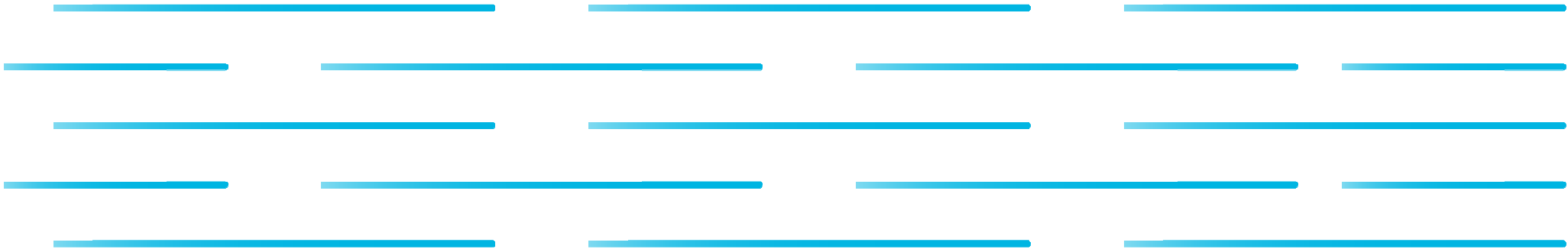




Vaccine downstream processing— an overview

DCVMN 10 March 2017



Outline

Vaccines overview

Demands on vaccine purification

Common techniques for vaccine purification

Example of a purification process

Summary



Vaccine overview

Vaccines and production

Vaccines

Bacteria based



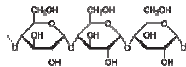
Virus based



Protein based



Polysaccharide based



The manufacturing process

Cell culture/
fermentation

Purification

Fill and finish

Analysis (QA/QC)

[E. Coli Bacteria](#) from NIH Image Library
[Influenza virus](#) by Kat Masback.



Demands on vaccine purification

Safety and quality is priority



- Regulatory requirements
- Vaccine with no or minimal adverse effects
- Effective dose
- Stability
- Process control
- Reproducible process

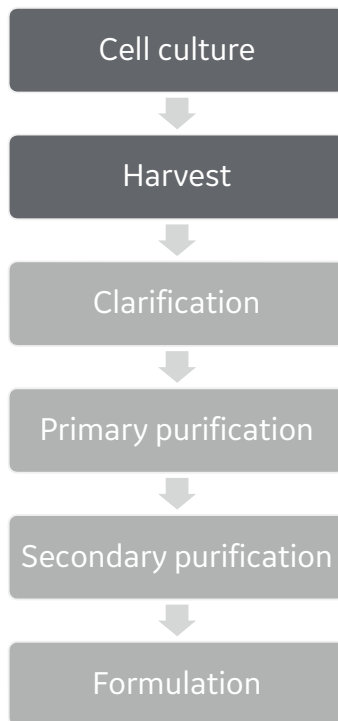
Vaccine image by www.torange-pt.com.



Available technologies for downstream purification of vaccines

Release of target molecules

Process flow



Lytic virus

Non-lytic virus

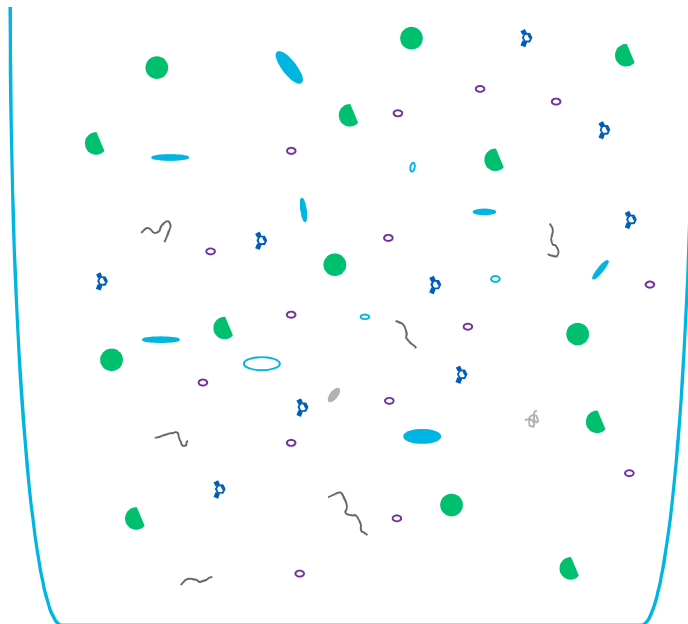
- Detergent
- Mechanical disruption/homogenization
- Osmotic shock
- Freeze-thaw

Yeast- and bacteria-based vaccines



Impurity challenges after lysis

After cell lysis



Antigen (e.g. virus) ●

Impurities

Process chemicals



DNA/RNA



Proteins



Cell membranes/organelles/lipids

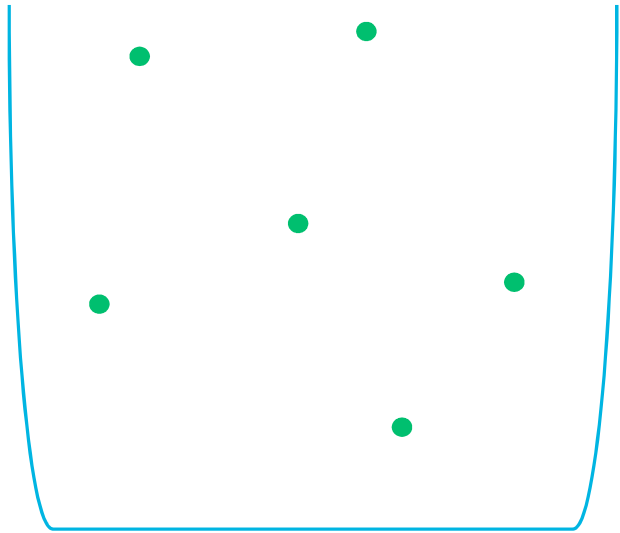


Antigen related impurities



Goal with purification

Purified sample

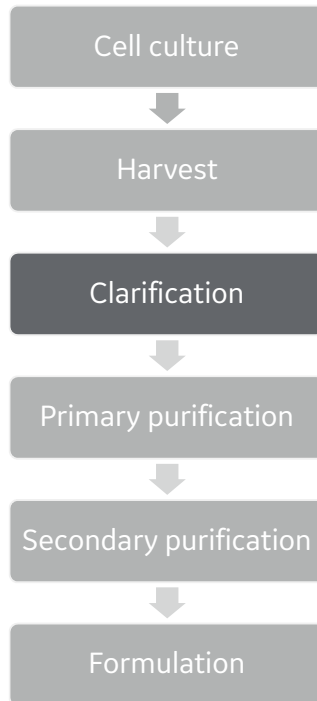


Antigen (e.g. virus) ●



Clarification

Process flow



Available techniques

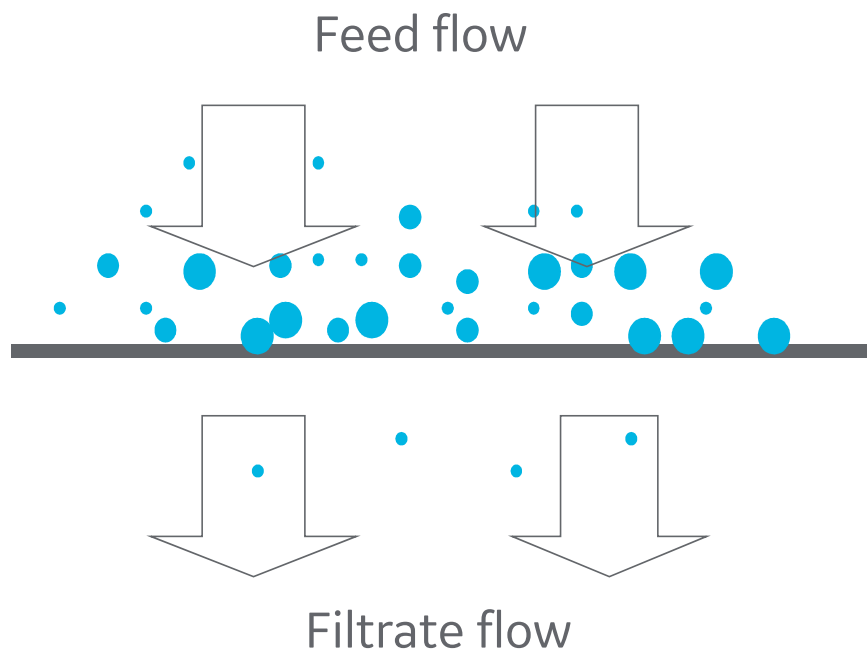
Filtration

- Normal flow (NFF)
- Tangential flow (TFF)

Centrifugation



Normal flow filtration (NFF)

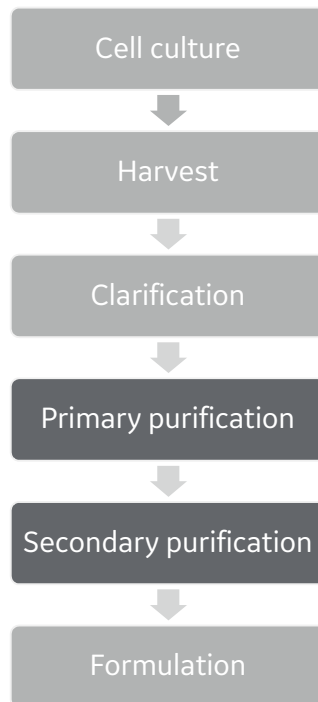


- Removal of cell debris and larger particulates
- Porosities from 0.2 to 20 μm
- Scalable
- Single-use technology
- Straightforward process set-up
- Not recommended for harvest with high particulate content



Purification

Process flow



Available techniques

TFF

Density gradient centrifugation

Selective precipitation

Chromatography

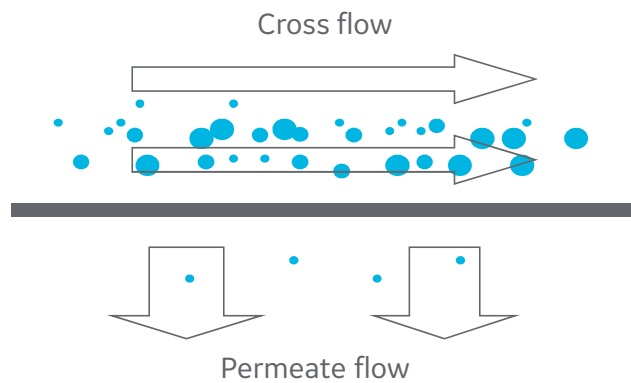
- IEX, MM, AC, HIC, SEC
- Bead format (packed bed)
- Membrane format (capsule)

Chromatography techniques:
AC = affinity, HIC = hydrophobic interaction
IEX ion exchange, MM = multi modal, SEC = size exclusion



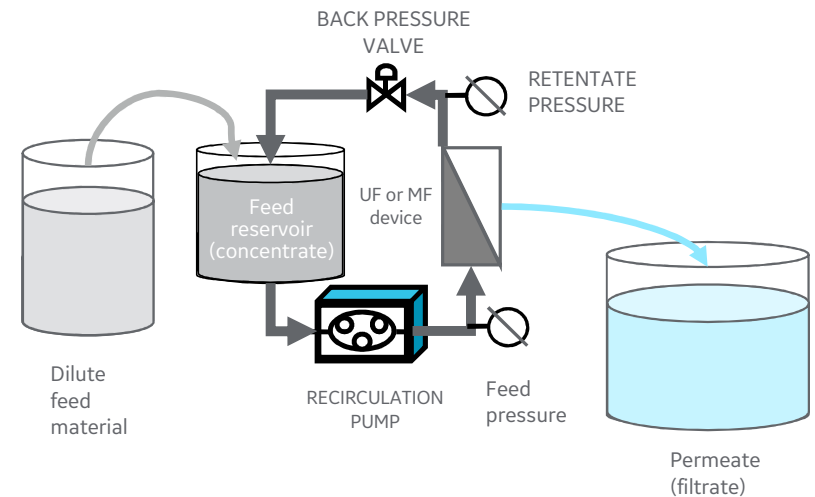
TFF

Basic principle



- Sweeping effect clean filter surface
- Allow greater throughput on smaller surface area

Schematic set-up



MF = micro filtration
UF = ultrafiltration



TFF

Flat sheet cassettes

Cassettes consists of sheet membranes

Concentration/diafiltration

Defined pore sizes

Reusable

Scalable



Hollow fiber filters

Hollow fiber cartridge consists of tubular fibers

Concentration/diafiltration

Microfiltration

Suitable for shear sensitive material

Possible to handle high particle loads
(e.g., cell harvest)

Defined pore sizes

Reusable

Scalable



Ion exchange chromatography (IEX)

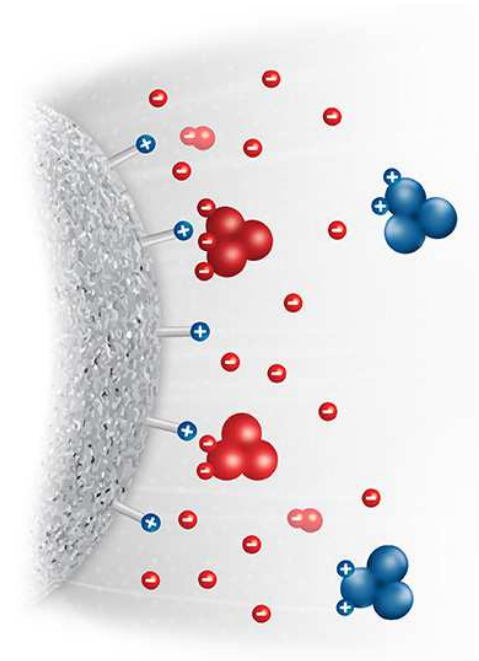
Anion exchange chromatography

(-) Negatively charged molecules binds to (+) positively charged ligands

Cation exchange chromatography

(+) Positively charged molecules binds to (-) negatively charged ligands

Principle



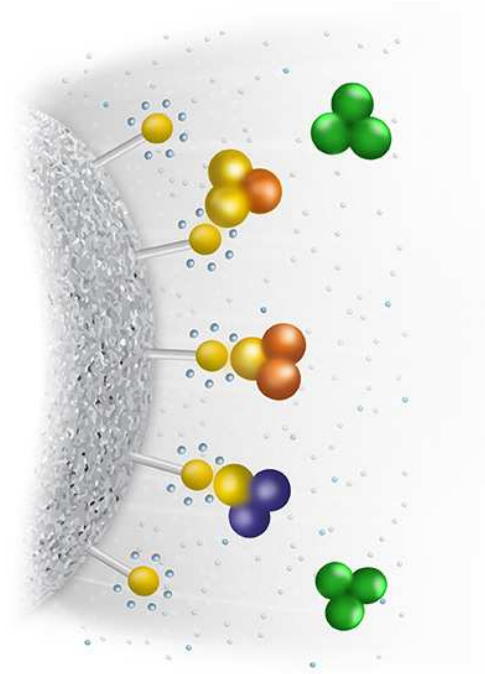
Hydrophobic interaction chromatography (HIC)

Separation by hydrophobicity

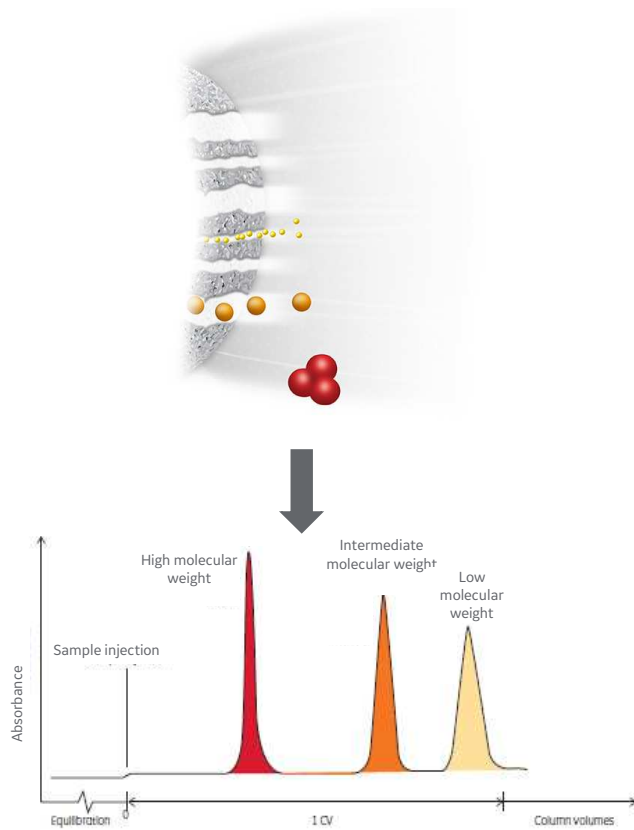
Hydrophobic surfaces of proteins interact with the ligand in presence of salts

High salt content enhance and low salt weakens the interaction

Principle



Size exclusion chromatography (SEC)



Excluded from pores



Enter a fraction of the pores



Enter all pores



Affinity chromatography

Specific binding

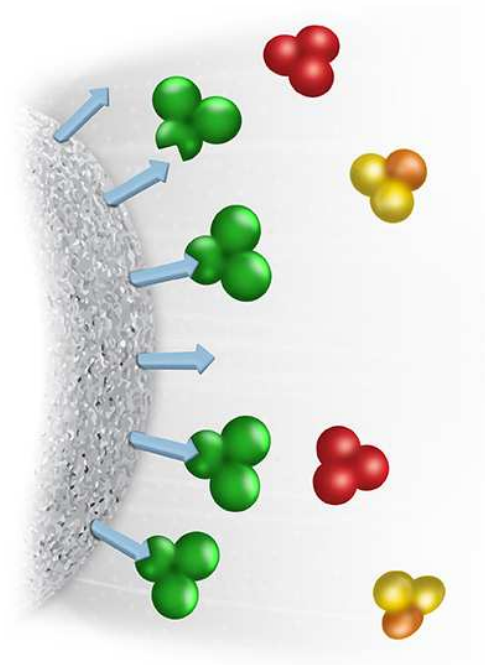
Few affinity resins available for vaccines

Agarose-based affinity resin for adeno associated virus

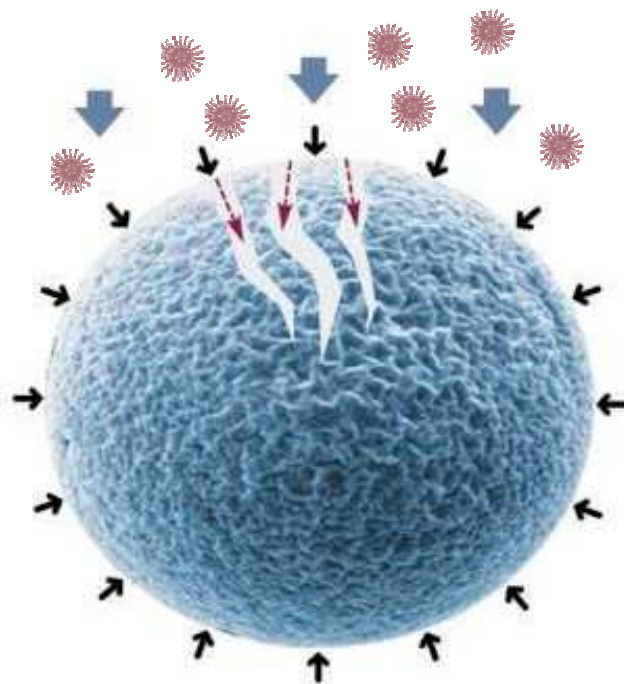
Pseudo affinity resins for influenza

- sulphated cellulose
- sulphated dextran

Principle



Chromatographic purification of large molecules can be challenging

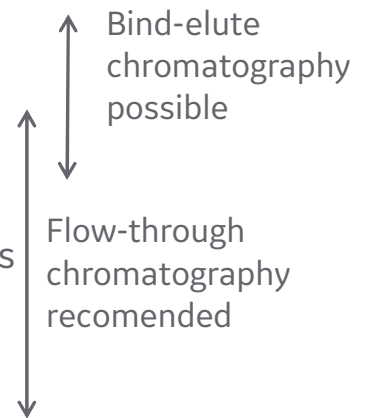


Pore size ~ 40-150 nm

- 1-7 nm proteins
- 25 nm polio
- 100 nm influenza virus

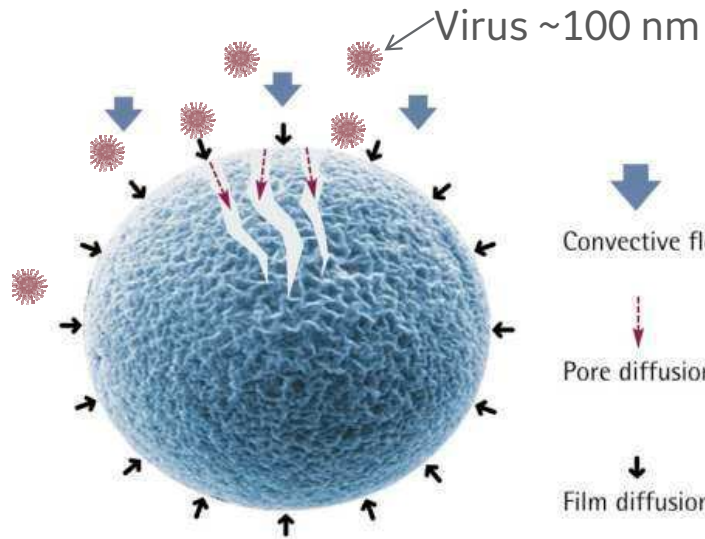


200 × 500 nm Pox virus



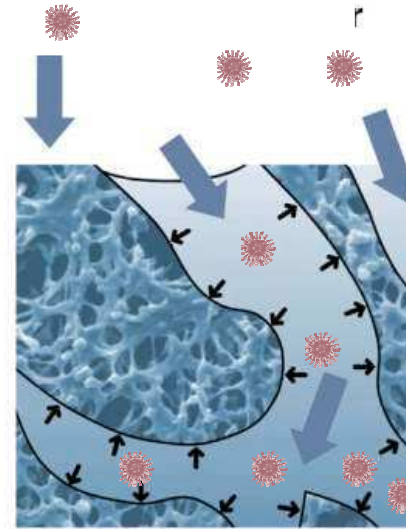
The pore size determines the properties

Chromatography beads
~ 2-6 min residence time



Pore size ~ 40-150 nm

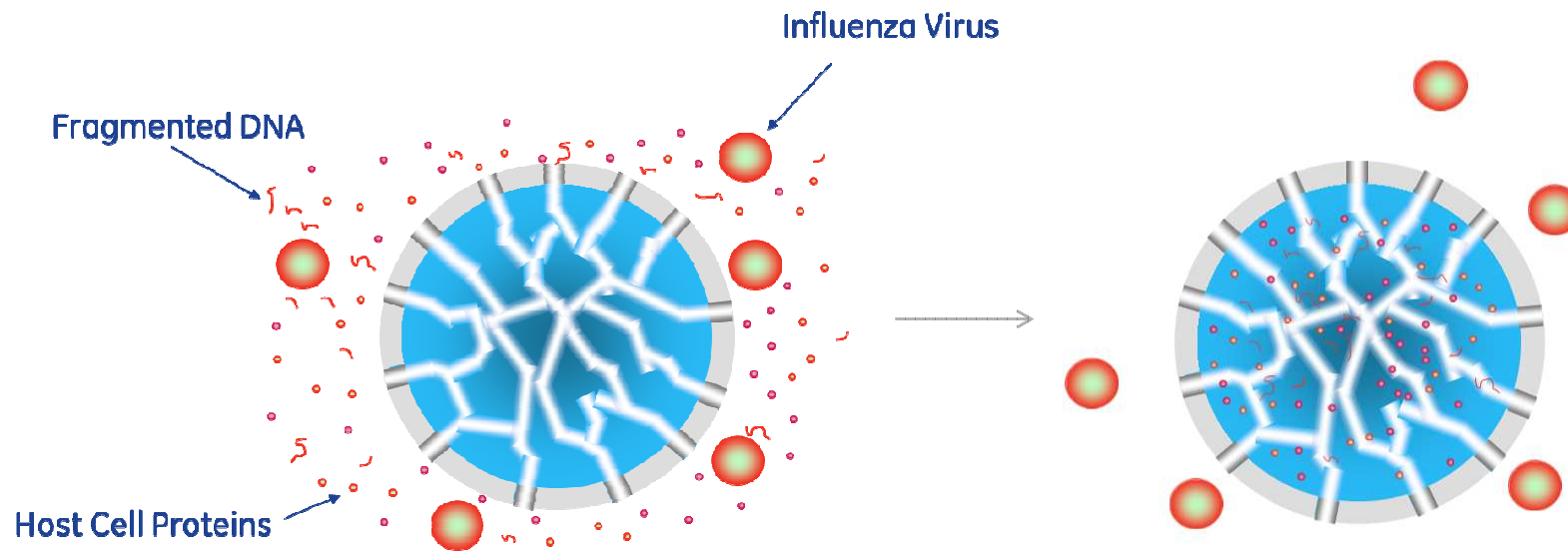
Membrane adsorbers
~ 2-60 s residence time



Average pore size 3-5 μm

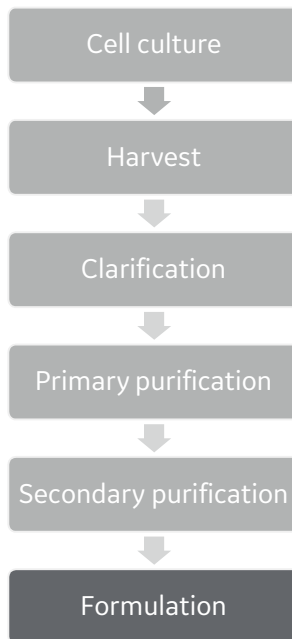


Core bead chromatography: host cell proteins and DNA fragments bind to the core and viruses stay in the void



Formulation

Process flow



Available techniques

Buffer exchange

TFF

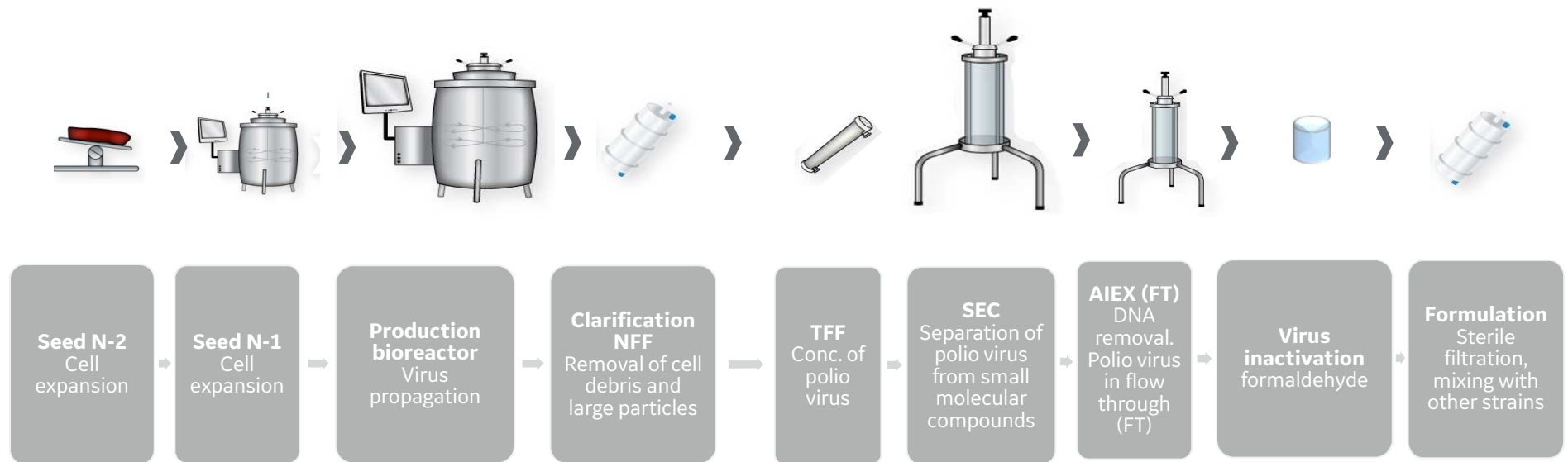
Chromatography

- SEC



Process example

Process example inactivated polio vaccine (IPV)



Summary: robust downstream process can ensure high quality

Most vaccines have unique purification processes

Preferably use scalable techniques when developing new processes

Purification of particles in binding mode can be difficult with classic chromatography

Core bead chromatography suitable for purification of particles > 700 kDa



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