Monolith-like Particle (MLP) for Next-Generation Virus Purification -Full/empty AAV Capsids Separation and Vaccine Production-

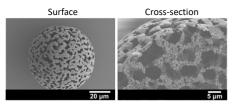
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Abstract

JNC has developed a novel cellulose-based monolith-like particle (MLP) with large through-pores. The MLP is an innovative bead architecture with high industrial applicability, optimized for the efficient separation of large targets such as viral particles. Structural analysis revealed that the MLP possesses a unique architecture with large through-pores, enabling easy access for large targets to reach the intraparticle chromatographic surface. This poster will provide detailed insights into the development of the MLP and explore their potential applications in biomolecule and virus purification processes.

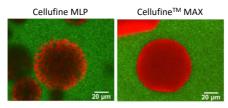
Introduction of Cellufine MLP

Fig. 1. SEM Morphology of Cellufine MLP.



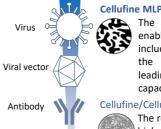
The through-pores of MLP were estimated to have a mode pore radius of 1.5 µm by using mercury porosimetry [1]. The average particle diameter of MLP is about 90 µm.

Fig. 2. CLSM observation of Cellufine MLP



Green fluorescent nanoparticles (100 nm) can diffuse into the intraparticle area of Cellufine MLP.

Target biomolecules



Small proteins

The large porous enables large biomolecules, including virus particles to access the intraparticle surface area, leading to high dynamic binding capacity.

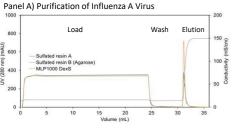
Cellufine/Cellufine MAX



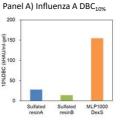
The resin with small pores shows high dynamic binding capacity due to its high specific surface area for small targets.

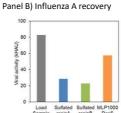
Cellufine MLP1000 DexS - Pseudo Affinity Bind and Elute Purification of Influenza A virus and SARS-CoV-2

Fig. 3. Bind and Elution Chromatogram of virus purification. Fig. 4. Results of purification process - Virus recovery and contaminants removal

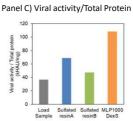


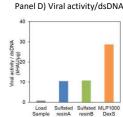




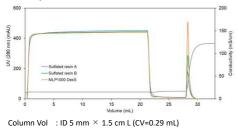


Panel E) SARS-CoV-2 recovery

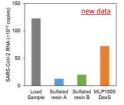


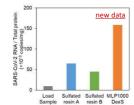


Panel B) Purification of SARS-CoV-2

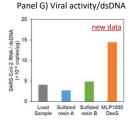


: 0.5 mL/min Load Sample: Panel A; Influenza A virus (H1N1) from MDCK cells : Panel B : SARS-CoV-2 XBB





Panel F) Viral activity/Total Protein



- ✓ MLP1000 DexS showed high dynamic binding capacity (DBC_{10%}) for influenza A virus and SARS-CoV-2.
- The purification efficiency leading to removal of contaminants by the MLP1000 DexS in bind and elute mode is higher than that of conventional sulfated resins.

Cellufine MLP1000 AEX resin for AAV empty/full particle — Comparison with benchmarks

Figure 5. Comparison of chromatograms between MLP1000 AEX prototype and benchmarks

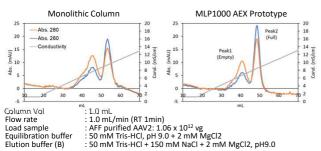


Table 1. Comparison of quantitative results

		260/280			
	*Full (%)	Peak1	Peak2 (peak top)	Peak2 (Peak area)	Resolution
Before purified	15.9	-	-	-	-
Commercial AEX	-	0.60	1.20	-	-
Monolithic Column	42.8	0.64	1.22	1.19	0.83
MLP1000 AEX Prototype	70.6	0.58	1.27	1.24	1.09

*Determined by Mass Photometry

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Conclusion

- A novel cross-linked Cellufine MLP beads with large pore structure allowing large biomolecules to access the intraparticle area have been developed.
- MLP1000 DexS exhibits a high dynamic binding capacity and high purification efficiency for whole virus particles.
- \checkmark MLP based AEX resin demonstrated high capacity and superior separation compared to benchmark.

- [1] K. Kadoi, E. Iwamoto, T. Nakama, Fabrication and characterization of a cellulose monolith-like particle for virus purification, Biochem Eng J. 192 (2023) 108849.
- [2] K. Kadoi, J. Toba, A. Uehara, N. Isoda, Y. Sakoda & E. Iwamoto, Enhanced sulfate pseudo-affinity chromatography using monolith-like particle architecture for purifying SARS-CoV-2, Vaccine. 53 (2025) 126951.





[√] The MLP1000 AEX showed superior separation of empty and full AAV2 particles compared to benchmarks.