

IMGT overview: the mouse T cell receptor alpha TRA genes

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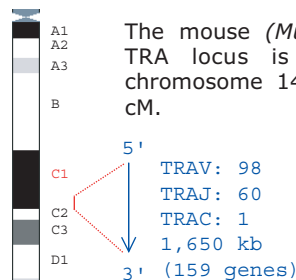


http://imgt.cines.fr

How many TRA genes?

Chromosome 14 C1

The total number of T cell receptor alpha TRA genes per haploid genome in *Mus musculus* laboratory mice is 159 (including 10 TRAV/DV: 8 rearranged either to TRAJ or to TRDD, 2 assigned by homology).



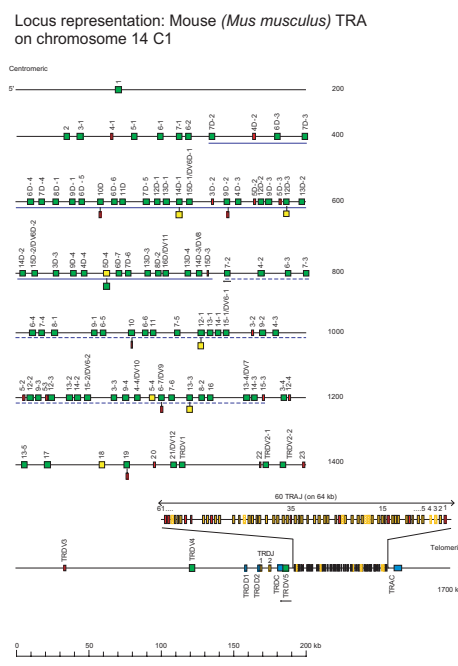
The mouse (*Mus musculus*) TRA locus is located on chromosome 14 C1 at 19.7 cM.

The mouse TRA locus spans 1,650 kb and comprises, from 5' to 3', 98 TRAV genes organized in 2 clusters upstream of 60 TRAJ genes and of a single TRAC gene.

Lefranc, M.-P. et al., *In Silico Biology*, 5, 45-60 (2005)

How are TRA genes organized?

TRA locus



Bosc, N. et al., *Dev. Comp. Immunol.*, 27, 465-497 (2003)

How many functional TRA genes?

Potential repertoire

The potential TRA repertoire per haploid genome comprises 112-123 functional genes (including 10 TRAV/DV): 73-84 TRAV (10 TRAV/DV), 38 TRAJ and 1 TRAC.

Overview		
	Number of genes	Functional genes
TRAV	98	73-84
TRAJ	60	38
TRAC	1	1
Total	159	112-123

The definitive IMGT nomenclature of the mouse TRAV genes and the correspondence with the provisional nomenclature have been established.

The mouse TRA genes and alleles and the corresponding IMGT reference sequences were provided to Mouse Genome Informatics MGD in July 2002 and are available in [IMGT/GENE-DB](#).

Giudicelli, V. et al., *Nucl. Acids Res.*, 33, D256-D261 (2005)

IMGT tools to analyse expressed variable genes

IMGT/V-QUEST IMGT/JunctionAnalysis

Sequence analysis

Analysis of the TRAV genes (germline or rearranged) can be performed by [IMGT/V-QUEST](#) and analysis of the V-J junctions by [IMGT/JunctionAnalysis](#).

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Results of IMGT/V-QUEST
Alignment for V-GENE
M73551 TRAV14-2*02 score 1198 CAGGACAGGTGAGCAAAATCCCAATCTC
L42255 TRAV14-2*02 1199 CAGGACAGGTGAGCAAAATCCCAATCTC
M54625 TRAV14-2*03 1199 CAGGACAGGTGAGCAAAATCCCAATCTC
L27248 TRAV14-2*03 1199 CAGGACAGGTGAGCAAAATCCCAATCTC
M54522 TRAV14-1*02 1099 CAGGACAGGTGAGCAAAATCCCAATCTC
A3311368 TRAV14-1*03 1099 CAGGACAGGTGAGCAAAATCCCAATCTC

Alignment for J-GENE
M73551 score CAGGCAATCCGGAAATCATCTTTGGACTGGGACACCTTACAGTGCACACACATCCGAA
M64239 TRAJ37*01 285 .....

Results of IMGT/JunctionAnalysis
Analysis of the JUNCTION
Input V name V-REGION N J-REGION J name Npc
#1 M73551 TRAV14-2*02 tggtcgacgaag...g...acaggaatccggaaactctttt TRAJ37*01 1/1

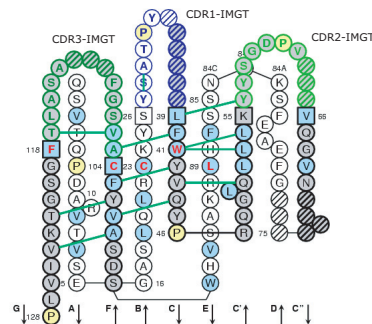
Translation of the JUNCTION
105 106 107 109 113 115 117 CDR3-IMGT
104 106 108 110 114 116 118 frame length
#1 M73551 tgi gca agg aca ggc aat acc gga aaa ctc att ttt + 11
C A A R T G N T T G K L I F
  
```

Giudicelli, V. et al., *Nucl. Acids Res.*, 32, W435-W440 (2004)

IMGT Colliers de Perles

2D representations

Mus musculus (Mouse) TRAV_1 V-DOMAIN from 2C (1g6r_A)
CDR-IMGT lengths [6.7.10]



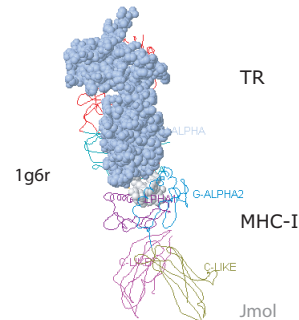
IMGT Colliers de Perles are according to the IMGT unique numbering for V-DOMAIN. Hydrogen bonds are shown as green lines.

Lefranc, M.-P. et al., *Dev. Comp. Immunol.*, 25, 55-77 (2003)
Lefranc, M.-P. et al., *Dev. Comp. Immunol.*, 29, 185-203 (2005)

IMGT/3Dstructure-DB IMGT/StructuralQuery

3D structures

Three-dimensional structures of 17 V-ALPHA domains encoded by rearranged TRAV-TRAJ genes are available in [IMGT/3D-structure-DB](#). Seven TR-ALPHA-BETA have been crystallised.



The ALPHA chain is in spacefill (the BETA chain and MHC-I are in wireframe).

Kaas, Q. et al., *Nucl. Acids Res.*, 32, D208-210 (2004)
Kaas, Q. and Lefranc, M.-P., *In Silico Biol.*, 5, 0046 (2005)

