

**Supplementary Material “*In silico* characterization of microRNAs-like sequences in the genome of *Paracoccidioides brasiliensis*”**

**Table S2 - Oligonucleotide sequences used in the present study**

<b>Genes</b>	<b>Forward sequences</b>	<b>Reverse sequence</b>
<i>dcr 1</i>	GGAGATTGAAGCTACTGAGAC	TCTGGCAGACACTATTTACAAC
<i>dcr 2</i>	GAGGGAGGCAACCAACTATC	TTAGAAACCACCTCGTCCTTG
<i>ago-1</i>	CCATGGCTGCAGTGTGAGTA	AAACACCATCGCGGAAGTAGT
<i>ago-2</i>	CGACTATTTTCAGACGCACATAT	GGGTTAAGCTTAGCATTGCC
<i>Act</i>	CGTCCTCGCCATCATGGTAT	TCTCCATATCATCCCAGTTCG
Super.:2.3:1128222-1128358(-)	GCAGTGTGCATGTGCAT	GGTCCAGTTTTTTTTTTTTTTTTCAC
Super.:2.5:587199-587317(+)	CGCAGAAATCACCTTCAC	GGTCCAGTTTTTTTTTTTTTTTGAAG
Super.:2.6:955763-955879(+)	GGCGGACGCGATG	GTCCAGTTTTTTTTTTTTTTTCCAC
Super.:2.12:21200-21275(-)	CAGCGCAGTAGGATTAGGAT	GGTCCAGTTTTTTTTTTTTTTTCT
Super.:2.14:171583-171670(+)	GGAGAGGGGGCCG	GGTCCAGTTTTTTTTTTTTTTTCAAC

(\*) all primers described in Table S2 were constructed for this work, except the actin primer.