



Enhancing Large Molecule Design by Early Integration of MAPPs using Defined Allele Antigen Presenting Cells

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Executive Summary

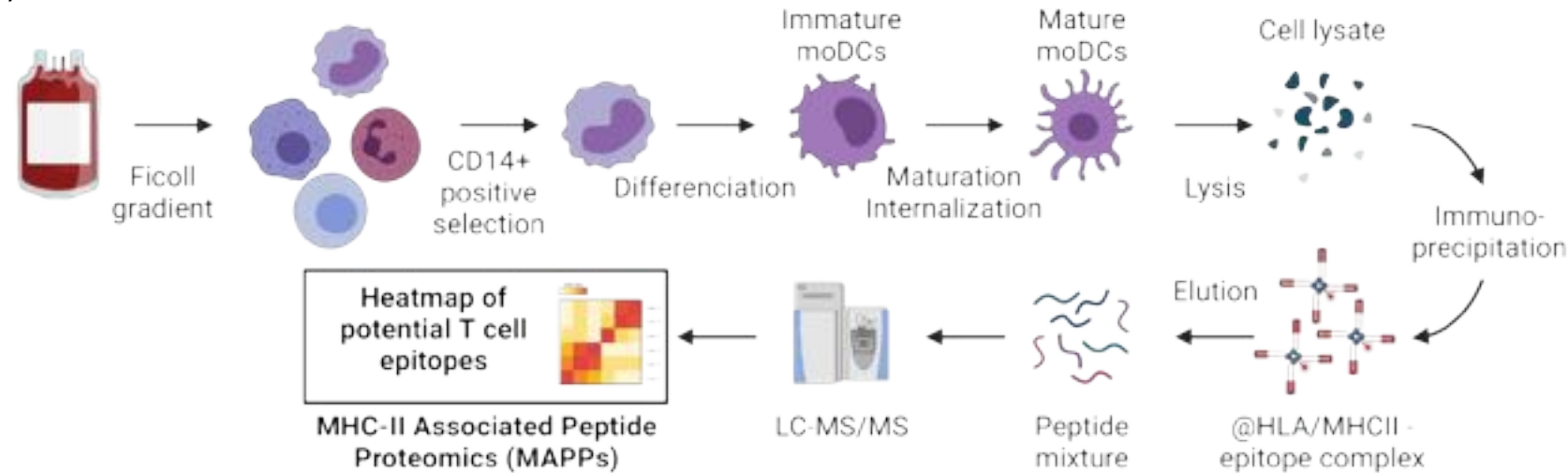
- The current MAPPs methodology faces challenges
- One factor is the biological variability for MHC haplotypes of the donor material
- Another the labor-intensive process of dendritic cell generation
- Overcoming these hurdles would allow for the earlier and more routine use of MAPPs e.g. in the space of improving large molecule design
- Here we present the development and a first PoC for a mono allelic (artificial) antigen presenting cell line



Introduction

Why Artificial Antigen Presenting Cells?

- APCs for MAPPs are derived from PBMCs of healthy donors
- DC generation adds (labor intensive / time consuming) working steps to MAPPs protocol
- Input material is expensive and needs continuous replenishment (cyroPBMCs) or supply (Buffy Coat)
- HLA-II set is not fully controllable

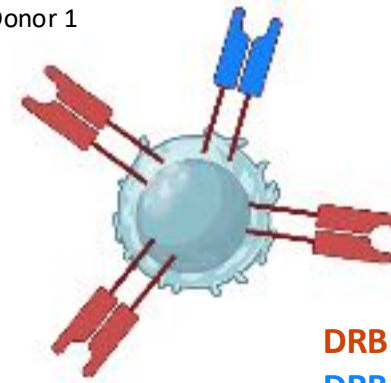


Introduction

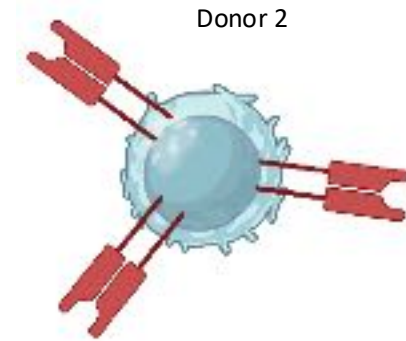
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Donor 1



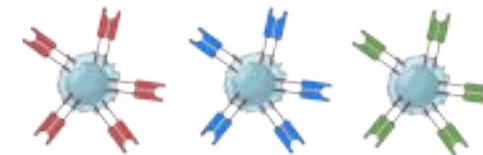
Donor 2



DRB1*15:01
DRB1*04:01
DRB1*07:01

Donors

DRB1*15:01
DRB1*04:01
DRB1*07:01



aAPCs

DRB1*15:01



DRB1*04:01



DRB1*07:01



superposition



Introduction

Why Artificial Antigen Presenting Cells?

- aAPCs allow for
 - unlimited amount of cells with regular cell culture expansion without the need for differentiation
 - defined HLA-II alleles

Virtual Population



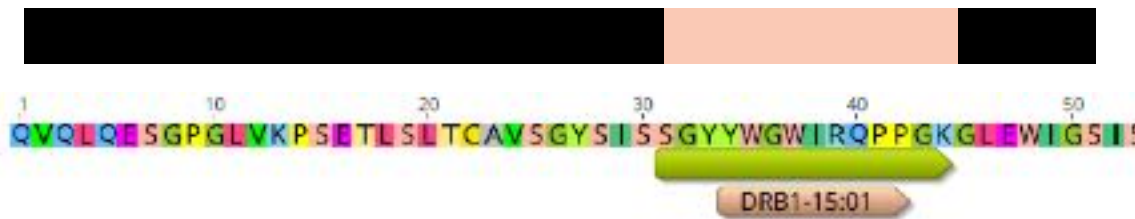
Table 4. HLA-DRB1 allele frequencies in worldwide pop

| DRB1 alleles | Northern Europe | | Western Europe | | |
|--------------|-------------------|-------------------------------|------------------------------|--------------------------------|------------------------------|
| | Denmark 2n=100 | Ireland (North) 2n=2000 | Portugal (North) 2n=92 | Portugal (Centre) 2n=300 | Portugal (South) 2n=98 |
| | *01 | 0.139 | 0.120 | 0.163 | 0.040 |
| *02 | 0.100 | 0.050 | 0.131 | 0.160 | 0.141 |
| *04 | 0.203 | 0.088 | 0.098 | 0.140 | 0.131 |
| *07 | 0.108 | 0.060 | 0.171 | 0.140 | 0.153 |
| *08 | 0.028 | 0.021 | 0.033 | 0.070 | 0.030 |
| *09 | - | 0.007 | 0.011 | 0.020 | 0.010 |
| *10 | 0.009 | 0.007 | - | - | 0.030 |
| *11 | 0.018 | 0.041 | 0.008 | 0.110 | 0.132 |
| *12 | 0.039 | 0.011 | 0.011 | - | 0.031 |
| *13 | 0.139 | 0.081 | 0.130 | 0.160 | 0.171 |
| *14 | 0.009 | 0.015 | 0.033 | 0.020 | 0.010 |
| *15 | 0.185 | 0.185 | 0.109 | 0.110 | 0.020 |

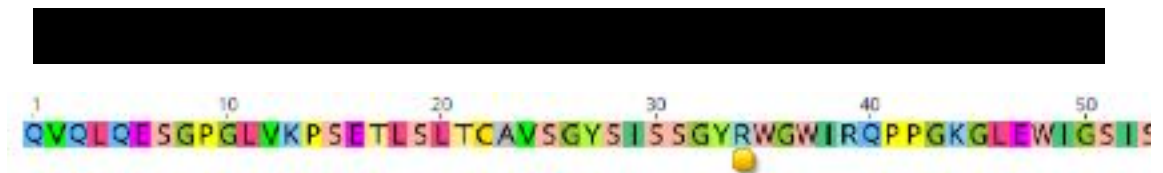
DG Brum et al. 2007

Lead Optimization

aAPCs
DRB1*15:01

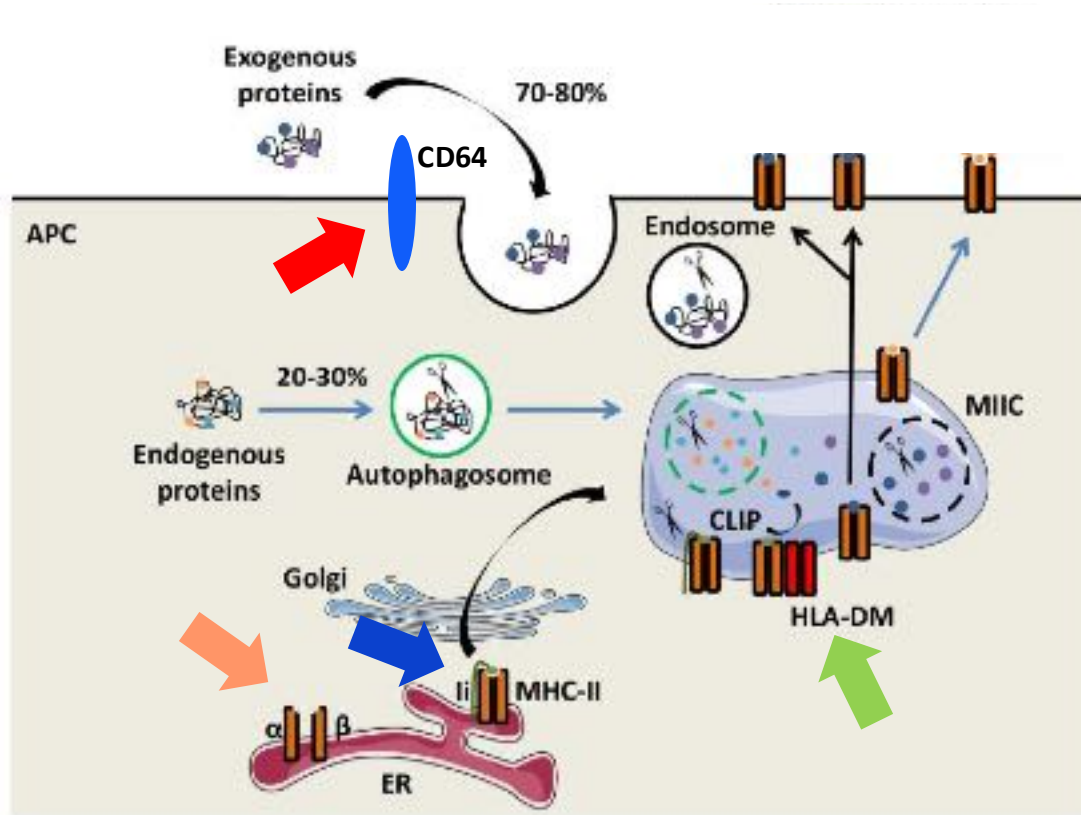


aAPCs
DRB1*15-01



Methods

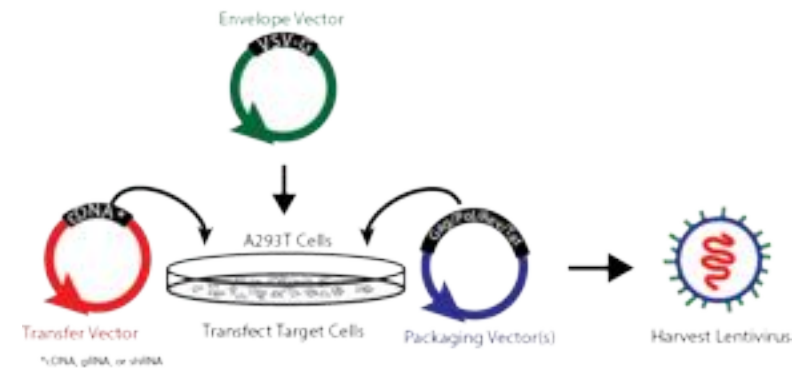
Components for aAPCs



Alexandre Couture et al. 2019

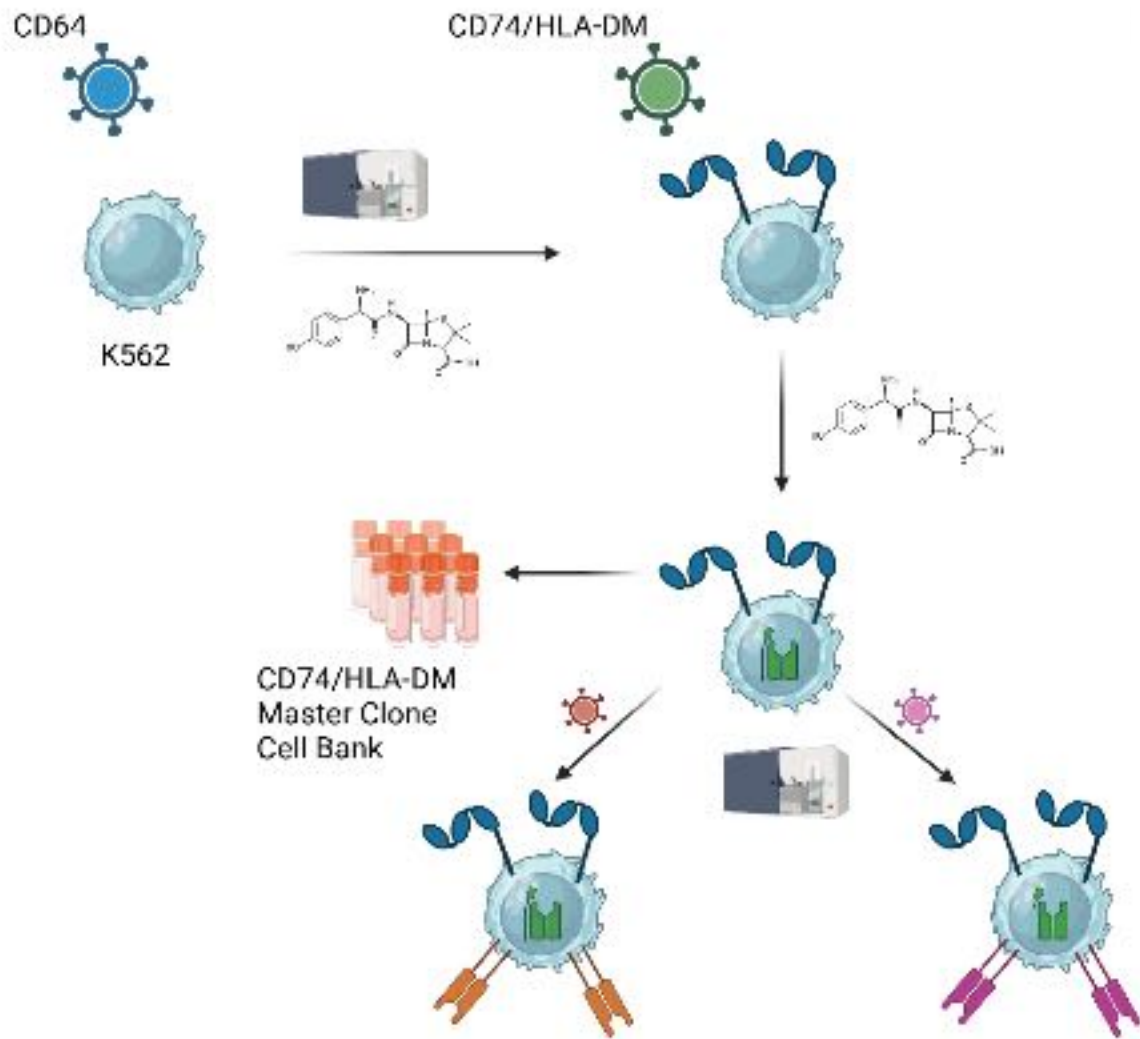
K562: HLA deficient B cell lymphoblast

- ➔ CD64: high affinity FcγRI for improved uptake
- ➔ CD74: li / CLIP for HLA stability
- ➔ HLA-DM: HLA loading
- ➔ HLA-D(R/P/Q): antigen presentation



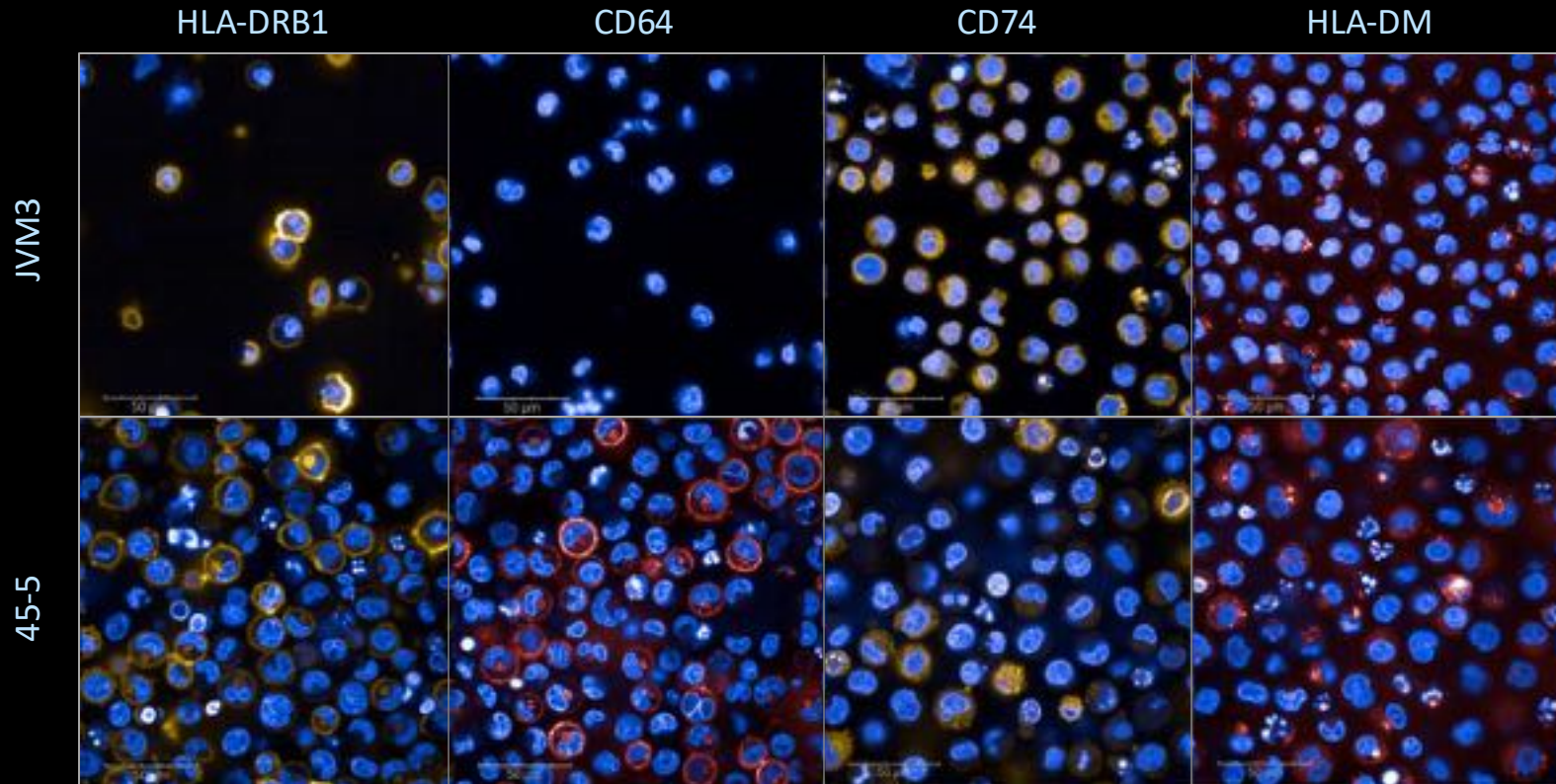
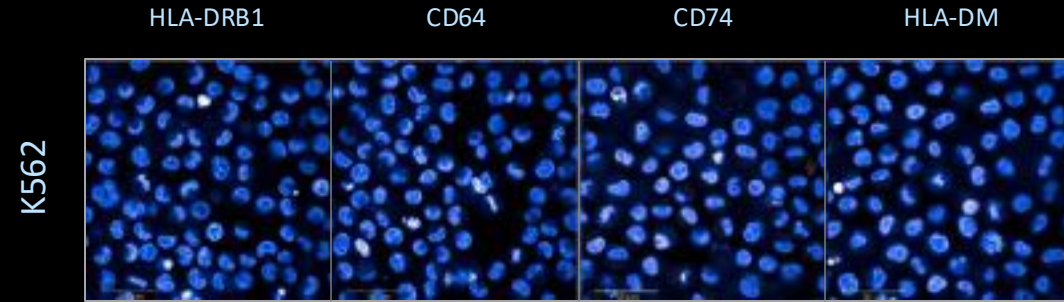
Methods

Transduction Strategy



Results

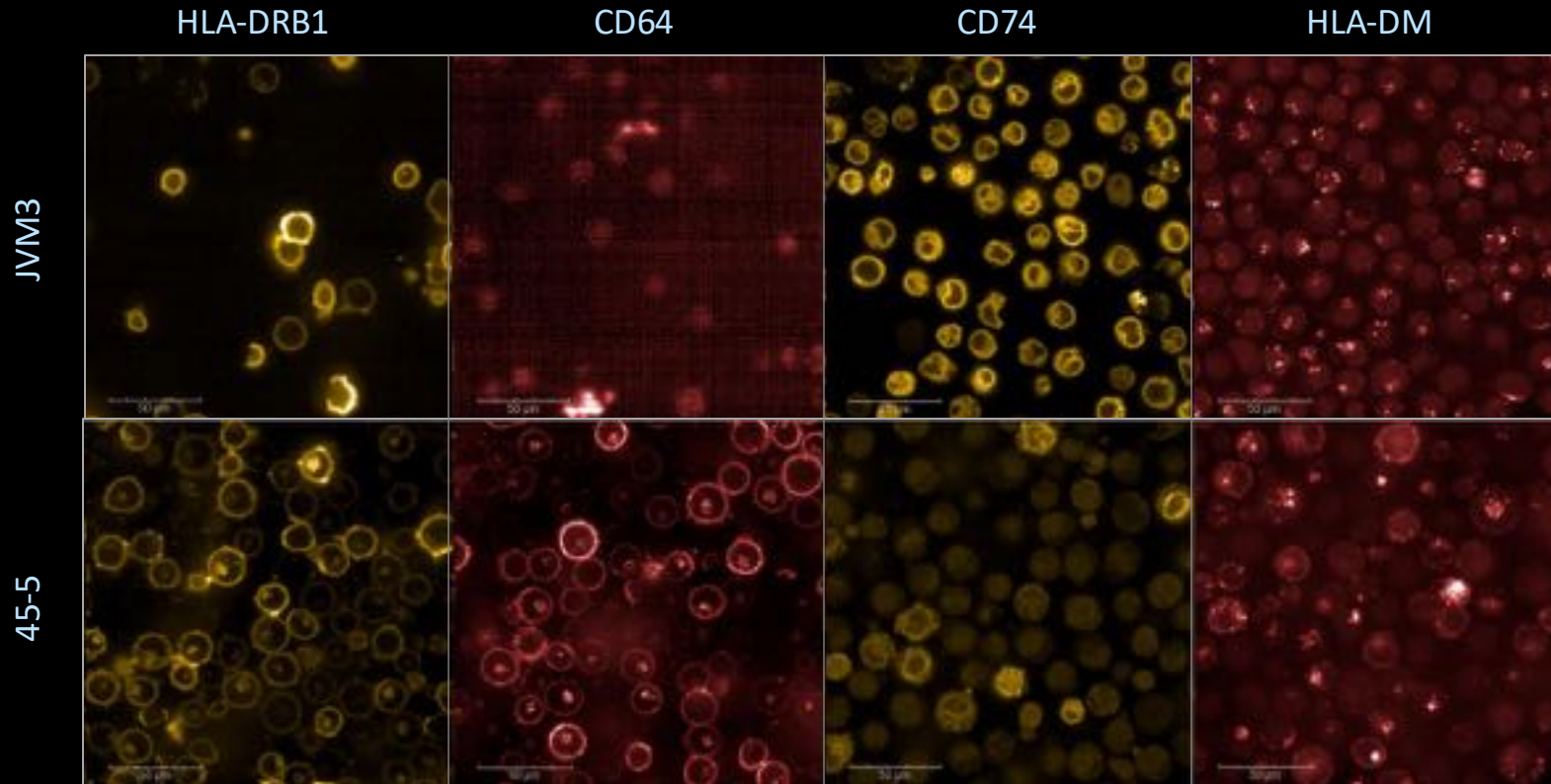
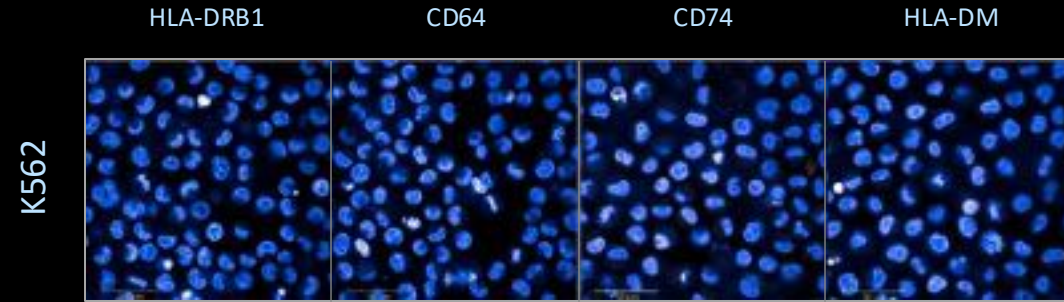
Expression Data - Imaging



- All four constructs are expressed
- Expression pattern / compartmentation comparable with JVM3
- Lower expression level for CD74 in aAPC

Results

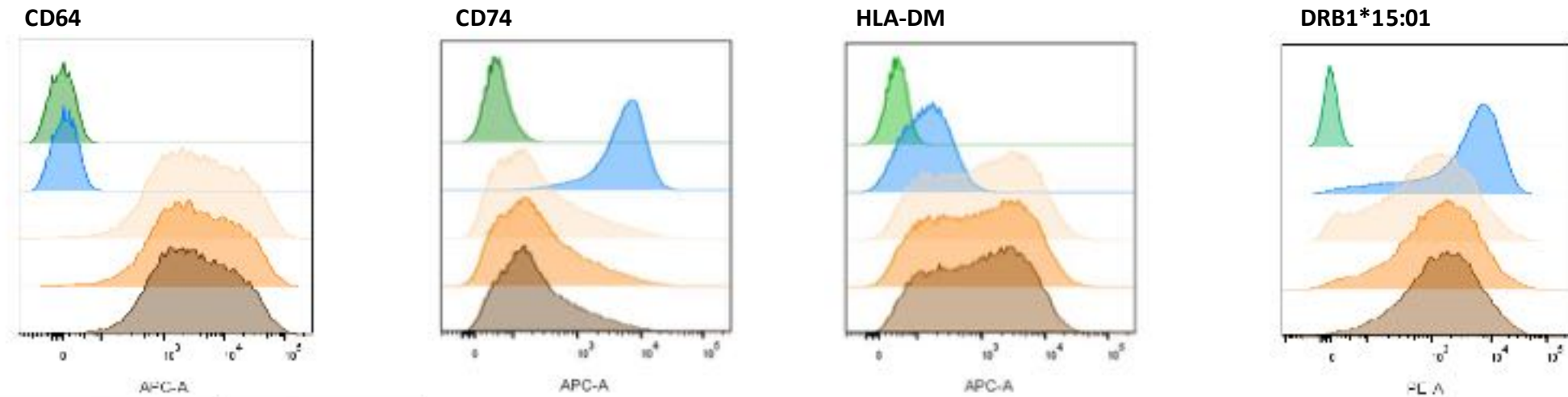
Expression Data - Imaging



- All four constructs are expressed
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Results

Expression Data – Flow Cytometry

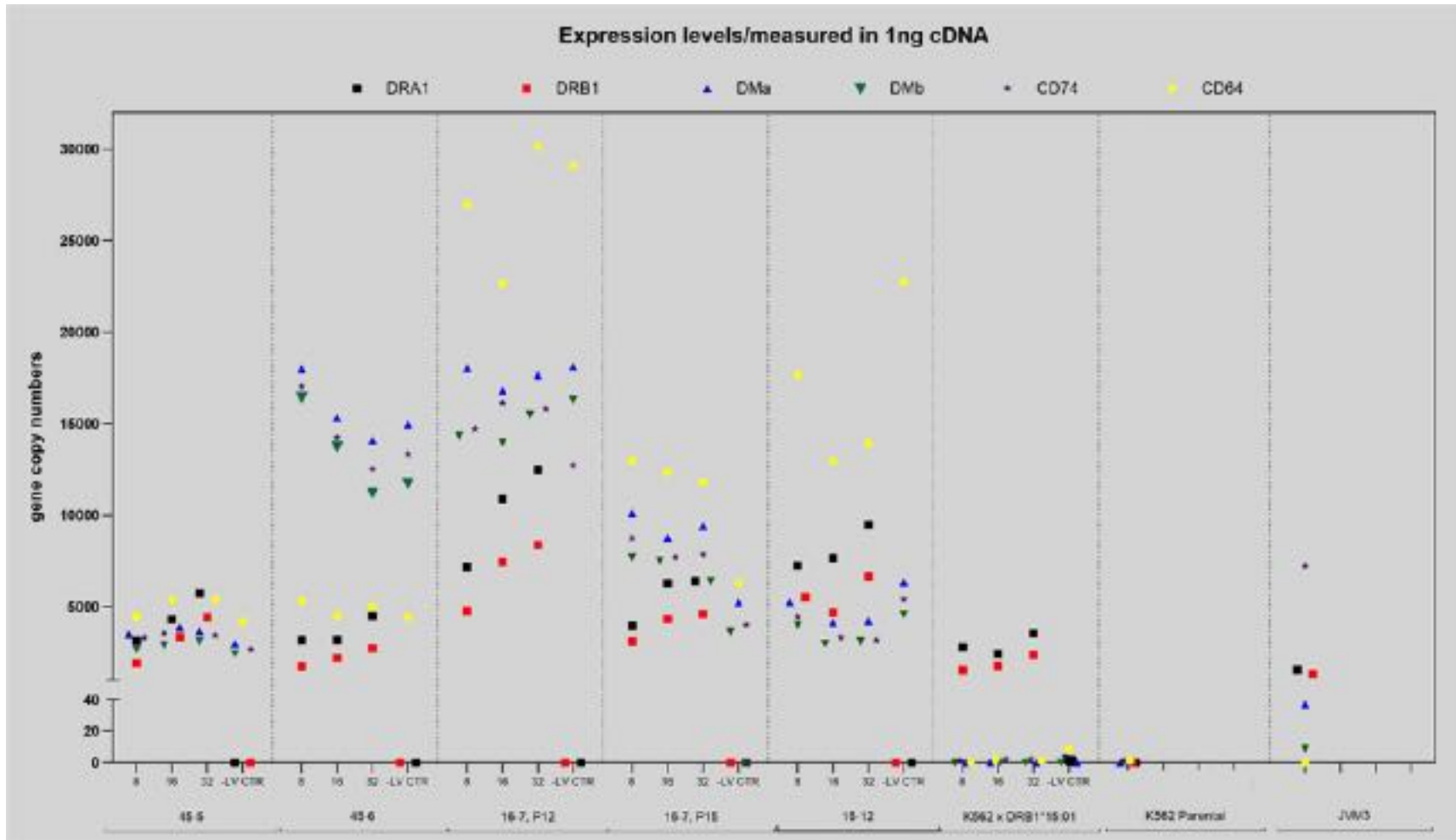


| Sample Name |
|-------------------|
| K562F_stained.fcs |
| JYMD_stained.fcs |
| MO10_stained.fcs |
| MO16_stained.fcs |
| MO82_stained.fcs |

- Expression data consistent with microscopy
- Lower expression level for CD74
- Higher expression of HLA-DM
- Comparable expression of DRB1

Results

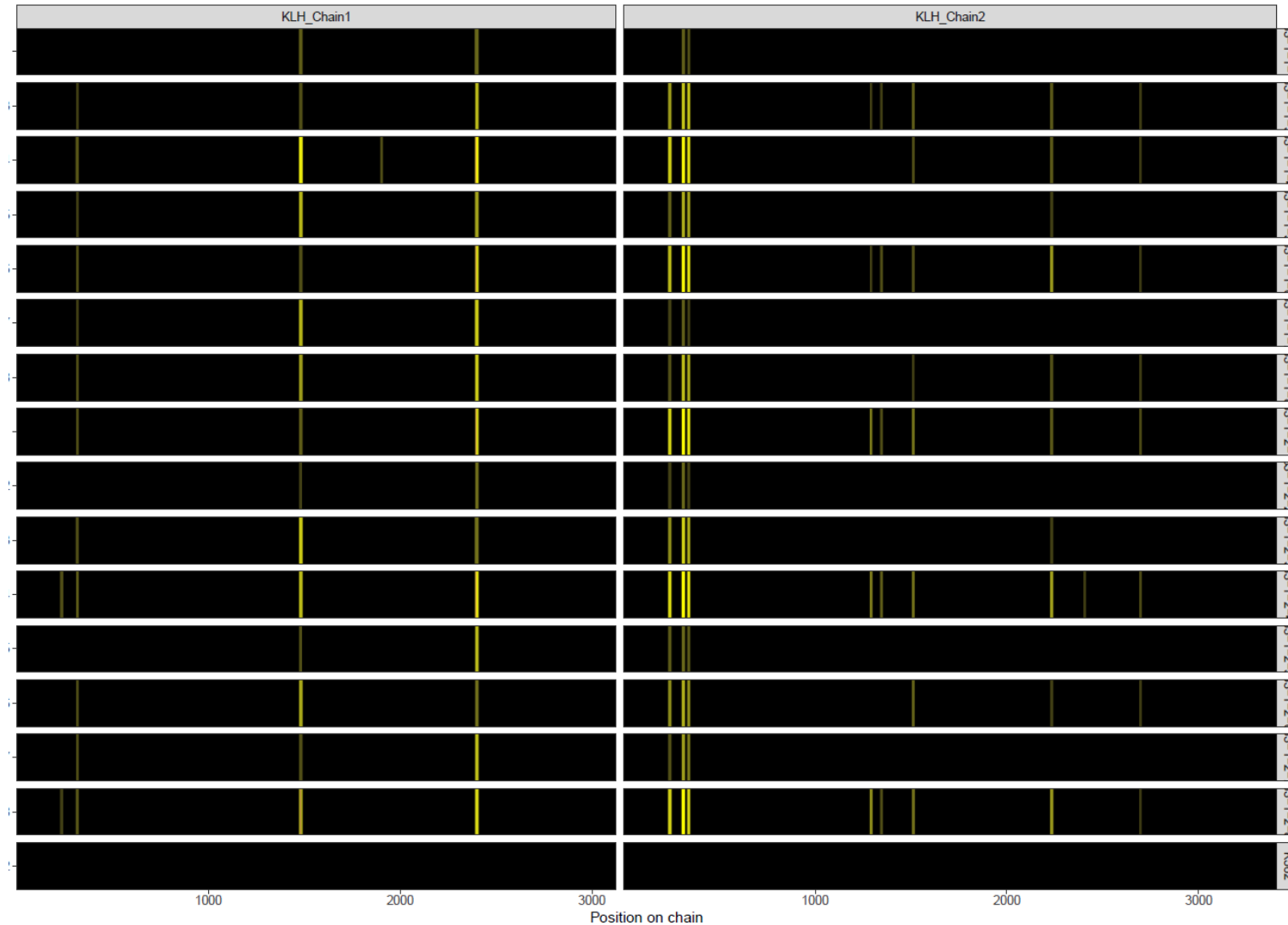
Expression Data – Genetic Copy Numbers



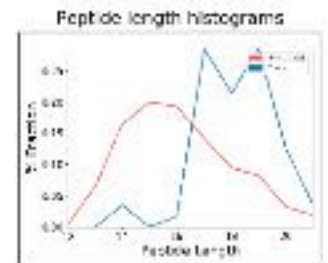
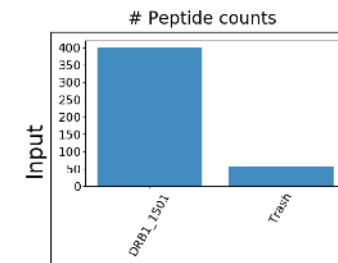
- Equal copy numbers for «paired» hetero dimeric components
- Copy numbers do not entirely reflect the observed expression levels

Results

MAPPs on PoC subClones 33 - KLH



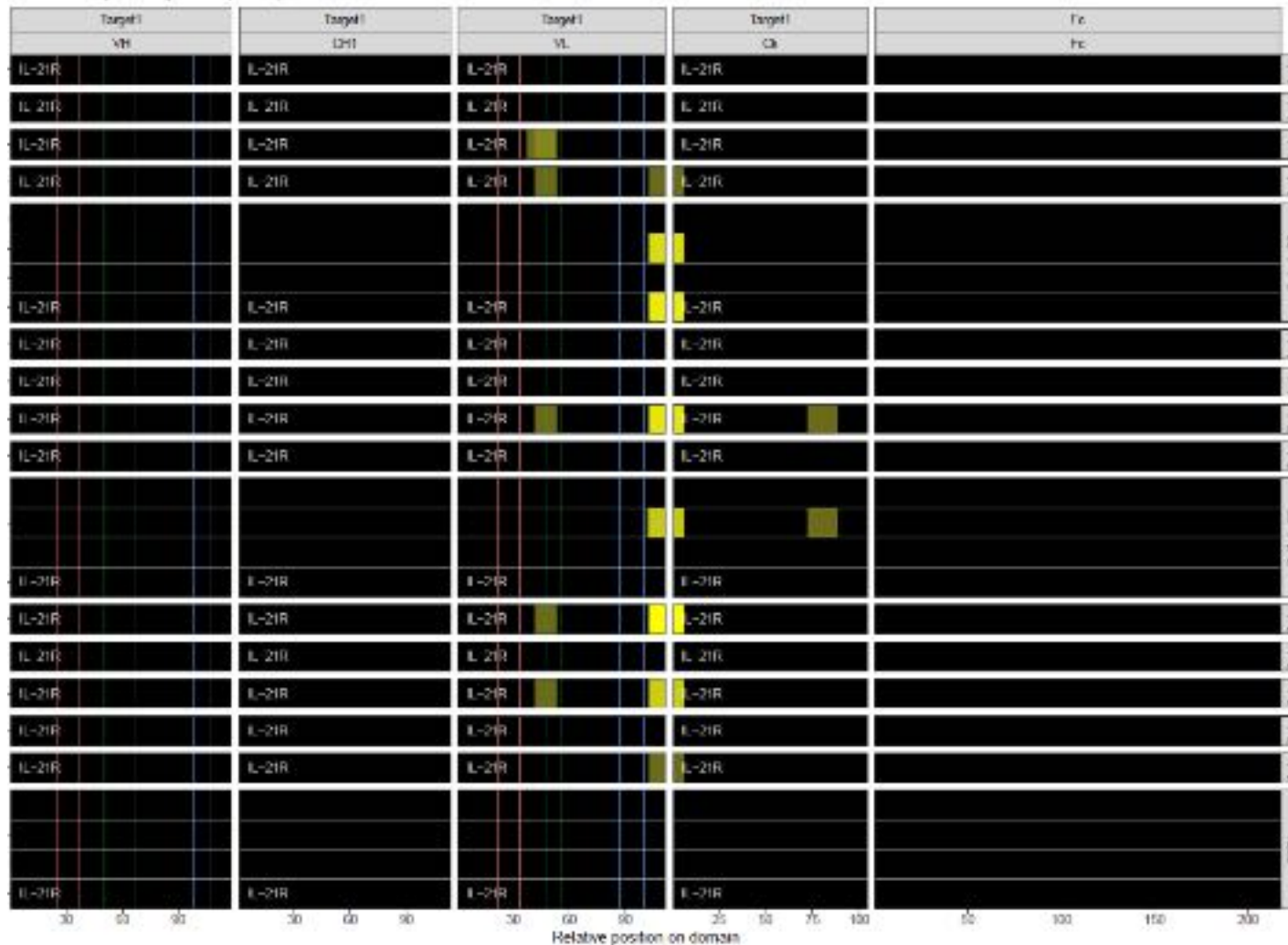
- Consistant pattern over all subclones
- No clusters detected for parental cell line K562
- Majority of peptides maps well on DRB1*15:01



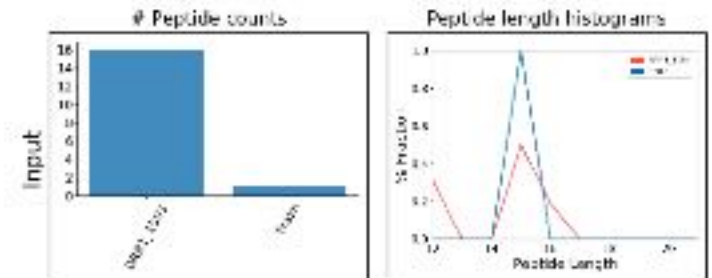
MHCMotifDecon - 1.2

Results

MAPPs on PoC subClones 33 – ATR-107



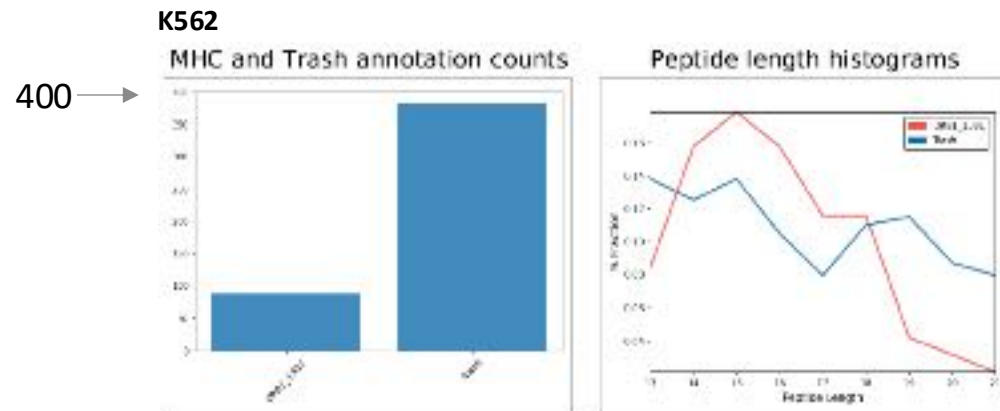
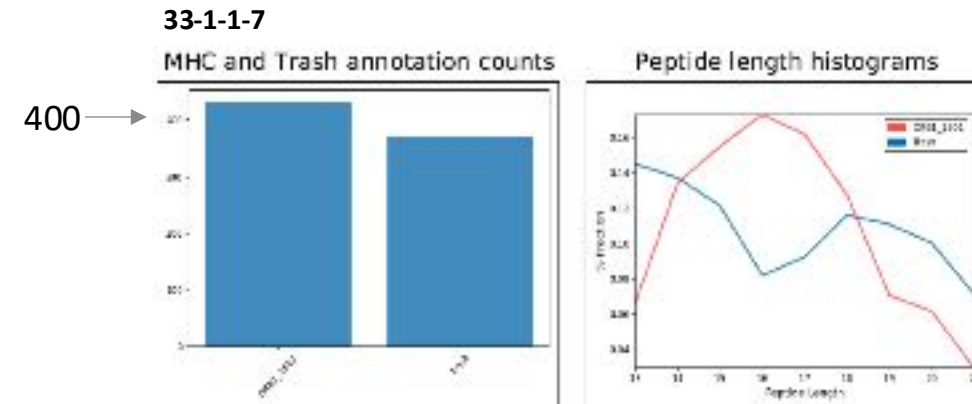
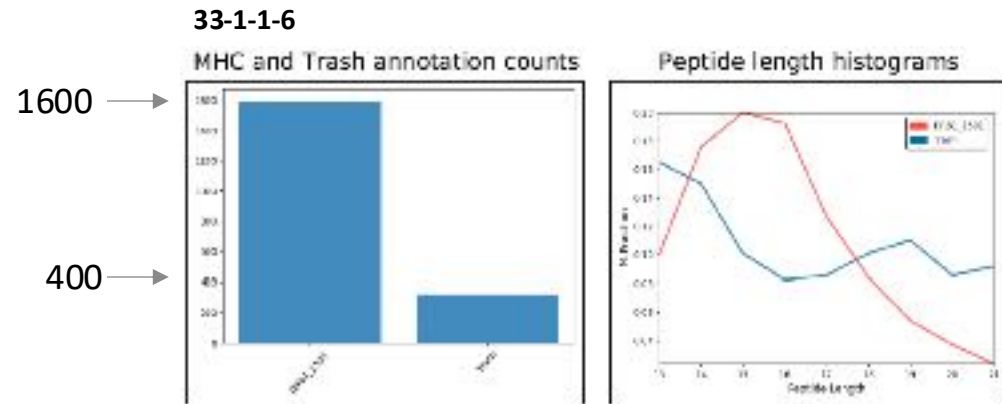
- Consistant pattern over all subclones (if clusters detected)
- No clusters detected for parental cell line K562
- Majority of peptides maps well on DRB1*15:01



MHCMotifDecon - 1.2

Results

MAPPs Fidelity on Endogenous Proteins



- Wide fidelity spread for endogenous proteins for different (sub)-clones
- Low fidelity (sub)-clones tend to present overall less
- Lowest/Zero fidelity for K562 cell line

Final Thoughts

- We established a first PoC for a mono allelic (artificial) antigen presenting cell line for DRB1*15:01
- Feasibility for more HLAs including DP and DQ will be assessed
- Generation and testing of oligo allelic panels and cell mixtures will be tested
- Integration of the system for MAPPs supported lead optimization and other suitable applications to be assessed

Acknowledgments

Merci Viel Mohl!

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Hana Glier

Thelma Lopes

MS – 360 Labs

Shahrzad Talavaei

CM120

Lilly von Muenchow

DTU

Morten Nielsen

Doing now what patients need next