

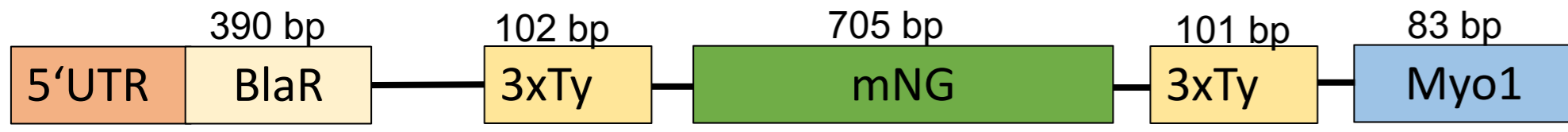
Characterisation of the two myosins of *T. brucei*

Sisco Jung*, Xenia Malzer*, Fabian Link, Mara Pöllmann, Monika Wieland, Antonia Konle, Brooke Morriswood

Department of Cell and Developmental Biology, Biocentre, University of Würzburg, Germany

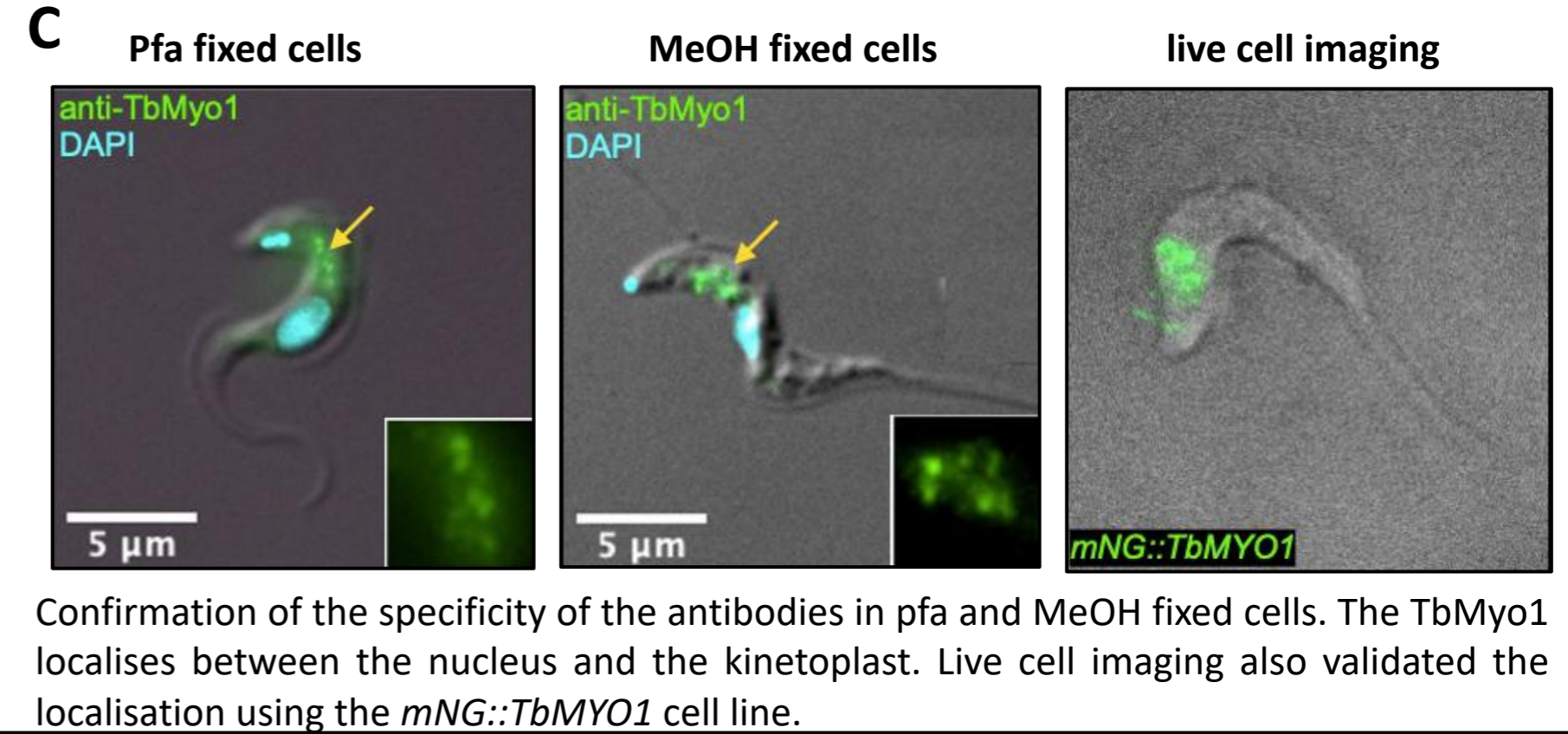
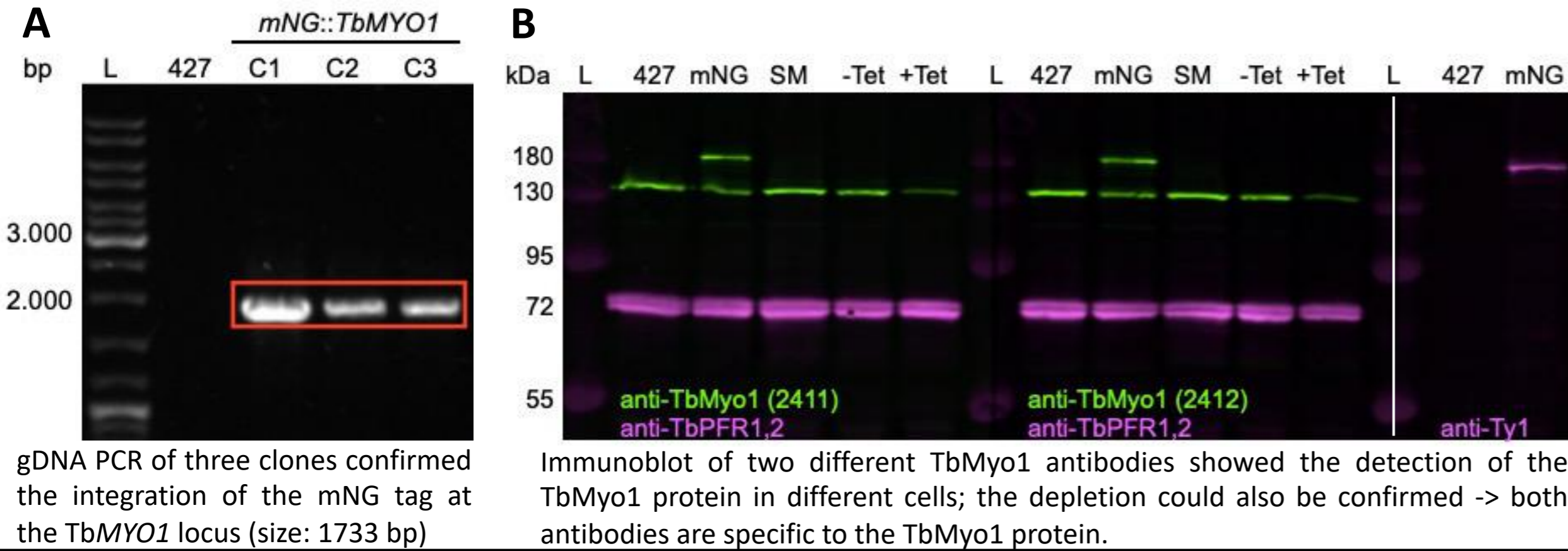
Abstract: Trypanosomes have a very reduced actomyosin system, and *Trypanosoma brucei* expresses only a single actin gene and two myosin genes. The retention of this rudimentary actomyosin system alongside the trypanosomes' massive investment in their microtubule system suggests that the actomyosin system performs certain functions that the microtubule cytoskeleton cannot recapitulate. Both of the *T. brucei* myosins have received little attention to date. The data here represent the first steps in a full characterisation of these two proteins in bloodstream form *T. brucei*.

1. Generation of an *mNG::TbMYO1* cell line

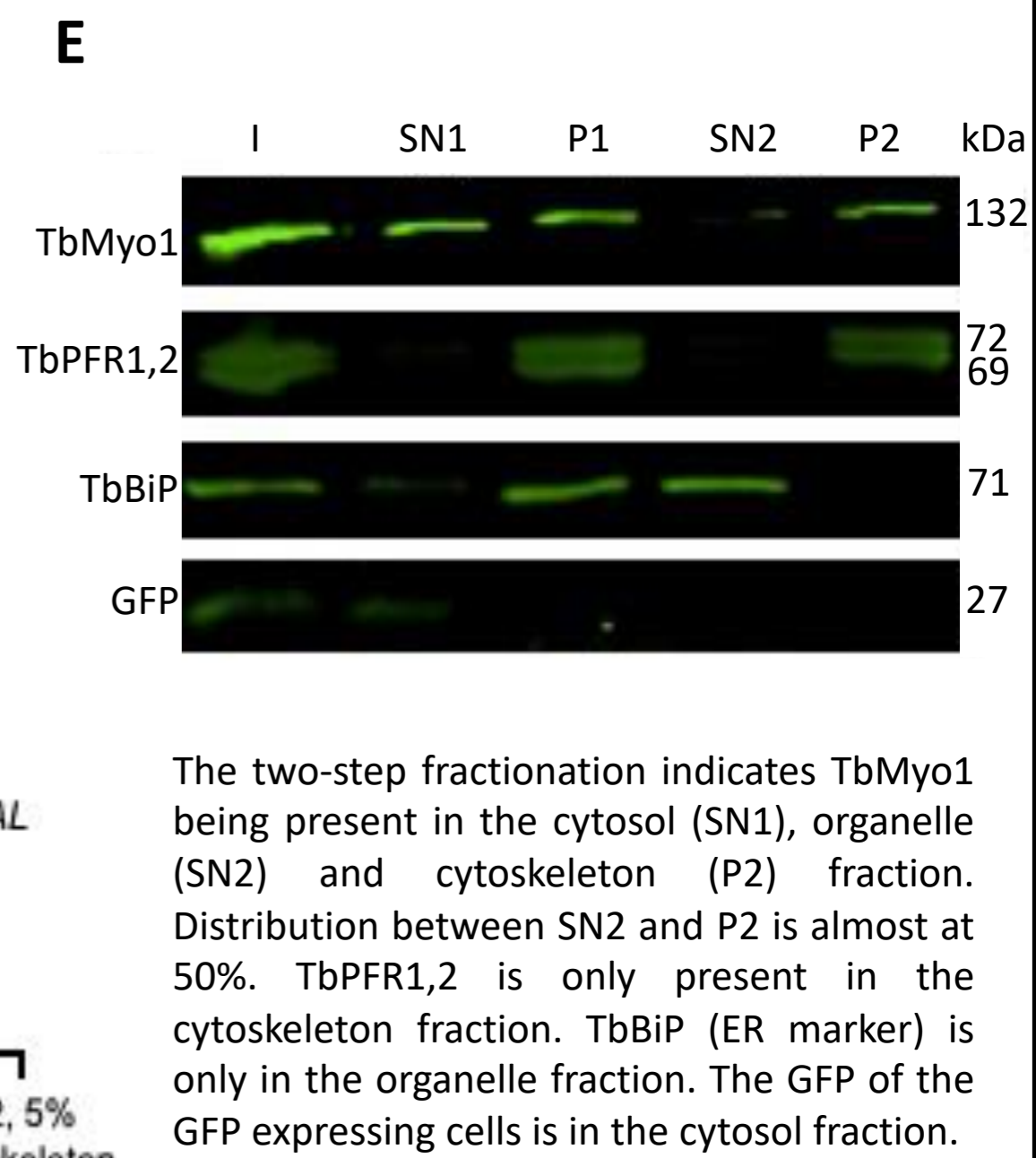
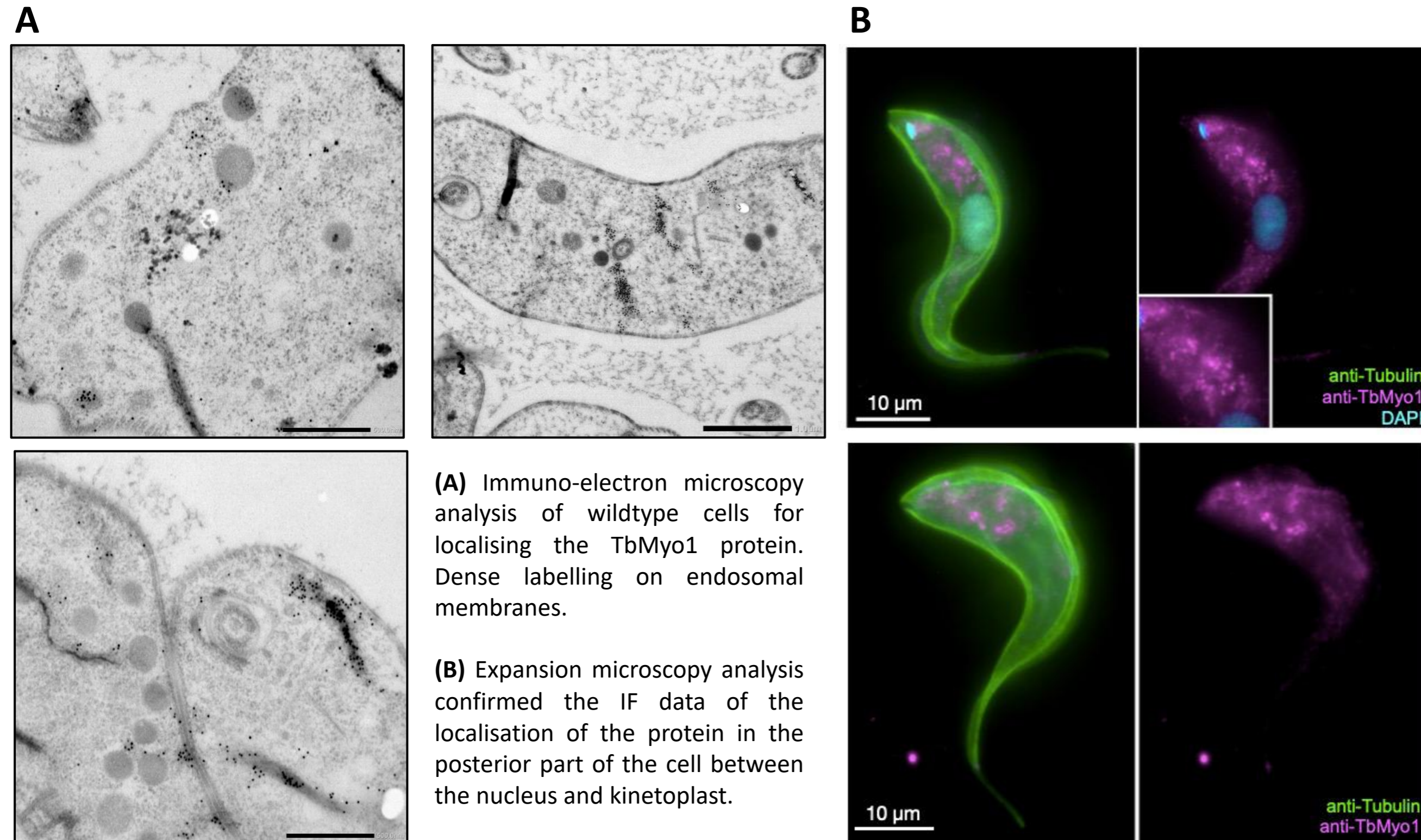


Schematic view of the *TbMyo1* in situ tagging construct. The wildtype strain 427 was transfected with the PCR tagging product. The tagging construct encoded a blasticidin resistance gene, a 3xTy1 tag, the mNG gene followed by a 3xTy1 tag.

2. Validation of the *TbMyo1* antibodies and localisation of the protein



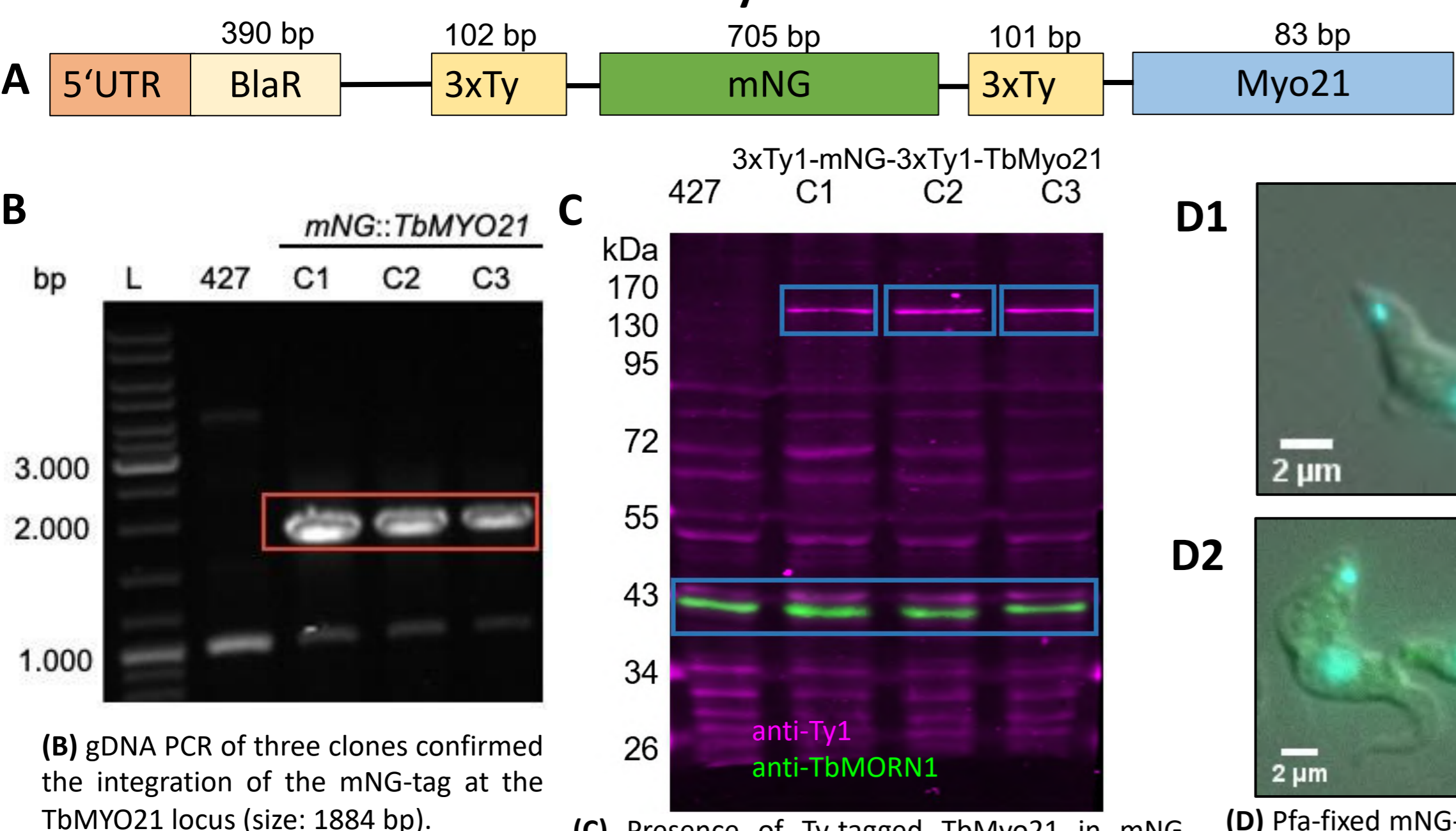
3. Ultrastructural analysis and biochemical localisation of the *TbMyo1* protein



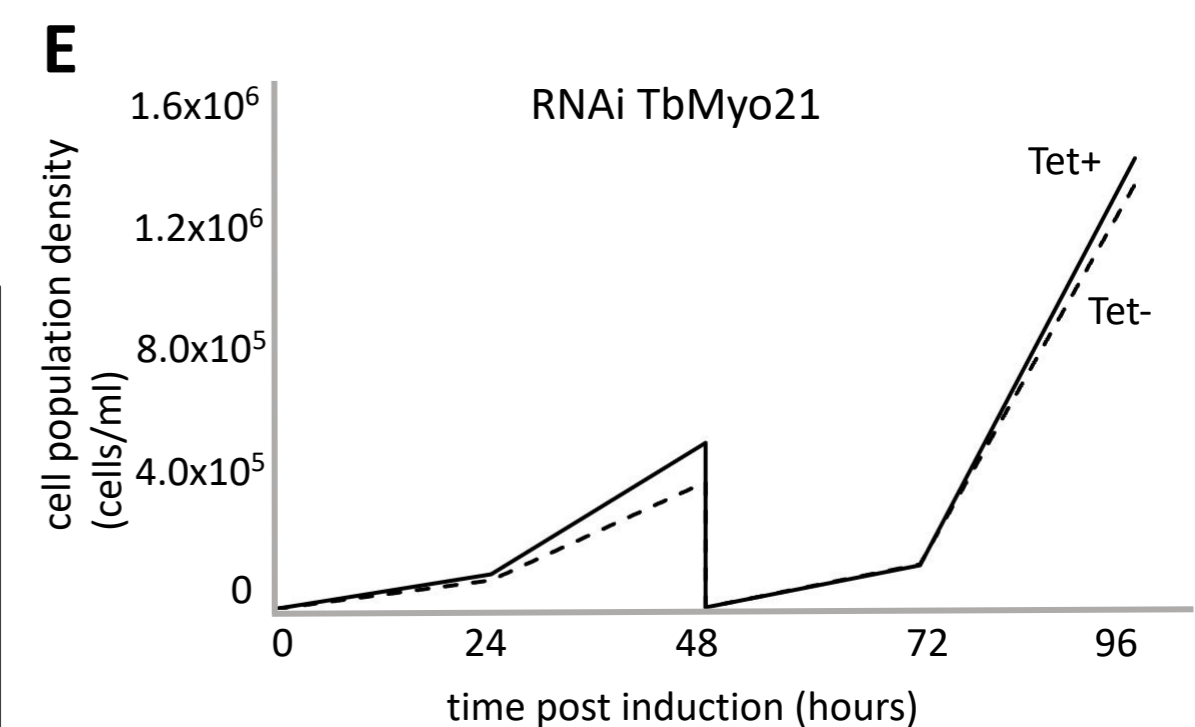
The two-step fractionation indicates *TbMyo1* being present in the cytosol (SN1), organelle (SN2) and cytoskeleton (P2) fraction. Distribution between SN2 and P2 is almost at 50%. *TbPFR1,2* is only present in the cytosol fraction. *TbBiP* (ER marker) is only in the organelle fraction. The GFP expressing cells is in the cytosol fraction.

Schematic view of the two-step fractionation.

4. Validation and localisation of *TbMyo21*



(A) Schematic view of the *TbMyo21* in situ tagging construct. The wildtype strain 427 was transfected with the shown PCR tagging product.



(E) The depletion of *TbMyo21* protein did not influence the cell viability of bloodstream form *T. brucei*, but had a small impact on the growth of the induced cells.

Outlook for *TbMyo21* localisation:

- Immunofluorescence of *TbMyo21* with anti-Myo21 antibodies
- Reverse transcription PCR of *TbMyo21*

5. Summary

TbMyo1 was localised to the posterior part of bloodstream form *T. brucei* cells using two different antibodies and an mNG tag. Ultrastructural analysis using immuno-electron microscopy and expansion microscopy suggested that *TbMyo1* is present on endosomes and vesicles. Biochemical fractionation indicated that there are cytosolic, organelle-associated and cytoskeleton-associated pools of *TbMyo1*. The expression level of *TbMyo21* in bloodstream form *T. brucei* appears to be much lower than that of *TbMyo1*. The exact localisation of *TbMyo21* remains unclear. Depletion of *TbMyo21* has a mild effect on population growth.