

## Flow Crossmatch Exchange Serum Results Upload Instructions

1. From the Serum Evaluation Report page, click on “[Upload Results File.](#)”

### UCLA Serum Evaluation Report

Exchange#:	31	Exchange Type:	CROSS MATCH
Center# :		Center Name:	

[Upload Result File](#)

Enter Results Serum #: 183 | 184 | 185 | 186

Class I  Class II

CLASS I 183 (green)

Locus	Positive Cutoff Value	Antigen Class I MFI
A		<a href="#">Enter/Edit Info Class I</a>
B		
C		
Class I		
Manufacturer		
Kit Type		
Lot No		
Comment		
Sera Pretreatment		
<a href="#">Enter/Edit Info</a>		

2. Download the sample file provided on the website to ensure your file is formatted correctly.

### Upload Results File

Exchange:	31	Exchange Type:	CROSS MATCH
Center #:		Center Name:	
Status:	Center Received		

Please select below the file containing the exchange results. You can download a sample file by clicking [here](#) (please preserve the column's order, keep the column's headers, and remove the example record from the sample file).

Results File Name: \*

 

The file should be a .csv file - comma-separated values file.

- a. Please preserve the column's order, keep the column's headers, and remove the sample record from the sample file.
- b. Enter the sample number, class (I, II), locus, antigen, and MFI under the corresponding column.
  - i. For C-locus, please include the letter "W" with the antigen name. For example, enter "C" under the "Locus" column, and "W12" under the "Antigen" column.
  - ii. For DPA1 and DQA1, please include an asterisk at the end of the locus name and a zero before the antigen number. For example, enter DPA1\* or DQA1\* under the "Locus" column and '01 or '04 under the "Antigen" column.
- c. File must be saved as a .csv file (comma separated values file).
- d. Results for all 4 sera may be reported on a single file.

3. **Choose a file** to upload then click "**Process File.**"

4. The processed file will appear on the bottom of the webpage. Errors will be shown in **RED**. Results will not be uploaded until errors have been corrected.

Process File

Go Back

The following records were readed from the uploaded file. The items marked in red were not possible to record them in the system, please review and solve the bullet points under the **Record Errors** column and upload them again in another file. You can download the displayed table by clicking on the button **Download Process Output**

Download Process Output

#	Serum Number	Class	Locus	Antigen	MFI	Record Errors
1	183	I	B	58	1757	
2	183	I	B	82	1920	
3	183	I	A	68	2115	
4	183	I	B	56	2426	
5	183	I	C	W10	2480	
6	183	I	A	69	2654	
7	183	II	DR	11	2528	
8	183	II	DR	1	2839	
9	183	II	DR	13	3396	
10	183	II	DQ	6	26786	
11	184	II	DPA1*	01	2343	
12	185	II	DQA1*	04	3071	
13	185	II	DQ	4	5673	
14	185	II	DQ	5	2050	
15	185	II	DQ	6	1479	

5. The displayed table may be downloaded by clicking on “[Download Process Output.](#)” An example of the downloaded file is below:

Exchange Type: CROSS MATCH						
Exchange: 31						
Center: 51						
Download Date and Time: 10/15/2024 12:28:54						
#	Serum Nu	Class	Locus	Antigen	MFI	Record Errors
1	183	I	B	58	1757	
2	183	I	B	82	1920	
3	183	I	A	68	2115	
4	183	I	B	56	2426	
5	183	I	C	W10	2480	
6	183	I	A	69	2654	
7	183	II	DR	11	2528	
8	183	II	DR	1	2839	
9	183	II	DR	13	3396	
10	183	II	DQ	6	26786	
11	184	II	DPA1*	01	2343	
12	185	II	DQA1*	04	3071	
13	185	II	DQ	4	5673	
14	185	II	DQ	5	2050	
15	185	II	DQ	6	1479	
16	185	II	DQ	8	4493	
17	185	II	DQA1*	05	4398	
18	185	II	DQA1*	06	4755	
19	185	II	DR	103	2828	
20	186	I	B	7	10666	
21	186	II	DR	11	4001	

6. Click on “[Go Back](#)” to return to the results entry page. If there are no errors, data from the file will populate the corresponding fields for each sample.

Class I
Class II

**CLASS I 183 (green)**

Locus	Positive Cutoff Value	Antigen	Class I MFI
A		A69	2654
B		CW10	2480
C		B56	2426
Class I		A68	2115
Manufacturer		B82	1920
Kit Type		B58	1757
Lot No		<a href="#">Enter/Edit Info Class I</a>	
Comment			
Sera Pretreatment			
<a href="#">Enter/Edit Info</a>			