# The key to Immunologic compatibility in the highly sensitized -The combined role of understanding Anti HLA antibody and HLA allelic prevalences.



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### **BACKGROUND**

- \* Finding suitable donors for highly sensitized transplant patients is challenging, even with advanced HLA typing and anti-HLA antibody detection methods.
- \* Highly sensitized patients develop antibodies against certain HLA antigens, which limits their donor options.
- ❖ While current tests can precisely identify these antibodies and HLA types, a clearer understanding of which HLA antibodies and antigens are most common could improve compatibility predictions.
- ❖ By recognizing these patterns, we may unlock ways to expand donor availability, allowing for more successful transplants for those who are highly sensitized.



#### AIM & OBJECTIVE

To evaluate the frequency of allele-specific anti-HLA antibodies and the prevalence of the corresponding HLA alleles to assess the clinical probability of finding an antigen negative donor.

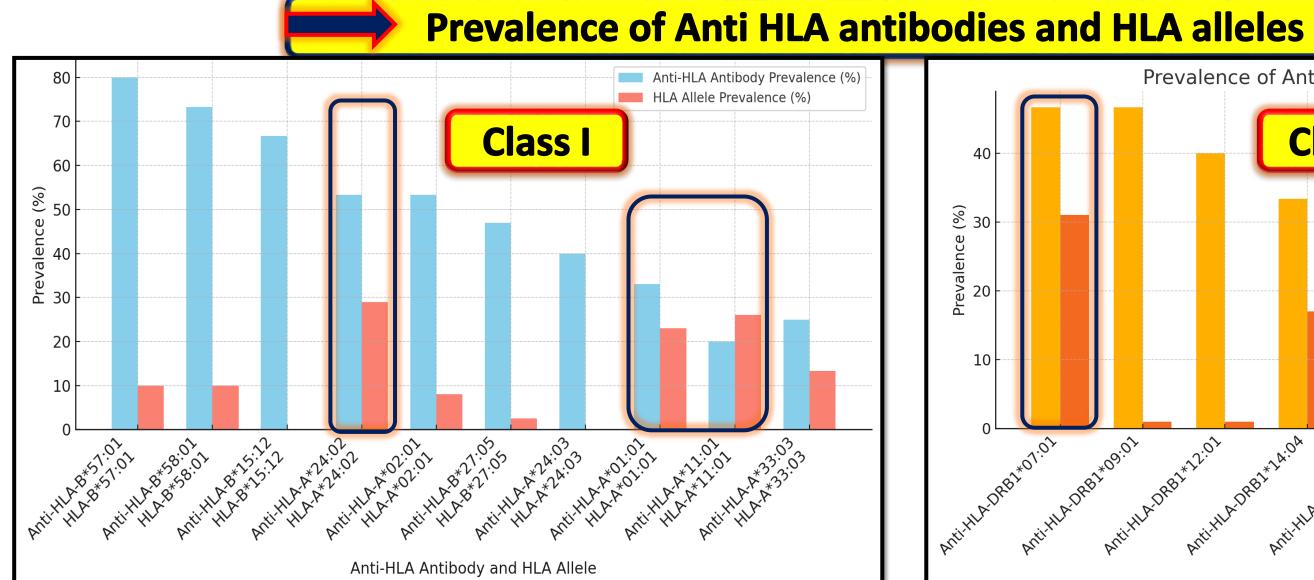


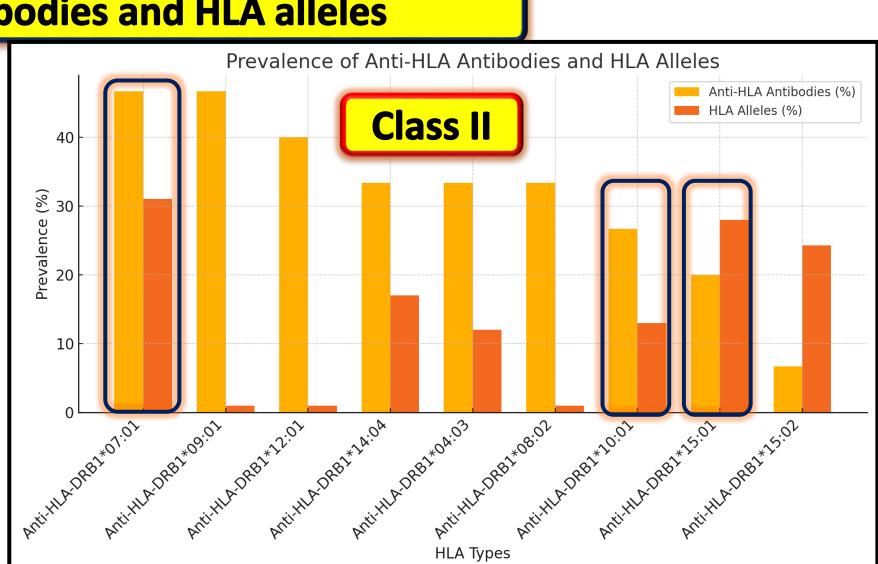
## **MATERIALS & METHODS**

- **☐** Study participants and recruitment:
- This retrospective study analysed the **frequency of antibody specificities** present in **15 highly sensitized** patients.
- ❖ A total of 3050 samples of High resolution HLA typing were analyzed to determine HLA allele frequency.
- ☐ Anti HLA Antibody detecting platform:
- \* Antibody testing was performed using the Luminex Single Bead Antigen Assay (SAB), with an MFI (Mean Fluorescence Intensity) > 1500 considered positive.
- ☐ HLA typing platform:
- \* High-resolution HLA typing was done with MIA FORA kits on the Illumina MiniSeq platform.
- ❖ HLA-A\*, HLA-B\*, HLA-C\*, HLA-DRB1\*, HLA-DQB1\* and HLA-DPB1\* genes were analyzed.
- ☐ Statistical tools —
- ❖ An in-house built software (Database-Driven HLA Matching Algorithm with Transaction-Safe CRUD Operations) designed by institutional IT team was used to determine the HLA allele frequency and the results were collated.



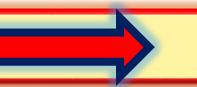
#### RESULTS





# **DISCUSSION & CONCLUSION**

- ❖ Our results highlight the various combinations of antibody frequency and allelic prevalence In Class I, HLA-A\*24:02 which shows high prevalence (29%) and , suggesting strong immune engagement.
- \* Conversely, HLA-B\*15:12, despite its high frequency (66.67%), has limited clinical relevance due to its very low prevalence (0.29%).
- ❖ In Class II, HLA-DRB1\*07:01 plays a significant role in immune recognition with both high frequency (46.67%) and prevalence (31%). On the other hand, HLA-DRB1\*09:01 and HLA-DRB1\*12:01, despite their high frequencies, have low prevalence (1%), indicating minimal immune significance.
- Understanding these various combinations in our population is essential for predicting the chances of finding a suitable donor in highly sensitized individuals.



# REFERENCES

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