

Swiss TPH



Swiss Tropical and Public Health Institute
Schweizerisches Tropen- und Public Health-Institut
Institut Tropical et de Santé Publique Suisse

Pascal Mäser

Parasite Chemotherapy Unit
Medical Parasitology and Infection Biology

Academic research case No. 3

African trypanosomes

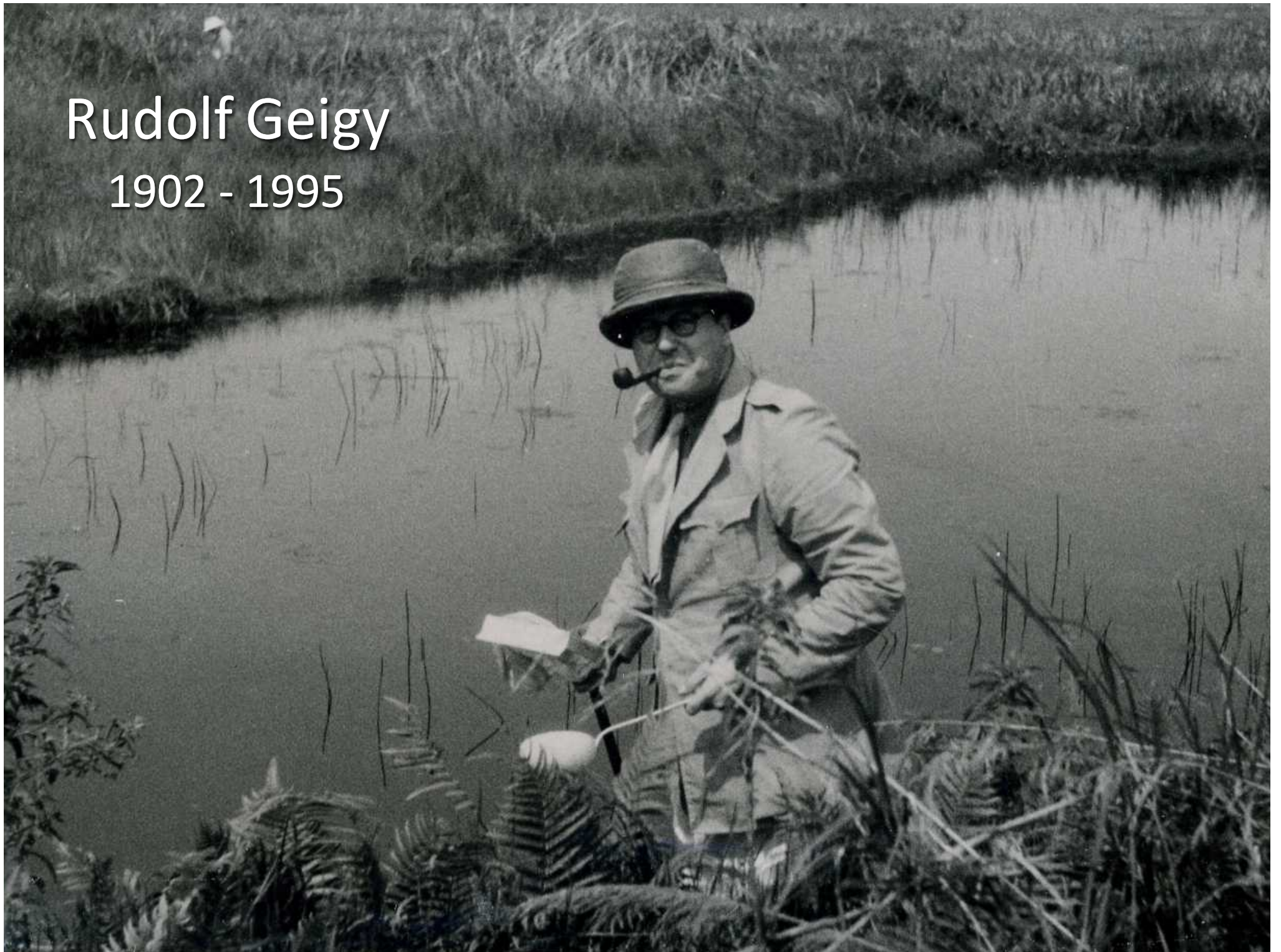




1943

Rudolf Geigy

1902 - 1995

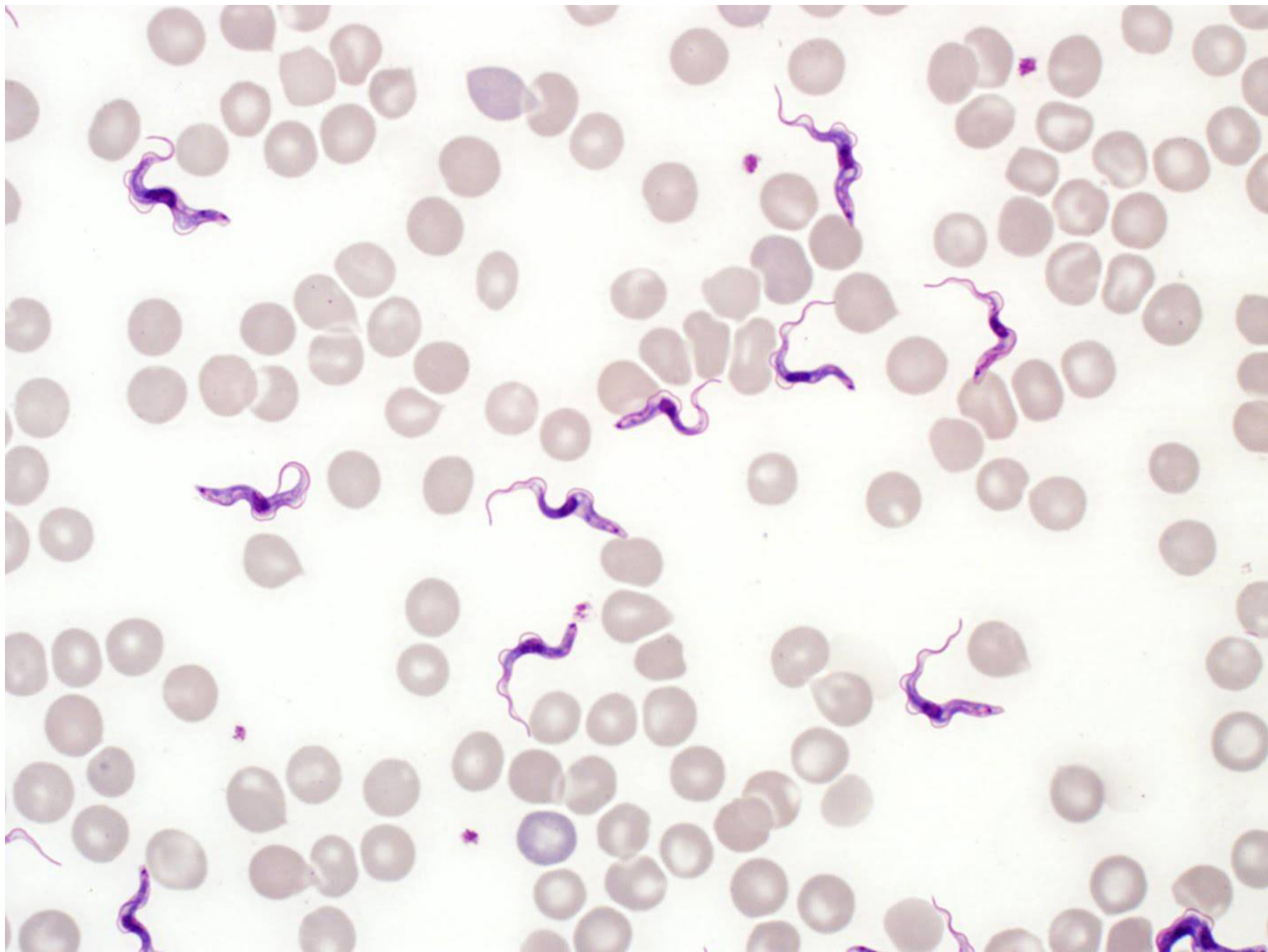


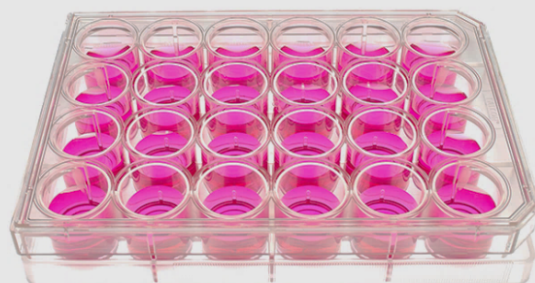
Sleeping Sickness Survey in the Serengeti Area 1971

R. Geigy & M. Kauffmann

Species	No. examined	No. infected
<i>Panthera leo</i>	43	39
<i>Crocuta crocuta</i>	31	23
<i>Kobus defassa</i>	1	1
<i>Alcelaphus buselaphus</i>	20	11









Susceptibility of Ugandan *Trypanosoma brucei* isolated from man and animal to melarsoprol, isometamidium and melarsoprol

Enoch Matovu^{1,2}, Monika Iten², John C. K. E. B. and Ronald Kaminsky²

¹ Livestock Health Research Institute, Tororo, Uganda

² Swiss Tropical Institute, Basel, Switzerland

³ Faculty of Veterinary Medicine, Makerere University

Summary

Thirty-six *Trypanosoma brucei* isolates from south-east Uganda were studied for sensitivity to melarsoprol *in vitro*. All stocks were sensitive to melarsoprol. In contrast, the remaining stocks were resistant. Isometamidium were required for the treatment of the remaining stocks. 0.01–0.20 ng/ml isometamidium showed reduced susceptibility. The results suggest that the population. Control of sleeping sickness is therefore, face serious problems. The disease is likely not be eliminated by re-

Comparative Genomics Reveals Multiple Genetic Backgrounds of Human Pathogenicity in the *Trypanosoma brucei* Complex

Mark Sistrom^{1,*}, Benjamin Evans¹, Robert Bjornson², Wendy Gibson³, Oliver Balmer^{4,5}, Pascal Mäser⁴, Serap Aksoy⁶, and Adalgisa Caccone¹

¹Department of Ecology and Evolutionary Biology, Yale University

²Department of Computer Science, Yale University

³School of Biological Sciences, University of Bristol, United Kingdom

⁴Swiss Tropical and Public Health Institute, Basel, Switzerland

⁵Zoological Institute, University of Basel, Switzerland

⁶Department of Epidemiology of Microbial Diseases, Yale School of Public Health, New Haven, CT

*Corresponding author: E-mail: mark.sistrom@yale.edu.

Accepted: September 30, 2014

Data deposition: This project has been deposited at NCBI under the accession 705162.

Abstract

The *Trypanosoma brucei* complex contains a number of subspecies with exceptionally variable life histories, including zoonotic subspecies, which are causative agents of human African trypanosomiasis (HAT) in sub-Saharan Africa. Paradoxically, genomic variation between taxa is extremely low. We analyzed the whole-genome sequences of 39 isolates across the *T. brucei* complex from diverse hosts and regions, identifying 608,501 single nucleotide polymorphisms that represent 2.33% of the nuclear genome. We show that human pathogenicity occurs across a wide range of parasite genotypes, and taxonomic designation does not reflect genetic variation across the group, as previous studies have suggested based on a small number of genes. This genome-wide study allowed the identification of significant host and geographic location associations. Strong purifying selection was detected in genomic regions associated with cytoskeleton structure, and regulatory genes associated with antigenic variation, suggesting conservation of these regions in African trypanosomes. In agreement with expectations drawn from meiotic reciprocal recombination, differences in average linkage disequilibrium between chromosomes in *T. brucei* correlate positively with chromosome size. In addition to insights into the life history of a diverse group of eukaryotic parasites, the documentation of genomic variation across the *T. brucei* complex and its association with specific hosts and geographic localities will aid in the development of comprehensive monitoring tools crucial to the proposed elimination of HAT by 2020, and on a shorter term, for monitoring the feared merger between the two human infective parasites, *T. brucei rhodesiense* and *T. b. gambiense*, in northern Uganda.



2nd edition

A Guide for Transboundary Research Partnerships

11 Principles

Swiss Commission for Research Partnerships
with Developing Countries (KFPE)

sc | nat 

Swiss Academy of Sciences
Akademie der Naturwissenschaften
Accademia di scienze naturali
Académie des sciences naturelles



1. Set the agenda together
2. Interact with stakeholders
3. Clarify responsibilities
4. Account to beneficiaries
5. Promote mutual learning
6. Enhance capacities
7. Share data and networks
8. Disseminate results
9. Pool profits and merits
10. Apply results
11. Secure outcomes

1. Set the agenda together
2. Interact with stakeholders
3. Clarify responsibilities
4. Account to beneficiaries
5. Promote mutual learning
6. Enhance capacities
7. Share data and networks
8. Disseminate results
9. **Pool profits and merits**
10. Apply results
11. Secure outcomes

Researchers cannot sell their results as a farmer would sell potatoes on the market.

Most research partnerships produce knowledge as a public good.

However, the situation becomes more difficult in cases where several of the parties involved lay claim to the same piece of the cake.

When property rights or patent rights are at stake, it is essential to set clear rules early on.