

CLONING OF *Babesia bigemina* AND *B. bovis* REPETITIVE DNA FOR SPECIES-SPECIFIC IDENTIFICATION OF THE PARASITES

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Genomic libraries of *Babesia bigemina* and *B. bovis* were constructed with the *Escherichia coli* EMBL<sub>4</sub> phage. <sup>32</sup>P-labelled parasite genomic DNAs were used as probes to screen the libraries, in high stringency and short time hybridization conditions. Several recombinant clones gave strong signals with the probes. Six *B. bigemina* clones, giving the strongest signals, were selected and upon back-hybridization all showed to have homologous repetitive DNA sequences. The same procedure was performed with *B. bovis* clones, with similar results. The cloned sequences do not cross-hybridize with host DNA and are species-specific. The probes can be used as alternative or complementary tools for the identification of two species of bovine *Babesia*.

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DISCOVERY OF A PUTATIVE 5.5 KBP DOUBLE-STRANDED RNA VIRUS PRESENT IN *Babesia bovis*

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*Babesia bovis* is an intraerythrocytic protozoa that causes a severe bovine disease. Nucleic acids from two *B. bovis* isolates presented in agarose gel electrophoresis, besides genomic DNA, a molecule of approximately 5.5 kbp characterized as double-stranded RNA. Sonicated *B. bovis* cells were centrifuged in a CsCl buoyant-density gradient and a fraction containing the double-stranded RNA was examined by electron microscopy. Virus-like particles were observed, which might be related to the 5.5 kbp RNA.

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