

Use of mobile genetic element PCR (MGE-PCR) as a tool to investigate the distribution and spread of parasitic diseases

> Professor Geoff Hide University of Salford, UK

Overview MGE-PCR: What is Mobile Genetic Element PCR? Examples: Spread of human sleeping sickness in Uganda Strain typing in Toxoplasma

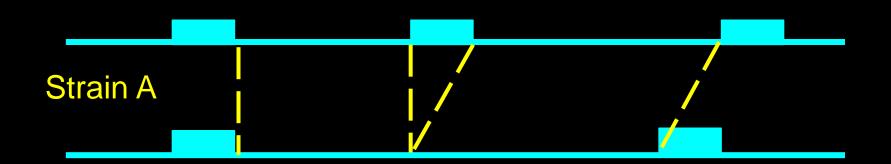
- Examples
- RAPDs
- RFLP
- Microsatellite genotyping (MS)

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- RAPDs
- RFLP
- Microsatellite genotyping (MS)
- All have specific uses

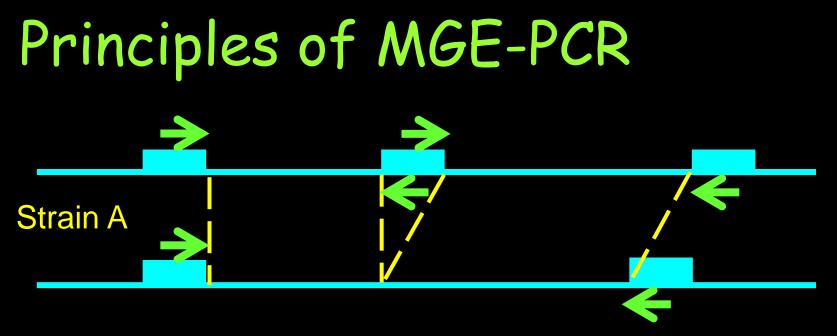
- limitations
- Most techniaues require several stages

- Aim
- To develop a single stage analytical tool that provides good discrimination between parasite strains

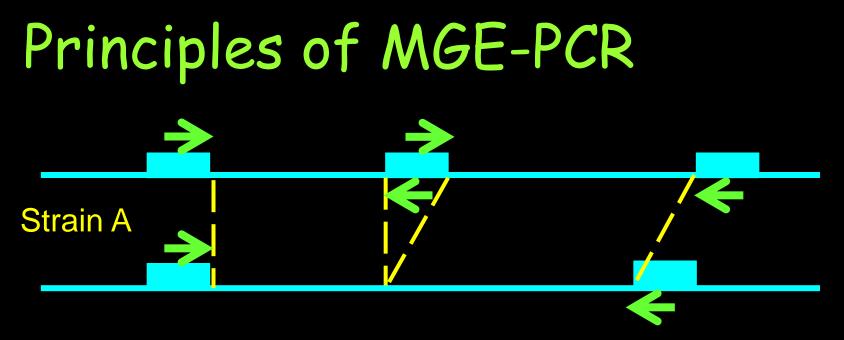
Principles of MGE-PCR



Strain B

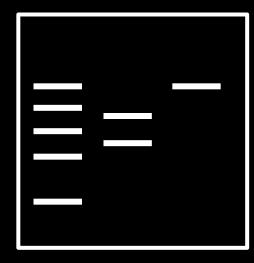


Strain **B**

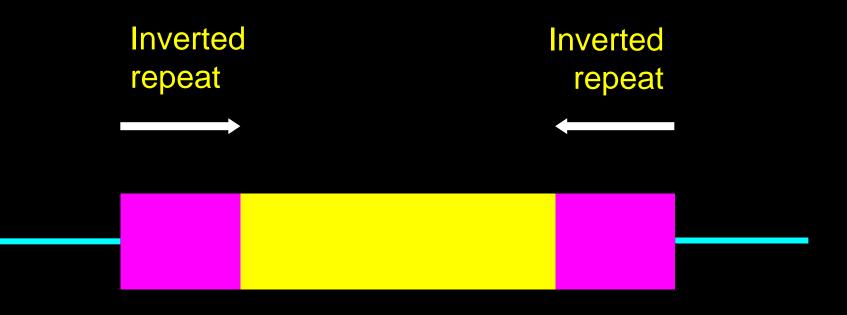


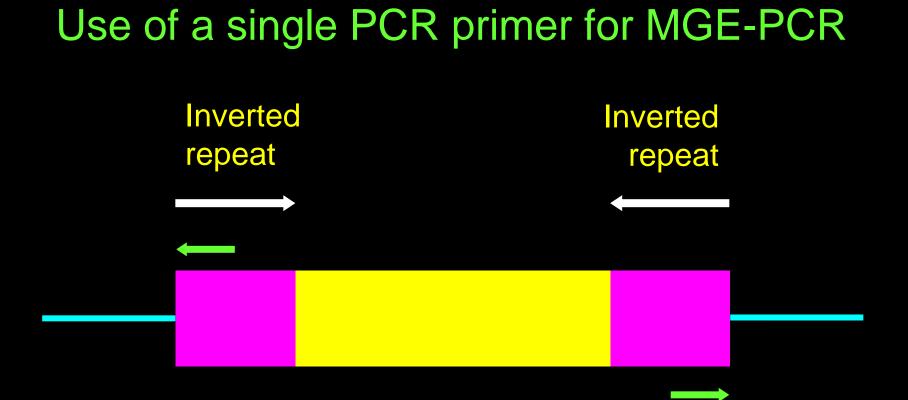
Strain **B**





Structure of a typical MGE





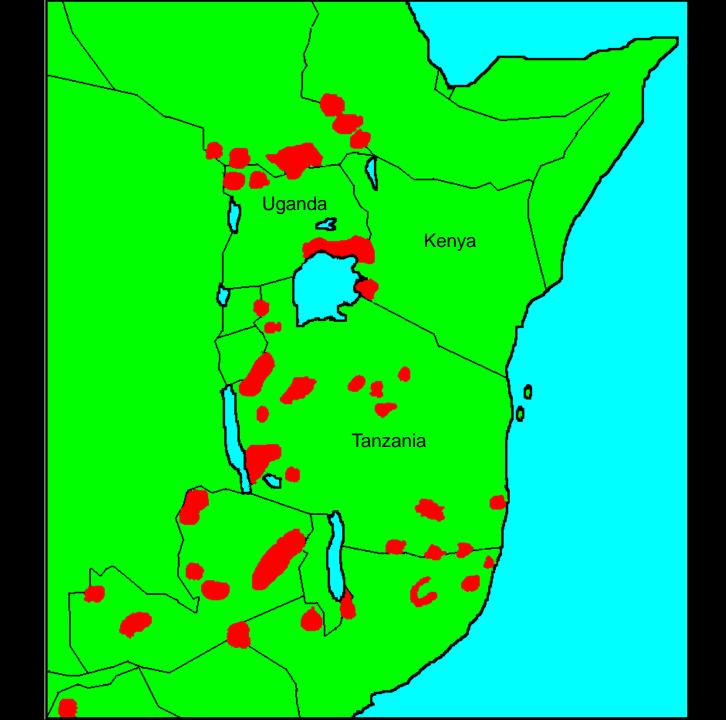
A single PCR primer is complementary to both inverted repeats Examples

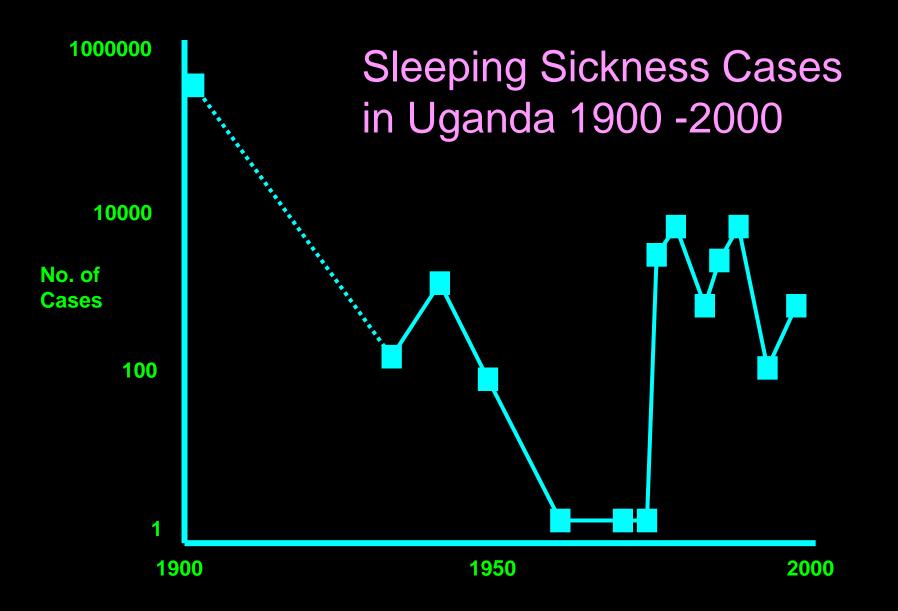
Examples

The origins of Human sleeping sickness epidemics

Human sleeping sickness in Uganda

Why are sleeping sickness foci where they are? What causes epidemics?



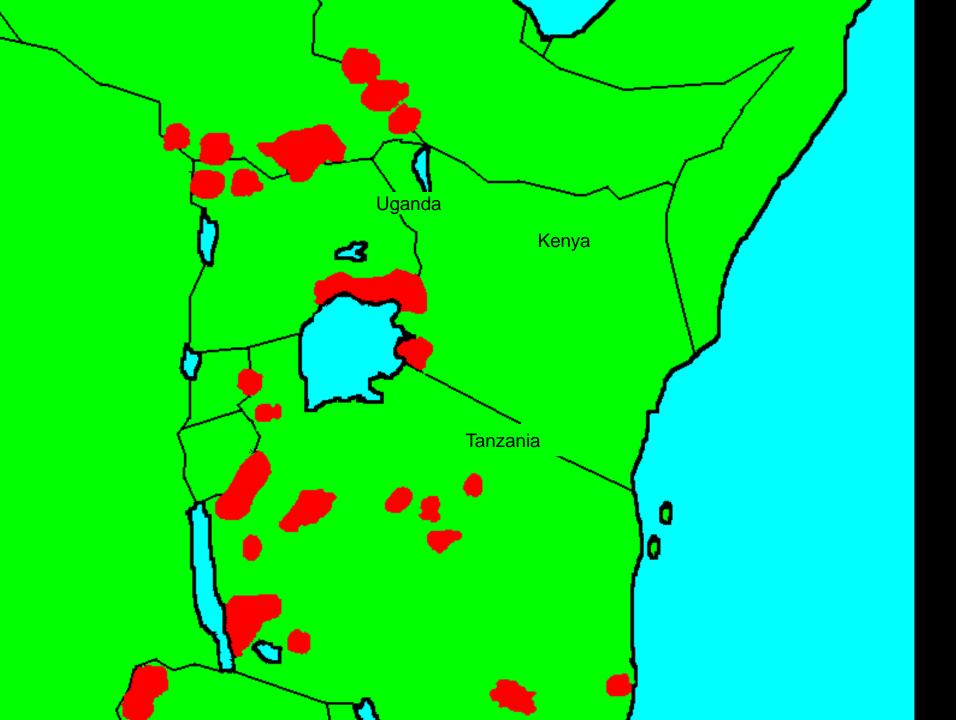


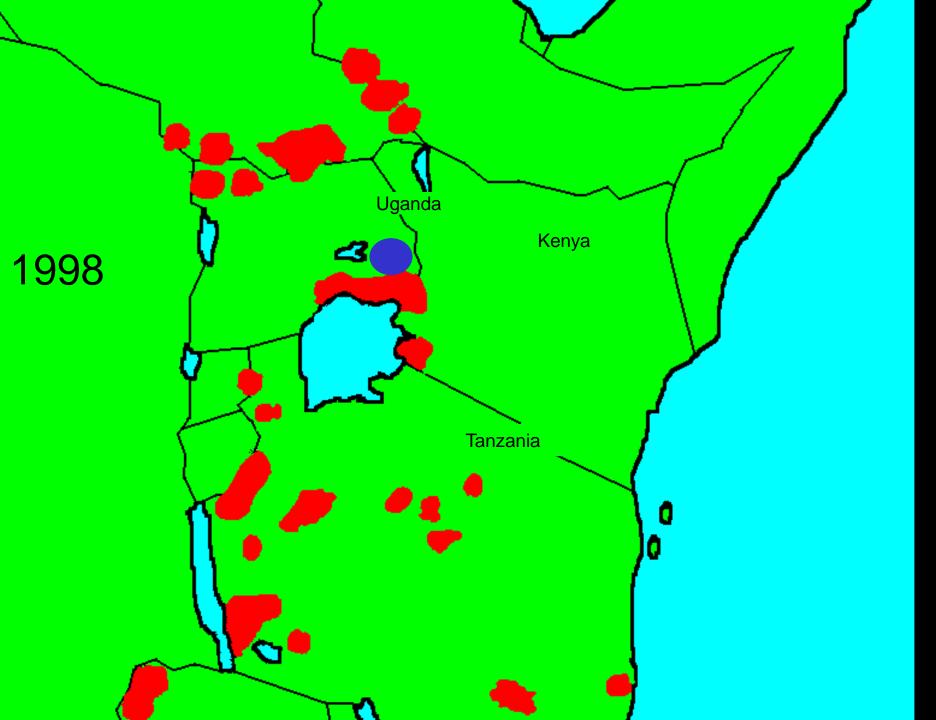
Patterns of Human Disease

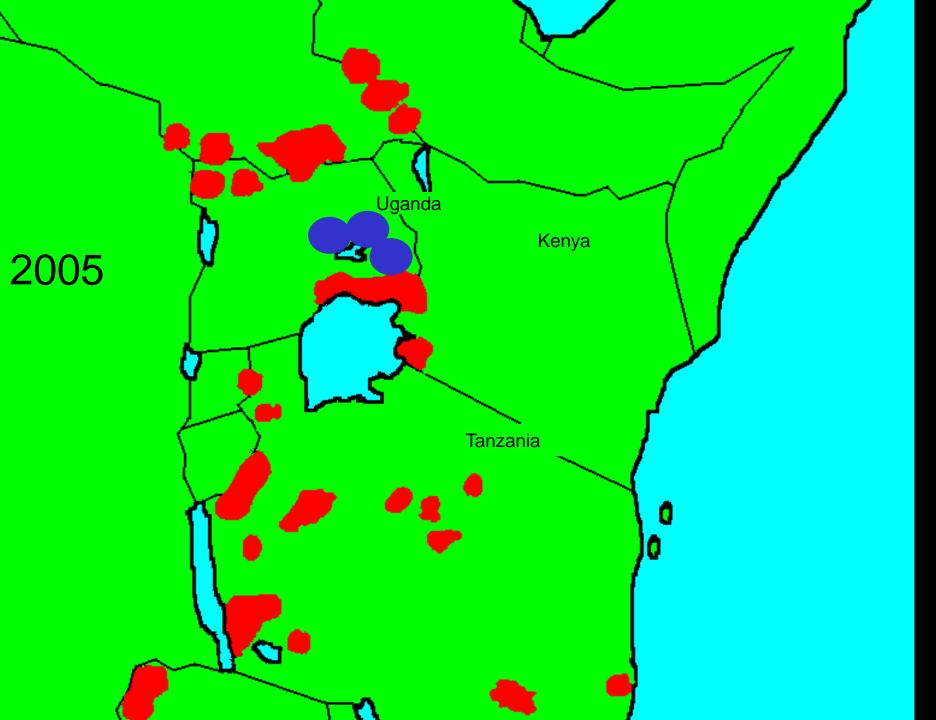
- Animal disease throughout
- Human disease localised to specific foci
- Human disease occurs as sporadic epidemics

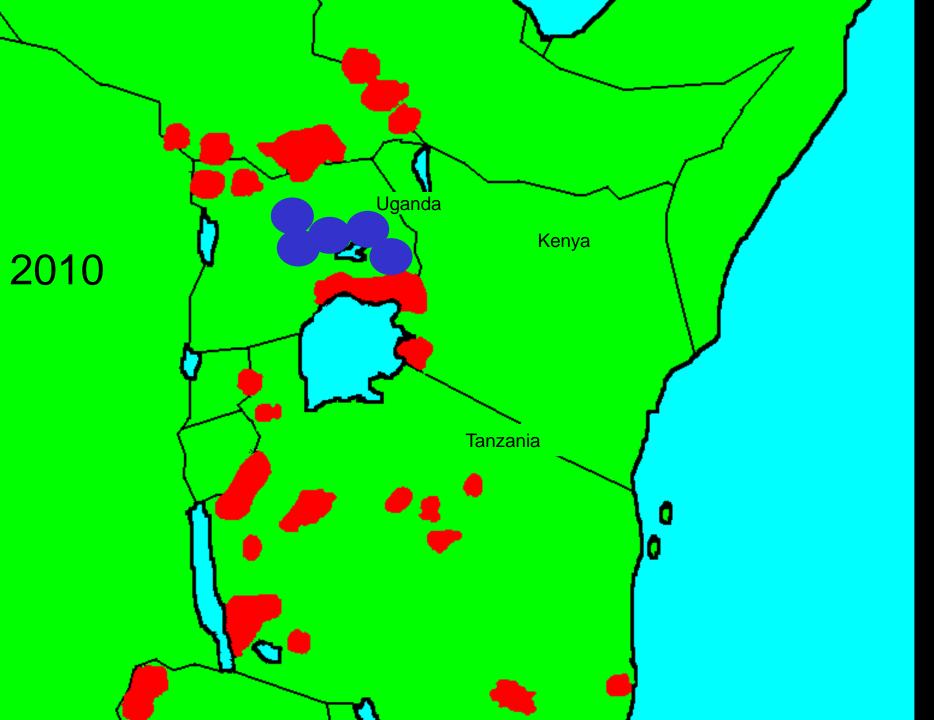
Patterns of Human Disease in Uganda

- Human focus is spreading
- North and West
- Merge with a *T.b.gambiense* focus??



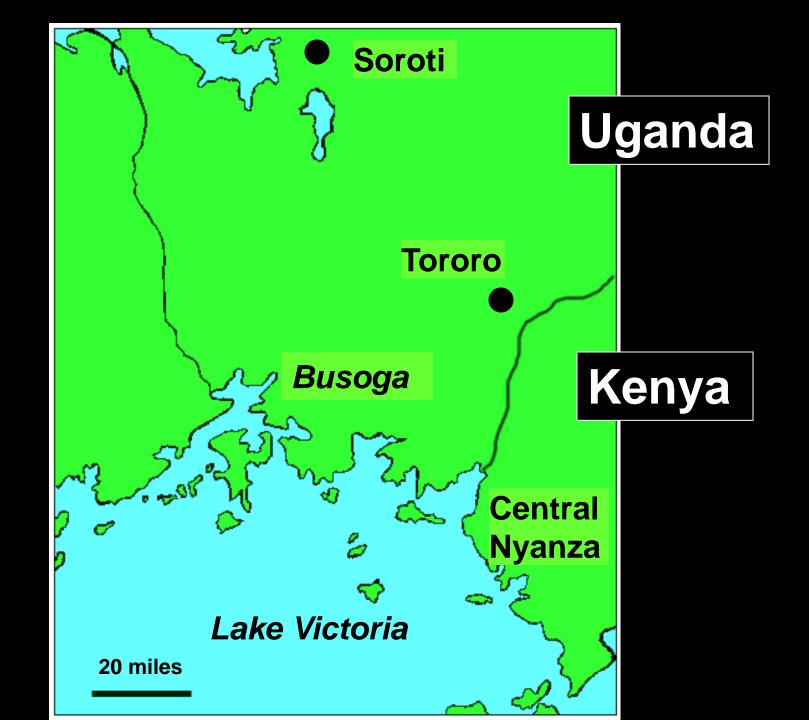






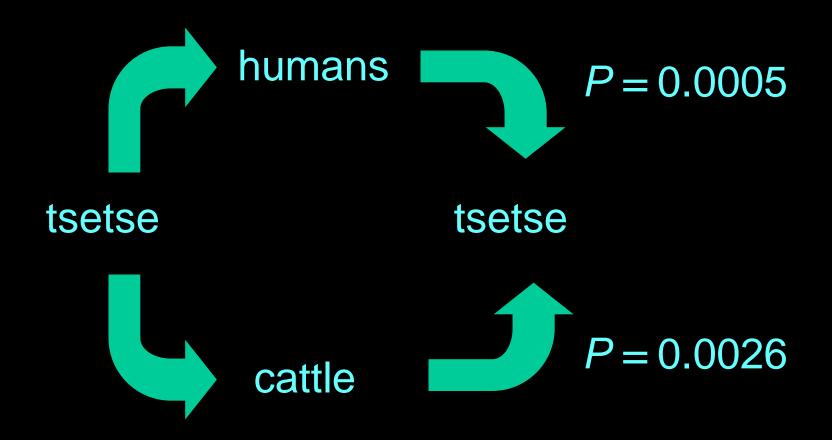
Recent Epidemics in Uganda

1988 - 1992 - Tororo, Uganda 200-300 cases per year 1998 to date - Soroti, Uganda 119 cases in first 18 months



Tororo Epidemic

- No cases prior to 1984
- Epidemic Peaked 1990
- Still a few cases

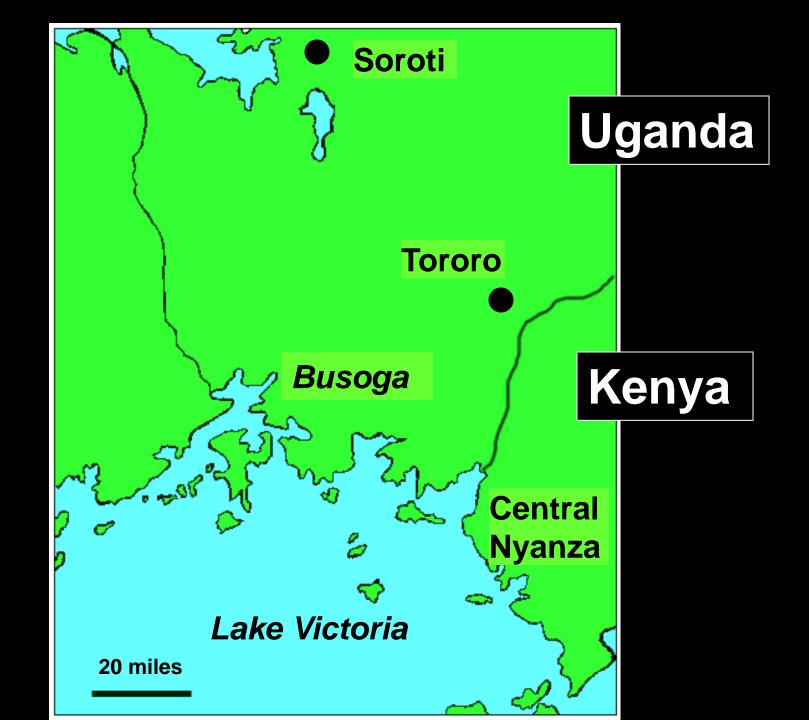


Main findings

- A Specific human strain
- Presence of human strain may determine extent of focus
- Cattle are a significant reservoir for the human strain

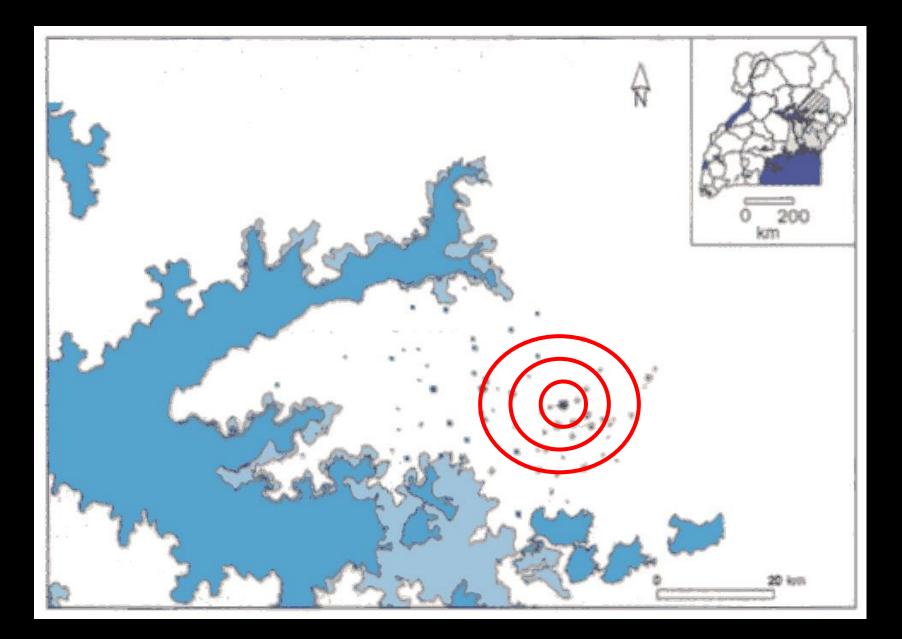
1998 Soroti Epidemic

- Never had sleeping sickness
- 31st Dec 1998 local nurse reported first case
- 119 cases by June 2000
- Recent movement of cattle from into the area



Location of disease associated with local cattle markets





Where did the Soroti epidemic come from?

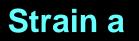
- Need a method of accurately identifying strains
- MGE-PCR as a tool

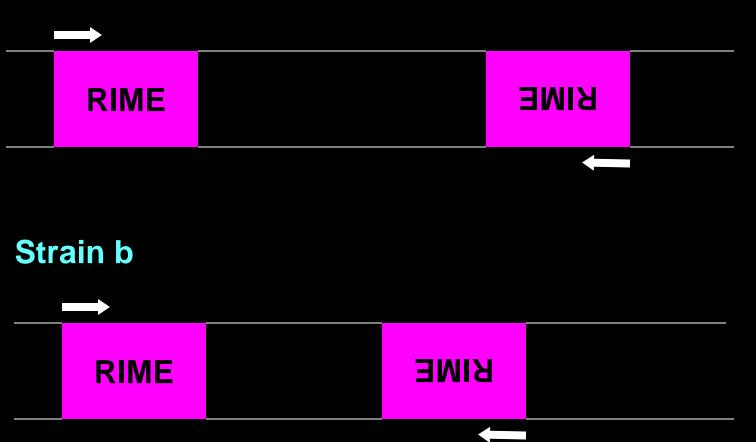
The trypanosomal mobile genetic element RIME

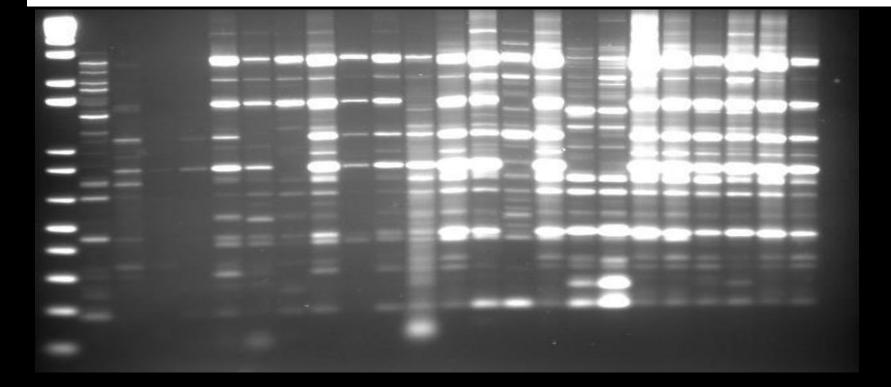
(b) RIME + Ingi MGE



Analysis variation in RIME position between trypanosome strains - single primer







Marker a Bumanda 14

a Bumanda 146

a UGL

c LIRI 16

c AKO C20

b KINU C21

b KINU P4

b BUWHTK

b BUG H1

b BUG H2

b KINU HBW

c LIRI 014

a UGK

a Mela 27

a Mela 32

a Mela 71

a Mela Pig1

a Mela Pig 2

a UGB 88

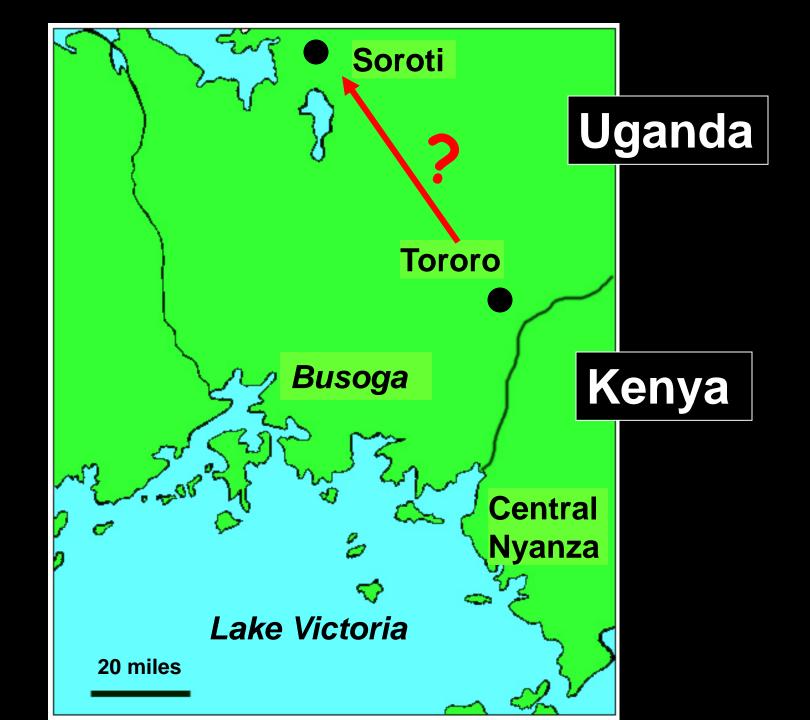
a UGC 88

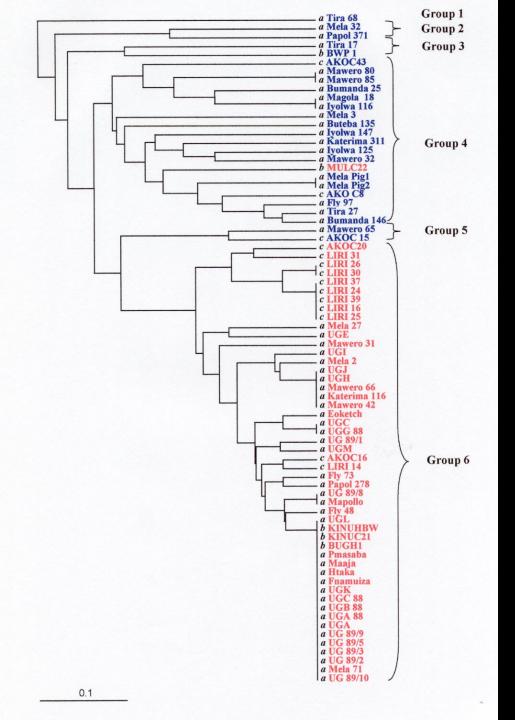
a UGE a UGE

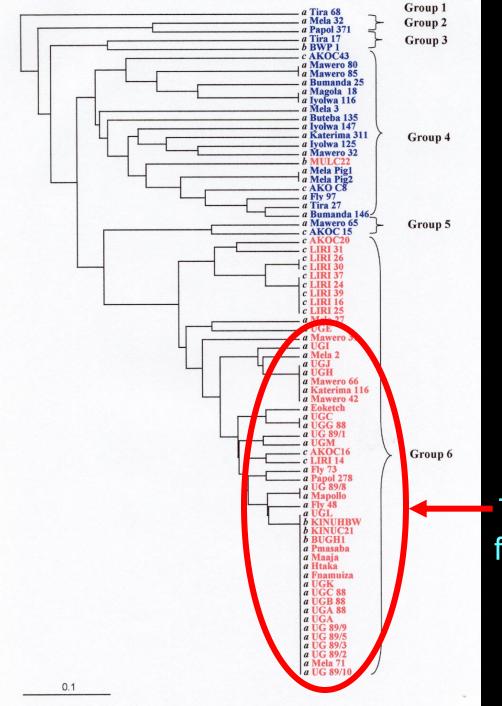
a UGG 88

a UGM

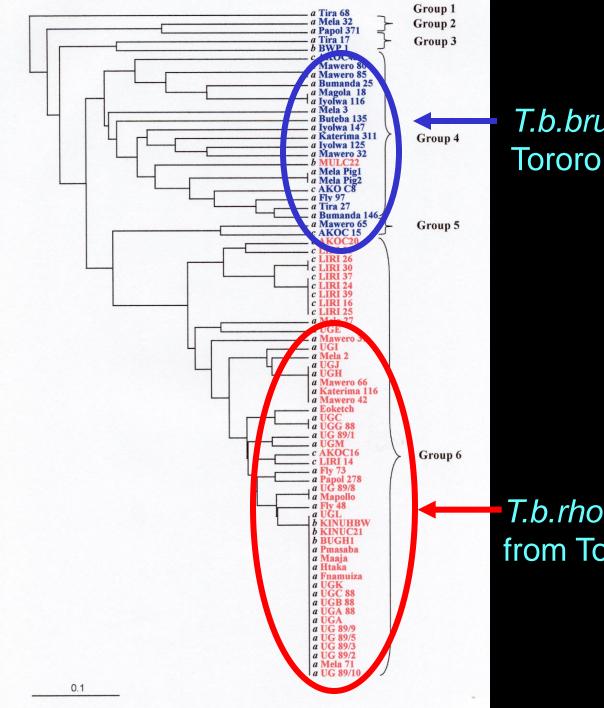
Water





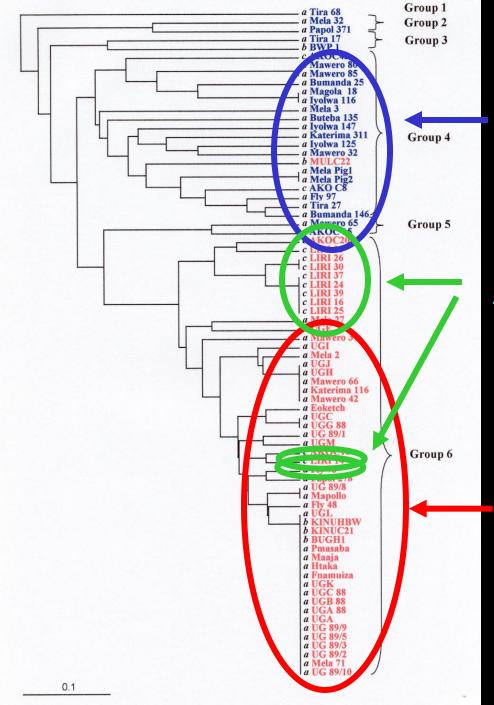


T.b.rhodesiense from Tororo



T.b.brucei from Tororo

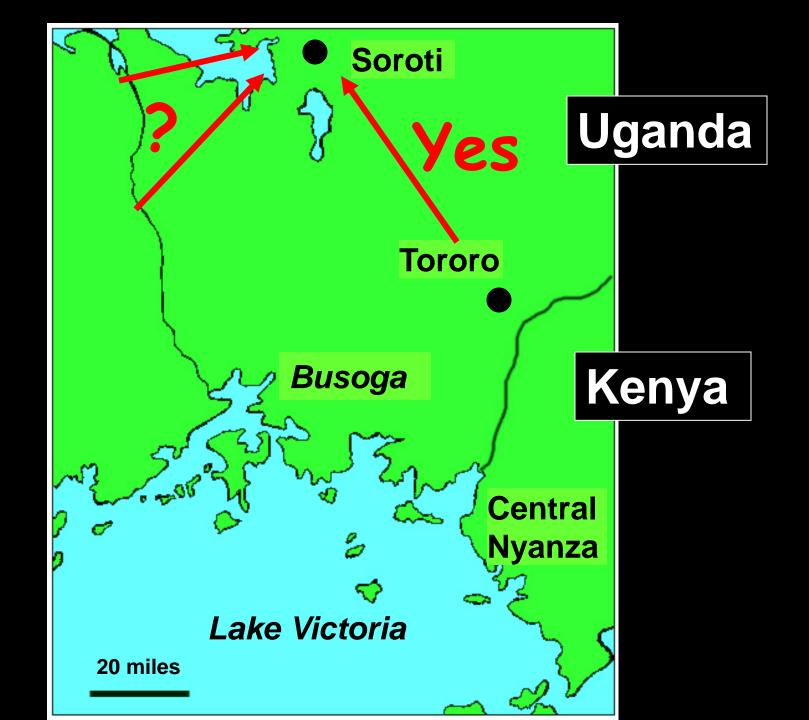
T.b.rhodesiense from Tororo



T.b.brucei from Tororo

T.b.rhodesiense from Soroti

T.b.rhodesiense from Tororo



Conclusions

- Two *T.b.rhodesiense* isolates in Soroti from Tororo (18%)
- Others from another source
- MGE-PCR a useful tool for tracking

Conclusions

- The origins of the Soroti epidemic may be complex
- Cattle important
- Need to identify risk areas and cattle movement

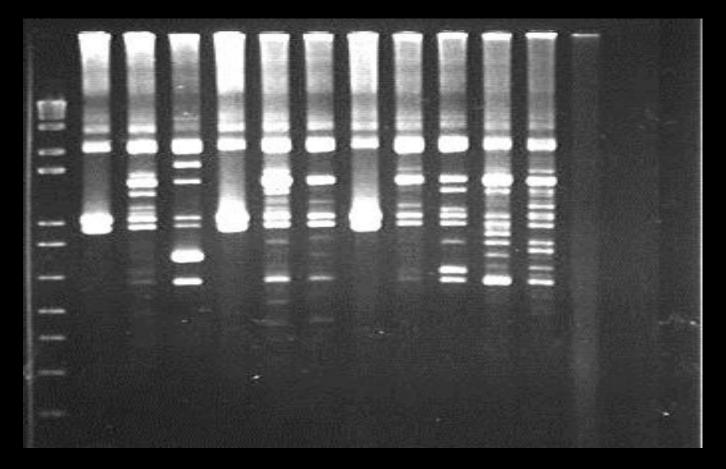
Examples

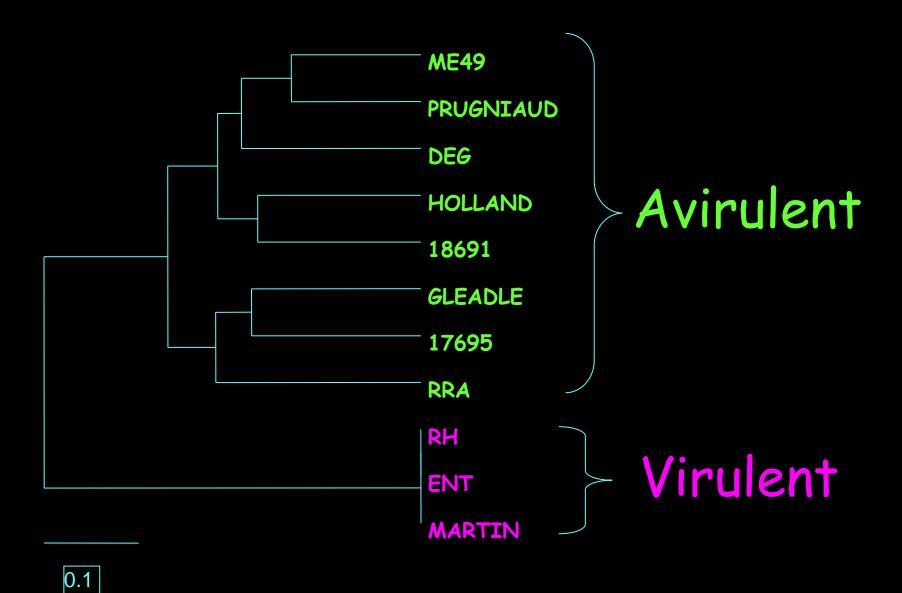
Markers for strain typing in *Toxoplasma gondii*

Markers for strain typing in *Toxoplasma gondii*

Traditional types:Type Ivirulent to miceType IIavirulent to miceType IIIavirulent to mice

1 2 2 1 2 2 1 2 2 2 2





Conclusions

 MGE is a useful tool for tracking parasite strains and understanding epidemiology Salford Prof Geoff Hide Dr Aimee Tilley Dr Rebecca Terry Dr Andy Cox Dr Jackie Hughes Edinburgh Prof Sue Welburn Dr Aimee Tilley Dr Eric Fevre Dr Kim Picozzi Prof Ian Maudlin

Uganda Dr Martin Odiit and colleagues