

<b>Organism</b>	<b>Number of analyzed sequences</b>	<b>Number of predicted IDPs</b>	<b>Percent of IDPs with predicted function</b>	<b>Percent of hypothetical IDPs</b>
<i>L. braziliensis</i>	7965 (95.8%)	5725 (68.9%)	28.9%	70.9%
<i>L. major</i>	8171 (97.2%)	5959 (70.8%)	29.1%	70.6%
<i>L. infantum</i>	7835 (95.3%)	5797 (70.5%)	28.7%	71%
<i>L. mexicana</i>	7952 (100%)	5774 (72.6%)	29.5%	70.4%
<i>L. tarentolae</i>	7465 (100%)	5483 (73.4%)	26.9%	72.5%
<i>T. cruzi</i>	8178 (79.2%)	5565 (53.9%).	38.5%	61.5%
<i>T. brucei</i>	9472 (95.7%)	5510 (55.6%)	32%	68%