

Criteria for Grading

1. B-cell line Exchange - class I/II molecular typing

- a. Consensus is based on a minimum of 10 labs reporting and ≥ 90% agreement among labs for HLA-A, B, C, DRB1, DQA1, DQB1, and DPA1 loci. For DRB3/4/5 and DPB1 loci, consensus is based on 80% agreement.
- b. Results falling within the consensus are graded as satisfactory.
- c. For loci presumed to be homozygous, the second field will be graded as satisfactory if either a blank is reported (A*02, -) or the antigen/allele is reported twice (A*02, A*02).

2. **Serum Exchange** – class I/II antibody identification

- a. Consensus is based on a minimum of 10 labs reporting and ≥ 90% agreement among labs.
- b. Assignment of ≥ 80% of antibody specificities falling within consensus will be graded as satisfactory.

3. KIR Exchange – KIR genotyping

- a. Consensus is based on a minimum of 10 labs reporting and ≥ 90% of labs reporting the positive/negative assignments for individual KIR loci.
- b. Results falling within the consensus are graded as satisfactory.
- c. Loci reported as not tested (NT) will not be included in the grading analysis.

4. **MICA Exchange** – MICA genotyping

- a. Consensus is based on a minimum of 10 labs reporting and ≥ 80% agreement among labs.
- b. Results falling within the consensus are graded as satisfactory.
- c. For loci presumed to be homozygous, the second field will be graded as satisfactory if either a blank is reported (MICA*002, -) or the allele is reported twice (MICA*002, MICA*002).

5. Single antigen, Flow and Virtual Crossmatch Exchange

- a. **Flow Crossmatch** T/B-cell Flow cytometric crossmatch
 - I. Consensus is based on a minimum of 10 labs reporting and 80% agreement among labs reporting positive/negative assignments.
 - II. Results falling within the consensus are graded as satisfactory. Results reported as borderline are not graded.

b. **Serum Evaluation** - class I/II antibody identification

- I. Consensus is based on a minimum of 10 labs reporting and 90% agreement among labs.
- II. Assignment of ≥ 80% of antibody specificities with ≥ 90% consensus will be graded as satisfactory.

c. Paper Challenge (educational exercise)

- I. Virtual predictions virtual prediction of physical crossmatch
 - 1. Consensus is based on a minimum of 10 labs reporting and 80% agreement among labs reporting positive/negative assignments.
 - Results falling within the consensus are marked as satisfactory. Results reported as borderline are not graded.

II. DSA Assignment – identification of donor specific antibodies

- 1. Consensus based on a minimum of 10 labs reporting and 90% agreement among labs.
- 2. Assignment of all antibody specificities with ≥ 90% consensus is required to be marked as satisfactory.

