Application note



Polishing mAb derived host cell proteins with HCPure[™] mixed-mode adsorbent

Protein A adsorbents are commonly used as the first chromatography step in downstream purification to capture monoclonal antibodies (mAbs) and fragments containing an Fc region. Intermediate and polishing steps are incorporated after the capture step to achieve the final purity required for the intended application. These chromatographic steps remove residual impurities, including host cell proteins (HCPs), host cell DNA, aggregates, leached Protein A, endotoxin, and a variety of viruses. This application note describes the optimal conditions for the use of HCPure[™] mixed-mode adsorbent for the polishing of mAb derived host cell proteins.

The primary capture of mAbs and fragments within this application note, use the GORE[®] Protein Capture Device, a composite PTFE membrane with a native-Protein A ligand. Use of intermediate washes with high salt, low pH, or arginine were evaluated for optimizing HCP removal on the GORE[®] Protein Capture Device. Post-affinity mAbs and fragments were then polished using HCPure[™], a novel adsorbent consisting of a proprietary mixed-mode ligand on beaded agarose.

HCPure[™]: A unique mixed-mode adsorbent

HCPure[™] primarily utilizes hydrogen bonding and hydrophobic interaction chromatography to bind host cell impurities while the target protein remains unbound. Figure 1 visualizes the chemical diversity of mixed-mode adsorbers, including HCPure[™] resin, by mapping ligand structures based on their functional group similarities. Each point represents a different resin, with its position determined by the properties of its ligand groups rather than a simple characteristic such as charge.

Clusters in the middle of the graph indicate resins with similar ligand structures, suggesting comparable binding behaviors. In contrast, HCPure[®] resin is positioned distinctly to the right, highlighting its unique ligand composition and different mode of action.

The distinct ligand chemistry of HCPure" resin, as shown in Figure 1, offers unique purification advantages compared to other resins with similar structures that may behave alike. This underscores the importance of selecting appropriate adsorbents for specific purification needs. This reinforces the value of a toolbox approach, allowing for resin selection based on specific process needs rather than a one-size-fits-all solution.



Figure 1: Chemical space diagram of commercial mixed-mode adsorbents.

Exploring the pH wash regime for Protein A capture

Clarified immunoglobulin G (IgG) produced in Chinese hamster ovary (CHO) cells was loaded onto a 1 mL GORE® Protein Capture Device with Protein A equilibrated with Phosphate-Buffered Saline (PBS). The device was washed with PBS, followed by a high-salt wash of PBS and 1.8 M NaCl. The salt was washed off with PBS before being eluted with 100 mM sodium citrate pH 3.4. An identical run was performed with a 100 mM sodium citrate pH 5 wash added prior to elution. IgG concentration was measured by UV, while aggregates, HCP levels, and HC DNA were assessed using SEC HPLC, ELISA, and PicoGreen, respectively. Table 1 summarizes the analytical results, showing that the run with the pH 5 wash had a similar impurity profile but a lower IgG yield.



| Reverse-phase HPLC | lgG recovery (mg) | lgG recovery (%) | Aggregate (%) | HCP concentration (ng/mg lgG) | HC DNA concentration (ng/mg lgG) |
|-----------------------|-------------------|------------------|---------------|----------------------------------|--|
| Control | 22 | 83 | 7 | 185 | 9 |
| With pH 5 wash | 12 | 44 | 4 | 184 | 11 |

Table 1: Comparison of IgG recovery and various impurity levels between the control capture step and the capture step with a pH wash.

Exploring an arginine wash regime for Protein A primary capture

An additional batch of CHO IgG was purified with two additional runs on a 9 mL GORE® Protein Capture Device. One of the runs eluted the IgG immediately after the PBS post-load wash, while the other was run with a high-salt wash, followed by an arginine wash prior to elution (Table 2). IgG concentration was measured by UV280, and HCP was measured by ELISA. High-salt and arginine washes reduced HCP impurities by approximately threefold while having minimal impact on IgG yield (Figure 2).

| Step | Buffer |
|----------------|------------------------------|
| Equilibration | PBS |
| Load | CHO harvest |
| PLW | PBS |
| High-salt wash | PBS + 1.8 M NaCl |
| EQ wash | PBS |
| Arginine wash | PBS + 0.1 M arginine pH 7.5 |
| Elution | 100 mM sodium citrate pH 3.4 |
| Strip | 0.1 M citric acid |
| CIP | 0.1 M NaOH |

 Table 2: Conditions for Protein A primary capture, including an arginine wash step.



Figure 2: Comparison of HCP content post-Protein A with highsalt and arginine washes.

Flow rate recovery of GORE[®] Protein Capture Device during cleaning

The same 9 mL device was used for more than 20 cycles of CHO IgG feedstock. The flow rate was automatically reduced to maintain the pressure below the limit (0.4 MPa), indicating an increase of pressure in subsequent runs (Figure 3). The flow rate was recovered after the extensive cleaning method (1), as indicated by the teal bars below (Figure 3).

Polishing host cell impurities with HCPure adsorbent

HCPure[™] is a tunable resin effective for post-affinity polishing of full-length IgG, IgG fragments, and bispecific antibodies from CHO, HEK293, *E. coli*, and Pichia cell lines. Optimal running conditions for HCPure[™] resin rely on the host cell and the target protein, therefore a screen of pH (4-9) and conductivities (6-18 mS/cm) should be performed for each target. Astrea Bioseparations offers 0.2 mL RoboColumns[®] and 1 mL pre-packed columns with HCPure[™] which are invaluable tools for screening.



Extended cleaning method recovers flow rate on GORE® Protein A device loading



| Buffer | CV | | | | |
|--|-----------|--|--|--|--|
| 200 mM arginine pH 2.3 | 5 | | | | |
| 50 mM Tris, 100 mM NaCl, 1% SDS, 10 mM DTT pH 10.4 | 10 | | | | |
| (10-minute hold) | | | | | |
| (10-minute hold) | | | | | |
| (10-minute hold) 200 mM arginine pH 2.3 | 10 | | | | |
| (10-minute hold) 200 mM arginine pH 2.3 0.2 M NaOH | 10 3.6 | | | | |

(1) GORE® cleaning regime

Figure 3: Effect on extended cleaning on the GORE^{\otimes} Protein A device flow rate.

Broad range screen for buffer conditions using HCPure[™] adsorbent

In another example, a post-Protein A elution fraction of V κ -overexpressed in *E. coli* was screened for HCP removal on HCPure[®] RoboColumns[®], using a factorial DOE with the following pH and conductivity conditions in 40 mM sodium phosphate:

- pH 6, 6 mS/cm
- pH 6, 18 mS/cm
- pH 7, 12 mS/cm
- pH 8, 6 mS/cm
- pH 8, 6 mS/cm

In this example, the DOE identified a significant improvement in HCP reduction at pH 6 compared to pH 8, from 1.6 log to 2 log (Figure 4). The V κ yield ranged from 75-96; however, no significant correlation was observed between the V κ yield and pH or conductivity parameters. Based on the predictions from Design-Expert[®] 13 software, optimal conditions of pH 6 and 18 mS/cm were expected to achieve a 2.19 log reduction in HCPs and an 89% yield of the V κ fragment. In another example, a post-Protein A elution fraction of V κ -overexpressed in *E. coli* was screened for HCP removal on HCPure[®] RoboColumns[®], using a factorial DOE with the following pH and conductivity conditions in 40 mM sodium phosphate:

- pH 6, 6 mS/cm
- pH 6, 18 mS/cm
- pH 7, 12 mS/cm
- pH 8, 6 mS/cm
- pH 8, 6 mS/cm

In this example, the DOE identified a significant improvement in HCP reduction at pH 6 compared to pH 8, from 1.6 log to 2 log (Figure 5). The V κ yield ranged from 75-96; however, no significant correlation was observed between the V κ yield and pH or conductivity parameters. Based on the predictions from Design-Expert® 13 software, optimal conditions of pH 6 and 18 mS/cm were expected to achieve a 2.19 log reduction in HCPs and an 89% yield of the V κ fragment.

These two examples demonstrate the importance of optimizing HCPure $\Tilde{\times}$ resin conditions based on the specific requirements of each target protein.

Screening pH and conductivity conditions for IgG recovery and HCP reduction on HCPure[™]



Figure 4: Broad range screen for HCP removal conditions from a CHO-produced IgG using HCPure[™] resin.





Figure 5: Heat map illustrating the impact of various pH and conductivity conditions on HCP removal efficiency for a V κ fragment produced in *E. coli*. The analysis suggests optimal conditions for both HCP reduction and V κ yield represented by the red regions. Sub-optimal conditions are highlighted by the blue regions.. These examples demonstrate the importance of optimizing HCPure^{**} conditions to the specific requirements of each target protein.

Polishing of HEK293 IgG using HCPure[™] adsorbent

A mAb produced in HEK293 cells was purified using a GORE® Protein Capture Device, followed by polishing with HCPure™ adsorbent. The column was equilibrated with 40 mM sodium phosphate, 160 mM NaCl, pH 6, 18 mS/cm. The Protein A elution fraction was dialyzed into the equilibration buffer before loading 30 mg on to a 1 mL HCPure™ pre-packed column. Unbound IgG was collected, and its concentration was measured by UV. HCP and Protein A leachate were measured by ELISA, HC DNA using PicoGreen, and endotoxin levels using a limulus amebocyte lysate (LAL) assay.

The IgG recovery was 80% with a significant reduction of HCP, dsDNA, Protein A leachate, and endotoxin impurities (Figure 6).

Conclusion

These studies demonstrate the effectiveness of HCPure[™] mixed-mode adsorbent as a polishing step across multiple expression systems, for the removal of HCP, Protein A leachate, and other impurities following Protein A capture. High-salt and arginine washes during Protein A chromatography significantly reduced HCP levels while maintaining IgG yield. Further purification with HCPure[™] resin efficiently reduced residual impurities, including HCP, host cell DNA, Protein A leachate, and endotoxin, bringing them within regulatory guidelines.

By utilizing a unique chemical space compared to other commercially available adsorbents, HCPure[™] offers a valuable addition to a purification toolbox. HCPure[™] resin's unique mixed-mode ligand chemistry provides a versatile and effective solution for polishing full-length IgG, IgG fragments, and bispecific antibodies across multiple expression systems, including CHO and HEK293. Its ability to selectively remove impurities while preserving target protein yield reinforces its value as a robust tool for achieving high-purity biologics. By screening each target protein and optimizing wash and polishing conditions, researchers and manufacturers can enhance process efficiency and meet stringent purity requirements for downstream applications.

References

- (1) GORE® application note: Cleaning protocols to reduce the effects of pressure rise over the lifetime of a GORE® Protein Capture Device
- (2) Molden R, et al. (2021). Host cell protein profiling of commercial therapeutic protein drugs as a benchmark for monoclonal antibody-based therapeutic protein development. mAbs, 13(1). https://doi.org/10.1080/19420862.2021.1955811



Figure 6: Impurity removal from HEK IgG with a HCPure[™] adsorbent polishing step.

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