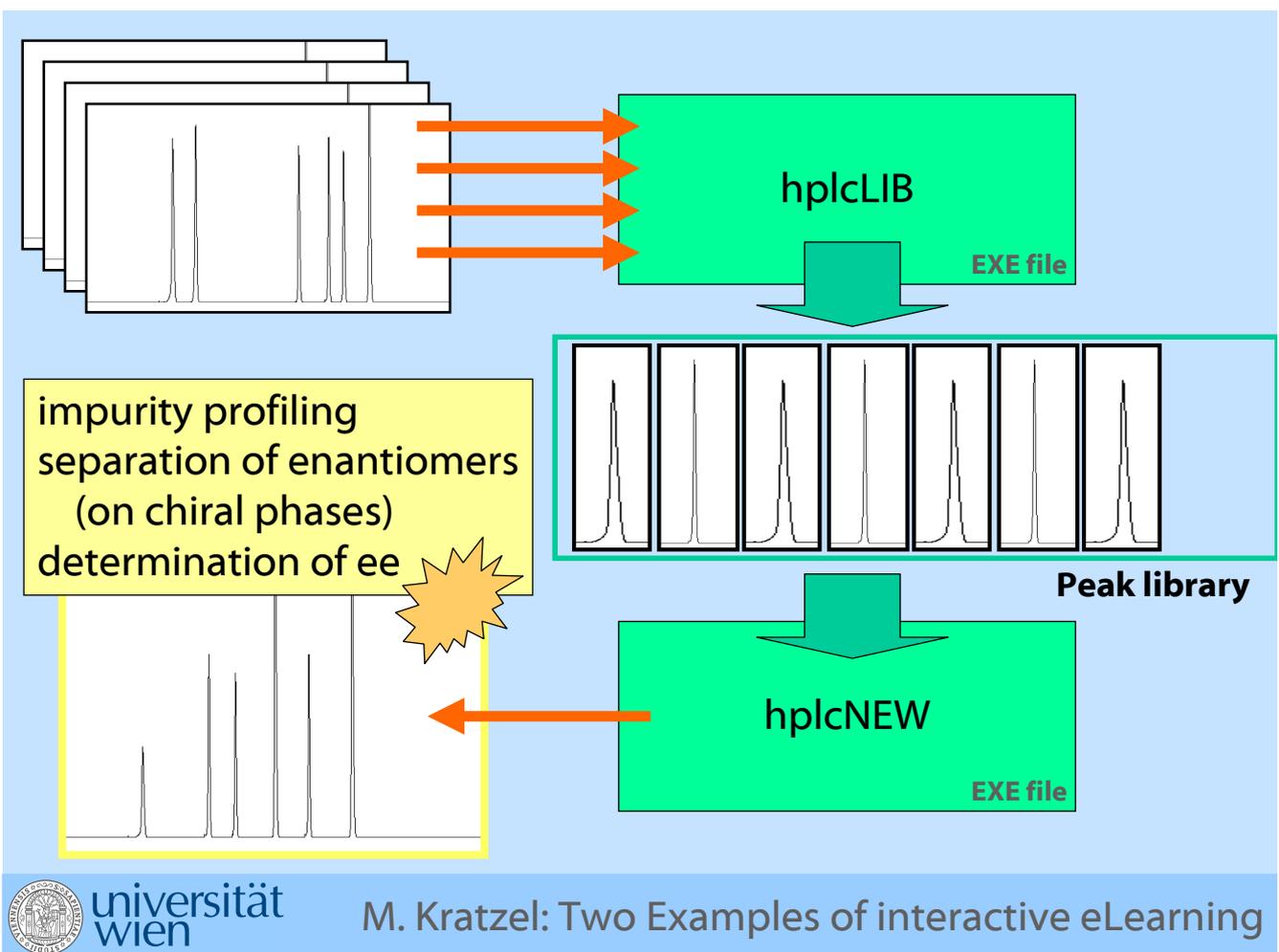
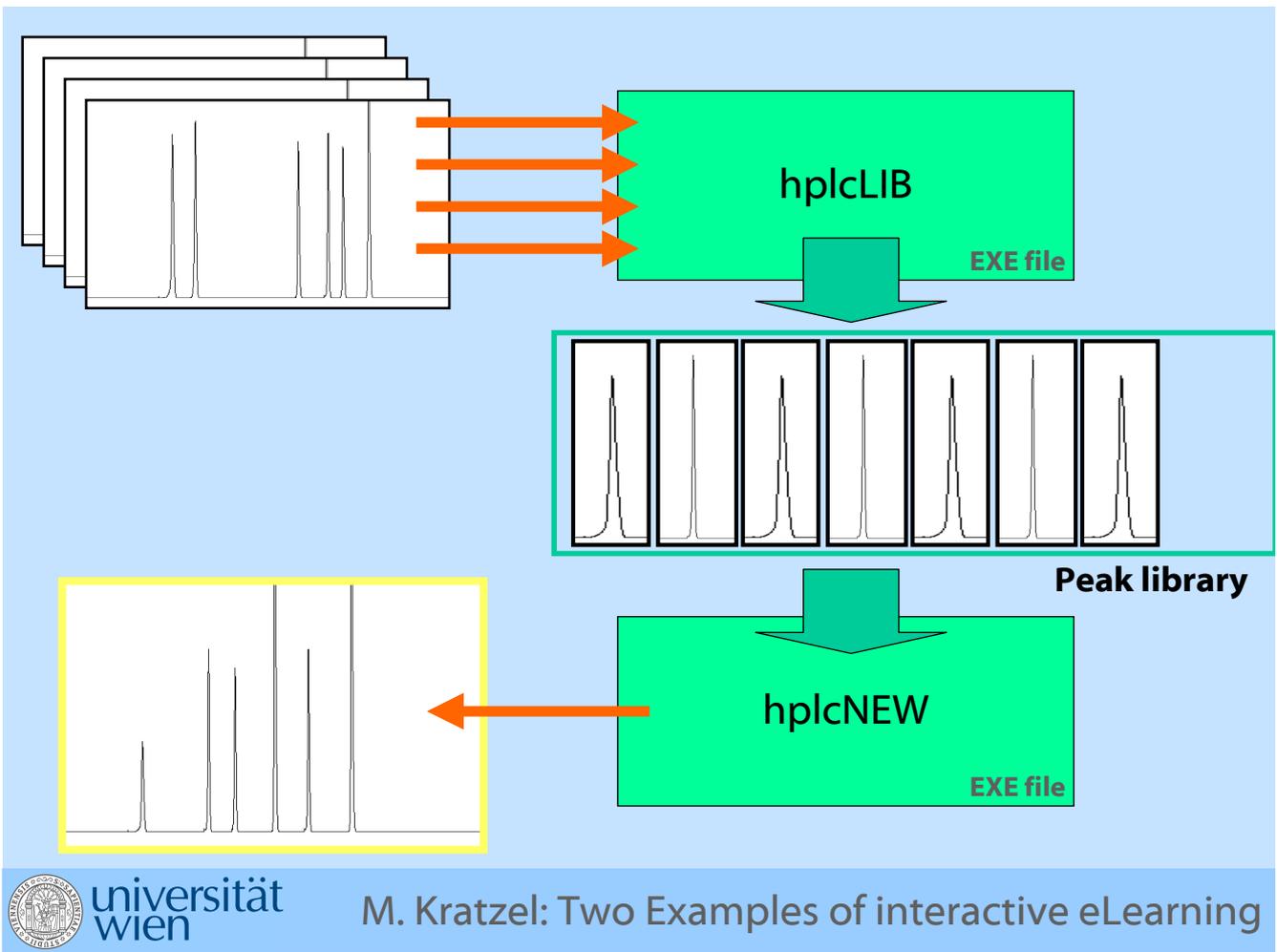


Peak library



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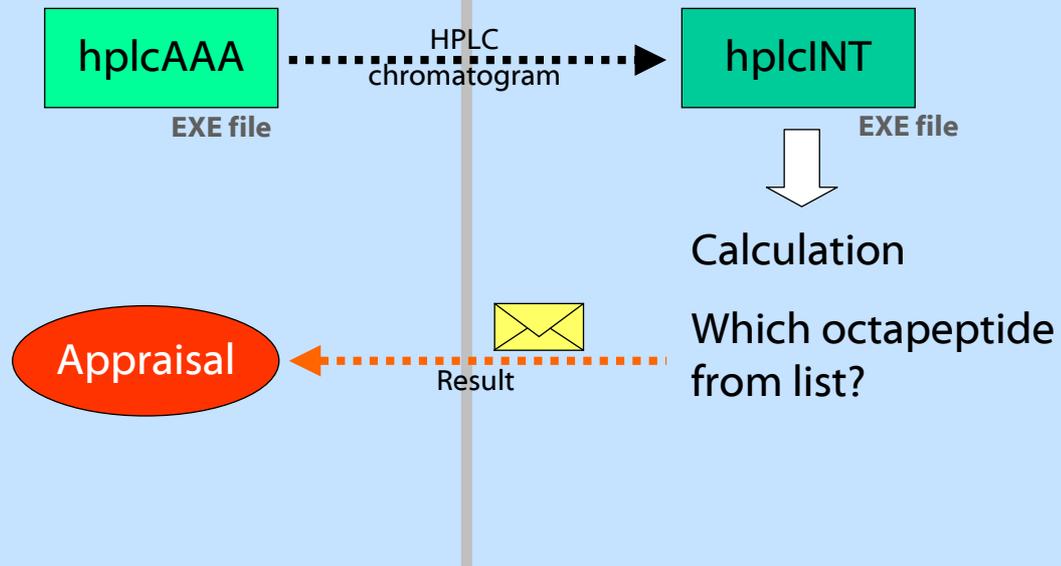
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# Flowchart - Amino acid analysis

Teacher's side

Student's side



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# Web pages - Amino acid analysis

<http://synthon.pch.univie.ac.at/px-te/simul/as001.htm>

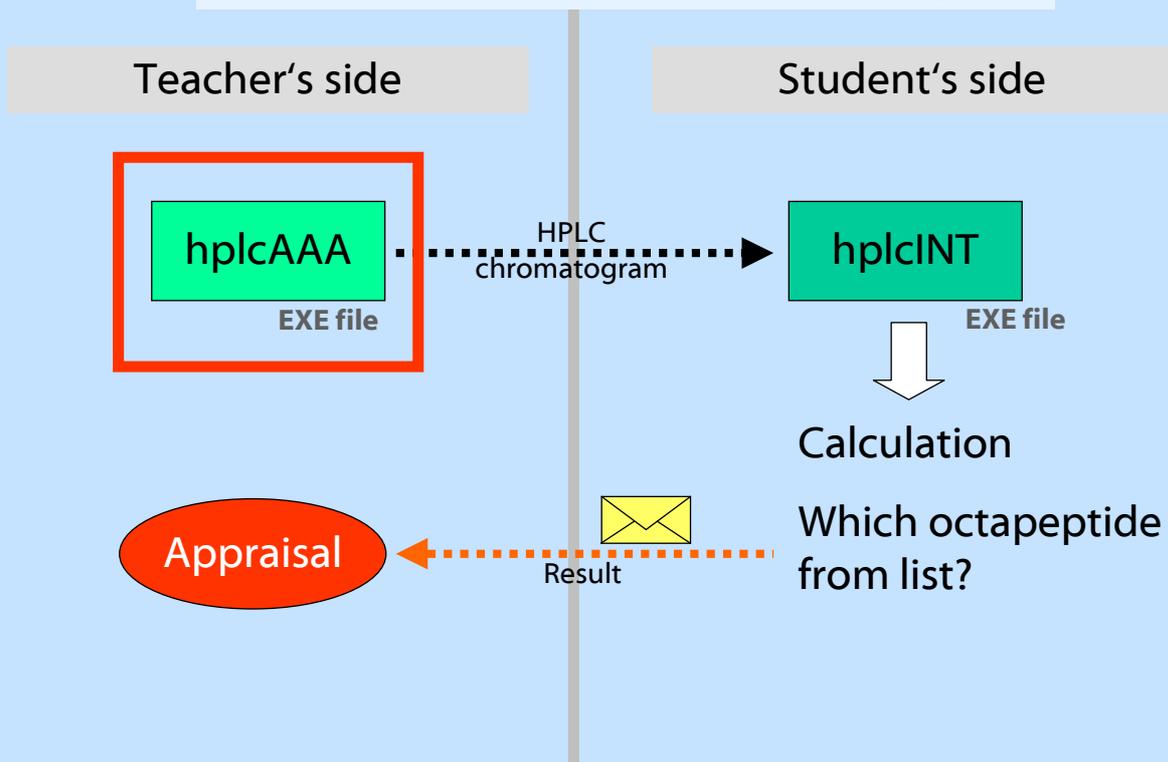
[http://www.univie.ac.at/ulg-pqm/aawe/programm\\_aawe\\_ws07.html](http://www.univie.ac.at/ulg-pqm/aawe/programm_aawe_ws07.html)



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# Flowchart - Amino acid analysis



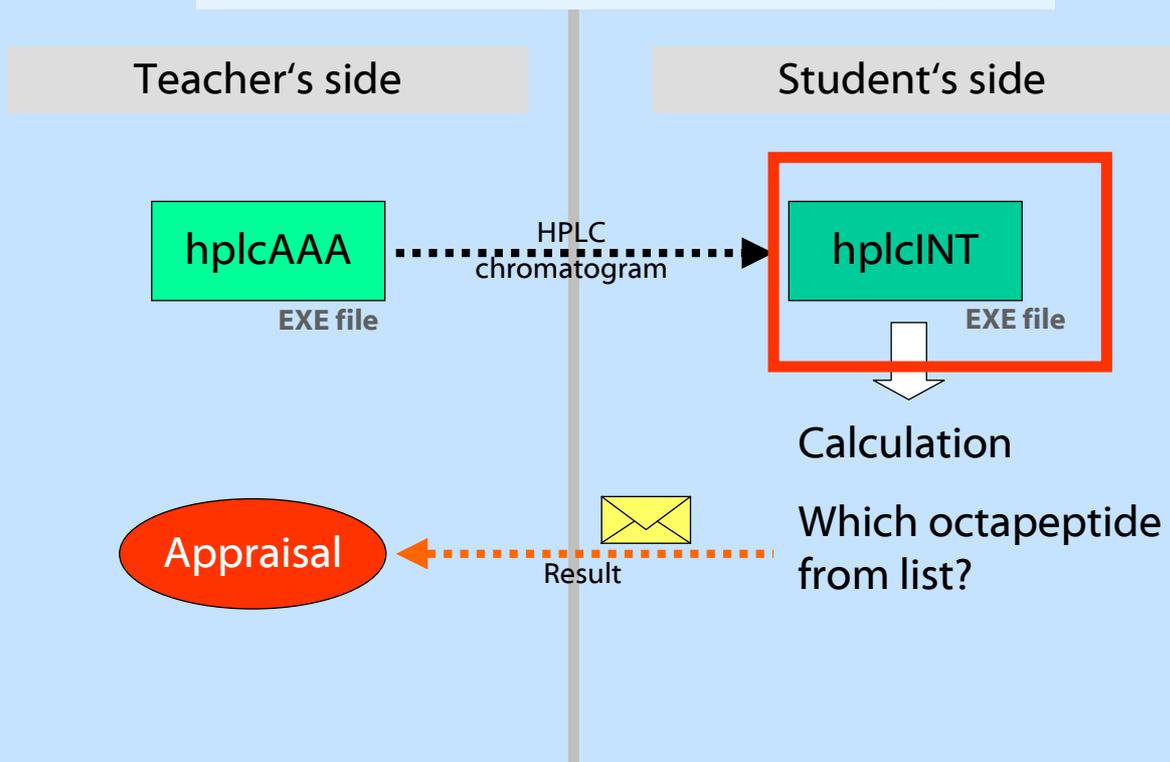
# Web pages - Sample download

<http://www.univie.ac.at/ulg-pqm/aawe/downloads.php>

[http://www.univie.ac.at/ulg-pqm/aawe/programm\\_aawe\\_ws07.html](http://www.univie.ac.at/ulg-pqm/aawe/programm_aawe_ws07.html)

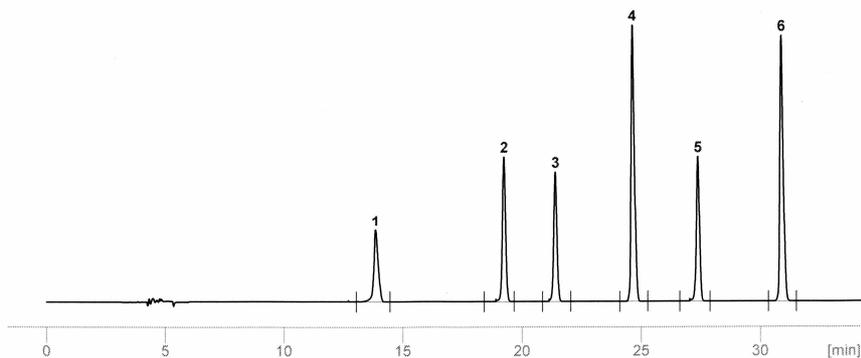


# Flowchart - Amino acid analysis



D:\eLearning\MEIER\_M\_3.HPD

hplcINT  
(C) M.Kratzel



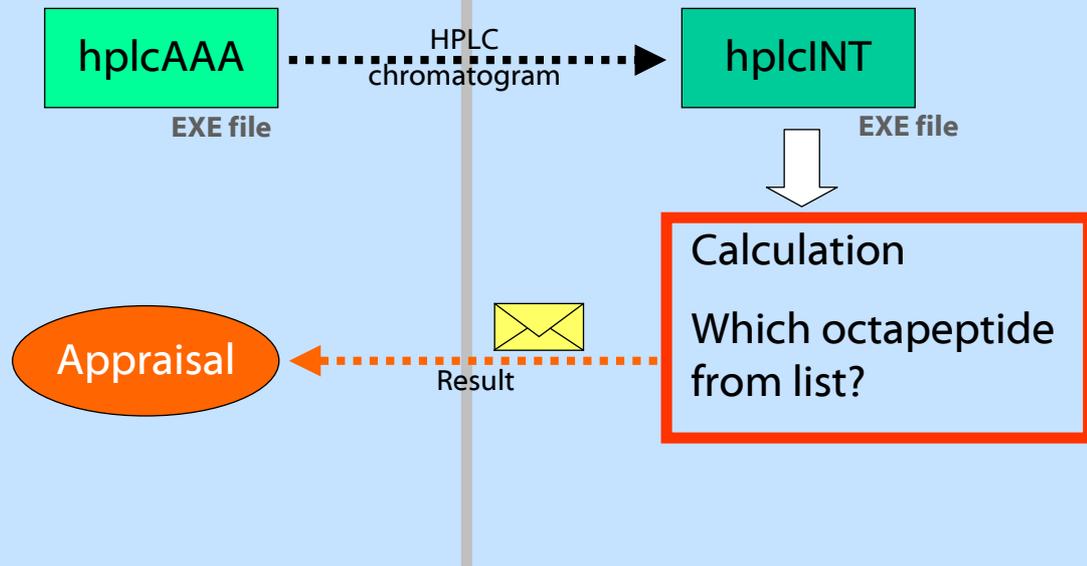
#	r. t.	PA	rel. %	
1	13.85	1001.19	9.09	Asp
2	19.24	1478.38	13.42	Arg
3	21.40	1319.32	11.98	Ala
4	24.62	2903.01	26.35	Val
5	27.37	1481.18	13.44	Ile
6	30.88	2833.77	25.72	Phe



# Flowchart - Amino acid analysis

Teacher's side

Student's side



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## Some (but no more) comments on evaluation:

1. The reference samples contain amino acids in equimolar amounts. Different peak areas refer to different UV absorption of PTC derivatives.
2. The result comprises the relative (molar) amounts of present amino acids.
3. And which octapeptide relates to your result?



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Microsoft Excel - CalcAAA.xls

Datei Bearbeiten Ansicht Einfügen Format Extras Daten Fenster ?

Arial 10 F X U

B23 = 1001,19

	A	B	C	D	E	F	G	H	I	J	K	L
1	AA	PA	Corr. Factor									
2												
3	<b>Standard 1</b>											
4												
5	Asp	1193,17	1,2562									
6	Glu	1215,98	1,2326									
7	<b>Ala</b>	<b>1498,86</b>	<b>1,0000</b>									
8	Ile	1667,46	0,8989									
9												
10												
11	<b>Standard 2</b>											
12												
13	Arg	1672,6	0,8966									
14	<b>Ala</b>	<b>1499,64</b>	<b>1,0000</b>									
15	Val	1700,34	0,8820									
16	Leu	1638,29	0,9154									
17	Phe	1651,64	0,9080									
18	Lys	2648,54	0,5662									
19												
20												
21	<b>Sample</b>		corr. PA									
22												
23	Asp	1001,19	1257,69									
24	Arg	1478,38	1325,50									
25	Ala	1319,32	1319,32									
26	Val	2903,01	2560,35									
27	Ile	1481,18	1331,42									
28	Phe	2833,77	2572,98									
29												
30												

Tabelle1 / Tabelle2 / Tabelle3 / **Tabelle4** / Tabelle5 / Tabelle6 / Tabelle7

Bereit



Microsoft Excel - CalcAAA.xls

Datei Bearbeiten Ansicht Einfügen Format Extras Daten Fenster ?

Arial 10 F X U

F14 =

	A	B	C	D	E	F	G	H	I	J	K	L
1	AA	PA	Corr. Factor									
2												
3	<b>Standard 1</b>											
4												
5	Asp	1193,17	1,2562									
6	Glu	1215,98	1,2326									
7	<b>Ala</b>	<b>1498,86</b>	<b>1,0000</b>									
8	Ile	1667,46	0,8989									
9												
10												
11	<b>Standard 2</b>											
12												
13	Arg	1672,6	0,8966									
14	<b>Ala</b>	<b>1499,64</b>	<b>1,0000</b>									
15	Val	1700,34	0,8820									
16	Leu	1638,29	0,9154									
17	Phe	1651,64	0,9080									
18	Lys	2648,54	0,5662									
19												
20												
21	<b>Sample</b>		corr. PA	<b>rel. %</b>								
22												
23	Asp	1001,19	1257,69	<b>12,13</b>	<b>1 x Asp (D)</b>							
24	Arg	1478,38	1325,50	<b>12,79</b>	<b>1 x Arg (R)</b>							
25	Ala	1319,32	1319,32	<b>12,73</b>	<b>1 x Ala (A)</b>							
26	Val	2903,01	2560,35	<b>24,70</b>	<b>2 x Val (V)</b>							
27	Ile	1481,18	1331,42	<b>12,84</b>	<b>1 x Ile (I)</b>							
28	Phe	2833,77	2572,98	<b>24,82</b>	<b>2 x Phe (F)</b>	<b>"=&gt; No. 23: DVFAVIFR</b>						
29												
30			<b>10367,27</b>									

Tabelle1 / Tabelle2 / Tabelle3 / **Tabelle4** / Tabelle5 / Tabelle6 / Tabelle7

Bereit



## II. Assay Validation



### ICH parameters versus type of validation

Type of Analytical Procedure	Identification	Impurity Testing		Assay
		Quantitative	Limit Tests	
Accuracy	No	Yes	No	Yes
Precision				
Repeatability	No	Yes	No	Yes
Interm. Prec.	No	Yes	No	Yes
Specificity	Yes	Yes	Yes	Yes
LOD	No	No	Yes	No
LOQ	No	Yes	No	No
Linearity	No	Yes	No	Yes
Range	No	Yes	No	Yes



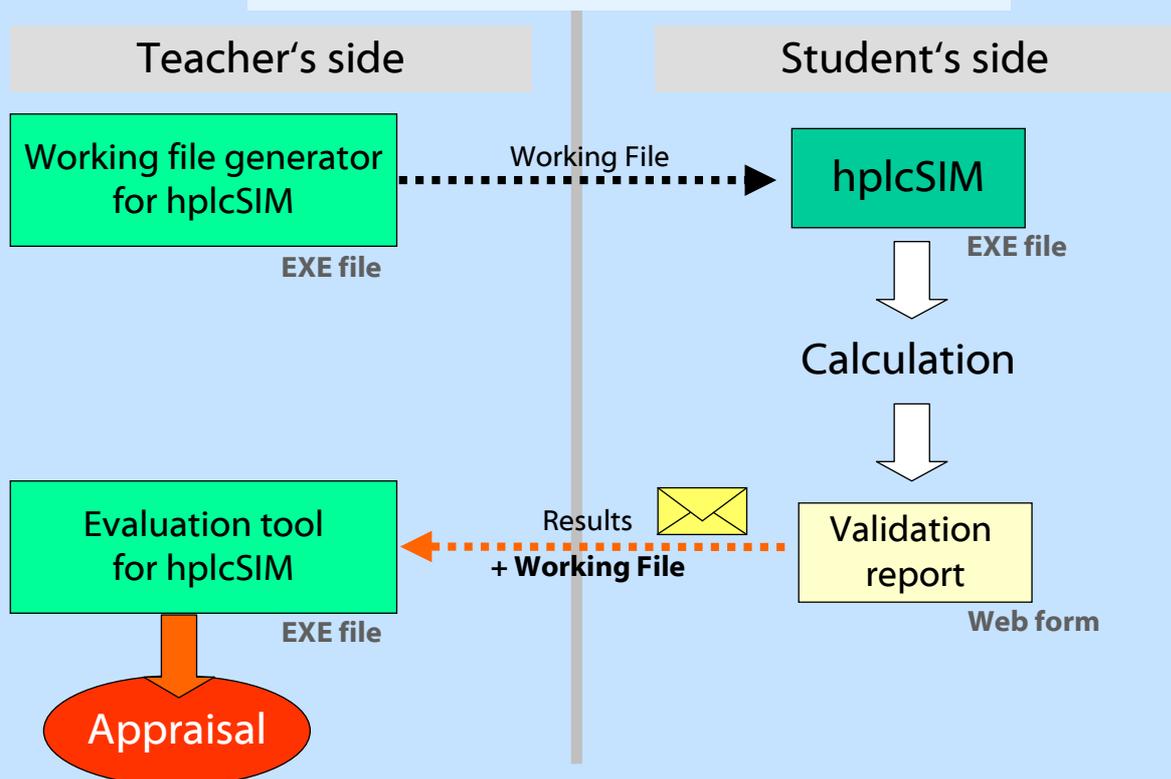
# Validation plan

## Assay of Nitrendipine by HPLC

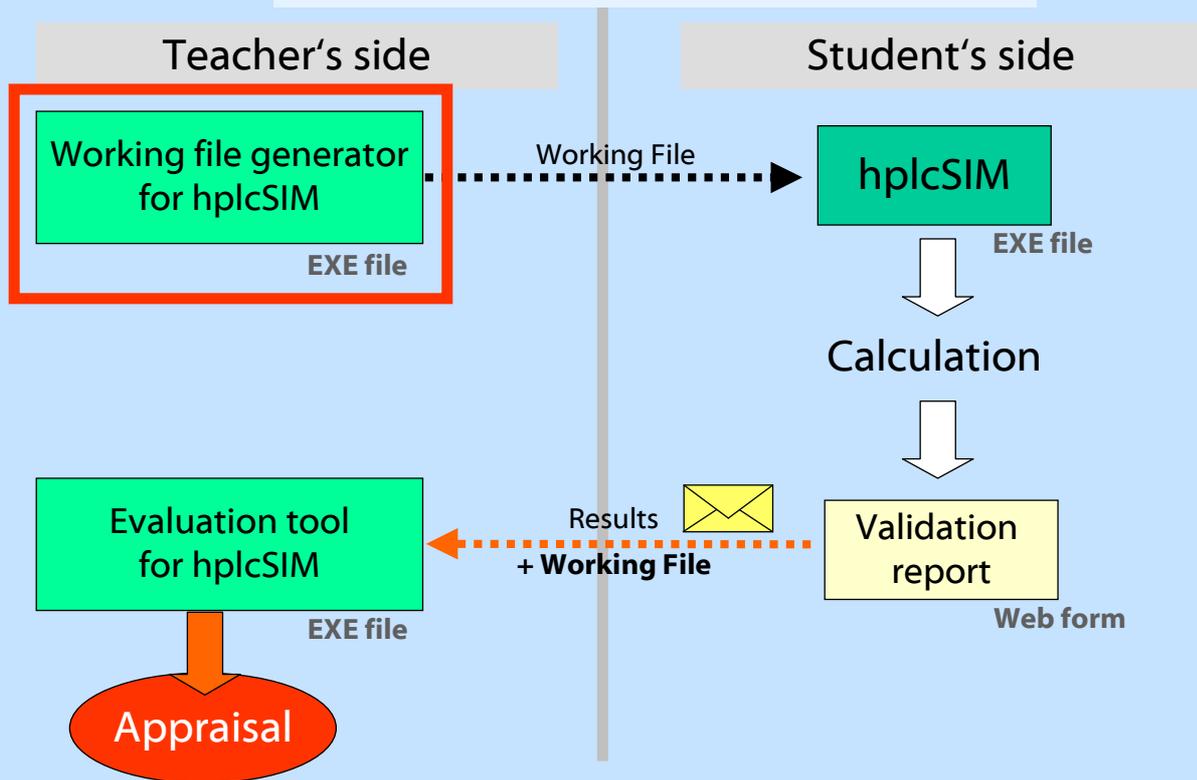
- I. System Suitability
- II. Specificity
- III. Precision
- IV. Linearity
- ~~V. LOD + LOQ~~
- (VI. Range)  $\Rightarrow$  included in IV. Linearity
- VII. Accuracy
- VIII. Precision - Repeatability
- IX: Precision - Intermediate precision



## Flowchart - Assay validation



# Flowchart - Assay validation



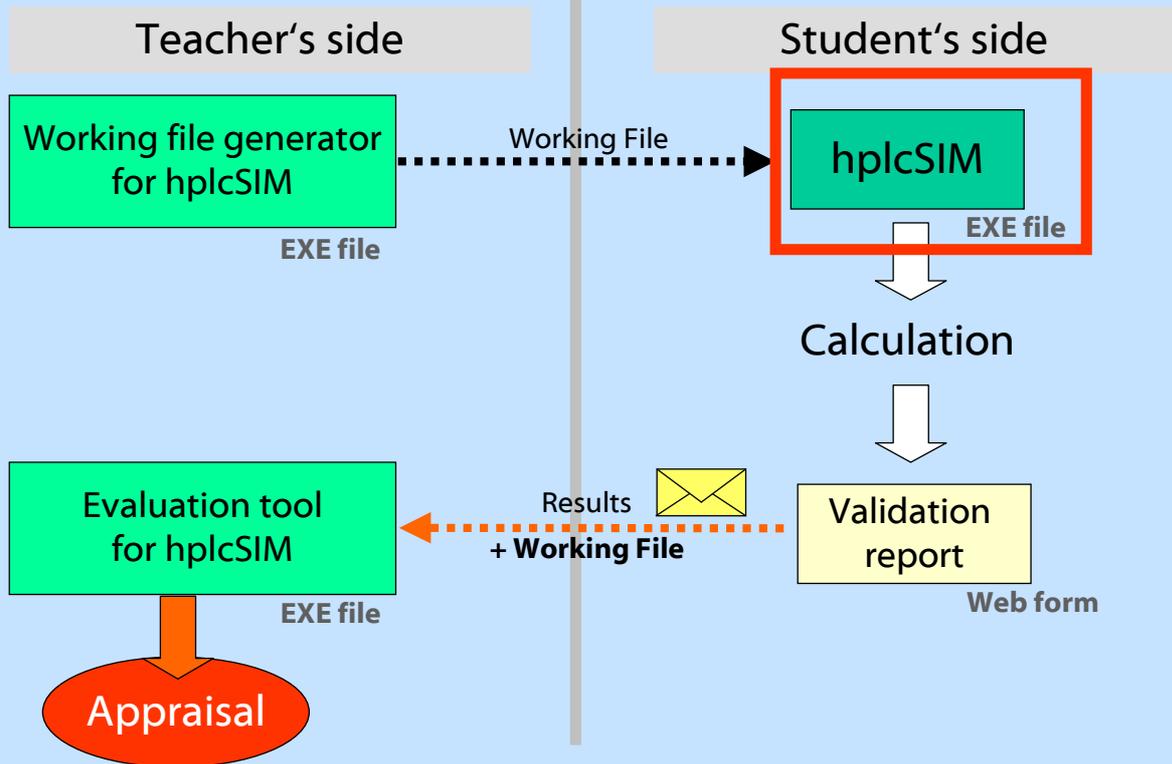
# Web pages - Assay validation

[http://synthon.pch.univie.ac.at/px-te/simul/valid\\_hplc\\_01.htm](http://synthon.pch.univie.ac.at/px-te/simul/valid_hplc_01.htm)

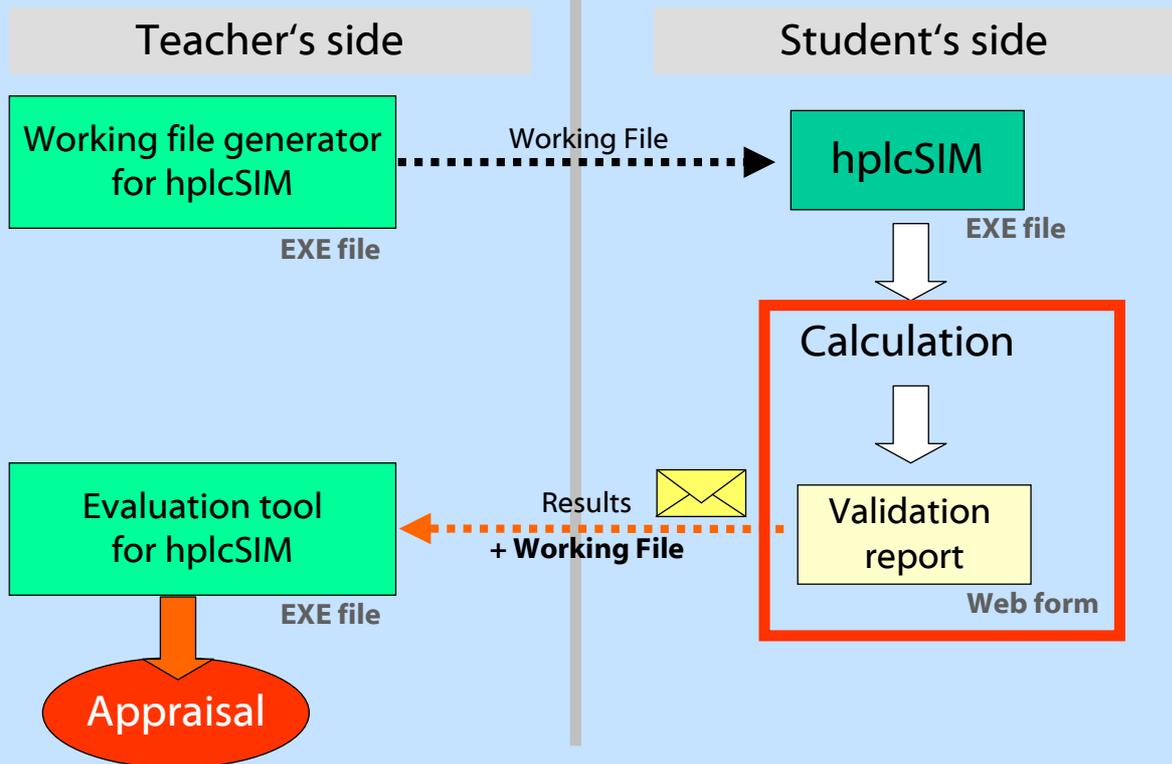
[http://www.univie.ac.at/ulg-pqm/aawe/programm\\_aawe\\_ws07.html](http://www.univie.ac.at/ulg-pqm/aawe/programm_aawe_ws07.html)



# Flowchart - Assay validation



# Flowchart - Assay validation



## Web form - Validation report

<http://synthon.pch.univie.ac.at/px-te/simul/abschlussbericht3.htm>

[http://www.univie.ac.at/ulg-pqm/aawe/programm\\_aawe\\_ws07.html](http://www.univie.ac.at/ulg-pqm/aawe/programm_aawe_ws07.html)

## Flowchart - Assay validation

