Minnesota Health

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HLA/Immunology Laboraotry

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ASHI: 08-4-MN-01-1

CLIA UMMC: 24D0688128 CLIA Outreach 24D0995276

TEST INFO

Sample Date: Amp Tech: Kit Lot #: **K3471-CR**Sample ID: **1804214** Analysis Tech: Test Date: **1/7/2019**

Patient ID: AFK3449 Review Tech:

SUMMARY

This session includes tests for A, B, C, DR, DQA, DQB, DPA and DPB

TEST	GENOTYPE	PHENOTYPE
HLA-A	A*02 A*03	A2 A3
HLA-B	B*14 B*57	B65 B57
HLA-DRB1	DRB1*07 DRB1*13	DR7 DR13
HLA-DQA1	DQA1*01 DQA1*02	DQA1*01 DQA1*02
HLA-DPA1	DPA1*01 DPA1*01	DPA1*01 DPA1*01

TEST	GENOTYPE	PHENOTYPE
HLA-C	C*06 C*08	Cw6 Cw8
Bw		Bw4 Bw6
HLA-DRB345	DRB3*03 DRB4*01:03:01:02N	DR52 DR53-Null
HLA-DQB1	DQB1*03 DQB1*06	DQ9 DQ6

EPITYPE	ALLELES	HVR-A	HVR-B	HVR-C	HVR-D	HVR-E	HVR-F
EDP04	DPB1* 04:01:01 (+109 rare alleles)	LFQG	EEFA	AAE	ILEEK	М	GGPM
EDP72	(1 rare alleles)	LFQG	EEFA	AAE	LLEEK	М	GGPM

One side of this epitype requires a RARE allele, review results to confirm assignment.

This epitype was MANUALLY chosen.

The HLA-DPB1 test result requires a RARE allele assignment.

(Notice: Cannot exclude other combinations. Review all results.)
All test results are based on IMGT/HLA release: version 3.25.0 - July 2016

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LABORATORY ASSIGNMENT

HLA-A: **A2 A3** HLA-B: **B65 B57** Bw: **Bw4 Bw6**

HLA-C: Cw6 Cw8 HLA-DRB1: DR7 DR13 HLA-DRB345: DR52 DR53-Null HLA-DQA1: DQA1*01 DQA1*02 HLA-DQB1: DQ9 DQ6 HLA-DPA1: DPA1*01 DPA1*01 HLA-DPB1: DPB1*04:01

DPB1*72:01

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HLA-A

GROUP	ALLELES	ANTIGE
A*02	A* 02:01:01:01, 02:01:04, 02:02:01:01, 02:04-02:05:01, 02:06:01:01, 02:07:01, 02:08-02:09, 02:11:01, 02:14, 02:17:01, 02:20:01, 02:22:01, 02:24:01; A* 02:12d-02:13d, 02:16d, 02:18d, 02:21d, 02:25d, 02:29d, 02:34d, 02:49d, 02:74:01d; A* 02:01:01:02Le-02:01:03e, 02:01:05e-02:01:13e, 02:01:15e-02:01:35e, 02:01:37e-02:01:122e, 02:02:01:02e-02:03e, 02:05:02e-02:05:06e, 02:06:01:02e-02:06:22e, 02:07:02e-02:07:08e, 02:11:02e-02:11:09e, 02:17:02e-02:17:03e, 02:20:02e, 02:22:02e, 02:24:02e, 02:31e, 02:35:02e, 02:41e-02:42e, 02:59e, 02:66e-02:68e, 02:70e-02:72e, 02:74:02e, 02:77e, 02:79:01e, 02:89e-02:90e, 02:97e, 02:105e	A2
	A* 02:03:05e	A203
	A* 02:15Ne, 02:32Ne, 02:43Ne, 02:82Ne-02:83Ne, 02:88Ne, 02:94Ne, 02:113Ne, 02:125Ne	Null
	A* 02:30:01, 02:60:01; A* 02:27d, 02:33d, 02:36d, 02:44d-02:45d, 02:58d, 02:63d-02:64:01d, 02:84d-02:86d, 02:96d, 02:119d, 02:123d, 02:131d, 02:137d; A* 02:01:14Qe, 02:26e, 02:28e, 02:30:02e, 02:37e, 02:40e, 02:47e, 02:51e, 02:54e, 02:56e-02:57e, 02:60:02e-02:62e, 02:69e, 02:75e-02:76e, 02:80e, 02:91e-02:92e, 02:99e-02:104e, 02:106e-02:109e, 02:111e, 02:115e-02:116e, 02:118e, 02:120e-02:121e, 02:126e-02:128e, 02:130e, 02:132e-02:134e, 02:138e-02:142e, 02:144e-02:147e, 02:149e-02:151e, 02:155e, 02:157e-02:170e, 02:172e-02:229e, 02:231e-02:241e, 02:243e, 02:246e-02:252e, 02:254e-02:257e, 02:259e-02:263e, 02:265e-02:278e, 02:283e-02:299e, 02:301Ne-02:303e, 02:305Ne-02:308e, 02:310e-02:314Ne, 02:316e-02:333e, 02:335e-02:337e, 02:340e-02:344e, 02:346e-02:354e, 02:356Ne-02:369e, 02:371e-02:399e-02:408e, 02:410e-02:411e, 02:413e-02:426e, 02:48e-02:430e, 02:432e-02:446e, 02:448e-02:452e, 02:455e-02:462e, 02:464e-02:465e, 02:467e-02:479e, 02:481e-02:504e, 02:506Ne-02:524:01e, 02:525Ne-02:526e, 02:528e, 02:530e-02:543e, 02:583e-02:556e, 02:558e-02:567e, 02:569e-02:570e, 02:572e-02:579e, 02:581e, 02:583e-02:593e, 02:632e	-
A*03	A* 03:01:01:01, 03:02:01, 03:05:01 ; <i>A* 03:01:01:03d, 03:01:02d</i> ; A* 03:01:01:04e-03:01:01:07e, 03:01:04e-03:01:18e, 03:01:20e-03:01:37e, 03:01:40e-03:01:58e, 03:02:02e-03:02:04e, 03:04e, 03:05:02e, 03:20e	А3
	A* 03:21Nd; A* 03:01:01:02Ne, 03:03Ne, 03:11Ne	Null
	A* 03:06d-03:09d, 03:22:01d, 03:26d, 03:50d, 03:108d; A* 03:12e-03:16e, 03:17:02e-03:19e, 03:22:02e, 03:23:02e, 03:25e, 03:27e-03:29e, 03:31e-03:35e, 03:37e-03:40e, 03:42e, 03:44e-03:49e, 03:51e-03:81e, 03:83e-03:106e, 03:109e-03:151e, 03:153e-03:166e, 03:168Ne-03:170e, 03:172e-03:185e, 03:187e-03:218e, 03:220e-03:230e, 03:232e-03:249e	-
	This genotype was MANUALLY chosen.	
	This genotype was chosen as the most likely; cannot exclude one or more rare genotypes.	

HLA-B

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GROUP	ALLELES	ANTIGEN
B*14	B* 14:06:02e	B14
	B* 14:02:01:01 ; B* 14:02:01:02e-14:02:14e	B65
	<i>B* 14:09d</i> ; B* 14:15e-14:17e, 14:20e, 14:22e, 14:24e-14:25e, 14:27e, 14:29e-14:31e, 14:34e-14:36e, 14:38e-14:39e, 14:41Ne, 14:43e-14:45e, 14:50e-14:52e	-
B*57	B* 57:01:01 ; B* 57:01:02e-57:01:08e, 57:01:10e-57:01:22e, 57:08e	B57
	B* 57:06e, 57:10e, 57:13e-57:15e, 57:18e-57:21e, 57:23e-57:27e, 57:29e-57:31e, 57:33e-57:38e, 57:40e-57:41e, 57:43e-57:44e, 57:47e-57:50e, 57:52e-57:56e, 57:59e-57:62e, 57:64e-57:65e, 57:68e, 57:72e-57:75e, 57:77e-57:79Ne, 57:81e	-

	ALLELES	ANTIGEN
Bw4	B* 57:01:01 ; B* 57:01:02e-57:01:08e, 57:01:10e-57:01:22e, 57:06e, 57:08e, 57:10e, 57:13e-57:15e, 57:18e-57:21e, 57:23e-57:27e, 57:29e-57:31e, 57:33e-57:38e, 57:40e-57:41e, 57:43e-57:44e, 57:47e-57:50e, 57:52e-57:56e, 57:59e-57:62e, 57:64e-57:65e, 57:68e, 57:72e-57:75e, 57:77e-57:79Ne, 57:81e	Bw4
Bw6	B* 14:02:01:01 ; <i>B* 14:09d</i> ; B* 14:02:01:02e-14:02:14e, 14:06:02e, 14:15e-14:17e, 14:20e, 14:22e, 14:24e-14:25e, 14:27e, 14:29e-14:31e, 14:34e-14:36e, 14:38e-14:39e, 14:41Ne, 14:43e-14:45e, 14:50e-14:52e	Bw6

HLA-C

GROUP	ALLELES	ANTIGEN
C*06	C* 06:02:01:01 ; <i>C* 06:02:01:02d</i> ; C* 06:02:01:03e-06:02:07e, 06:02:09e-06:02:26e, 06:02:28e-06:02:45e	Cw6
	C* 06:16Ne	Null
	<i>C*</i> 06:07d-06:09d, 06:27d; C* 06:03:02e, 06:04:02e, 06:10e-06:15e, 06:17e-06:26e, 06:28e-06:29e, 06:32e-06:34:01e, 06:36e-06:43e, 06:45e-06:71e, 06:73e-06:76:01e, 06:77e-06:100e, 06:102e-06:117e, 06:119e-06:123e, 06:125e-06:126e, 06:128Ne-06:131e, 06:133e-06:135e, 06:137e, 06:139e-06:142e, 06:145e-06:146e, 06:148e-06:152Ne, 06:154e-06:167e, 06:169e-06:187e	-
C*08	C* 08:02:01:01, 08:04:01; C* 08:02:01:02e-08:02:03e, 08:02:05e-08:02:06e, 08:02:08e-08:02:10e, 08:02:12e-08:02:14e, 08:04:02e-08:04:03e	Cw8
	<i>C*</i> 08:05d, 08:12d; C* 08:07e, 08:17e-08:19e, 08:23e, 08:28e, 08:30e, 08:32e, 08:33:03e-08:35e, 08:37e, 08:43e, 08:45e, 08:48e-08:49e, 08:52Ne-08:55Ne, 08:62e-08:63e, 08:66e-08:71e, 08:73e-08:77e, 08:90e, 08:92e-08:93e, 08:100e, 08:103e-08:104e, 08:107e-08:108e, 08:110e-08:112e, 08:114e-08:116e, 08:120e, 08:125e-08:126e, 08:132e, 08:134e	-

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HLA-DR

HLA-DRB1

GROUP	ALLELES	ANTIGEN
DRB1*07	DRB1* 07:01:01:01 ; <i>DRB1* 07:03d</i> ; DRB1* 07:01:01:02e-07:01:19e, 07:04e, 07:09e	DR7
	DRB1* 07:10Ne	Null
	<i>DRB1* 07:05d</i> , <i>07:07d</i> , <i>07:11d</i> , <i>07:13d</i> ; DRB1* 07:06e, 07:08e, 07:12e, 07:14e-07:49e, 07:51e-07:70e	-
DRB1*13	DRB1* 13:01:01:01, 13:02:01; DRB1* 13:01:03d, 13:08d, 13:16d, 13:19d, 13:36d; DRB1* 13:01:01:02e-13:01:02e, 13:01:04e-13:01:16e, 13:02:02e-13:02:12e, 13:52e, 13:57e	DR13
	DRB1* 13:31d, 13:39d-13:40d, 13:59d, 13:76d; DRB1* 13:28e, 13:32e, 13:34e-13:35e, 13:51e, 13:64e-13:65e, 13:68e-13:69e, 13:72e-13:74e, 13:79e-13:80e, 13:83e-13:84e, 13:87e, 13:91e-13:93e, 13:97:01e, 13:98e-13:99e, 13:102e-13:103e, 13:105e-13:109e-13:110e, 13:112e-13:113Ne, 13:117e, 13:121e, 13:123e-13:128e, 13:130e-13:131e, 13:135e, 13:137Ne-13:143e, 13:145e, 13:147e-13:148e, 13:153e, 13:155e, 13:160e, 13:165e-13:166e, 13:168e, 13:171:02e, 13:173e, 13:177e, 13:182e, 13:184e-13:187e, 13:190e, 13:200Ne-13:202e, 13:204e-13:205e, 13:207e-13:213e, 13:215e, 13:218e, 13:220e	-

HLA-DRB345

GROUP	ALLELES	ANTIGEN
DRB3*03	DRB3* 03:01:01; DRB3* 03:01:02e, 03:01:03e, 03:01:05e, 03:02e	DR52
	DRB3*03:03e, 03:04e, 03:05e, 03:06e, 03:07e, 03:08e, 03:09e, 03:11e	-
DRB4*01:03: 01:02N	DRB4* 01:03:01:02N	Null

Cannot exclude RARE alleles from another locus.

HLA-DQA1

GROUP	ALLELES	ANTIGEN
DQA1*01	DQA1* 01:01:01:01, 01:01:02, 01:02:01:01, 01:03:01:01, 01:04:01:01, 01:05:01; DQA1* 01:02:02d; DQA1* 01:01:01:02e-01:01:01:03e, 01:01:03e, 01:02:01:02e-01:02:01:07e, 01:02:03e-01:02:04e, 01:03:01:02e-01:03:01:06e, 01:04:01:02e-01:04:02e, 01:05:02e-01:13e	-
DQA1*02	DQA1* 02:01:01:01; DQA1* 02:01:01:02e	-

HLA-DQB1

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GROUP	ALLELES	ANTIGEN
DQB1*03	DQB1* 03:01:13e, 03:01:15e, 03:01:21e, 03:01:25e	DQ7
	DQB1* 03:10e	DQ8
	DQB1* 03:03:02:01; DQB1* 03:03:02:02e-03:03:13e	DQ9
	DQB1* 03:12e, 03:15e-03:16e, 03:19:02e-03:20e, 03:23e, 03:26e, 03:30e-03:31e, 03:33e-03:34e, 03:38e-03:41e, 03:43e, 03:55e, 03:60e, 03:65e, 03:77e, 03:79e, 03:86e-03:91Qe, 03:95Ne-03:99Qe, 03:101e, 03:103e-03:105e, 03:108e, 03:111e-03:112e, 03:117e, 03:123e-03:124e, 03:126e, 03:130e, 03:136e-03:137e, 03:141e, 03:145e, 03:149e, 03:155e-03:156e, 03:168e, 03:172e, 03:176e-03:177e, 03:180e, 03:183e, 03:186e-03:187e, 03:195e, 03:200e, 03:202e, 03:209e, 03:212e, 03:217e, 03:222e, 03:227e, 03:230e	-
DQB1*06	DQB1* 06:11:01d; DQB1* 06:11:02e-06:12e	DQ1
	DQB1* 06:02:01, 06:03:01, 06:04:01, 06:08:01, 06:09:01; DQB1* 06:05:01d; DQB1* 06:01:14e, 06:02:02e-06:02:25e, 06:03:02e-06:03:22e, 06:04:02e-06:04:10e, 06:05:02e, 06:08:02e-06:08:03e, 06:09:02e-06:09:06e, 06:14e, 06:25e	DQ6
	DQB1* 06:26Ne	Null
	DQB1* 06:10d; DQB1* 06:07e, 06:13e, 06:15e-06:22e, 06:24e, 06:27e-06:34e, 06:36e-06:44e, 06:46e-06:52e, 06:57e-06:60e, 06:62e-06:98e, 06:106e-06:107e, 06:109e-06:119e, 06:121e-06:123e, 06:125e-06:131e, 06:133e-06:138e, 06:141e-06:148e, 06:150e-06:152e, 06:154e-06:156e, 06:158Ne-06:178e-06:180e, 06:182e-06:200e, 06:202e-06:204e, 06:206e-06:208e	-
	This genotype was MANUALLY chosen.	
	This genotype was chosen as the most likely; cannot exclude one or more rare genotypes.	

HLA-DPA1

GROUP	ALLELES	ANTIGEN
DPA1*01	DPA1* 01:03:01:01, 01:04 ; DPA1* 01:03:01:02e-01:03:05e, 01:05e-01:07e, 01:09e-01:10e, 01:12e	-
DPA1*01	DPA1* 01:03:01:01, 01:04 ; DPA1* 01:03:01:02e-01:03:05e, 01:05e-01:07e, 01:09e-01:10e, 01:12e	-

HLA-DPB1

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EPITYPE	ALLELES	HVR-A	HVR-B	HVR-C	HVR-D	HVR-E	HVR-F
EDP04	DPB1* 04:01:01:01; DPB1* 04:01:01:02e-04:01:12e, 04:01:14e-04:01:32e / DPB1* 99:01e / DPB1* 120:01Ne / DPB1* 121:01e / DPB1* 128:01e / DPB1* 134:01e / DPB1* 149:01e / DPB1* 169:01e / DPB1* 174:01e / DPB1* 175:01e / DPB1* 180:01e / DPB1* 181:01e / DPB1* 192:01e / DPB1* 180:01e / DPB1* 195:01e / DPB1* 192:01e / DPB1* 194:01e / DPB1* 195:01e / DPB1* 212:01e / DPB1* 213:01e / DPB1* 224:01e / DPB1* 225:01e / DPB1* 228:01e / DPB1* 231:01e / DPB1* 232:01e / DPB1* 240:01e / DPB1* 253:01e / DPB1* 255:01e / DPB1* 262:01e / DPB1* 272:01e / DPB1* 275:01e / DPB1* 276:01e / DPB1* 281:01e / DPB1* 320:01e / DPB1* 283:01e / DPB1* 298:01e / DPB1* 330:01e / DPB1* 336:01e / DPB1* 334:01e / DPB1* 350:01e / DPB1* 336:01e / DPB1* 341:01e / DPB1* 350:01e / DPB1* 356:01e / DPB1* 378:01e / DPB1* 396:01e / DPB1* 376:01e / DPB1* 415:01e / DPB1* 418:01e / DPB1* 425:01e / DPB1* 453:01e / DPB1* 455:01Ne / DPB1* 434:01e / DPB1* 464:01e / DPB1* 455:01Ne / DPB1* 459:01e / DPB1* 464:01e / DPB1* 465:01e / DPB1* 479:01e / DPB1* 475:01e / DPB1* 466:01e / DPB1* 479:01e / DPB1* 475:01e / DPB1* 550:01e / DPB1* 523:01e / DPB1* 553:01e / DPB1* 552:01e / DPB1* 534:01e / DPB1* 553:01e / DPB1* 554:01e / DPB1* 556:01e / DPB1* 559:01e / DPB1* 559:01e	LFQG	EEFA	AAE	ILEEK	M	GGPM
EDP72	DPB1* 72:01d	LFQG	EEFA	AAE	LLEEK	М	GGPM

One side of this epitype requires a RARE allele, review results to confirm assignment. This epitype was MANUALLY chosen.

This genotype was selected.

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