

Characterization of lentiviral vectors using capillary electrophoresis platform technology

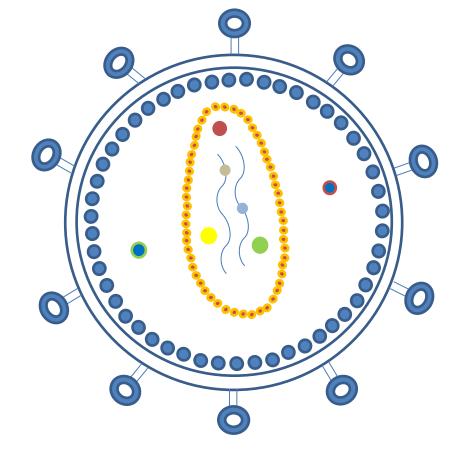
Aaron Shafer, PhD St. Jude Children's Research Hospital





- Lentiviral vector (LVV) background and analytical characteristics
- Platform methods for analyzing LVV based products LVV proteome analysis
 - Titer determination from p24 protein
 - Lentiviral vector protein profiling
 - LVV genome analysis
 - Transgene sizing and impurity screening
- Summary

Why is lentivirus proteome/genome analysis important?



Lentivirus

Structural proteins:

- Nucleocapsid proteins: p7
- Capsid proteins: p24
- Matrix proteins: p17
- Env. proteins: GP120, GP41

Non-structural proteins:

- Enzymes (4)
- Gene Regulatory Proteins(2)
- Accessory Proteins (5)
- HCPs
- etc.

Proteome

- Envelope Proteins: The glycoprotein complexes enable the virus to attach to and fuse with target cells to initiate the infectious cycle
- P24 protein is typically used for lentiviral titer determination, a critical quality attribute (CQA) for lentivirus production from lab scale to GMP manufacturing
- Protein profiling of the lentiviral vector by CE can provide enhanced specificity for titer determination as well as the assessment of multiple structural and non-structural proteins

Genome

- Product-related impurities can cause reduced transduction efficiency and efficacy:
 - Empty LVV particles
 - LVV containing a truncated transgene or incomplete 3' end
- Process-related impurities can be toxic to target cells and reduce transduction efficiency:
 - Residual host cell DNA and/or plasmid DNA
 - Increases the risk for innate immune response
- Lentiviral genome integrity analysis by CE offers high resolution insight into the quality of your therapeutic transgene



Analysis of Lentiviral Proteins

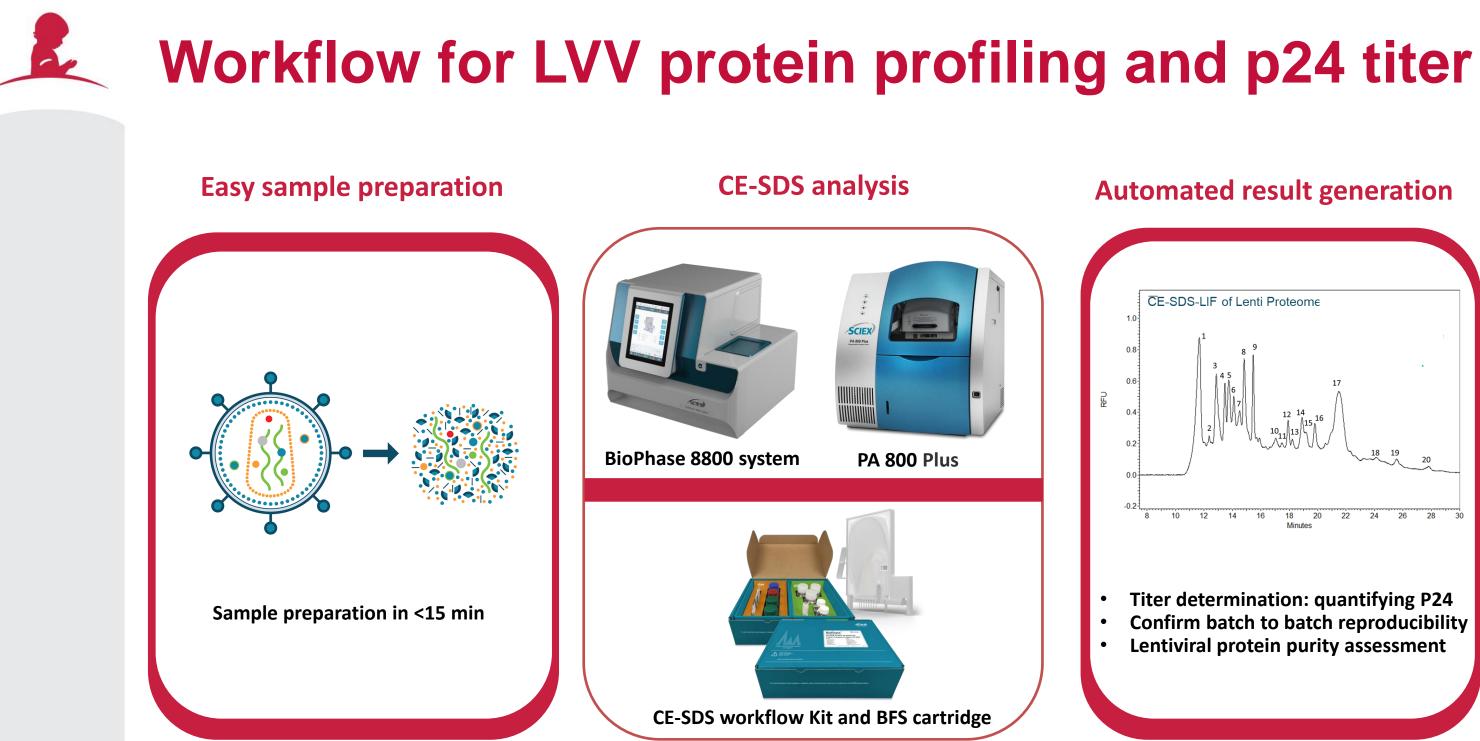
- Titer determination from p24 protein
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Analysis of Lentiviral Genome

Genome sizing and impurity screening

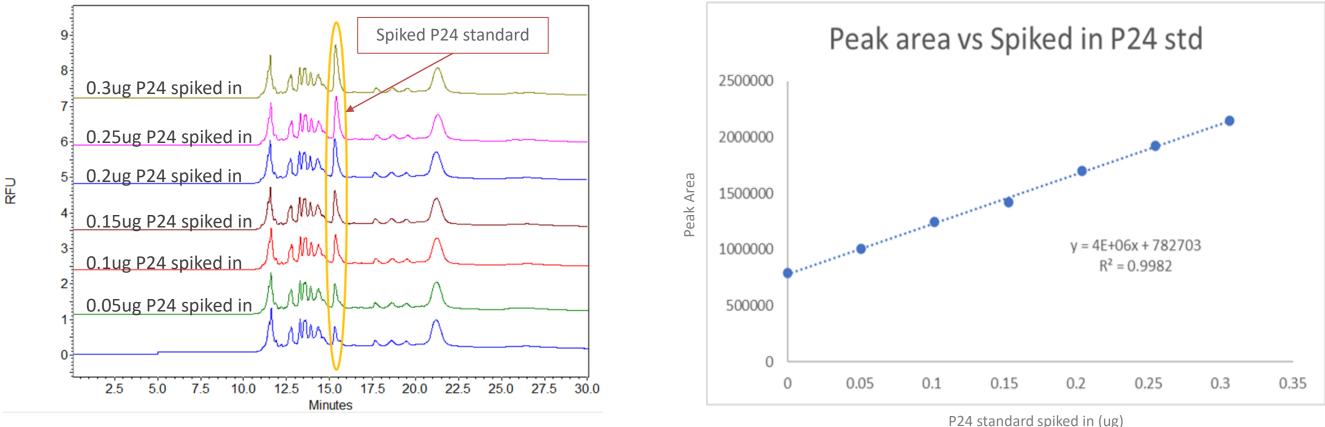






Identification of the P24 peak

P24 protein standard addition method



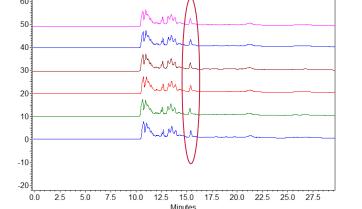
Identification of the P24 peak by standard addition to lentivirus sample of 1X10⁹ TU/mL



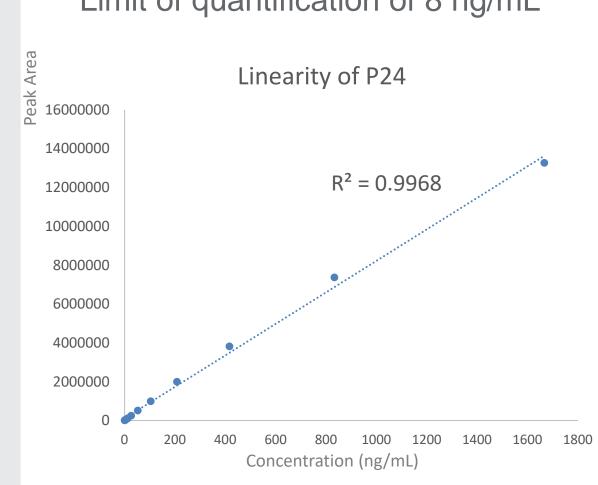
Linearity and LOQ

Repeatability assessment

Six consecutive injections 60 Man RFU Man Much

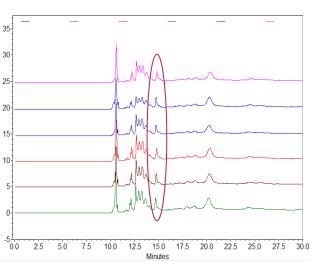


Injection	MT (Min)	Peak area
1	15.57	3594174
2	15.60	3615001
3	15.57	3621318
4	15.61	3594266
5	15.62	3565088
6	15.65	3535125
RSD%	0.20	0.91



Limit of quantification of 8 ng/mL

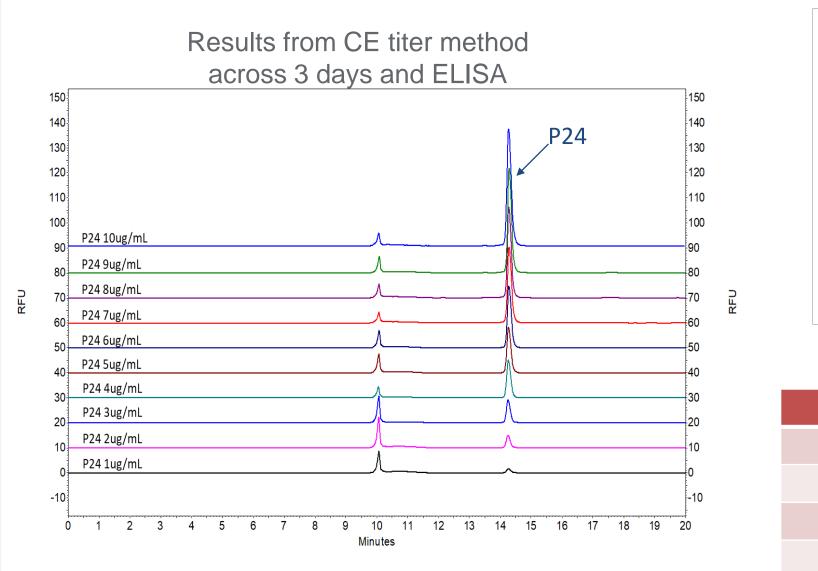
Six sample preparations

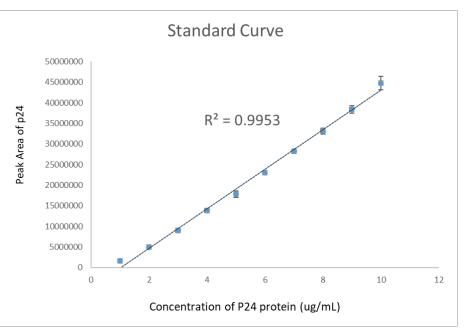


Injection	MT (Min)	Peak area
1	14.62	1874688
2	14.62	2145261
3	14.68	2013849
4	14.69	1974002
5	14.73	1965482
6	14.76	1996383
RSD%	0.39	4.41

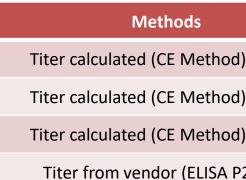
Calibration Curve and P24 titer determination

P24 standard curve from 3 sample preparations





Comparison between LVV P24 titer methods



- on 3 separate days
- P24 titer by CE is comparable to the p24 titer by ELISA .

	Titer (TU/mL)
d) - Day 1	1.45 X 10 ⁹
d) - Day 2	1.45 X 10 ⁹
d) - Day 3	1.46 X 10 ⁹
P24 kit)	1.5 X 10 ⁹

CE quantitation demonstrated consistency across 3 preparations



Analysis of Lentiviral Proteins

- Titer determination from p24 protein
- Lentiviral vector protein profiling

Analysis of Lentiviral Genome

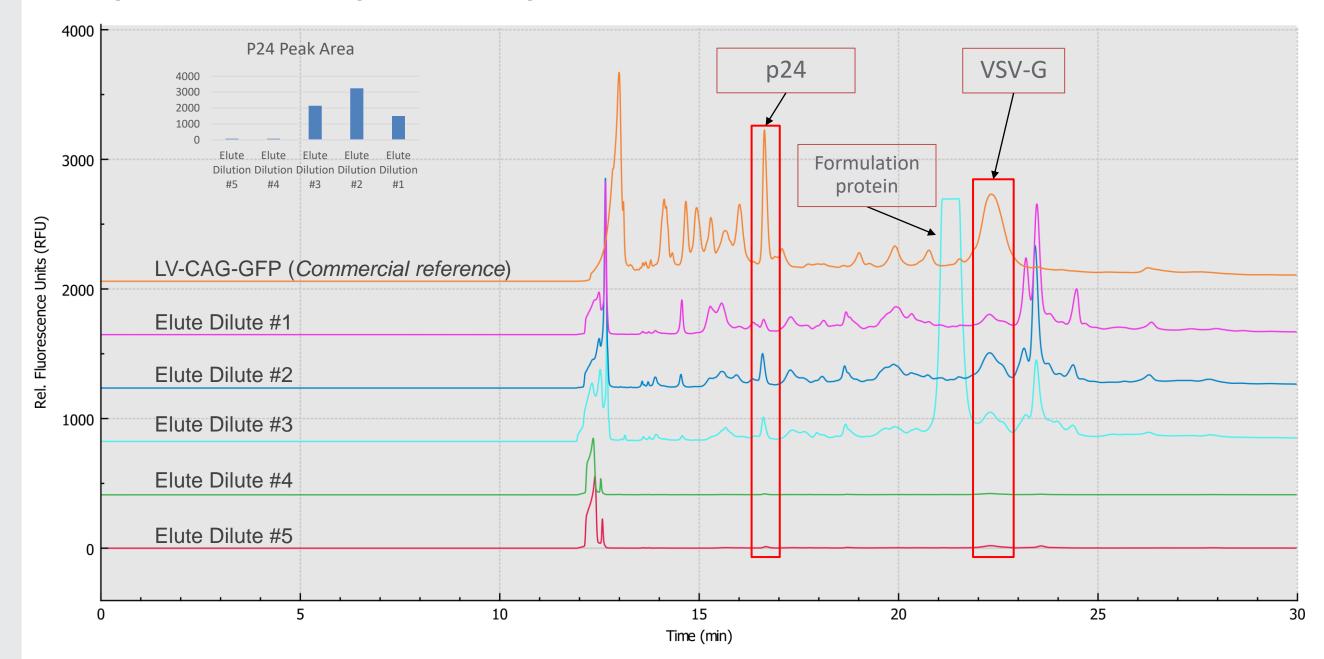
Genome sizing and impurity screening





Protein profiling to assess in-process samples

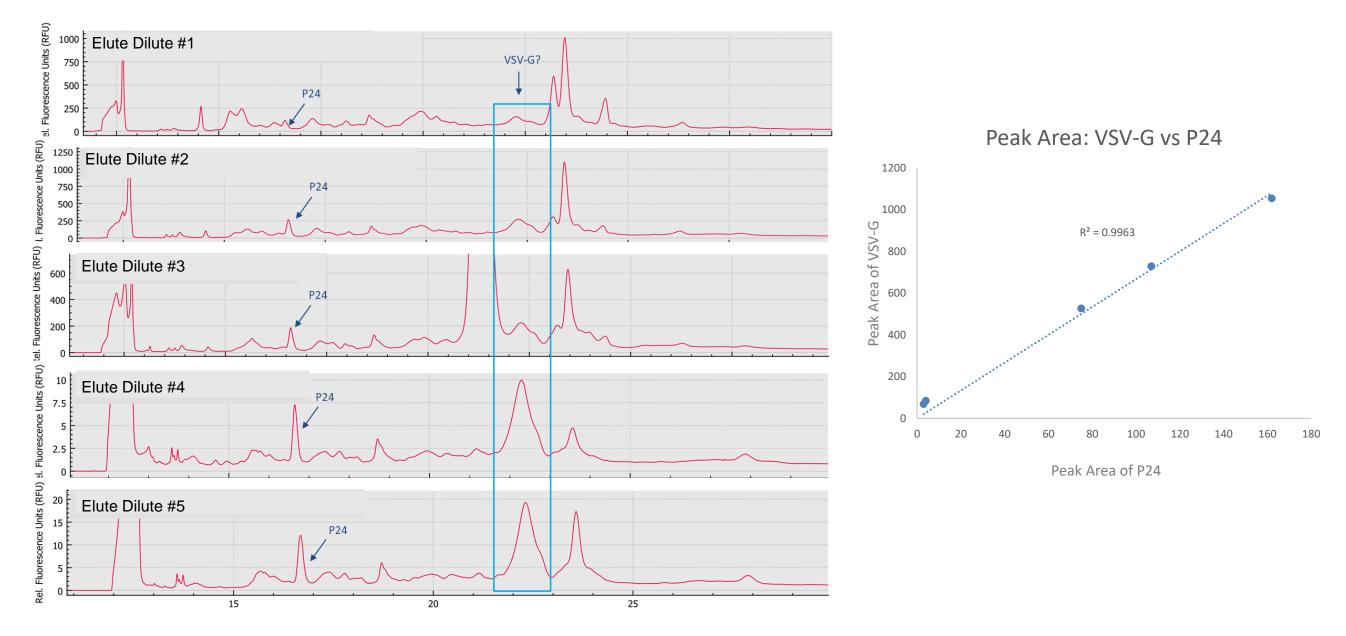
In-process samples compared to a commercial reference material





Protein profiling to assess in-process samples

Linear correlation of VSV-G to the P24 LVV proteins







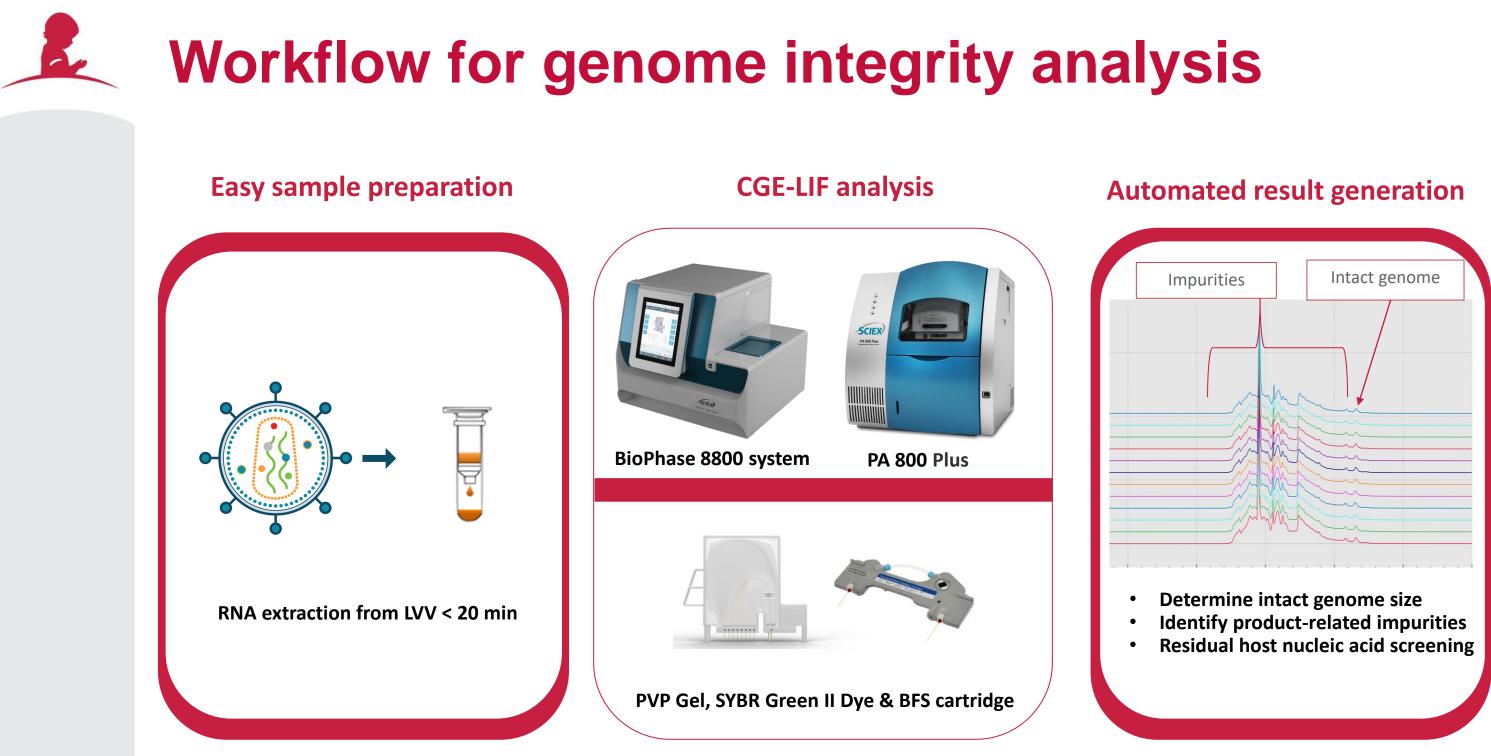
Analysis of Lentiviral Proteins

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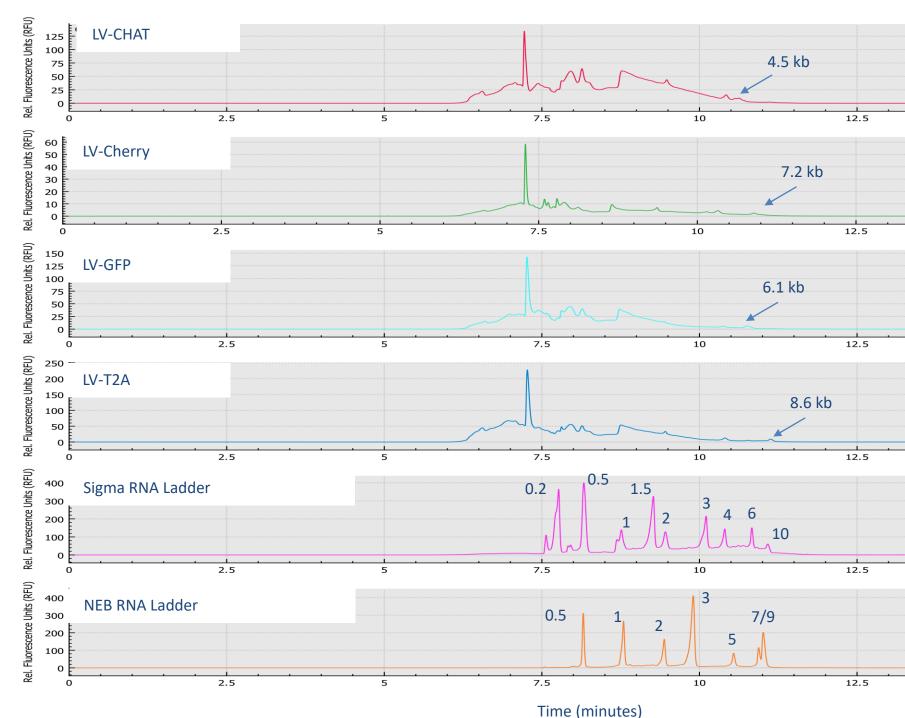
Analysis of Lentiviral Genome

Genome sizing and impurity screening



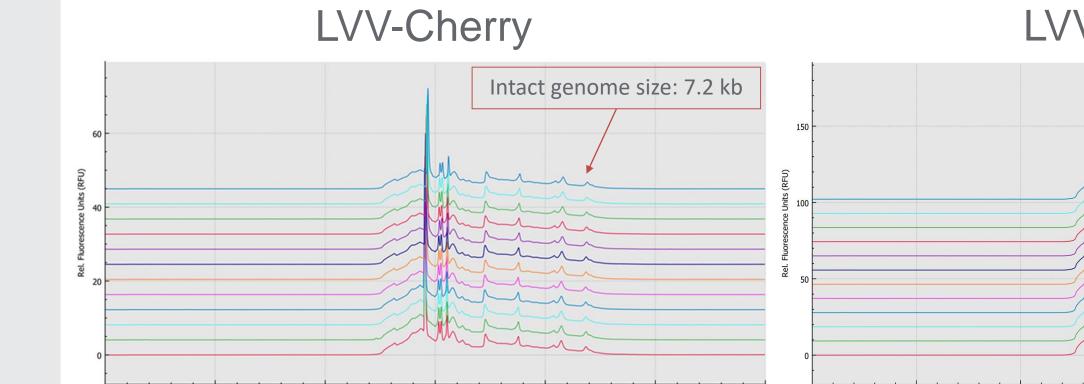


Analysis of LVVs with various genome sizes





Genome integrity workflow consistency



10

7.5

Time (min)

5

2.5

Consistent migration time for the intact genome peak across 12 consecutive injections of 2 commercial samples

12.5

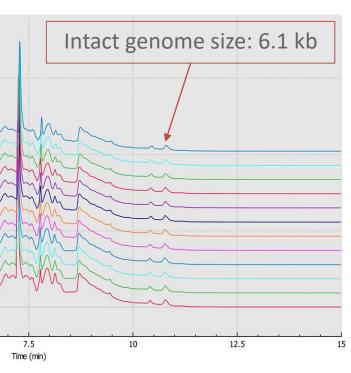
15

2.5

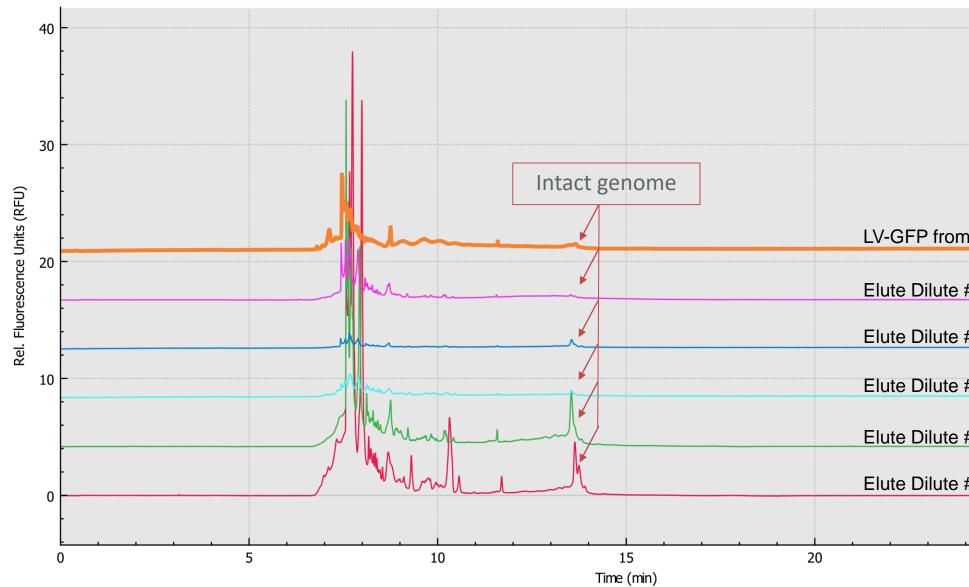
5



LVV-GFP

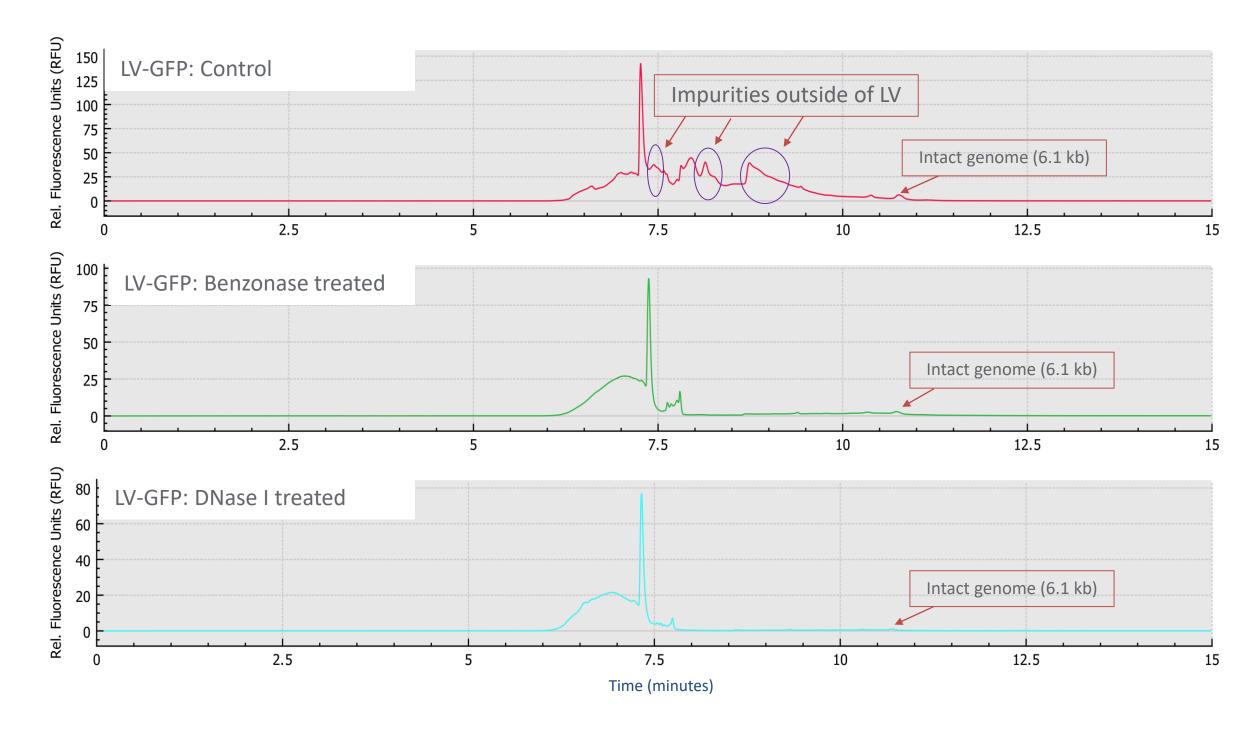


Genome integrity analysis of in-process samples



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Residual nucleic acids screening workflow





Conclusion

The CE platform technology demonstrates unique capabilities for quality attribute monitoring of lentiviral vectors

- Sensitive titer determination based on precise quantification of the separated p24 protein peak with 4 magnitude orders of detection linearity of $r^2 = 0.9968$ and LOQ of 8 ng/mL
- Easy to assess batch-to-batch reproducibility and protein purity based on the resulting proteome profile
- Good linear relationship of p24 and VSV-G indicates CE titer method is not measuring free p24.
- High sensitivity of CE-LIF enables genome integrity analysis of lentiviral vectors at a titer of 1 x 10⁹ TU/ml without amplification, providing high resolution of intact LV genome from various impurities with good repeatability

Acknowledgement

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