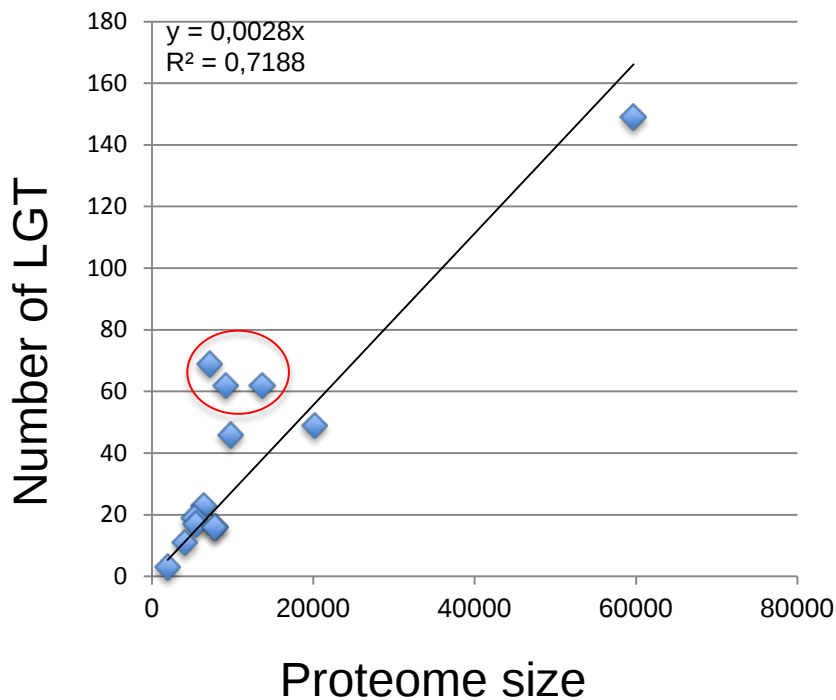


Additional File 9.



Taxa	No of annotated genes	LGT
<i>Trichomonas vaginalis</i>	59681	149
<i>Leishmania major</i>	7111	69
<i>Dictyostelium discoideum</i>	13605	62
<i>Entamoeba histolytica</i>	9090	62
<i>Trypanosoma cruzi</i>	20184	49
<i>Trypanosoma brucei</i>	9750	46
<i>Giardia lamblia</i>	6394	23
<i>Plasmodium falciparum</i>	5258	19
<i>Plasmodium vivax</i>	5393	17
<i>Plasmodium yoelii yoelii</i>	7813	16
<i>Toxoplasma gondii</i>	7793	16
<i>Cryptosporidium parvum</i>	4074	11
<i>Encephalitozoon cuniculi</i>	1918	3

Additional File 9. Numbers of LGTs detected compared to proteome size. The number of LGTs per individual annotated proteome was plotted against the proteome size. A linear regression forced over the point 0:0 and calculated for the 13 available points is shown. The three most obvious outliers are circled and highlighted in red in the table below which also contains all relevant numbers.