A

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                                  [   ] HAE III
L00635 ACCCCATAAACAAATAGGTTTGGTCCTAGCCTTTCTATTAGCTCTTAGTAAGATTACACATGCAAGCATCCCC H00708
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       .........................................................................
       .........................................................................
       .........................................................................
       ....c....................................................................
       ...........................G.............................................
       .T........................................................T..............
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        [         ]                            HINC II
L13257 AGTCAACTAGGACTCATAATAGTTACAATCGGCATCAACCAACCACACCTAGCATTCCTGCACATCTGTACCCACGCCTTCTTCAAAGCCATACTATTTATGTGCTCCGGGTCCATCAT H13377
       .....G............-....................................................................................................
       .....G.................................................................................................................
       .....G.................................................................................................................
       .....G....R................................................N.......K.......K..................................T........
       .....G.........................................................................................A.....................
       .....G.................................................................................................................
       ...............R....................................................A................................R.................
       ......................................................................................................................C
       .....................................................................................................A.................
       ....................................................................................G..................................
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                                                       ALU I                                                                   [    ]
L05054 TACAACCCTAACATAACCATTCTTAATTTAACTATTTATATTATCCTAACTACTACCGCATTCCTACTACTCAACTTAAACTCCAGCACCACGACCCTACTACTATCTCGCACCTGAAACAAGCTAACATGACT H05189
       ..................................................................................................................C...................
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       ......................................................................................................................................
       ......................................................................................................................................
       .........................................W............................................................................................
       .....T................................................G..T............................................................................
       .....T...................................................T....................T.......................................................
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B

223

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L16209                         223                                                                         298
CCCCATGCTTACAAGCAAGTACAGCAATCAACCCTCAACTATCACACATCAACTGCAACTCCAAAGCCACCCCTCACCCACTAGGATACCAACAAACCTACCCACCCTTAACAGTACATAGTACATAAAGCCA H16303
                    .............T...G......................................................................C....
                    .............T......................G...........................................T.......C..T.
                    .............T..........................................................................C....
                    .............T......................................S.B.Y...............................C....
                    .............T.......................................B..................................C....
                    .............T..........................................B...................................
                    .............T................................T.....................................G...C....
                    .............T..........................................................................C....
                    .............T.............................G............................................C....                    325 327
                                                                        16287 CACTAGGATACCAACAAACCTACCCACCCTTAACAGTACATAGTACATAAAGCCATTTACCGTACATAGCACATTACAGTCAAATCCCTTCTCGTCCCCATGGATGACCCCCCTCAGATAGGGGTCCCTTG H16379
                                                                                                  ..........C..........................C.T...................................................
                                                                                                  ..........C..........................C.T...................................................
                                                                                                  ....-.....C..........................C.T...................................................
                                                                                                  .....T....C..........................C.T...................................................
                                                                                                  ..........C............C.............C.T...........T.......................................
                                                                                                  ......T.T.......C..............................T...........................................
                                                                                                  ....T......................................................................................
                                                                                                  ............................................T..............................................
                                                                                                  ...........................................................................................
                                                                             Oxford direct        ..........C..........................C.T...................................................
                                                                                                 |..........C..........................C.T....................................T..............
                                                                                                 |..........C.G.....................T..C.T...................................................
                                                                                                 |..........C..........................C.T...................................................
                                                                             Oxford clones ------|..........C..........................C.T...................................................
                                                                                                 |..........C................................................................................
                                                                                                 |..........C..........................C.T...................................................
                                                                                                 |..........C..........................C.T...................................................
                                                                                                 |..........C..........................C.T...................................................
                                                                                                 |..........C.....G......A.G..........GC.TG..................G................T..............
                                                                                                 |..........C..........................C.T....................................TT.............
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C

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                                                                                               16S
L02828 AACAGAAAATCCTCCGAGCGATTTTAAAGACTAGACCAACAAGTCAAACCAAACCATCGCTTATTGATCCAAAAACTTGATCAACGGAACA H02922 - Ovis
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       CAGAACCCAACCTCCGAGCAGTACATGCTAAGACTTCACCAGTCAAAGCGAACTACTATACTCAATTGATCCAATAACTTGACCAACGGAACA      - Homo
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       ...................................................T......................A.......T..........
       ......................G............................G.........................................
       ....M..-...-.......................................R.........................................
       ....-..-...Y--..........................................C....................................
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                                                                                               12S
L01091 GCTTAGCCCTAAACACAAATAATTATAAAAACAAAATTATTCGCCAGAGTACTACCGGCAACAGCCCGAAACTCAAAGGACTTGGCGGTGCTTTATACCCTTC H01195 - Ovis
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       ........................................................S..............................................
       GCTTAGCCCTAAACCTCAACAGTTAAATCAACAAAACTGCTCGCCAGAACACTACGAGCCACAGCTTAAAACTCAAAGGACCTGGCGGTGCTTCATATCCCTC        - Homo
       .......................................................................................................
       ...........................................T...........................................................
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