# Immuno-qPCR analysis and immunogenicity assessment of gene therapeutics and their targeted delivery molecules

EBF YSS

Fanni Suomi – Syrinx Bioanalytics, Finland





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# Who am I?

(Can someone please tell me?)



Master's studies

Molecular Biotechnology and diagnostics,

University of Turku



B.Sc.

Biochemistry

Sept. 2018

# Background

- Nucleic acid-based biotherapeutic products are gaining interest, but bioanalytical methods and guidelines are yet to be defined.
  - Preclinical and clinical safety evaluation methods need to adapt to the new needs of the industry.



# Background

- Anti-sense oligonucleotides (ASOs) are primarily administrated by parental injection.
  - After administration ASOs are tranferred into tissues predominantly by endocytotic uptake.
- It has been suggested that peptide conjugates can enhance cell penetration and improve the targeting of antisense agents.
  - Pharmacokinetic analysis
- Short oligonucleotides are not likely to elicit immune responses. However, polypeptide conjugates may have immunogenic properties.
  - Immunogenicity assessment



### Aims

Development of a pharmacokinetic analysis method for nucleic acid-based biopharmaceuticals and their carrier molecules.

< Immuno-qPCR

- Anti-drug antibody (ADA) assay for the assessment of immunogenicity.
  - ADA Bridging assay





#### **MATERIALS AND METHODS**





#### Test molecules

- Production of a fusion protein containing an antigen-binding fragment (Fab) and SpyCatcher in *E.Coli*.
- Sioconjugation of Fab-SpyCatcher with SpyTag-oligonucleotide





### ADA bridging assay







# qIPCR







# qIPCR











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#### RESULTS





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### ADA bridging assay - Capture and tracer optimization

#### Signal-to-noise ratios with different Bio-Fab-ON/Eu-Fab-ON amounts.

| Eu-Fab-ON (ng/reaction) | 70  | 100 | 125 | 150 | 175 | 200 |                  |
|-------------------------|-----|-----|-----|-----|-----|-----|------------------|
| 50                      | 4.3 | 5.2 | 5.8 | 6.6 | 7.2 | 7.1 | Eu <sup>3+</sup> |
| 100                     | 4.1 | 4.6 | 5.3 | 5.9 | 6.4 | 6.5 |                  |
| 150                     | 3.8 | 4.3 | 4.9 | 5.3 | 5.9 | 6.1 |                  |
| 200                     | 3.4 | 4   | 4.4 | 4.8 | 5.4 | 4.9 |                  |



#### ADA bridging assay - Sample incubation



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#### ADA bridging assay - Tracer incubation





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#### ADA bridging assay



Polyclonal antibody was tested as positive control due to low signals with monoclonal antibody



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# Cut point analysis

- Validated Excel spreadsheet based on Devanarayan V, Smith WC, Brunelle RL, Seger ME, Krug K, Bowsher RR. Recommendations for Systematic Statistical Computation of Immunogenicity Cut Points. AAPS J. 2017 Sep;19(5):14871498
  - ✓ 50 individual rat serum samples and 10 samples from a rat serum pool were measured.
  - Data was normally distributed.
  - Cut point 1.16 (mean + 1.645 x SD) at 5 % false positive rate.
  - ✓ Sensitivity 241 ng/mL (Cut point + 3 x SD).



#### qIPCR - bead and capture antibody optimization

#### Cq cycles of 31.3 µg/mL samples with different Dynabead/pAb ratios

|           | Bio-Anti-Human-IgG pAb |        |        |        |        |        |  |
|-----------|------------------------|--------|--------|--------|--------|--------|--|
| Dynabeads | 75 ng                  | 100 ng | 125 ng | 150 ng | 175 ng | 200 ng |  |
| 2.5 μg    | 12.2                   | 12.9   | 11.95  | 11.45  | 13.95  | 10.5   | Contraction of the second seco |
| 5 µg      | 11.8                   | 11.7   | 12.5   | 11.6   | 11.5   | 11.1   |  |
| 7.5 μg    | 12.1                   | 11.25  | 10.8   | 10.2   | 11.9   | 11.9   |  |
| 10 µg     | 11.0                   | 11.6   | 12.0   | 11.4   | 11.0   | 11.4   |  |



#### qIPCR – bead dissolving volume

# %CV's of samples with the same amount of beads dissolved in 20 $\mu L$ vs. 50 $\mu L$ per reaction

|                             | Concentration (ng/mL) |             |             |             |  |  |
|-----------------------------|-----------------------|-------------|-------------|-------------|--|--|
| Bead dissolving volume (µL) | 1467                  | 146.7       | 14.67       | 1.467       |  |  |
| 50                          | <u>37.3</u>           | 1.3         | <u>95.8</u> | <u>22.1</u> |  |  |
| 20                          | 10.7                  | <u>27.7</u> | 2.0         | 16.9        |  |  |
| <u>Underlined:</u> >20 %    |                       |             |             |             |  |  |





### qIPCR – primer length



#### Cycles of quantification with different primer sets: Overhanging primers and Overlapping primers

|                     | Standard concentration (ng/mL) |        |       |      |      |      |  |
|---------------------|--------------------------------|--------|-------|------|------|------|--|
|                     | 147 000                        | 14 700 | 1 470 | 147  | 14.7 | 0    |  |
| Overhanging primers | 10.6                           | 13.8   | 16.2  | 20.9 | 24.6 | 33.1 |  |
| Overlapping primers | 14.8                           | 17.0   | 19.2  | 24.9 | 30.3 | 31.0 |  |





qIPCR – final results







#### CONCLUSIONS





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# ADA bridging assay

- SpyCatcher fusion protein might block mAb from binding the Fab-Oligonucleotide molecule.
- Quasi-quantitative assay.
  - In practice anti-drug antibodies vary between individuals.
  - Results above cut point are considered positive.
- Assay meets the regulations for preclinical testing.
  - Sensitivity 241 ng/mL at 5 % false positive rate.



# qIPCR

- Implementation of qIPCR assay was successful.
- qIPCR offers a super sensitive method for oligonucleotide-protein conjugate detection.
  - Less than 2 ng/mL of test molecule was detectable
- Problems with deviation.
  - Immunoassay phase-derived.
  - Beads may increase deviation.
- Changing the magnetic beads to a different solid phase might solve the deviation problem.



### Future aspects

- Autumn: oligonucleotide extraction method development.
- Future: bioanalysis of oligonucleotide-based biotherapeutic products.



### Acknowledgements

- Supervisor Nina Sirkka, Syrinx Bioanalytics
- Scientific Director Timo Piironen, Syrinx Bioanalytics
- Pasi Virta and Antti Äärelä, Department of Chemistry, University of Turku
- Tuomas Huovinen and Anastasiia Kushnarova-Vakal, Biotech. unit, Department of Biochemistry, University of Turku



For futher questions, don't hesitate to contact me (e.g. via LinkedIn)!

#### **THANK YOU!**



