

1- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCGACAA  
2- ACCA-TCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
3- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
4- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
5- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
6- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
7- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
8- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
9- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
10- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
11- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
12- ACCA-GCCAGCTGAGCCAATTCATGGACCAGAACAACCCGCTGTCGGGGTTGACCCACAA  
13- ACCGCGTCGTGTATGACTCTGTATACACAGAGGAGTCACGGCGCGCGTGGTGGTCTCCAT

**Figure 3 DNA sequence analysis alignment showing the difference between nucleic acid of *Mycobacterium tuberculosis* complex (samples 1-12) and *Mycobacteria* other than *M. tuberculosis* (sample 13)**

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