Supplementary Figure 1.

Ovine and bovine MAIT cell TCR α chain nucleotide sequences and location of primers.
In red: primers used for cloning and sequencing of bovine and ovine MAIT TCR; in blue: primers used in MAIT V region and α chain constant region qPCR; in pink: qPCR probes; in grey: CDR regions.

Supplementary Figure 2
Alignment of MR1 nucleotide sequences of different species.
Alignment of full-length bovine (bo) (accession number FJ028657.1), ovine (ov) (FJ423039), human (hu) (AJ249778.1), and murine (mu) (AF010448.1) MR1 nucleotide sequences.
* Indicates a position with a fully conserved residue.

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        ┌Leader                                                    ┌Extracellular domain
BoMR1   -----------ATGATGCTCCTATTGCCTCTCATCATTGTATTAATGATGAAGCTCAGCGATGCTCGGACTCACTCTCTGAGATATTTT
OvMR1   -----------ATGATGCTCCTATTGCCTCTCATCATTGTATTAATGATGAAGCTCAGCGATGCTCGGACTCACTCTCTGAGATATTTT
HuMR1   ATGGGGGAACTGATGGCGTTCCTGTTACCTCTCATCATTGTGTTAATGGTGAAGCACAGCGATTCCCGGACGCACTCTCTGAGATATTTT
MuMR1   -----------ATGATGCTCCTGTTACCTCTGCTCGCTGTATTCTTGGTGAAGCGAAGCCATACTCGGACCCACTCGCTGAGATATTTT
                   *** * **** ** ***** ** *** ** ** ****** *** ** * ***** ***** ***********

BoMR1   CGCCTGGGCATTTCAGAGCCTGGCTATGGGATCCCTGAATTTATTTCAGCCGGGTATGTGGATTCTCATCCCATCACCATGTACAACAGT
OvMR1   CGCCTGGGCATCTCAGAGCCTGGCTATGGGATCCCTGAATTTATTTCAGCTGGGTATGTGGATTCTCATCCCATCACCATGTACAACAGT
HuMR1   CGCCTGGGCGTTTCGGATCCCATCCATGGGGTCCCTGAATTTATTTCGGTTGGGTACGTGGACTCGCACCCTATCACCACATATGACAGT
MuMR1   CGTCTGGCTGTTTCCGATCCTCGGTCCCGTCGTCCCTGAATTTATCTCTGTTGGGTATGTGGACTCACACCCTATCACTACATACGACAGT
        ** ****   * ** ** **        *    ************ ** **  ***** ***** ** ** ** ***** *  ** *****

BoMR1   GTGTCCCAGCTGAAGGAGCCGCGGGCCCCTGTGGATGGAGGAAAACCTGGCACCTGATCACTGGGAGAGGTACACACAGCTGCTGCGGGGC
OvMR1   GTCTCCCAGCTGAAGGAGCCACGTGCCCCATGGATGGCGGAAAACCTGGAACCTGATCACTGGGAGAGGTACACACAGCTGCTGCGGGGC
HuMR1   GTCACTCGGCAGAAGGAGCCACGGGCCCCATGGATGGCAGAGAACCTCGCGCCTGATCACTGGGAGAGGTACACTCAGCTGCTGAGGGGC
MuMR1   GTCACTCGACAGAAGGAGCCGAAAGCTCCATGGATGGCAGAGAACCTGGCACCTGATCACTGGGAGAGGTACACTCAGCTGCTAAGGGGC
        **  * *  * ********     ** *  ******* ** *****  *  **************************** ********  *****

BoMR1   TGGCAGCAGGCCGTTCAAGGTGGAACTGAAGCAGCTGCAGCATCACTACAATCACTCAGGGTTTCACACTTACCAGAGAATGATTGGCTGT
OvMR1   TGGCAGCAGGCGTTCAAGGTGGAACTGAAGCAGCTGCAGCATCACTACAATCACTCAGGGTTTAACACTTACCAGAGAATGATTGGCTGT
HuMR1   TGGCAGCAGATGTTCAAGGTGGAACTGAAGCGCCTACAGAGGCACTACAATCACTCAGGGTCTCACACTTACCAGAGAATGATTGGCTGT
MuMR1   TGGCAGCAGACATTCAAGGCGGAGCTGAGGCACCTACAGAGGCACTACAACCACTCAGGGCTTCACACCTACCAGAGAATGATTGGCTGT
        *********  ******* *** **** ** ** *** **  ** ***    * **** ************************

BoMR1   GAGTTACTGGAGGATGGAAGCATCACGGGGATTTCTCCAATATGCATATGATGGACAGGATTTCCTTATCTTCAATAAAGATACCCTGTCC
OvMR1   GAGTTACTGGATGATGGCAGCACCACGGGGATTTCTCCAATATGCGTATGATGGACAGGATTCATTATCTTCAATAAAGATACCCTCTCC
HuMR1   GAGCTGCTGGAGGATGGAAGCACCACAGGGATTTCTGCAGTATGCATATGACGGGCAGGATTTCCTGATCTTCAATAAAGACACCCTCTCC
MuMR1   GAGTTGCTAGAAGATGGCAGCACCACAGGGTTTCTCCAGTATGCATATGATGGACAAGATTTCATCATCTTCAATAAAGACACCCTCTCC
        *** *  ** ** ** ***** **** *** ***** **  ***** ***** *  ** ** ** ************* **** ***

BoMR1   TGGATGGCCATGGATAATGTGGCTGACATCATCAGGCGGGTATGGGAGGCCAATCGGCATGAGTTACAATATCAGAAGAATTGGCTGGAA
OvMR1   TGGATAGCCATGGATAATGTGGCTAACATCATCAGGCGGGCATGGGAGGCCAATCGGCATGAGTTACAATATCAGAAGAATTGGCTGGAA
HuMR1   TGGCTGGCTGTAGATAATGTGGCTCACACCATCAAGCAGGCATGGGAGGCCAATCAGCATGAGTTGCTGTATCAAAAGAATTGGCTGGAA
MuMR1   TGGTTGGCCATGGATTATGTGGCTCACATCACCAAGCAAGCATGGGAGGCCAACCTGCATGAGTTGCAATACCAAAAGAACTGGCTGGAA
        *** * **  * *** ******* *** ** ** *  * ************ * ********* *  ** ** ***** *********

BoMR1   GAAGAATGCATTGCTTGGTTAAAGAGATTCCTGGAGTATGGGAAAGATGCCTTACAAAGAACAGAGCCCCCCAAAGTCAGAGTCAATCAC
OvMR1   GAAGAATGTATTGCTTGGTTAAAGAGATTCCTGGAGTATGGGAAAGATACCTTACAAAGGACAGAGCCCCCCAAAGTCCGAGTCAATTAC
HuMR1   GAAGAATGTATTGCCTGGCTAAAGAGATTCCTGGAGTATGGGAAAGACACCCTACAAAGAACAGAGCCCCCCACTGGTCAGAGTAAATCGC
MuMR1   GAAGAATGCATTGCCTGGCTAAAGAGGTTCTTGGAATATGGAAGAGATACCCTAGAAAGAACAGAGCATCCAGTGGTAAGAACAACTCGA
        ******** ***** ***** *** ***  *** **** ** * ****  * ***  ** ** **** *******   **   **   **    * *

BoMR1   AAAGAAACTTTCCCAGGGATTACAACTCTTTACTGCAGAGCTTATGGCTTTTACCCCCCAGAAATTTCCATTAACTGGATGAAAAACGGG
OvMR1   AAAGAAACTTTCCCAGGGATTACAACTCTTTACTGCAGAGCTCATGGCTTTTACCCCCCAGAAATTTCCATAAACTGGATGAAAAACGGG
HuMR1   AAAGAAACTTTTCCAGGGGTTACAGCTCTTCTGCAAAGCTCAGCTTTTACCCCCCAGAAATTTCATGACATGGATGAAAAACGGG
MuMR1   AAGGAAACTTTTCCAGGGATTACAACTTTCTTCTGCAGAGCTCATGGCTTCTACCCACCAGAAATTTCCATTGACATGGATGAAAAATGGG
        ** ******** ****** ***** **  ** ** * ***** ****  ** **** ********** *** **  ********** ***

BoMR1   GAAGAAATTTTCCAAGATACCGATTATGGAGGCATTCTTCCCAGTGGGGATGGGACCTATCAGACTTGGGTGTCTGTTGAGCTGGATCCT
OvMR1   GAAGAAGTTGTCCAAGACACCAATTATGGAGGCATTCTTCCCAGTGGGGATGGGACCTATCAGACTTGGGTGTCTGTTGAGCTGGATTCT
HuMR1   GAAGAAATTGTCCAAGAAATTGATTATGGAGACATTCTTCCCAGTGGGGATGGAACCTATCAGGCGTGGGCATCAATTGAGCTTGATCCT
MuMR1   GAAGAAATTGCCCAAGAAGTGGATTACGGAGGGGGTACTTCCCAGCGGGGATGGAACCTATCAAACGTGGCTGTCAGTTAATCTGGATCCT
        ****** **  ******    **** ****     **** ***  * ******** ******** ** *******  * **  ** * ** *** **

BoMR1   CAGAATGGTGACATTTACTCCTGTCACGTGGAGCATGGCGGAGTCCACATGGTTCTTCAGGGTTTCCAGGAATCGGAAACTATACTTCTC
OvMR1   CAGAACGGTGACATTTACTCCTGTCACGTGGAGCATGGCGGCGTCCACATGGTTCTTCCGGGTTTCCAGGATTCGGAAACTATGCTTCTG
HuMR1   CAGAGCAGCAACCTTTACTCCTGTCATGTGGAGCACTGCGGTGTCCACATGGTTCTTCAGGTCCCCCAGGAATCAGAAACTATCCCTCTT
MuMR1   CAGAGCAATGATGTTTATTCTTGTCATGTGGAGCACTGTGGTCGCCAAATGGTTCTGGAGGCCCCTCGGGAATCAGGGGACATCCTTCGA
        ****        *  **** ** ***** ******* *  ** *   *** ******** **     *** ***** *    ** * **

        Transmembrane                                             ┌Cytoplasmic domain
BoMR1   GTGGTGAAAGCTGT---TGGGTTCATTGTCCTCGCCATTGCCCTGGCTGGAGTTGGCATCCTAGCCTGGCGAAGAGGCCCCGAGGGAAA
OvMR1   GTGTGATGAAAGCTGT---TGGGTTCATTGTCCTCGCCATTGCCCTGGCTGGAGTTGGCATCTTAGCCTGGCGAAGGAAGCCCCGAGGGGAA
HuMR1   GTGTGATGAAAGCTGTCTCTGGGTCCATTGTCCTTGTCATTGTGCTGGCTGGAGTTGGTGTTCTAGTCTGGAGAAGAAGGCCCCGAGAGCAA
MuMR1   GTGAGCACGATCTC---TGGAACCACAATTCTCATCATCGCCCTGGCTGGAGTTGGTGTTCTGATCTGG---AGAAGGTCACAAGAACTA
        ***  *           ***   **    * **   *** *  ************* *  *   **** * *  * * * **        *

BoMR1   AACAAAGTCATCTGCCTTTCTACACCAGAACACTGA-----------------
OvMR1   AACAAAGTCATCTACCTTTCTACACCAGAACACTGA-----------------
HuMR1   AATGGAGCCATCTACCTTCCAACACCAGATCGATGA-----------------
MuMR1   AAAGAAGTCATGTACCAACCCACCCAAGTGAATGAGGGGAAGCTCTCCCTCTTAG
        **   ** *** * **   * ** * **
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