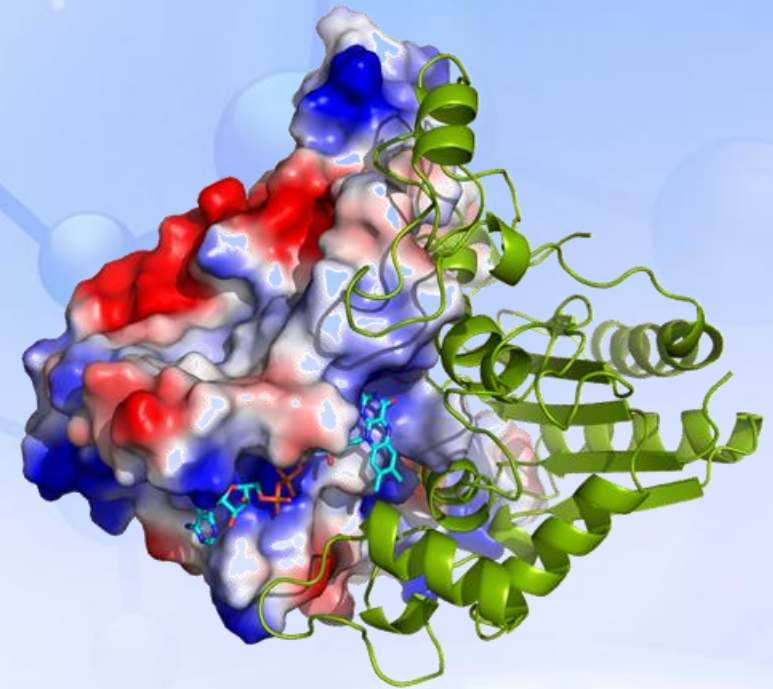




Scientists for Scientists: Protein Purification At Biortus

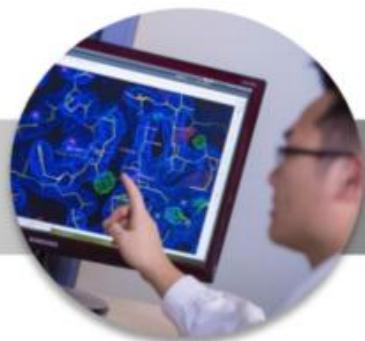
February, 2025

Web: en.biortus.bio



Guided Design from the Beginning

- Published Literature & Structures
- Structure Prediction Tools



Evaluation & Design

Construct & Condition Screening

- High-throughput parallel screening
- Shakers, Fermenters, WAVE
- *E. coli*, Insect Cell, Mammalian, Silkworm, etc.



Expression

An Extension of Your Lab

- Membrane, Soluble, & Refolded Proteins
- Standard & Specialty Columns



Purification



Quality Control

Soluble, Consistent, Functional

- Standard QC: SDS-PAGE, LC-MS, aSEC
- Custom QC: nanoDSF, SPR, Negative Stain, etc.

4-8 weeks

Our facilities and instruments available

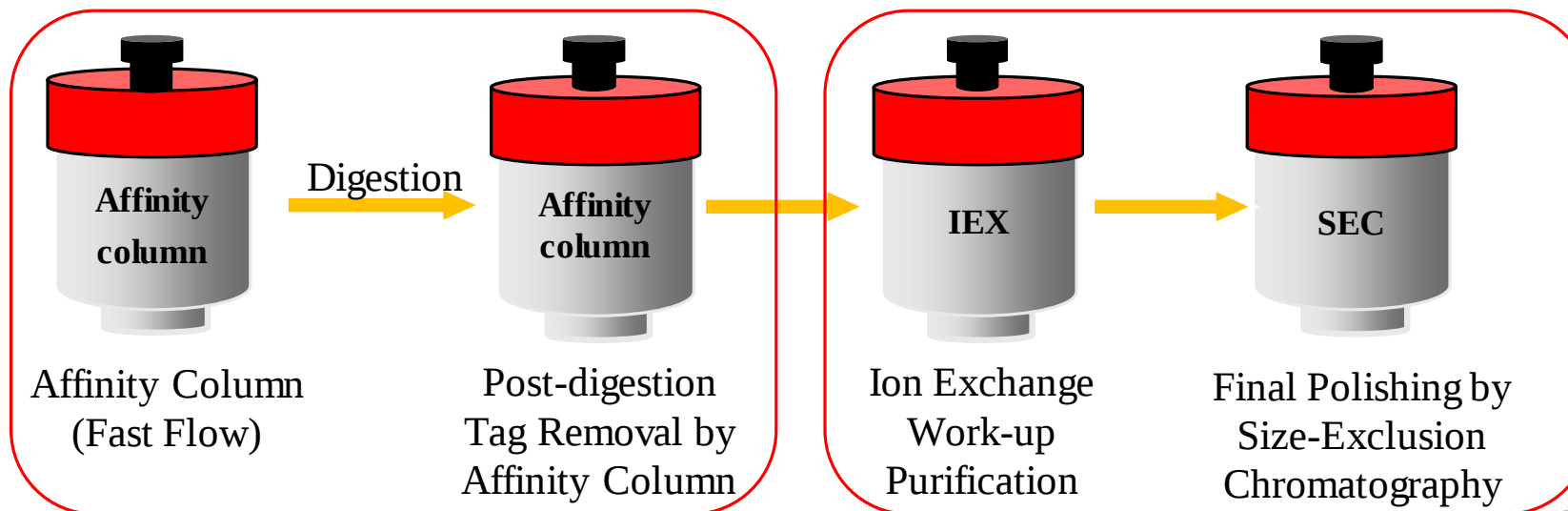
- >50 AKTAs
- >250 employees in protein expression and purification department
- Dedicated membrane protein purification team
- Over 200 batches of purified protein per week
- Endotoxin removal available
- Standard columns and specialty columns available
 - Nickel (NTA, Excel, etc.), GST, MBP, Flag, Protein A/G, IEX, SEC, and more.



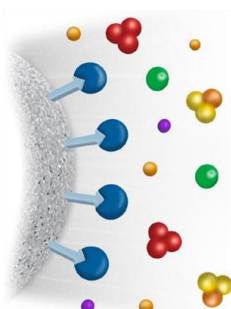
AKTAs At Biortus (>50 Instruments In Total)



Core Steps in the Purification Workflow

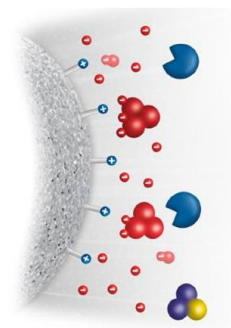


Affinity chromatography (AC)

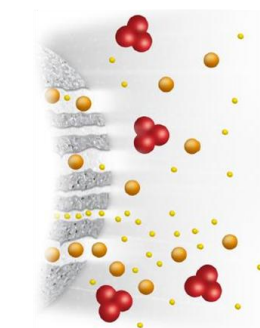


- Bind – elute principle
- Requires specific elution conditions
- Concentrating effect

Ion exchange chromatography (IEX)



Size exclusion chromatography (SEC)



- Diffusion – no binding
- Any elution conditions
- Diluting effect



Protein Preparation Platform-Purification (Cell Lysis)



For cell pellet

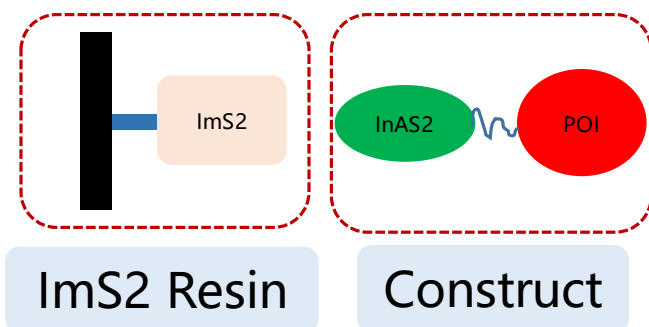


For culture medium

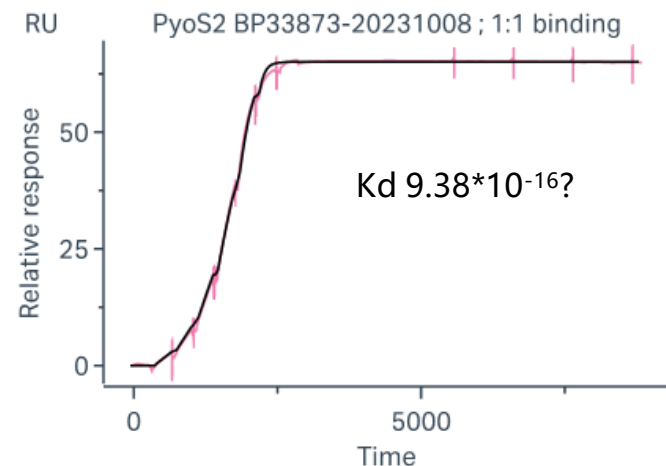
Primary Purification: Affinity Chromatography

Tag	Sequence	Residues	Affinity K_D (M)	Loading NaCl (M)	Capacity (mg/ml)	Affinity beads
His	HHHHHH	6	$10^{-8/-9}$	---	20-40	His FF
Strep	WRHPQFGG	8	10^{-6}	---	---	Strep-Tactin
StrepII	WSHPQFEK	8	10^{-9}	---	10-15	Strep-Tactin XT
Flag	DYKDDDK	8	$10^{-8/-9}$	< 0.2	0.5-1	Anti-Flag G1
HA	YPYDVPDYA	9	10^{-9}	< 0.2	---	4B2 antibody
ALFA	SRLEEELRRRLTE	13	10^{-11}	< 0.2	---	ALFA nanobody
Twin StrepII	SHHPQFEK-(GGGS)2-GGSAWSHPQFEK	28	10^{-12}	---	10-15	Strep-Tactin XT
CL7	TFWEEVSKDPELSKQFSRNNNDRI	130	$10^{-14/-17}$	---	15-20	IM7
GST	VLDIRYGVSRIAYSKDFETLKVDFL	218	$10^{-6/-7}$	< 0.2	10	GST FF
MBP	KYDIKDVGVNAGAKAGLTFLVD	370	10^{-6}	< 0.2	6-10	MBP FF
SpyTag	AHIVMVDAYKPTK	13	Covalent	---	---	Catcher-Protein
Biotinylated-Avi	Biotin+GLNDIFEAQKIEWHE	15	$10^{-12/-13}$	---	10-15	Strep-Tactin XT
HiBiT	MVSGWRLFKKIS	11	10^{-9}	---	---	LgBiT

- Preferred construct design: **His-StrepII-TEV-GG-POI**, Screened by FSEC+His Lite methods.



ImS2/InAS2



- Both the affinities of Im7/CL7 and ImS2/InAS2 are beyond the limit of the S200 SPR instrument, we termed as “Infinite Affinity (InA)” .

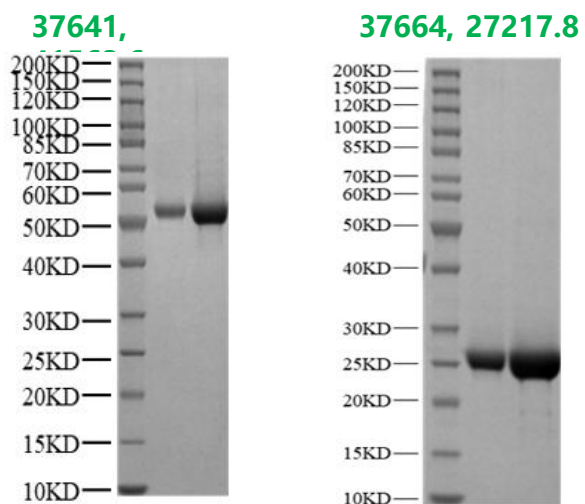
Beads	Binding Affinity(M)	Purity	Binding Capacity (mg/mL)	Recycle
His	10 ⁻⁸ -10 ⁻⁹	>80%	40	5
Flag	10 ⁻⁸ -10 ⁻⁹	>90%	1	2
GST	10 ⁻⁶ -10 ⁻⁷	>90%	5	5
StrepII	10⁻¹²	>90%	14	10
ImS2	Very high	>90%	70	10-100
Im7	Very high	>90%	40-60	10-100

❑ Features:

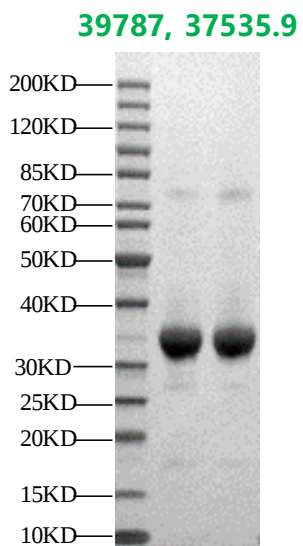
- can be attached to either N- or C-terminus of POI
- can be expressed in *E.coli*, insect cell and mammalian systems
- can be used for either cellular or secretion expression

Biortus: One Step Infinite Affinity (InA) Purification

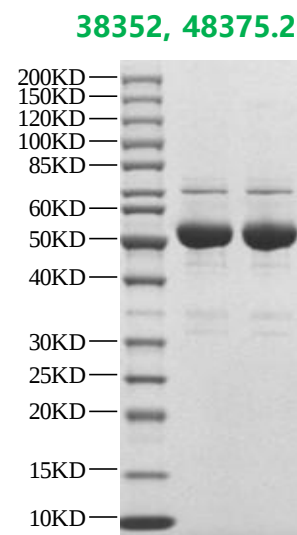
E.coli system



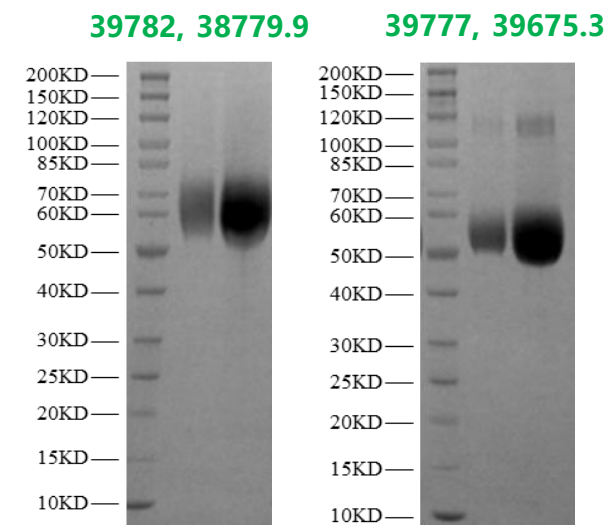
Insect, intracellular



Mammalian, intracellular



Mammalian, secreted



Introduction Self-developed Bio-AC™

Bio-AC™ is a fast protein affinity purification system that takes up very little laboratory space and can automate sample affinity chromatography with a simple program setup. It can help experimentalists reduce the probability of human error and the time of experimental operation.



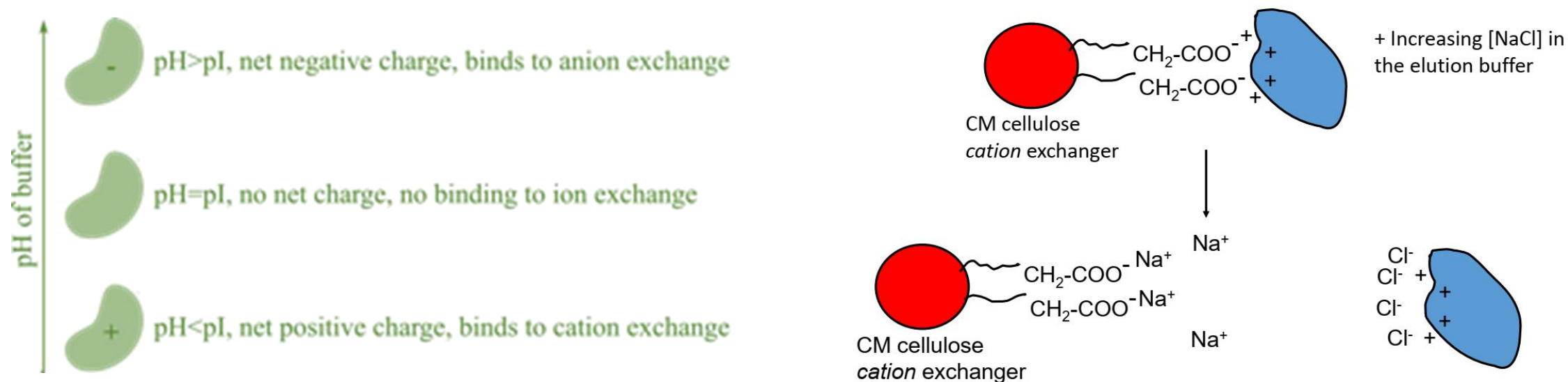
Comparison item	Bio-AC
Pump	Double plunger pump
Air sensor	+(Ultrasonic wave)
UV detector	-
Collector	Eight channel outlet valve
Control system	Touch screen
Column value	+(2 column position)
Pressure detector	+
Flow rate (ml/min)	0.1-20
Max pressure (MPa)	0.7
Price	Low price





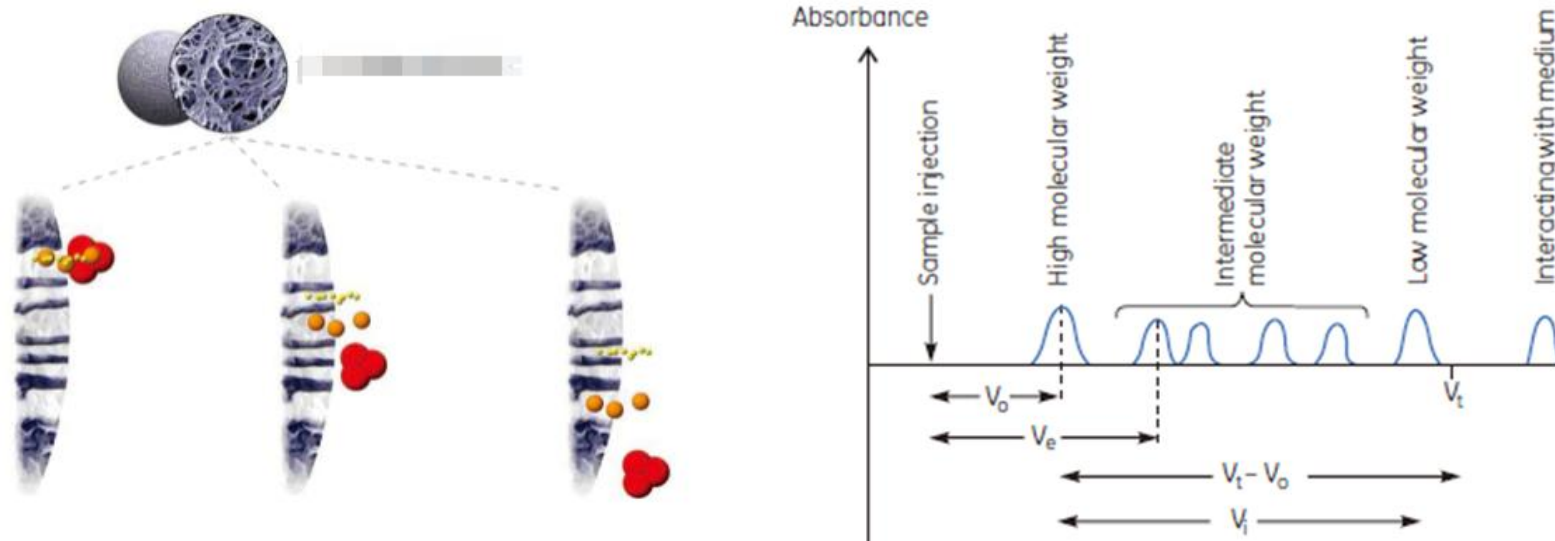
Polishing Purification: Ion Exchange Chromatography (IEC)

Group	Abbreviation	Exchange Type	Structure
sulphonate	S	strong cation	$-\text{CH}_2\text{SO}_3^-$
quaternary amine	Q	strong anion	$-\text{CH}_2\text{N}^+(\text{CH}_3)_3$



IEX	Mono Q 10/100 GL, Mono S 10/100 GL	8ml
	Resource Q, Resource S	6ml
	Source 15Q 4.6/100 PE, Source 15S 4.6/100 PE	1.7ml
	HiTrap Q HP / FF; HiTrap SP HP / FF	1ml, 5ml

Polishing Purification: Size-Exclusion Chromatography (SEC)

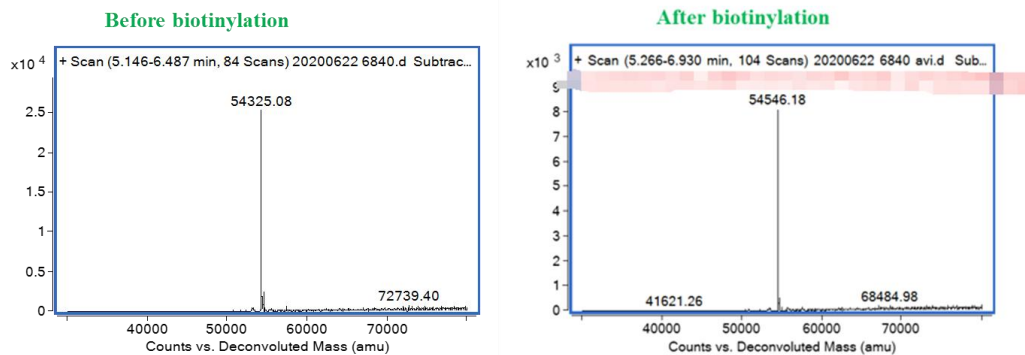


Column	Specification	Separation range	Sampling volume	Sampling volume	Flow rate	Pressure
Superdex 200 Increase 5/150 GL	3ml	QC, or small volume	<50 μ l		0.3ml/min	
Superdex 200 Increase 10/300 GL	24ml	10~600KD	<0.5ml	<10mg	0.5ml/min	3.5MPa
Superdex 75 Increase 10/300 GL	24ml	3~70KD	<0.5ml	<10mg	0.5ml/min	3.5MPa
Superose 6 Increase 10/300 GL	24ml	5~5000KD	<0.5ml	<10mg	0.5ml/min	3.5MPa
HiLoad 16/600 Superdex 200 pg	120ml	10~600KD	<5ml	<60mg	1.0ml/min	0.5MPa
HiLoad 16/600 Superdex 75 pg	120ml	3~70KD	<5ml	<60mg	1.0ml/min	0.5MPa
HiLoad 16/600 Superdex 30 pg	120ml	3~30KD	<5ml	<60mg	1.0ml/min	0.5MPa
HiLoad 26/600 Superdex 200 pg	320ml	10~600KD	<13ml	<120mg	1.5ml/min	0.5MPa
HiLoad 26/600 Superdex 75 pg	320ml	3~70KD	<13ml	<120mg	1.5ml/min	0.5MPa

Protein Preparation platform-Reaction of modification

➤ Biotinylation

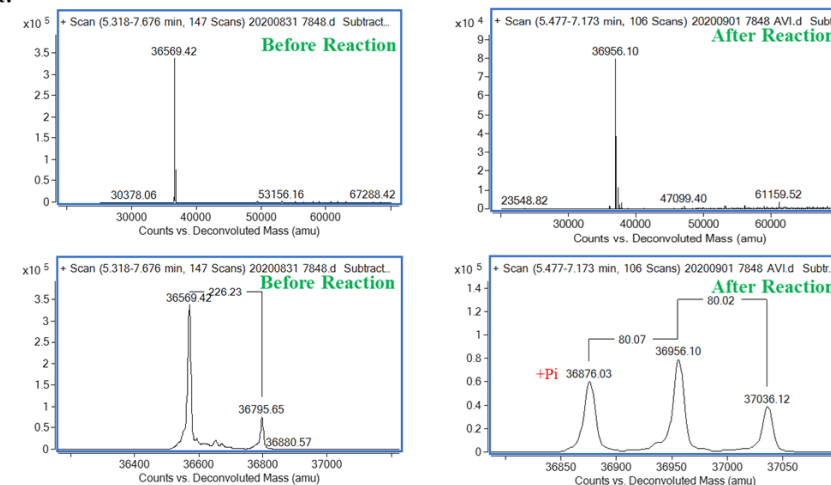
Commonly used enzymes: BirA



➤ Phosphorylation

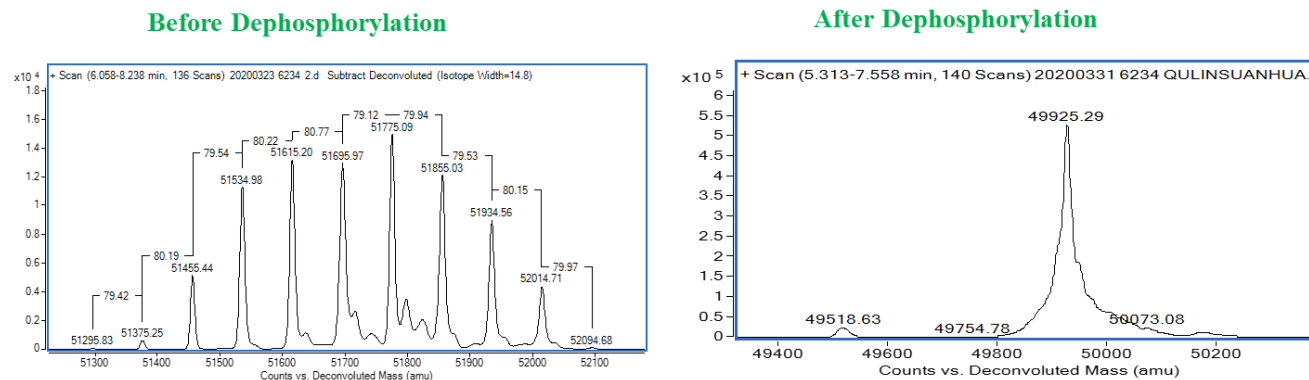
Commonly used enzymes: CAK, SRC

Result:

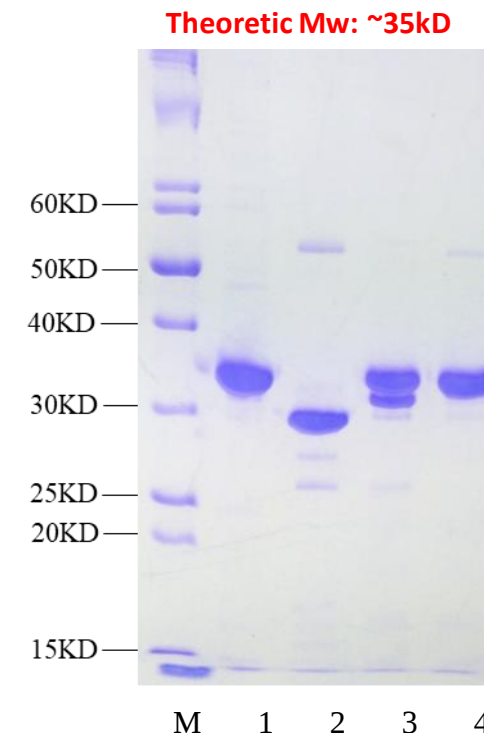
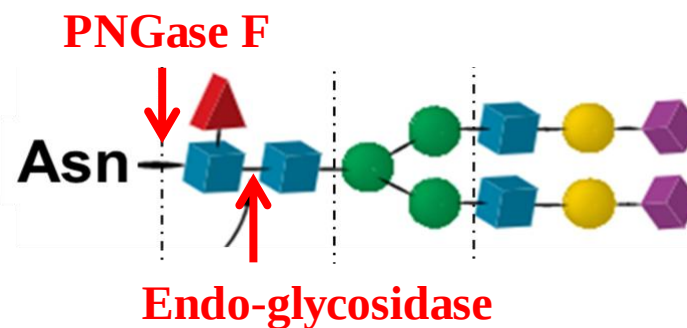
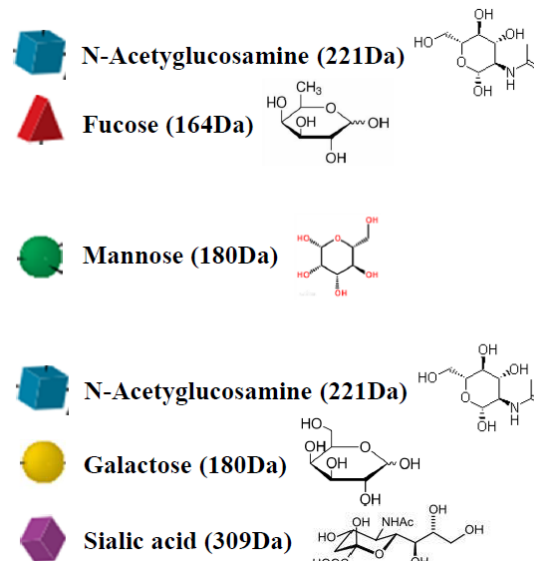
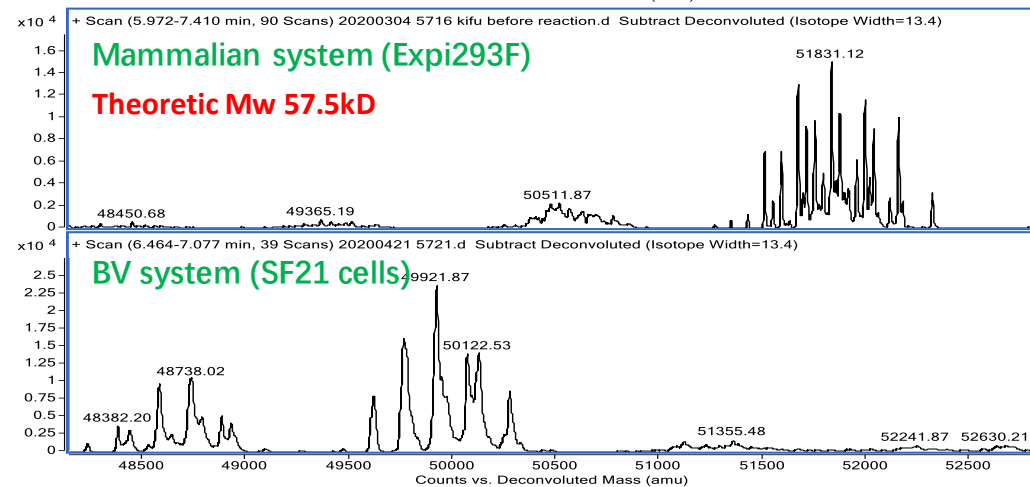
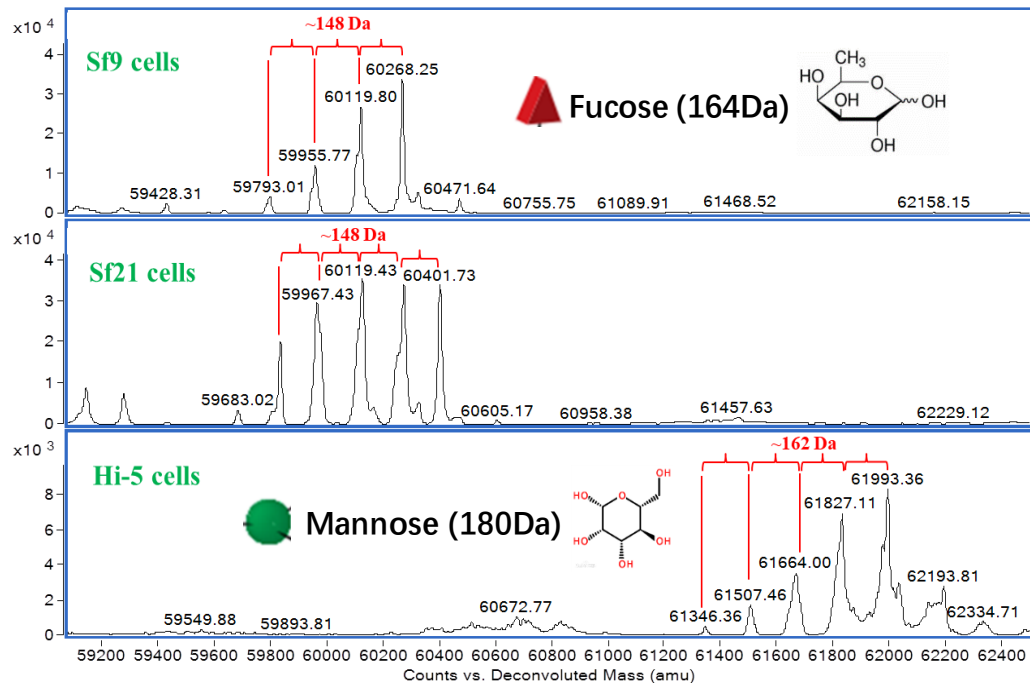


➤ Dephosphorylation

Commonly used enzymes: LamdaPP



Glycosylation in insect cell and mammalian cell expression



M = Marker
Lane 1 = before reaction
Lane 2 = Endo F1-overnight
Lane 3 = Endo F2-overnight
Lane 4 = Endo F3-overnight

(Re)Labeling with Sortase

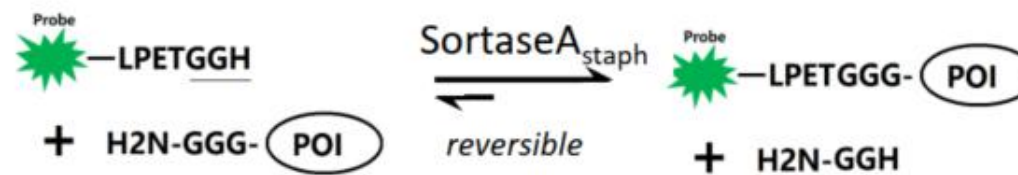
- Post-purification Tag addition
- Convenient & Specific Labeling with High-Efficiency
- Conjugate Dyes

For your pipeline

- Time saved: No need to start over for new affinity tags
- Common applications: Biotinylation for SPR, Fluorescent labeling

All constructs at BiorTus integrated

- His-StrepII-TEV-GG-POI → GGG-POI
- Natural Substrate for Sortase



Example: Fluorescent labeling with FAM & Alexa647

Item	No.	Amount of G	MW(Da)	Peptide	Label efficiency	Time in RT	Time in 4°C
1	6076	G	33851.39	FAM-Ahx-LPETG(2-hydroxyacetic acid)-GH	~100%	2h	2~4h
2	8168	GGG	25573.50	FAM-Ahx-LPETG(2-hydroxyacetic acid)-GH	~100%	1h	1~2h
3	8198	GGG	17948.03	FAM-Ahx-LPETG(2-hydroxyacetic acid)-GH	~100%	1h	1~2h
4	7109	G	17711.75	FAM-Ahx-LPETG(2-hydroxyacetic acid)-GH	~10%	4h	4h

Standard QC Package



SDS-PAGE

Purity
Stoichiometry



LC- MS

MW verification
Protein
Validation
PTMs



Analytical SEC

Solubility
Multimerization
Complex
Integrity

Custom QC Options



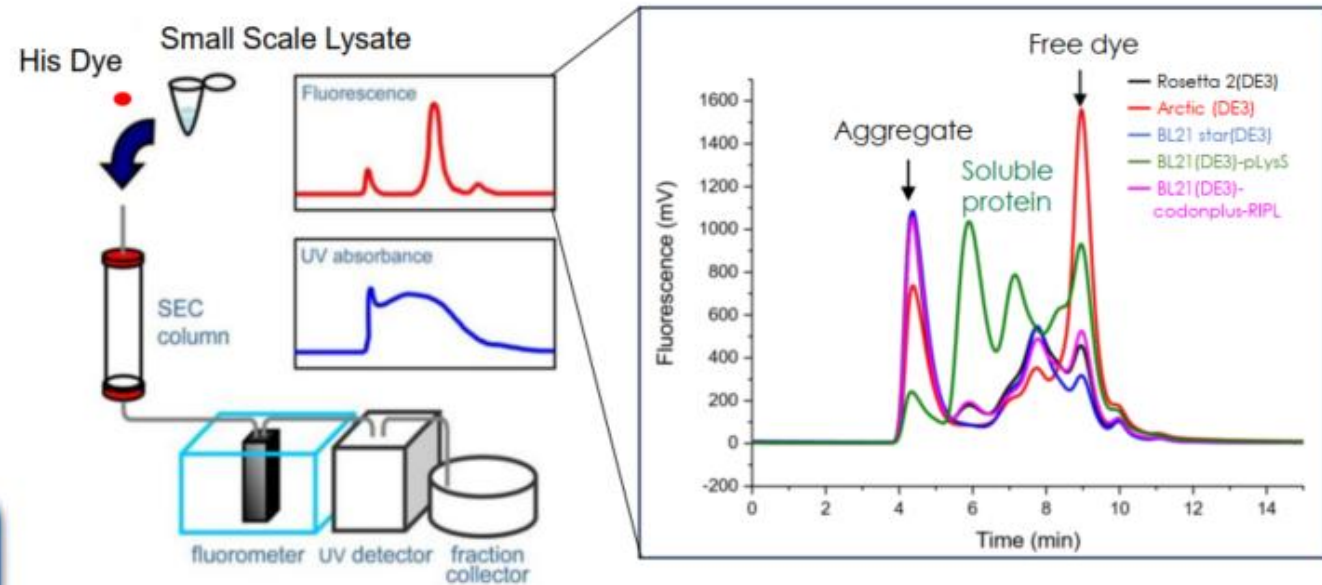
- Functional biochemical & biophysical assays
- SEC-MALS: oligomerization and solubility
- De-glycosylation
- Complex formation (gel-shift)
- Neg-stain EM
- Mass photometry
- DSF: T_m determination/stability

Through the Bottleneck: FSEC

- In parallel testing and monitoring
- Input – Small Scale Expression Lysate
- Visualization: His-Dye, GFP, Conjugated Antibody
- Up to 100 conditions or constructs /day /instrument (5 Total)

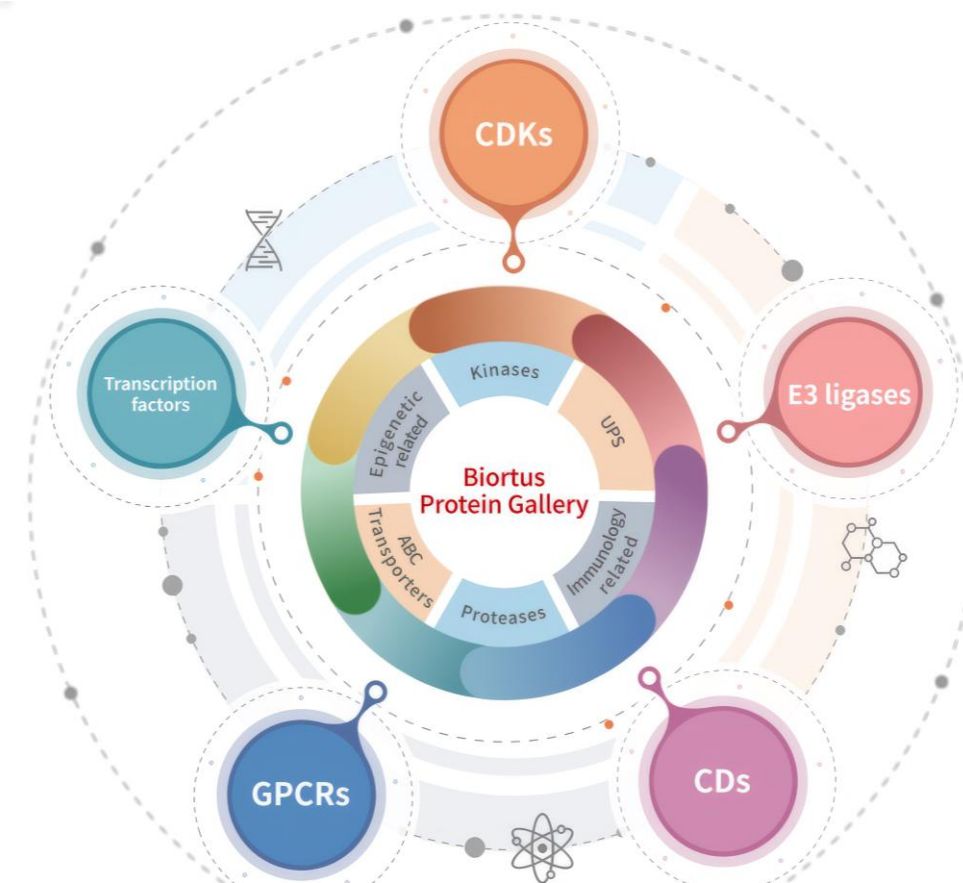
For your pipeline

- FL vs. Truncations vs. Mutations
- Expression Host System & Conditions
- Buffer & Solubility Screening
- Complex Formation



Specialize in Drug Targets

- Current inventory at over 2500 and expanding
- Focus on Drug Targets (E1, E2, E3, DUB, Ras, CDK, GPCR, SLC, etc.)
- High standards for QC
- >600 proteins, 17 assay kits are available on VWR and Chem-space and more products are coming.



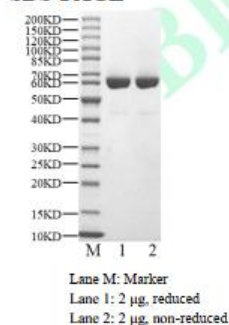
Delivery standard

STAT3

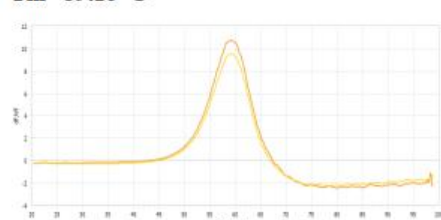
Category No.	BP39735-00A
Lot #	0001
Species	<i>Homo sapiens</i>
Uniprot ID/NCBI	P40763
AA sequence	G127-I722
Tags & Cleavage sites	No tag
Expression System	<i>E. coli</i>
92.6	67.9 KDa
Concentration	2.07 mg/ml **
Storage Temperature	-80°C
Storage Buffer	50 mM HEPES (pH7.5), 200 mM NaCl, 20% glycerol, 1 mM DTT
Notes	Please use rapid thawing with running water to thaw the protein samples. For complete recovery, mix well and spin before use. Product must not be store in diluted solutions. Aliquots below 10µl are not advisable. Avoid repetitive freeze-thaw cycles!

**Measured using UV absorption

SDS-PAGE



T_m = 59.16 °C

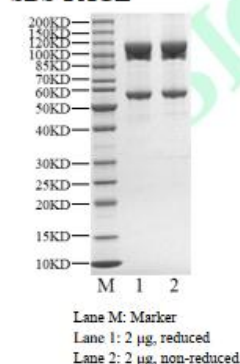


CRBN/DDB1

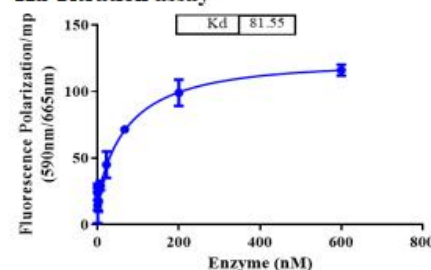
Category No.	BP12360-01A
Lot #	0001
Species	<i>Homo sapiens</i>
Uniprot ID/NCBI	Q96SW2/Q16531
AA sequence	M1-L442 end(CRBN) M1-H1140 end(DDB1)
Tags & Cleavage sites	N-His-TEV-Avi (CRBN) No tag (DDB1)
Expression System	Insect cell
MW	55.3 KDa/126.9 KDa
Concentration	1.71 mg/ml **
Storage Temperature	-80°C
Storage Buffer	50 mM Tris-HCl (pH7.5), 200 mM NaCl, 20% glycerol, 1 mM DTT
Notes	For complete recovery, mix well and spin before use. Product must not be store in diluted solutions. Aliquots below 10µl are not advisable. Avoid repetitive freeze-thaw cycles!

**Measured using UV absorption

SDS-PAGE



Kd Titration assay

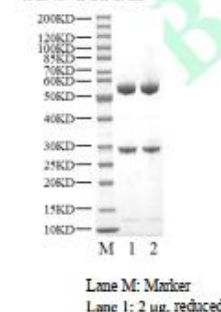


CDK2/Cyclin A2

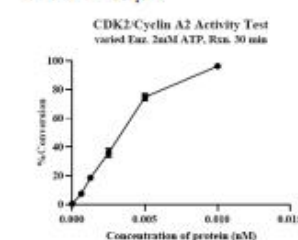
Category No.	BP469A2-21A
Lot #	0002
Species	<i>Homo sapiens</i>
Uniprot ID/NCBI	P24941/P20248
AA sequence	M1-L298(CDK2) E174-L432(Cyclin A2)
Tags & Cleavage sites	N-GST-3C(CDK2) N-6His-TEV(Cyclin A2)
Expression System	Insect cell
MW	60.4 KDa/32.8 KDa
Concentration	2.29 mg/ml **
Storage Temperature	-80°C
Storage Buffer	20 mM HEPES (pH 7.5), 200 mM NaCl, 5% glycerol, 1 mM DTT.
Notes	Please use rapid thawing with running water to thaw the protein samples. For complete recovery, mix well and spin before use. Product must not be stored in diluted solutions. Aliquots below 10µl are not advisable. Avoid repetitive freeze-thaw cycles!

**Measured using UV absorption

SDS-PAGE



Caliper assay ATP-K_m = 65 µM



Thanks!



Biortus Biosciences Co., Ltd.
<https://en.biortus.bio/>

For custom: info@biortus.bio

For off-the-shelf: order@biortus.bio