

## SUPPLEMENTARY MATERIAL No.9

## Characterization of the mRNA untranslated regions [UTR] of the *Trypanosoma cruzi* LYT1 isoforms derived by alternative *trans*-splicing

**Suppl. 9.** Multiple alignment of the complete 3´ UTR-II sequence from *T. cruzi* isolates listed in Table 1 and those sequences obtained in this study from 058PUJ (GenBank ID: KU973678) and D.A. (GenBank ID: KU973679) isolates. Sequences were aligned with CLUSTAL Omega [1.2.4] program. Numbers on the right correspond to the nucleotide position in the sequence. An asterisk means that nucleotides are identical in all sequences show in the alignment. The motifs of RNA are indicated in bold. The polymorphisms present in the RNA motifs are shown in red.

AGUCUUUAGCAUUAGGGAUGACAAUC**AUAGU**CG-UUUGCAGCUUCGUUGAGGUUGGUACC

AGUCUUUAGCAUUAGGGAUGACAAUC <b>AUAGU</b> CG-UUUGCAGCUUCGUUGAGGUUGGUACC	59
AGUCUUUAGCAUUAGGGAUGACAAUC <b>AUAG</b> UCG-UUUGCAGCUUCGUUGAGGUUGGUACC	59
AAUCCUUAGCAUUAGGGAUGACAAUC <b>AUAG</b> UCG-UUUGCAGCUUCGUUAAGGUUGGUACC	59
AGUCUUUAGCAUUAGGGAUGACAAUC <b>AUAG</b> UCGUUUUGCAGCUUCGUUGAGGUUGGUACA	60
AGUCUUUGGCAUUAGGGAUGACAAUC <b>AUAG</b> UCGUUUUGCAGCUUCGUUGAGGUUGGUACA	60
AAUCCUUAGCAUUAGGGAUGACAAUC <b>AUAG</b> UCG-UUUGCAGCUUCGUUAAGGUUGGUACC	59
AGUCUUUGGCAUUAGGGAUGACAAUC <b>AUAG</b> UCGUUUUGCAGCUUCGUUGAGGUUGGUACA	60
AGUCUUUGGCAUUAGGGAUGACAAUC <b>AUAG</b> UCGUUUUGCAGCUUCGUUGAGGUUGGUACA	60
GGUGUUUAGCAUUAGGGAUGACAAUC <b>AUAG</b> UCG-UUUGCAGCUUUGUUGACUUUGUUAUC	59
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UUCAGAGAAUUGUAUCUGCACCCUGUGCAGUGAUACACACAC	111
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UUCAGAGAAUUGUAUCUGCACAGGGUGCAAUGAUACACACAC	105
*** ******* **** **** ***** ***** *****	105
	1.77
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AUACAUGUACACUUAUGUCUUUUGCUAUCUACCAGGAGGGGCGAGGAGUGAGU	171
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AGCAAGCGGGUGAAAGCUUUCUUCCUCACAAUCUCCCGGCGACGCGAAGAGCAGGAGGAG	231
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AGCAAGCGGGUGAAAGCUUUCUUCCUCACAAUCUCCCGGCGACGCGAAGAGCAGGAGGAG	231
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AGCAAGCGGGUGAAAGUUUUCUUCCUCACAGUCUCCCGGCGACGCGAAGAGCAGGAGAAU	238
AACAGGCGCGUGAAAGUUUUCUUCCUCACAGCUUCCCGGCGACGCGAAGAGCAGGAGAAA	225
* ** *** ****** ******** * ********* * *	
GGAGGUACGAUGCGAGAAGAAGAAGAAAAAAAAAAAAA <b>GAAAGAAGUAA</b>	286
GGAGGUACGAUGCGAGAAGAAGAAGAAGAAG-AAAAAAAA <b>GAAAGAAGUAA</b>	287
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GGAGGUACGAUGCGAGAAGAAGAAGAAGAAG-AAAAAAAA <b>GAAAGAAGUAA</b>	289
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GGAAGUACAAUGCGAGAAGAAGAAGAAGAAG-AAAGGAAAAAAAAAA	291
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GGAGGUACAAUGCGAGAAGAAGAAGAAGAAGAAAAAAAAA	289
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GGAAGUACAAUGCGAGAAGAAGAAGAAGAAGAAG-AAGGAAAAAAAAAA	295
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Sylvio\_X10/1 KU973678.1 KU973679.1 CLBrenerEL Esmeraldo Y CLBrener CLBrener Tula.C12

Sylvio\_X10/1 KU973678.1 KU973679.1 CLBrenerEL Esmeraldo CLBrenerNEL CLBrener Tula.Cl2 Marinkellei JR\_cl.\_4 Dm28c Sylvio X10/1 KU973678.1 KU973679.1 CLBrenerEL Esmeraldo CLBrenerNEL CLBrener Tula.C12 Marinkellei JR\_cl.\_4 Dm28c Sylvio X10/1 KU973678.1 KU973679.1 Esmeraldo CLBrenerNEL CLBrener Tula, C12 Marinkellei JR cl. 4 Dm28c Sylvio X10/1 KU973678.1 KU973679.1 CLBrenerEL Esmeraldo CLBrenerNEL Tula, Cl2 Marinkellei