## REPORT OF THE 39<sup>th</sup> INTERNATIONAL MICA EXCHANGE MAY 11, 2020

MICA 157 - 162

For the 39<sup>th</sup> MICA Exchange, 6 DNA samples (MICA #157 - MICA#162) were shipped to 19 laboratories worldwide. MICA typing results were received

from 18 laboratories. Results are summarized on Table 1 and individual laboratory results are listed on tables 2 - 7.

**MICA #157**. MICA\*002 and MICA\*027 was the consensus type for this cell from a Caucasian donor. MICA\*002 was reported by 11 labs, with 8 assigning MICA\*002:01. A number of labs (n = 7) were unable to resolve MICA\*002 from MICA\*020 and MICA\*055. MICA\*002 is identical to MICA\*020 and MICA\*055, except for the number of GCT repeats in its transmembrane domain (exon 5). MICA\*002 has 9 GCT repeats, where as MICA\*020 has 10 GCT repeats and MICA\*055 has 8 GCT repeats.

MICA\*027 was reported by 11 labs, while MICA\*027/\*048 was reported by 7 labs. MICA\*048 differs from MICA\*027 by a single nucleotide substitution in exon 5 at codon 316 (GAG  $\rightarrow$  GAT). The substitution results in an amino acid change from glutamic acid in MICA\*027 to aspartic acid in MICA\*048.

**MICA #158**. MICA\*008 and MICA\*011 was the reported consensus type for this sample from an African American donor. MICA\*008 was reported by 16 labs, with 7 labs assigning MICA\*008:01. A number of labs (n = 4) were unable to resolve MICA\*008:01 from MICA\*008:03, MICA\*008:04 and/or MICA\*088N. MICA\*008:03 and MICA\*008:04 each differ from MICA\*008:01 by a single nucleotide substitution. For MICA\*008:03 the substitution in exon 4 at codon 193 (GCC  $\rightarrow$  GCA), and for MICA\*008:04 its in exon 1 (TTC  $\rightarrow$  TTT). Each results in a synonymous substitution. MICA\*011 was reported in complete consensus.

**MICA #159.** MICA\*001 and MICA\*004 was the reported consensus type for this sample a Hispanic donor. Each were well assigned by 100% of labs.

**MICA #160**. MICA\*012 was the reported consensus type for this sample from an Asian donor. MICA\*012 was reported in complete consensus, with 11 labs reporting MICA\*012:01. Interestingly, only 11 labs reported this sample as MICA\*012 homozygous. Five labs assigned the second allele as MICA\*012/\*018, and 2 reported it as MICA\*018. Two nucleotide substitutions distinguish MICA\*018:01 from MICA\*012:01. One is in exon 3 at codon 151 (CTC  $\rightarrow$  CAC) where leucine is replaced with histidine in MICA\*018:01. The second substitution is in exon 6 at codon 360 (GCC  $\rightarrow$  ACC), which results in an amino acid change from alanine to threonine in MICA\*018:01.

**MICA #161.** MICA\*010 and MICA\*015 was the reported consensus type for this sample from an African American donor. MICA\*010 was reported by 11 labs, with 8 labs assigning MICA\*010:01. MICA\*010/\*069 was reported by 6 labs and 1 lab reported MICA\*010:01/\*065/\*069. MICA\*069 differs from MICA\*010:01 by a single nucleotide substitution in exon 5 at codon 350 (GCT  $\rightarrow$  GAT), which results in alanine being replaced by aspartic acid in MICA\*069. MICA\*015, was reported in complete consensus.

**MICA #162**. MICA\*027 was reported in good agreement (> 80%) in this sample from a donor of unknown ethnicity. Ten labs assigned MICA\*027 and 5 labs assigned MICA\*027/\*048. No consensus was achieved for Allele-1 in this sample, as 5 labs assigned MICA\*009, 5 assigned MICA\*049, and 5 were unable to resolve MICA\*009 from MICA\*049. As result, this second allele was not graded. MICA\*049 differs from MICA\*009 by a single nucleotide substitution in exon 6 at codon 333 (ACG  $\rightarrow$  ATG), in which threonine is replaced by methionine in MICA\*049.

**NEXT MAILING DATE: August 5, 2020** 

Arlene Locke, David Gjertson, Qiuheng Zhang, and Elaine F. Reed

Table 1. Summary of 39th MICA Exchange #157-#162

MICA#15	7				
18 labs					
Allele-1	%(n)				
*002:01	44(8)				
*002:01/*002:02	6 (1)				
*002	11(2)				
*002:01/*020/*055	11(2)				
*002/*020/*055	22(4)				
*002/*020/*027/*048	6(1)				
18 labs					
Allele-2	%(n)				
*027	61(11)				
*027/*048	27(5)				
*008:03/*027/*048	6 (1)				
*027/*048/*055	6 (1)				

MICA#158	
18 labs	
Allele - 1	%(n)
*008:01:01	6 (1)
*008:01	33(6)
*008:01/*008:04	11(2)
*008	33(6)
*008:01/*008:03/*008:04	6 (1)
*008:01/*008:04/*088N	6 (1)
*008/*088N	6 (1)
18 labs	
Allele - 2	%(n)
*011	100(18)

MICA#16	60
18 labs	
Allele-1	%(n)
*012:01	61(11)
*012:01/*012:04	6 (1)
*012	33(6)
18 labs	
Allele-2	%(n)
*012:01	50(9)
*012:01/*012:04	5 (1)
*012	5 (1)
*012:01/*018:01	5 (1)
*012/*018	22(4)
*018:01	5 (1)
*018	5 (1)

MICA#161	
18 labs	
Allele-1	%(n)
*010:01	44(8)
*010	17(3)
*010:01/*069	11(2)
*010/*069	22(4)
*010:01/*065/*069	6 (1)
18 labs	
Allele-2	%(n)
*015	100(18)

MICA	<b>#159</b>
18 la	bs
Allele - 1	%(n)
*001	100(18)
18 la	he
Allele - 2	%(n)
*004	100(18)

MICA#162	
17 labs	
Allele-1	%(n)
*009:01	17(3)
*009	12(2)
*049	29(5)
*009:01/*049	12(2)
*009/*049	17(3)
*009:01/*049/*076	6 (1)
*009/*027/*048	6 (1)
17 labs	
Allele-2	%(n)
*027	56(10)
*027/*048	28(5)
*027/*048/*049	6 (1)
*008:03/*027/*048/*024	6 (1)

Table 2. MIC	Table 2. MICA typing results reported by participating laboratories.						
MICA# 157	CTR	Allele-1	Allele-2	Others	Method		
(Caucasian)	234	*002:01/*020/*055	*027/*048		SSO		
	278	*002:01	*027	*020/*048/*055	SSO		
	733	*002:01	*027		SBT		
	762	*002:01	*027		NGS		
	1426	*002:01/*020/*055	*027/*048		SSO		
	3753	*002/*020/*055	*027/*048		SSO		
	3798	*002/*020/*055	*027/*048		SSO		
	3966	*002	*027	*048/*055	SSO		
	4337	*002/*020/*055	*027		SSP		
	4345	*002/*020/*055	*027/*048				
	5133	*002:01	*027		NGS		
	8035	*002/*020/*027/*048	*027/*048/*055		SSO		
	8047	*002:01/*002:02	*008:03/*027/*048		NGS		
	8073	*002:01	*027		NGS		
	8080	*002:01	*027		NGS		
	8086	*002:01	*027		SBT		
	8105	*002:01	*027		NGS		
	8108	*002	*027	*020/*048/*055	SSO		

Table 3. MI	able 3. MICA typing results reported by participating laboratories.					
MICA# 158	CTR	Allele-1	Allele-2	Others	Method	
(Black)	234	*008:01/*008:04	*011		SSO	
	278	*008:01	*011	*008:04/*088N	SSO	
	733	*008:01:01	*011	*008:01:02/*008:04	SBT	
	762	*008:01/*008:04	*011		NGS	
	1426	*008:01/*008:04/*088N	*011		SSO	
	3753	*008	*011		SSO	
	3798	*008	*011		SSO	
	3966	*008	*011		SSO	
	4337	*008	*011		SSP	
	4345	*008	*011			
	5133	*008:01	*011		NGS	
	8035	*008/*088N	*011		SSO	
	8047	*008:01/*008:03/*008:04	*011		NGS	
	8073	*008:01	*011		NGS	
	8080	*008:01	*011		NGS	
	8086	*008:01	*011		SBT	
	8105	*008:01	*011		NGS	
	8108	*008	*011	*088N	SSO	

Table 4. MI	Table 4. MICA typing results reported by participating laboratories.					
MICA# 159	CTR	Allele-1	Allele-2	Others	Method	
(Hispanic)	234	*001	*004		SSO	
	278	*001	*004		SSO	
	733	*001	*004		SBT	
	762	*001	*004		NGS	
	1426	*001	*004		SSO	
	3753	*001	*004		SSO	
	3798	*001	*004		SSO	
	3966	*001	*004		SSO	
	4337	*001	*004		SSP	
	4345	*001	*004			
	5133	*001	*004		NGS	
	8035	*001	*004		SSO	
	8047	*001	*004		NGS	
	8073	*001	*004		NGS	
	8080	*001	*004		NGS	
	8086	*001	*004		SBT	
	8105	*001	*004		NGS	
	8108	*001	*004		SSO	

Table 5. MI	Table 5. MICA typing results reported by participating laboratories.					
MICA# 160	CTR	Allele-1	Allele-2	Others	Method	
(Asian)	234	*012:01	*018:01		SSO	
	278	*012:01	*012:01	*018:01	SSO	
	733	*012:01			SBT	
	762	*012:01			NGS	
	1426	*012:01	*012:01/*018:01		SSO	
	3753	*012	*012/*018		SSO	
	3798	*012	*012/*018		SSO	
	3966	*012	*018		SSO	
	4337	*012:01	*012:01		SSP	
	4345	*012	*012/*018			
	5133	*012:01	-		NGS	
	8035	*012	*012/*018		SSO	
	8047	*012:01/*012:04	*012:01/*012:04		NGS	
	8073	*012:01			NGS	
	8080	*012:01	*012:01		NGS	
	8086	*012:01	*012:01		SBT	
	8105	*012:01			NGS	
	8108	*012	*012	*018	SSO	

Table 6. MI	Table 6. MICA typing results reported by participating laboratories.					
MICA# 161	CTR	Allele-1	Allele-2	Others	Method	
(Black)	234	*010:01/*069	*015		SSO	
	278	*010:01	*015	*069	SSO	
	733	*010:01	*015	*065/*069	SBT	
	762	*010:01	*015		NGS	
	1426	*010:01/*069	*015		SSO	
	3753	*010/*069	*015		SSO	
	3798	*010/*069	*015		SSO	
	3966	*010	*015	*069	SSO	
	4337	*010	*015		SSP	
	4345	*010/*069	*015			
	5133	*010:01	*015		NGS	
	8035	*010/*069	*015		SSO	
	8047	*010:01/*065/*069	*015		NGS	
	8073	*010:01	*015		NGS	
	8080	*010:01	*015		NGS	
	8086	*010:01	*015		SBT	
	8105	*010:01	*015		NGS	
	8108	*010	*015	*069	SSO	

CA# 162	CTR	Allele-1	Allele-2	Others	Method
known)	234	*009:01/*049	*027/*048		SSO
,	278	*009:01	*027	*048/*049	SSO
	733	*009:01	*027	*048/*049	SBT
	762	*049	*027	*024/*076	NGS
	1426	*009:01/*049	*027/*048		SSO
	3753	*009/*049	*027/*048		SSO
	3798	*009/*049	*027/*048		SSO
	3966	*009	*027	*049/*048	SSO
	4337	*049	*027		SSP
	4345	*009/*049	*027/*048		
	5133	*049	*027	*024/*076	NGS
	8035	*009/*027/*048	*027/*048/*049		SSO
	8047	*009:01/*049/*076	*008:03/*027/*048/*024		NGS
	8073	no amplicon			NGS
	8080	*009:01	*027	*049/*048	NGS
	8086	*049	*027		SBT
	8105	*049	*027		NGS
	8108	*009	*027	*048/*049	SSO