



**Supplementary Table S3.** Read alignment to the *Leishmania infantum* genome

Sample	Total reads	Total mapped reads	Mapping rate (%)
BCN R1	21413648	8182463	38.21
BCN R2	22326356	8500058	38.07
BCN R3	20144002	7761730	38.53
BOS R1	20096416	7566644	37.65
BOS R2	18457056	6840244	37.06
BOS R3	19497644	7480189	38.36