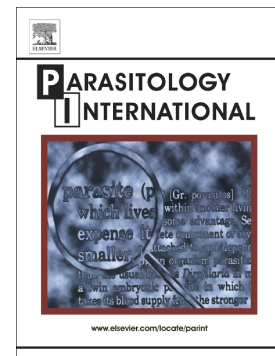


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RNA-seq analysis reveals differences in transcript abundance between cultured and sand fly-derived *Leishmania infantum* promastigotes

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Short communication

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