

Getting started with automated two-step protein purification

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Benefits of automated two-step purification

Typically, manual intervention is needed Step 1 Step 2

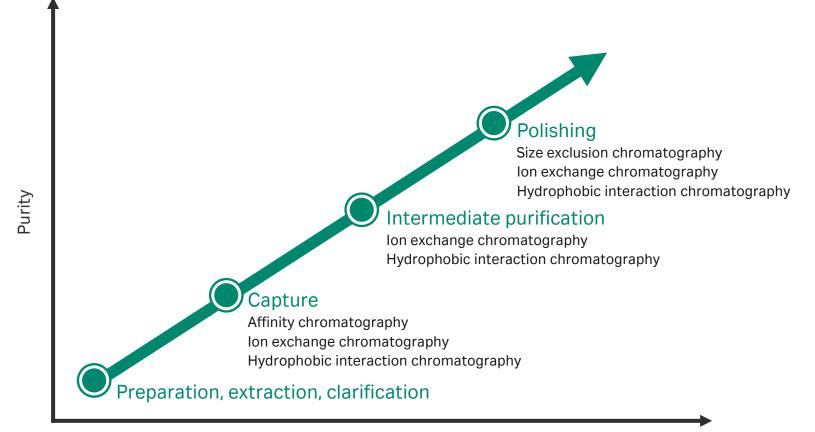
Reduce manual interactions

- Increase throughput
- Reduce labor
- Improve process consistency

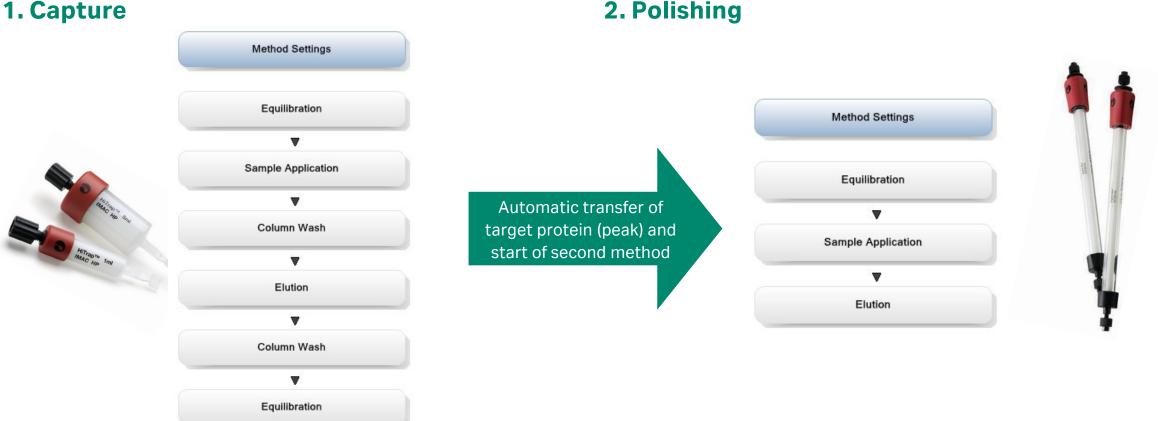
Eliminate hold steps

- Ensure product stability
- Eliminate or reduce hold containers

Apply multistep purification on a wide range of purification schemes



Example application: Automated two-step purification using predefined methods



2. Polishing

Ways of working

- 1. Select purification protocol
- 2. Set up ÄKTA[™] pure with components to support selected protocol
- 3. Set up capture method in UNICORN[™] Method Editor
- 4. Set up polishing method in UNICORN Method Editor
- 5. Set up Method queue
- 6. Run the methods first with buffer and then with sample
- 7. Evaluate results

The following slides will demonstrate a workflow for a two-step purification with intermediate loop collection.

1. Select purification protocol



Select purification protocol

- Discuss with colleagues
- Web searches
- Reference literature
- Discuss with suppliers

Set up the protocol to combine a capture and a polishing step in one run.

2. Set up ÄKTA pure with components to support selected protocol using intermediate loop collection

The flexible system configuration allows for an optimized flow path

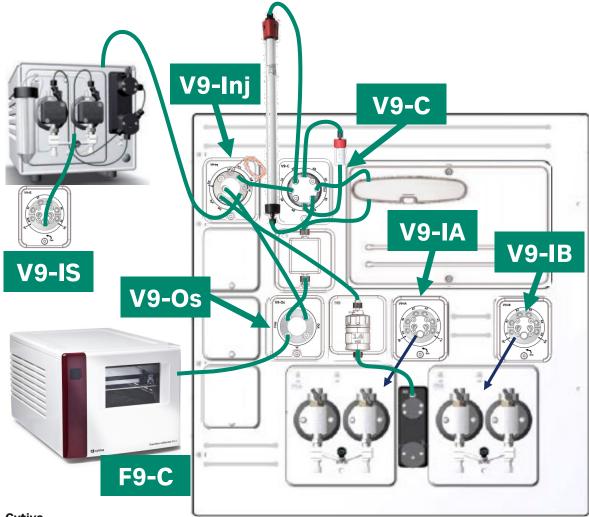
Components needed	Name
Sample inlet valve	V9-IS [‡]
Inlet valve	V9-IAB
Inlet valves*	V9-IA [‡] / V9-IB [‡]
Injection valve	V9-Inj [‡]
Column valve	V9-C [‡]
UV monitor 1	U9-M [‡] /U9-L
Conductivity	C9n [‡]
Outlet valve	V9-O V9-Os‡
Versatile Valve [†]	V9-V
Fraction collector	F9-R [‡] /F9-C
Sample pump	P9S [‡]

* Enables additional inlet buffers

[†] Use a versatile valve in front of injection valve if syringe position is needed for manual loop filling

[‡] Used in the example described in the following slides

2. Set up ÄKTA pure with components to support selected protocol (cont.)



Optimized flow path with intermediate loop collection used in the application example

A sample pump facilitates sample loading in capture step.

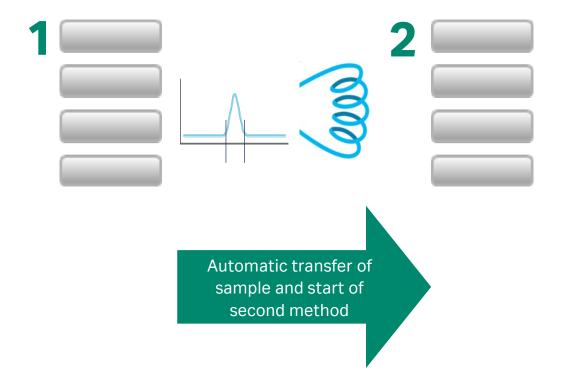
Connect outlet position from Outlet valve to Syringe port of injection valve.

When using V9-Os a fraction collector must be used.

Note: As an alternative, the sample can be loaded using the system pump and a mixer by-pass valve.

3. Method setup: Automated two-step purification with intermediate loop collection

Include first and second method in a *Method queue*



- Set up your methods for the two steps as individual methods using the predefined protocols
 - Capture step

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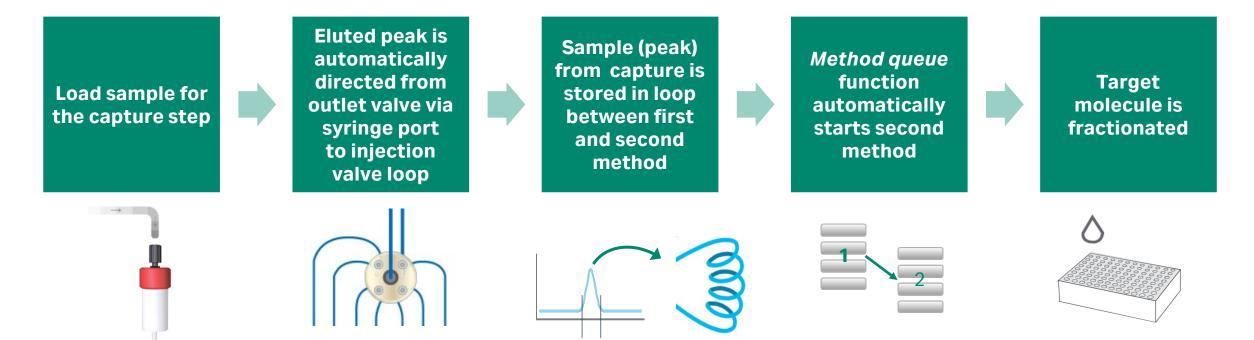
Use peak fractionation (via the outlet valve) to direct the eluted peak from the capture step to the loop

- **Polishing step** Fractionate normally to Fraction collector or V9-O
- 2. Create **Method queue** and start the run in System Control

3. Method setup

Method 1: Capture step

Method 2: Polishing step



3. Method setup: Capture Select adsorption technique

Select predefined method in *Method editor*:

New Method	×
System:	
test ~	
Create a new method by using the:	
Predefined Method:	
Affinity Chromatography (AC) $\qquad \qquad \lor$	
C Empty Method:	
Method Description	1
After equilibration and sample application, the protein of interest is adsorbed to the column ligand. After a wash to remove unbound sample, elution is performed either by using a buffer containing a competitor to displace the protein of interest, or by changing the pH or ionic strength. Finally, the column is re- equilibrated with start buffer.	
OK Cancel	

Different phases (steps) will be displayed in *Method outline*

Method Settings
Equilibration
*
Sample Application
Column Wash
Elution
Column Wash
*
Equilibration

3. Method setup: Capture Adjust **Phase properties**

Define parameters for each phase in *Phase properties* tab

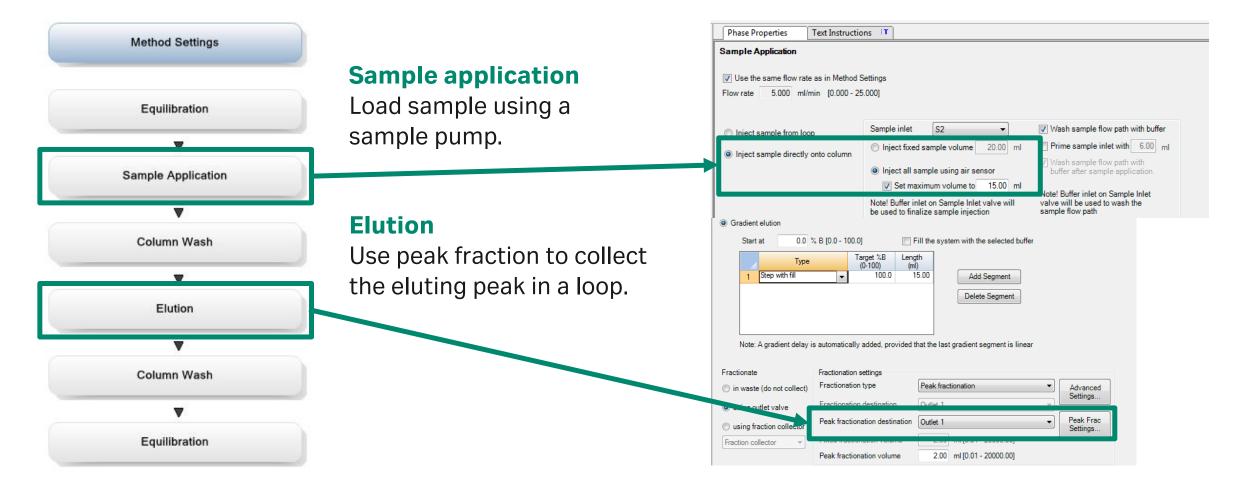
Phase Proper	ies Text Instructions	
Method Settings		
Column selection		Result Name & Location
Show by technique Affinity ~		Start Protocol
Column		Method Notes
Column Pressure Press	only suggested columns Column Properties volume 0.962 ml e limit pre-column 0.50 MPa [0.02 - 20.00] ure limit delta-column 0.30 MPa [0.02 - 20.00]	Unit selection Method Base Unit CV ~ Flow Rate Unit ml/min ~ Monitor settings
Column position	Bypass ~	UV variable wavelengths UV 1 280 [190 - 700] nm UV 2 254 [190 - 700] nm
Flow rate 1.000 ml/min [0.000 - 25.000]		UV 3 214 [190 - 700] nm Note! UV monitors with fixed
Inlet A Inlet B	A1 ~ B1 ~	wavelength are not presented in this view ☐ Enable pH monitoring
		Enable air sensor alarm
		✓ Inlet B

Settings available in *Phase properties*

- Flow rate and pressure limits for selected column
- Column position
- Inlets to be used
- Base unit
- Monitor settings



3. Method setup: Capture **Sample application** and **Elution** properties



4. Method setup: Polishing Select **SEC*/Desalting** in **Method editor**

Select predefined method: SEC*/Desalting in Method editor

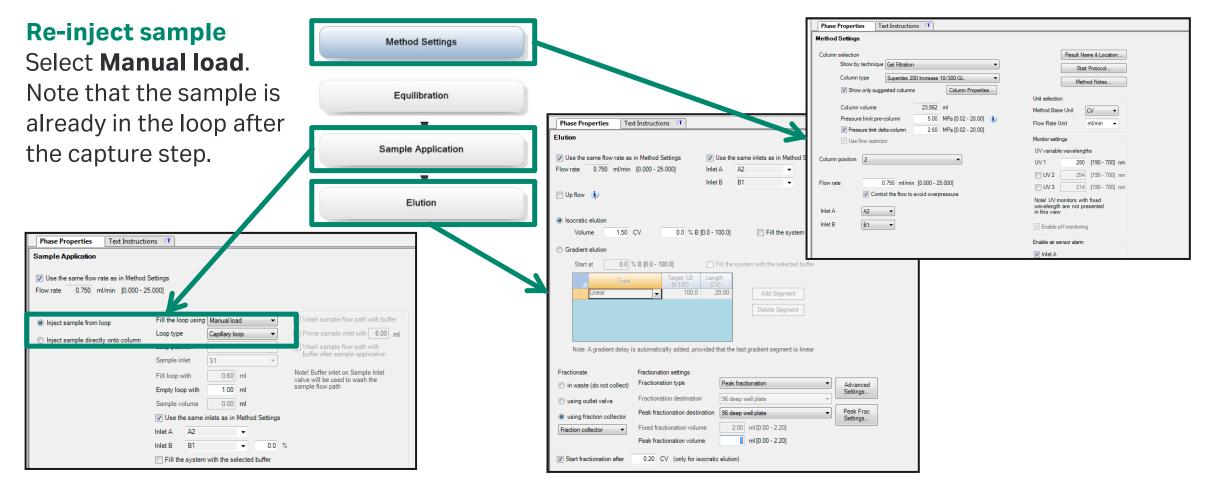
lew Method System:	Concentration of the local sectors of the local sec
1991976	
Create a new method by us	sing the:
Predefined Method:	-
Gel Filtration (GF)	
Empty Method: Method Description	
After equilibration and sa elute according to their s	ample application, proteins separate and size (largest first).
	OK Cancel
	UK Cancel

Method will be displayed in Method outline

_	Method Settings
	Equilibration
	•
	Sample Application
	Elution

*Size exclusion chromatography, also called gel filtration (GF)

4. Method setup: Polishing **Method settings**, **Sample application**, and **Elution** properties

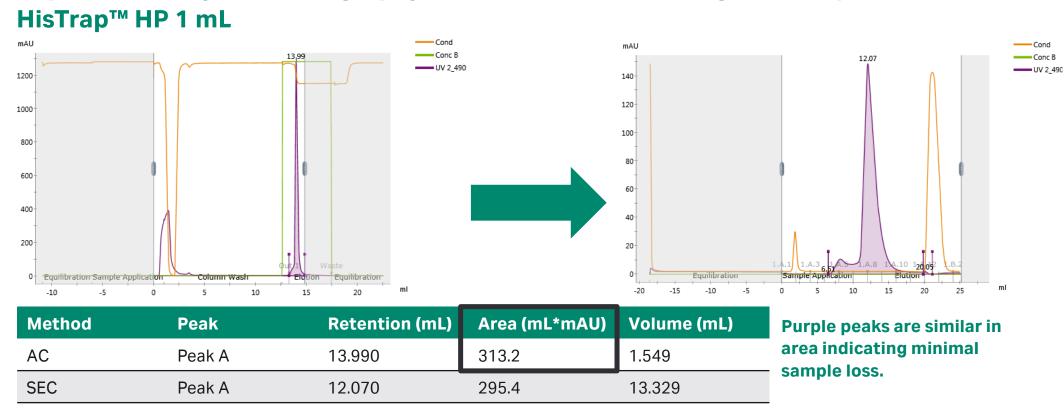


5. Method setup: Create **Method queue** Include the capture and polishing step in **Method queue**

Method queue is set in UNICORN™ Method editor

Met	thod Queue			×
Num	ber of included systems: 1	~		
Syst	tem test	8	×	Move Up
	Method		Start Condition	Move Down
1	Method One AC		At queue start	move bown
2	Method Two SEC		Immediately after the previous method has ended	V Insert Row
3				 ✓
4				✓ Delete
		Capture step	Polishing step	d.

6. Running the methods Automated two-step purification, example 1

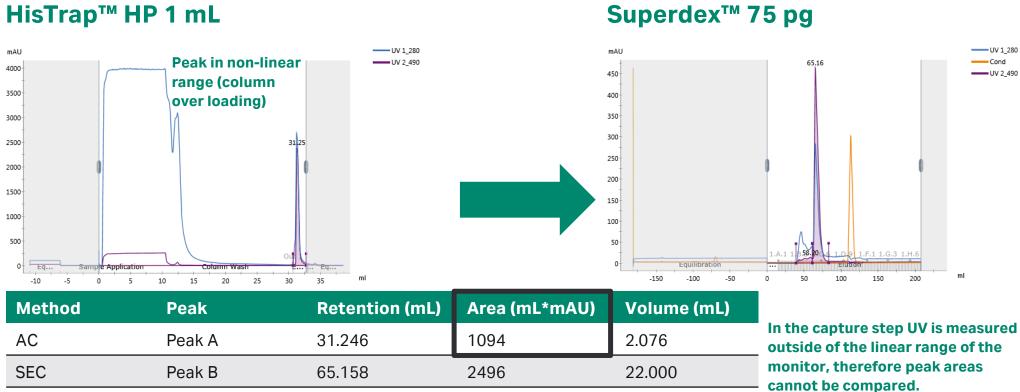


Polishing: SEC* Superdex™ 75 Increase

Sample: GFP-His (clarified homogenate, 5.3 mg GFP-His/g *E. coli* cell pellet) *Size exclusion chromatography, also called gel filtration (GF)

Capture: Affinity chromatography with

6. Running the methods Automated two-step purification, example 2



Polishing: SEC* HiLoad[™] 16/600

Capture: Affinity chromatography with HisTrap[™] HP 1 mL

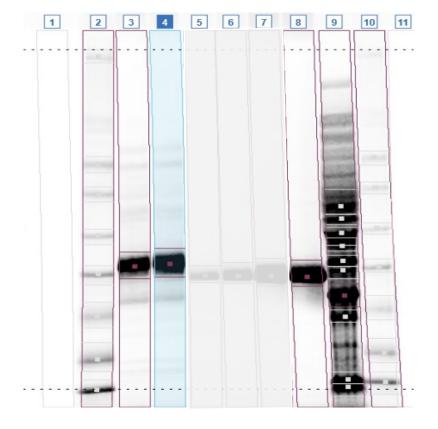
Sample: GFP-His (clarified homogenate, 5.3 mg GFP-His/g *E. coli* cell pellet) *Size exclusion chromatography, also called gel filtration (GF)

7. Evaluation Integration results from example 1 and 2

Example 2 Peak area Sample loading amount up to column capacity ("scaleup"). Difference in area between steps mL*mAU due to area from eluting peak from Method 1 is in the nonlinear region of the UV monitor. 2500 2000 1500 **Example 1** Sample loading amount below column capacity 1000 to show complete transfer between steps and reproducibility 500 0 Method 1: Method 2: Method 1: Method 2: Method 1: Method 2: HisTrap[™] HP Superdex[™] 75 Increase 10/300 HisTrap HP, duplicate Superdex 75 Increase 10/300 GL, HisTrap HP, sample load HiLoad[™] 16/600 Superdex 75 pg duplicate GL

7. Verification Reference analysis with electrophoresis

Results form electrophoresis



Reference analysis to verify identified peaks

Sample ID:

- 1. Blank
- 2. Amersham[™] WB molecular weight markers
- 3. Two-step purification protocol AC-GF HisTrap[™] 1 mL + Superdex[™] 75 Increase 10/300 GL
- 4. Two-step purification protocol AC-GF HisTrap 1 mL + Superdex 75 Increase 10/300 GL, duplicate
- 5.-7. Blanks
- 8. Two-step purification protocol AC-GF HisTrap 1 mL + HiLoad[™] 16/60 Superdex 75 pg, scaleup
- 9. Start material His-GFP, *E. coli*
- 10. Amersham WB molecular weight markers

Utilize standard hardware and predefined UNICORN methods for automated two-step purification

ÄKTA™ systems in automated multistep purification



Combine different chromatography steps in automated multistep purification to save time and add consistency.

Set up automated, two-step purification using predefined methods and **Method queue** function to quickly get started.

Store peak from step one in loop before automatically continuing to second step.

ÄKTA pure was used in the example application shown. ÄKTA avant can also be used in the same way.





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