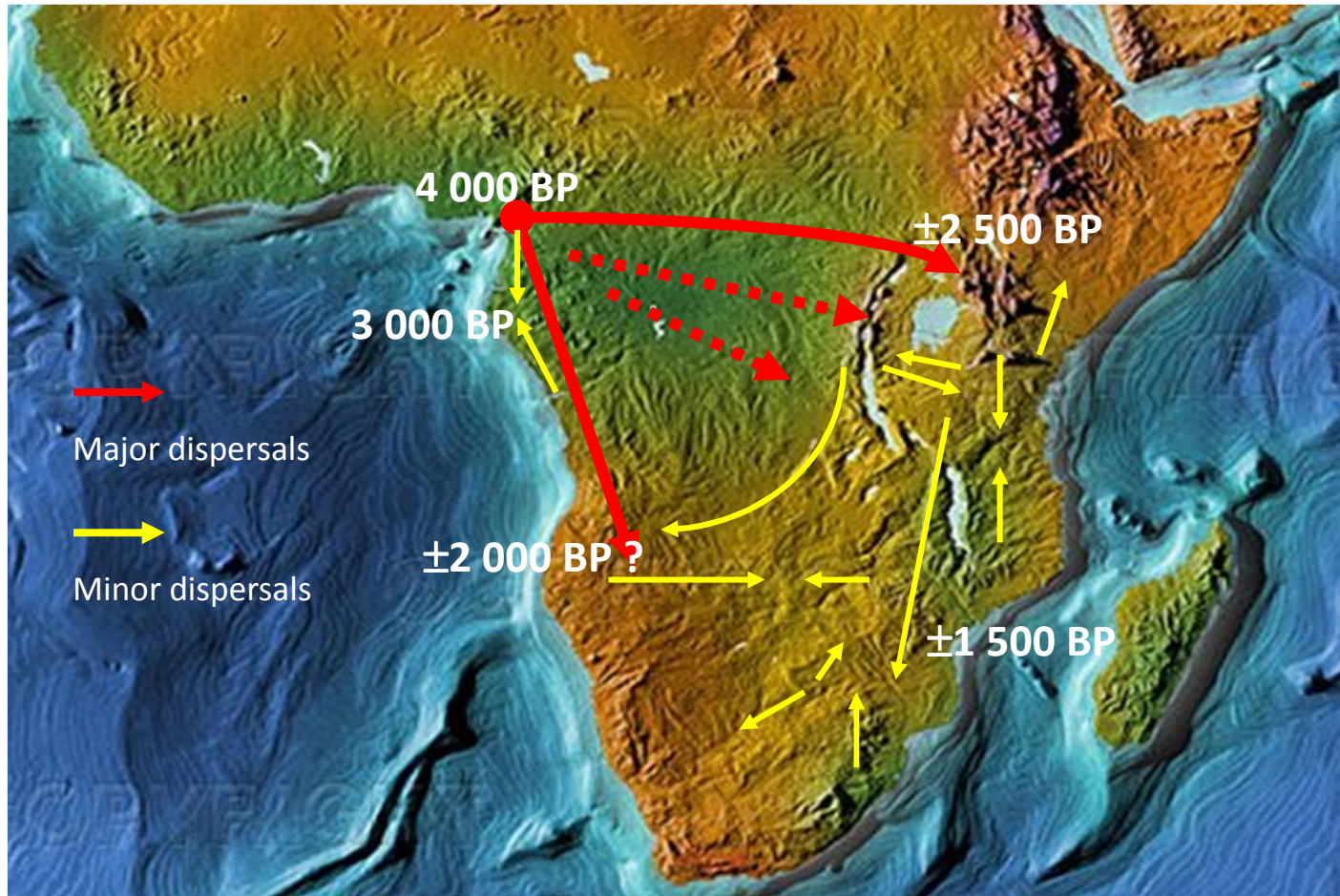


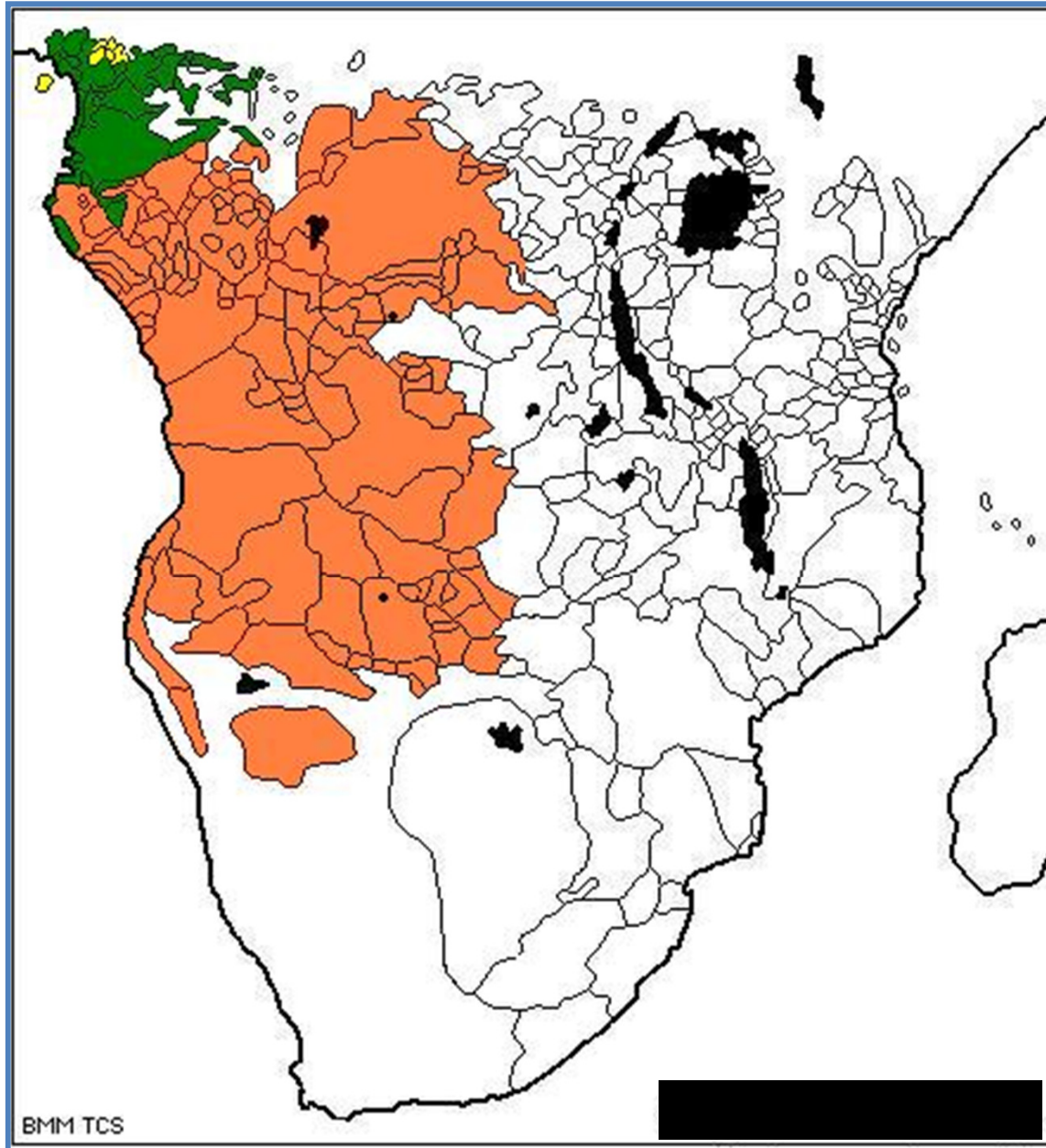
Linguistic implications of contacts between (agriculturalists) Bantu and Hunter-Gatherers

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Dynamique du Langage, Lyon, France
Jean-Marie.Hombert@univ-lyon2.fr

Methodology in Linguistics Prehistory Workshop,
Berlin, October 15-16, 2011

Bantu migrations

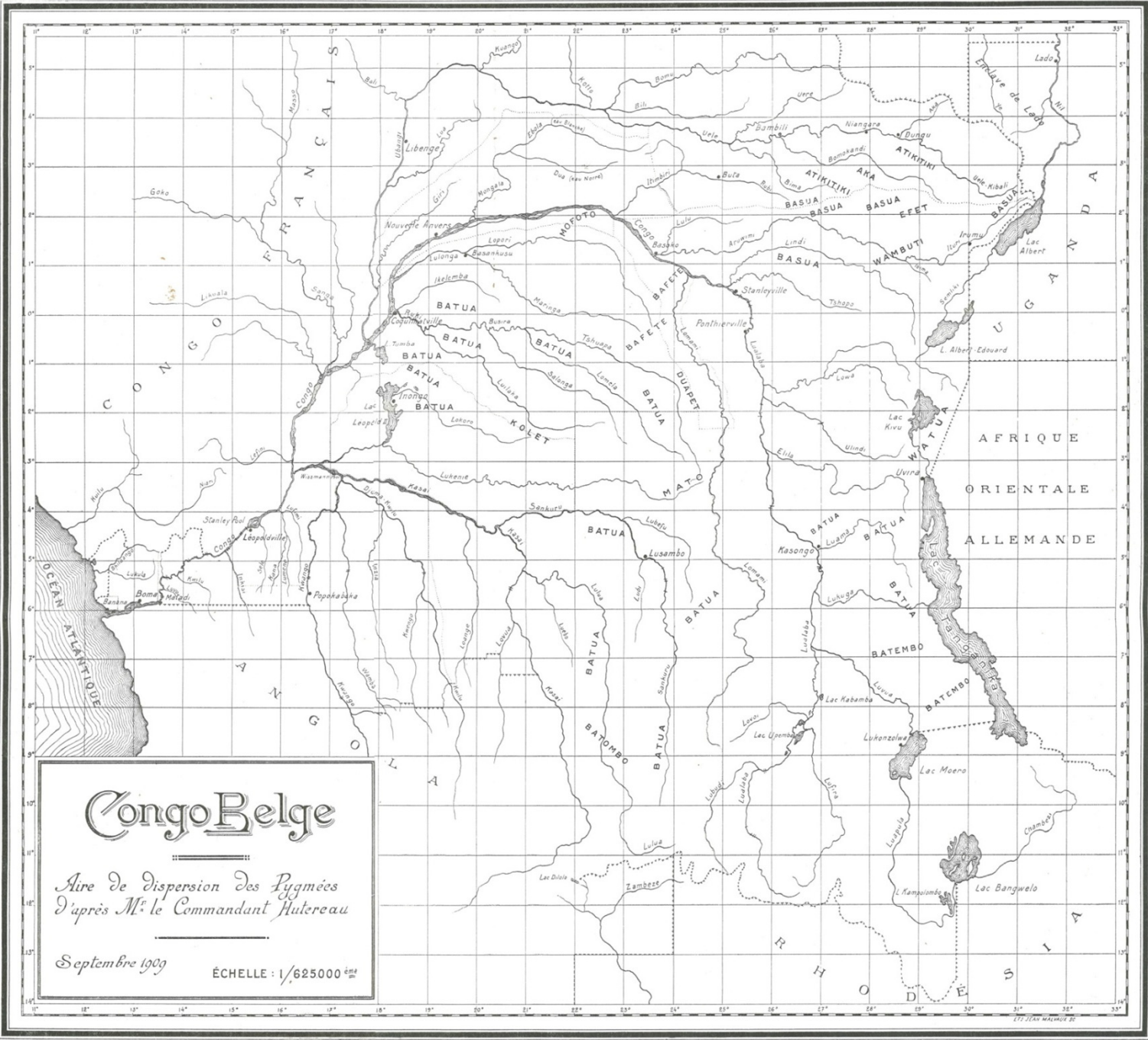


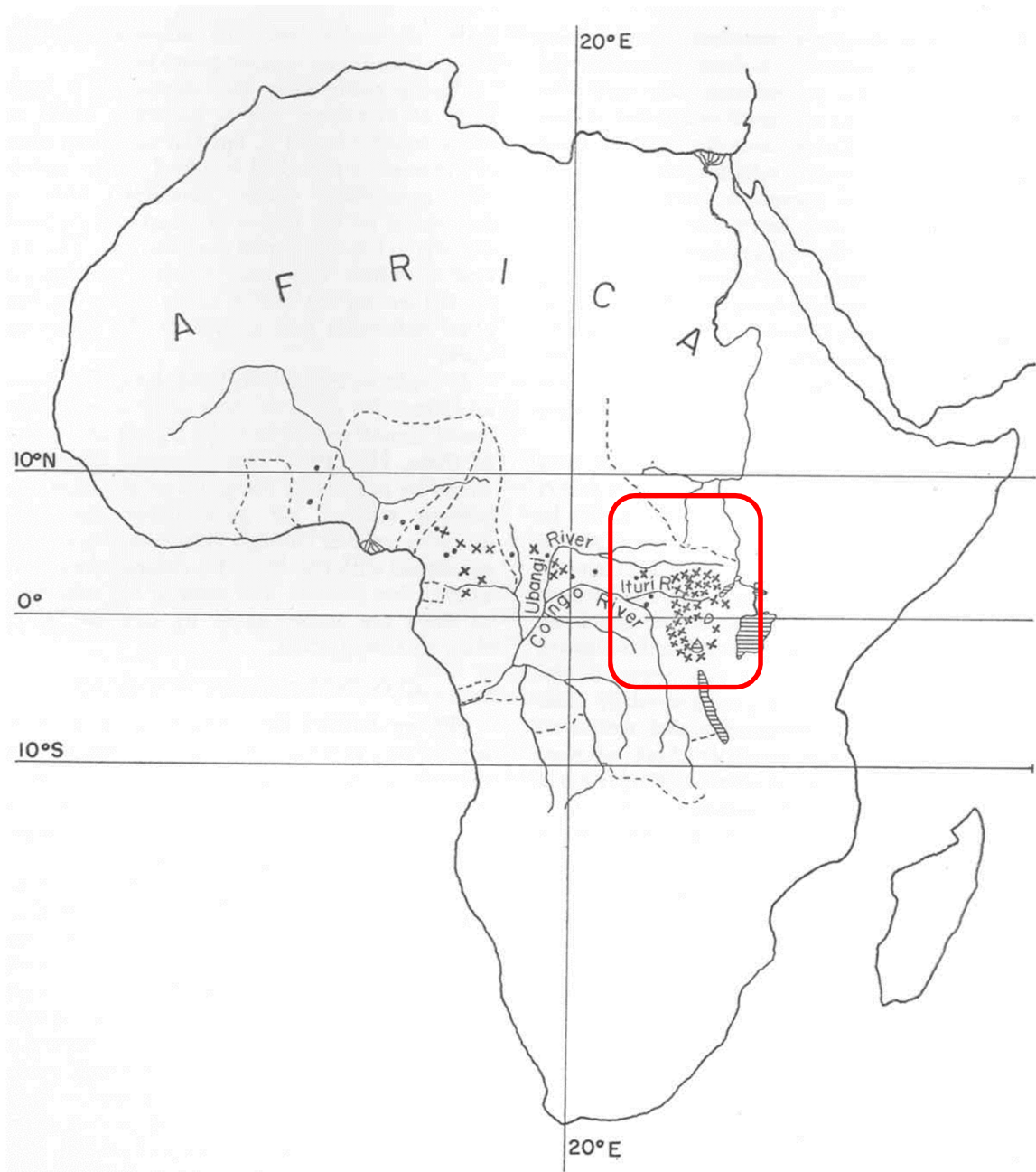


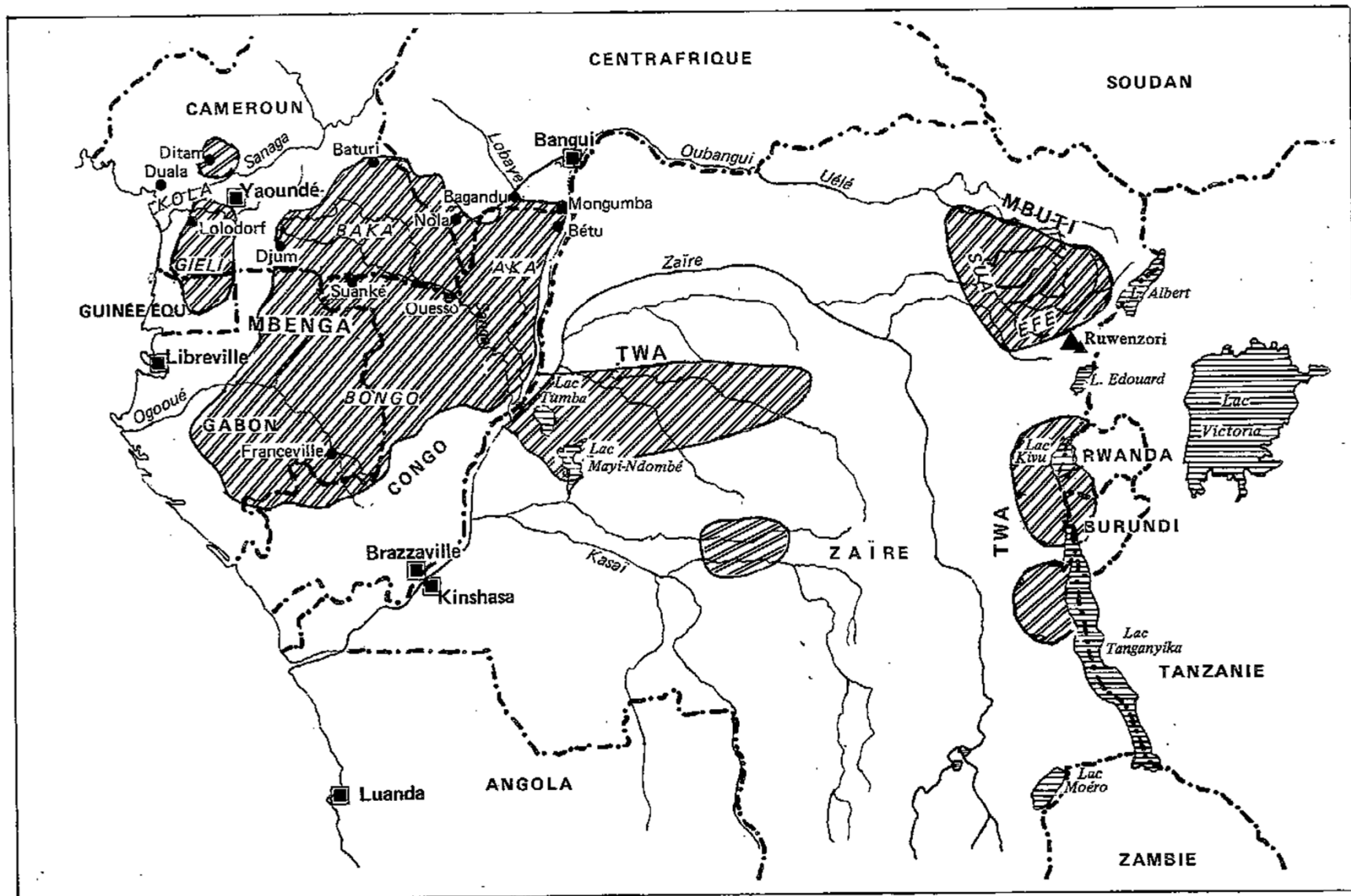
**Major Bantu subdivisions
Bastin and Piron 1999**

Hunter-gatherers

- How many different groups?
- Pygmies?
- Khoe San ?
- Other groups?

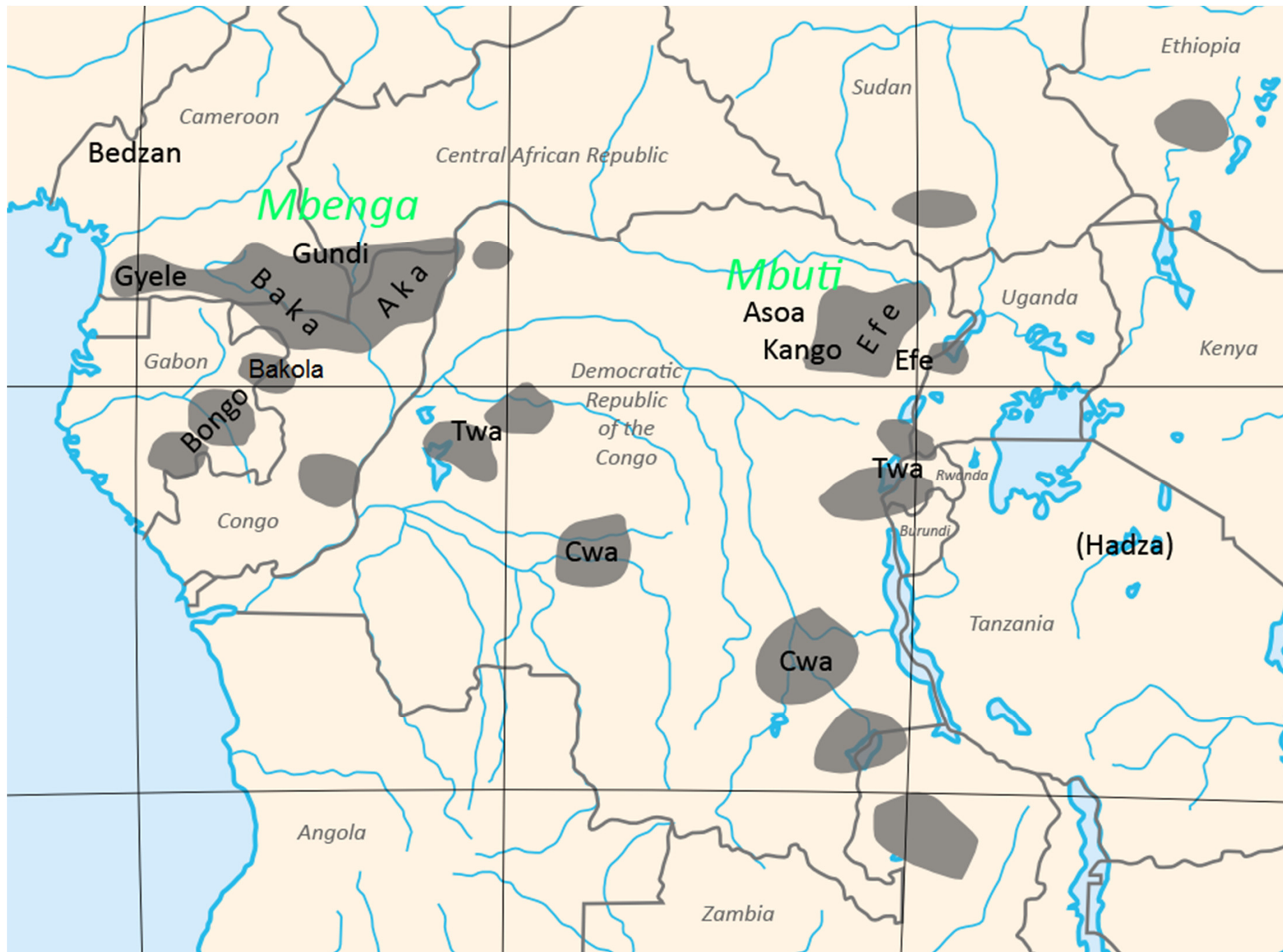


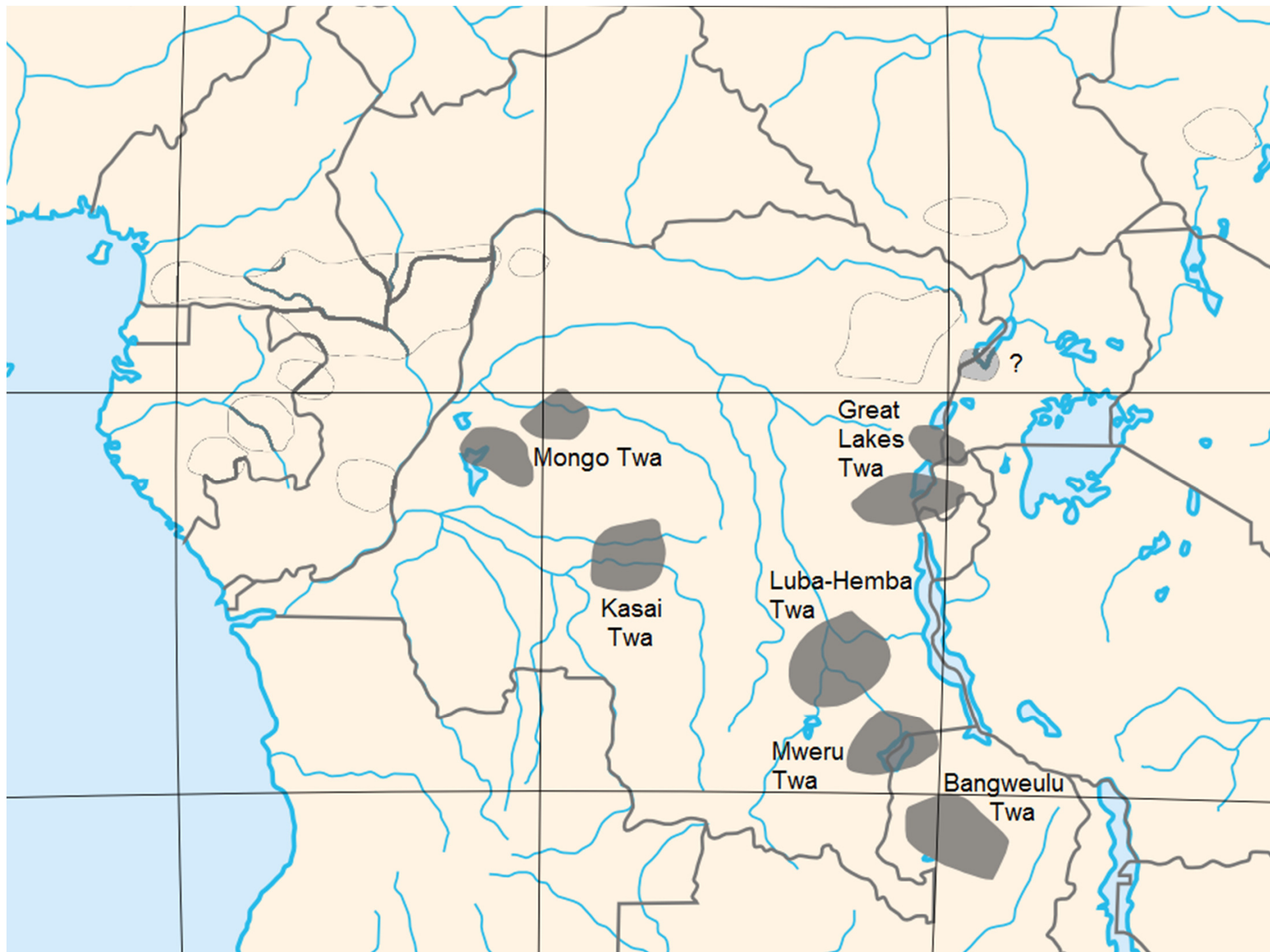


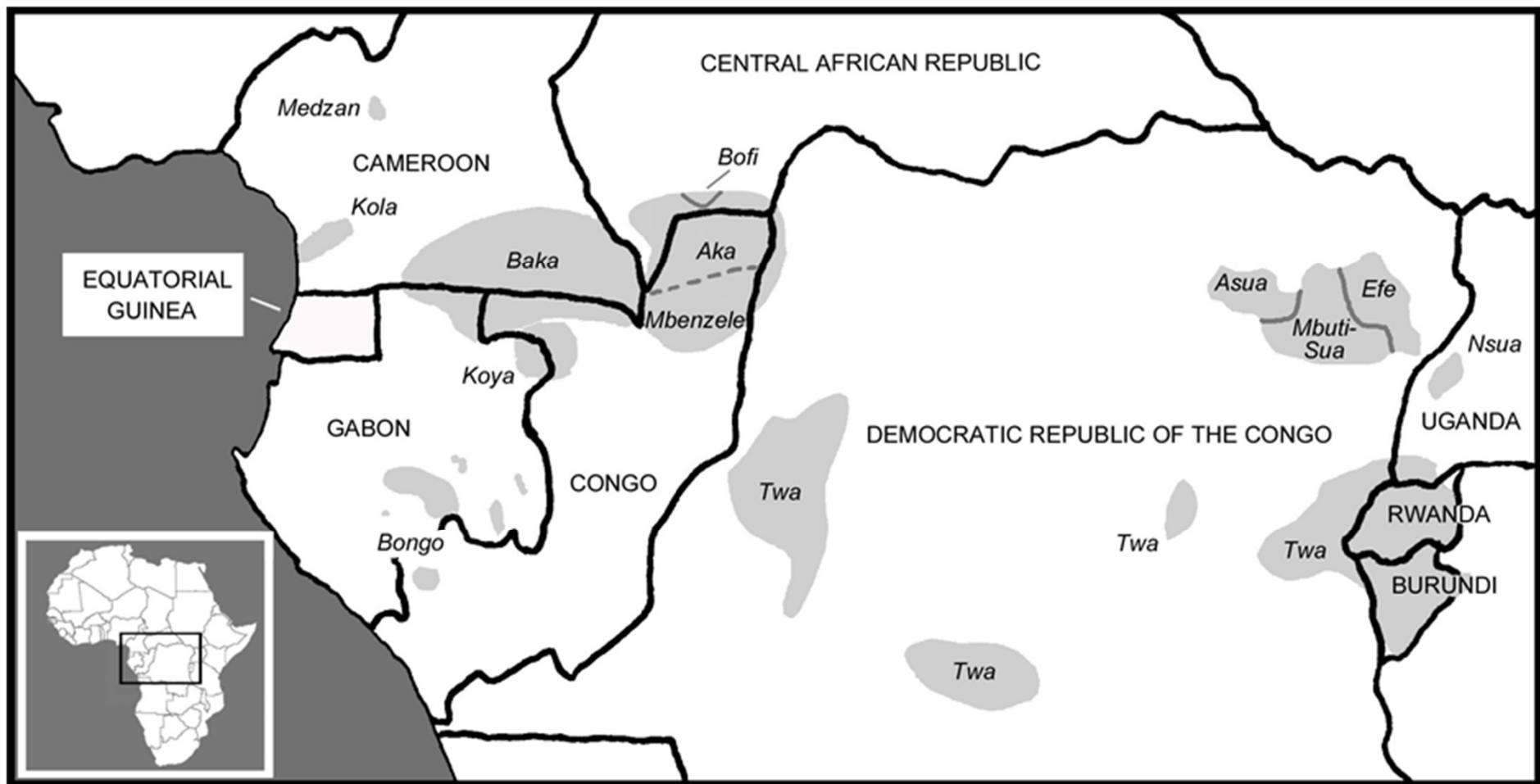


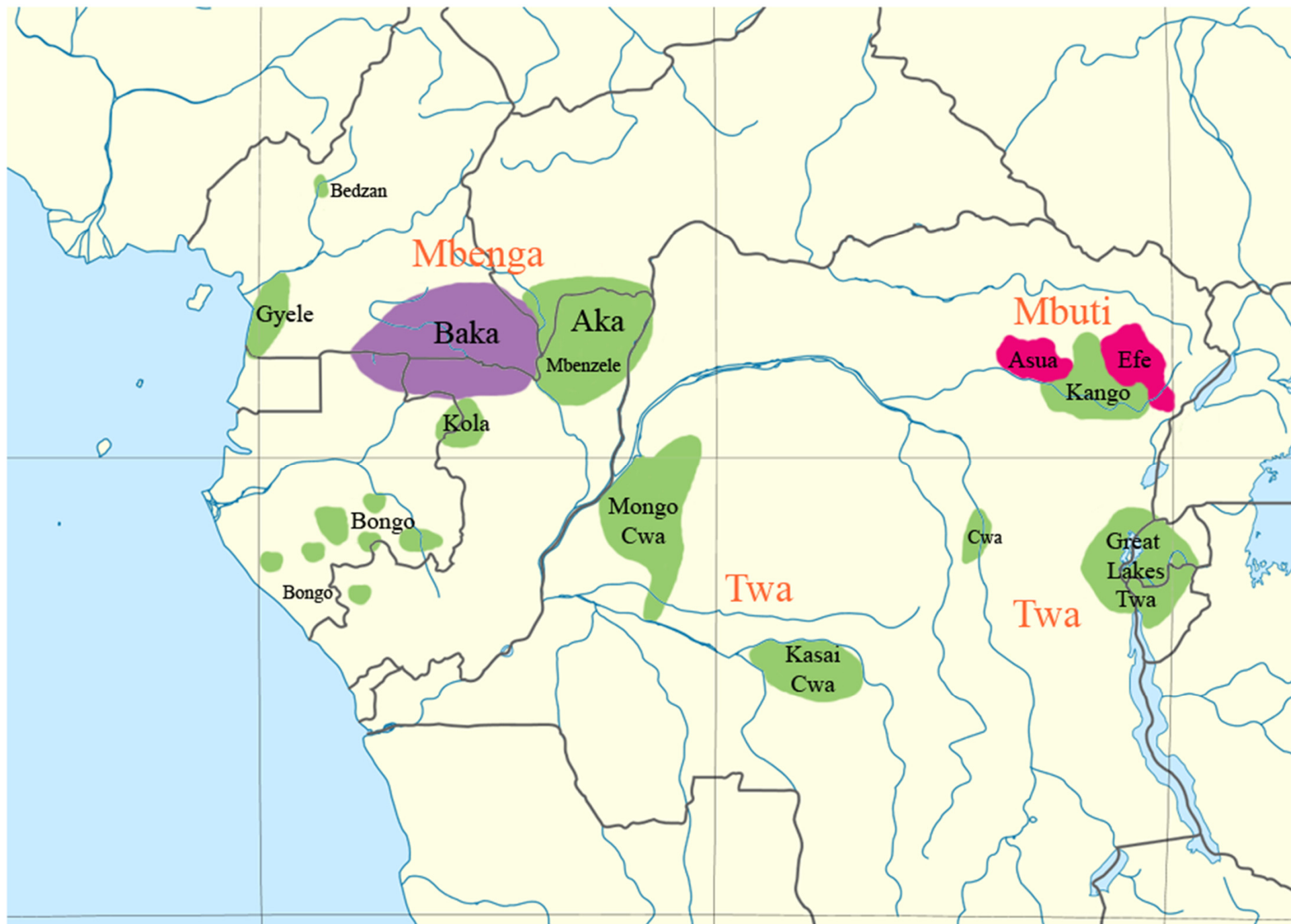
PB *túá «Pygmée»

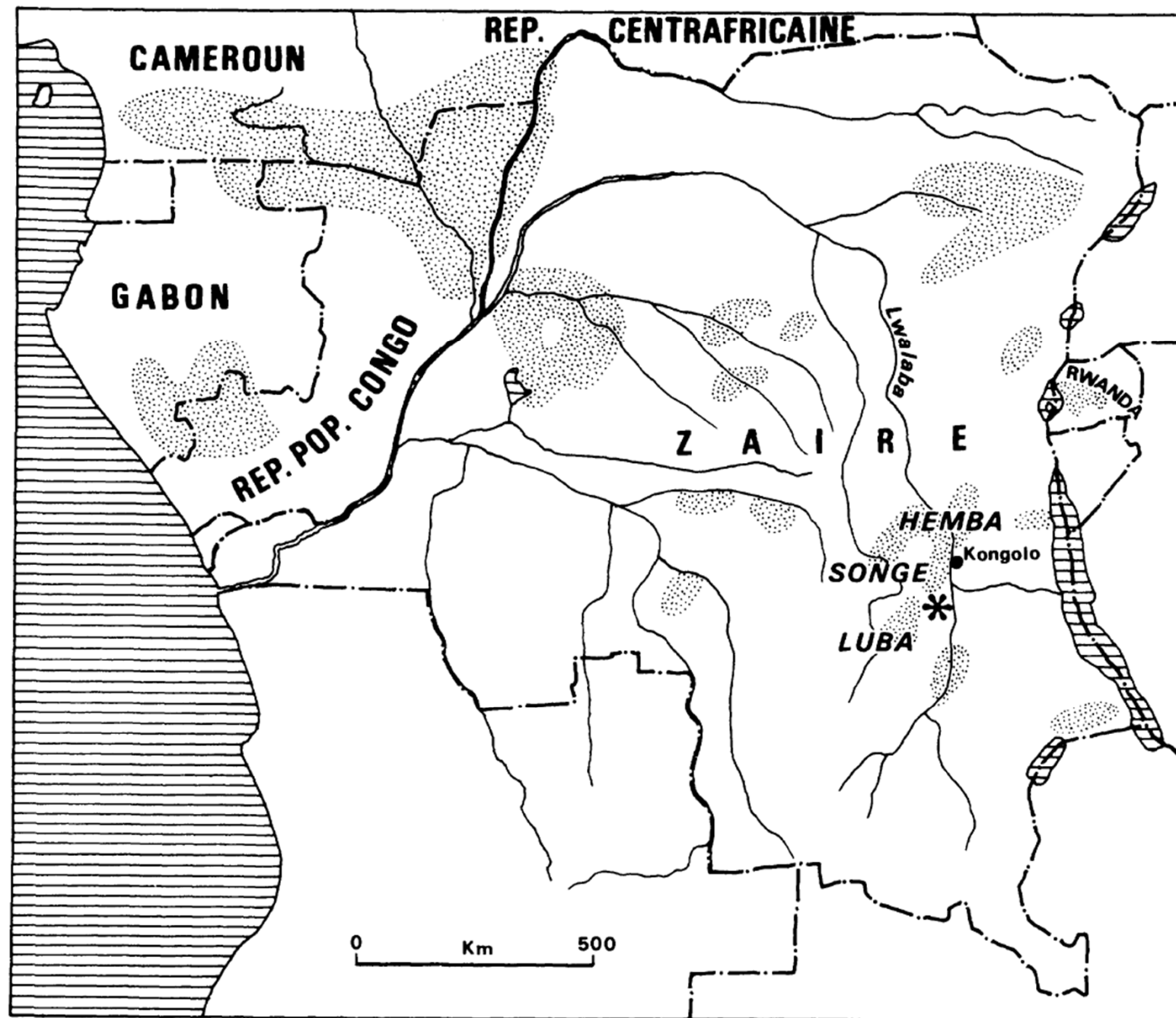
CARTE II — LES PYGMÉES (d'après H. GUILLAUME et d'autres, modifiée)





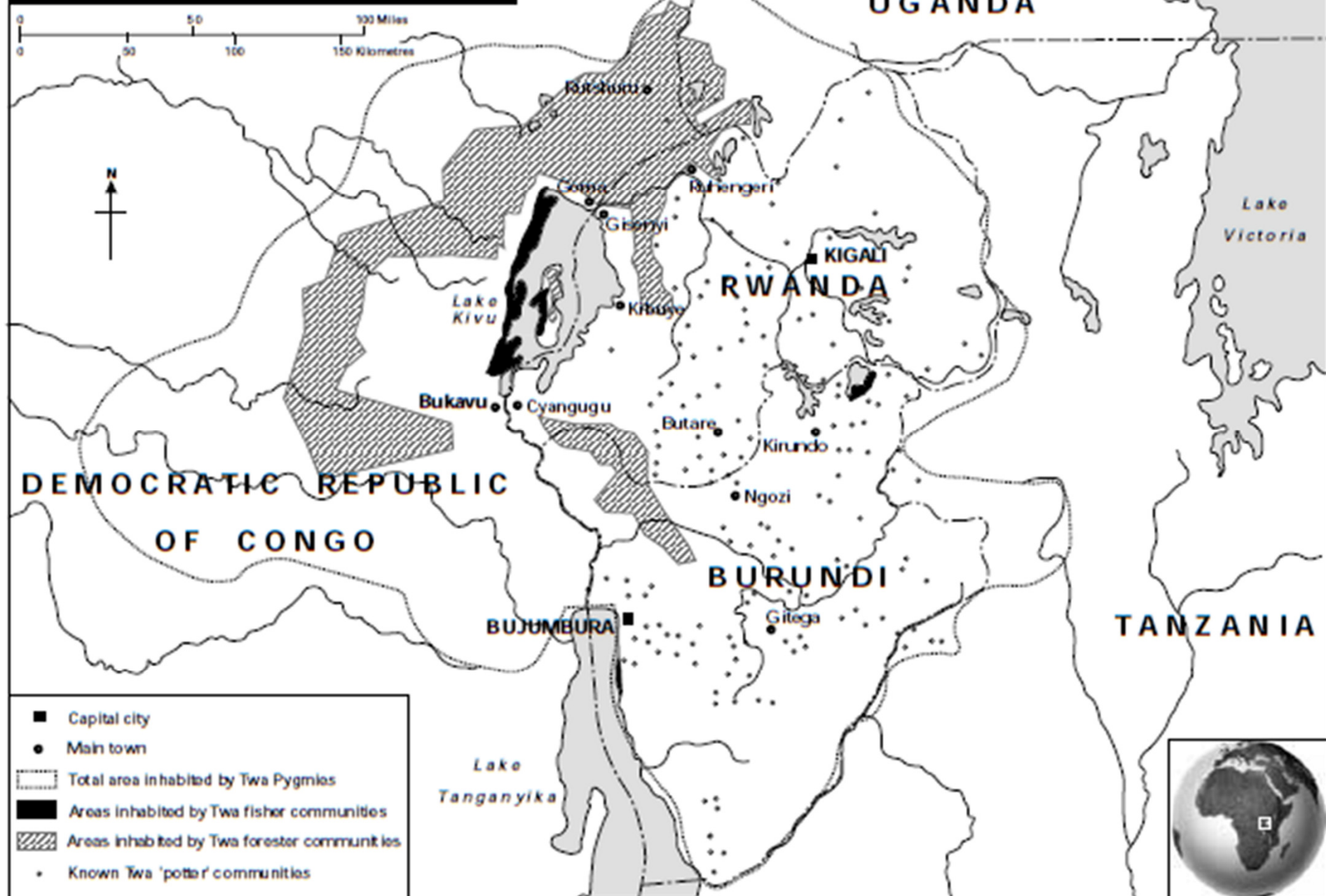


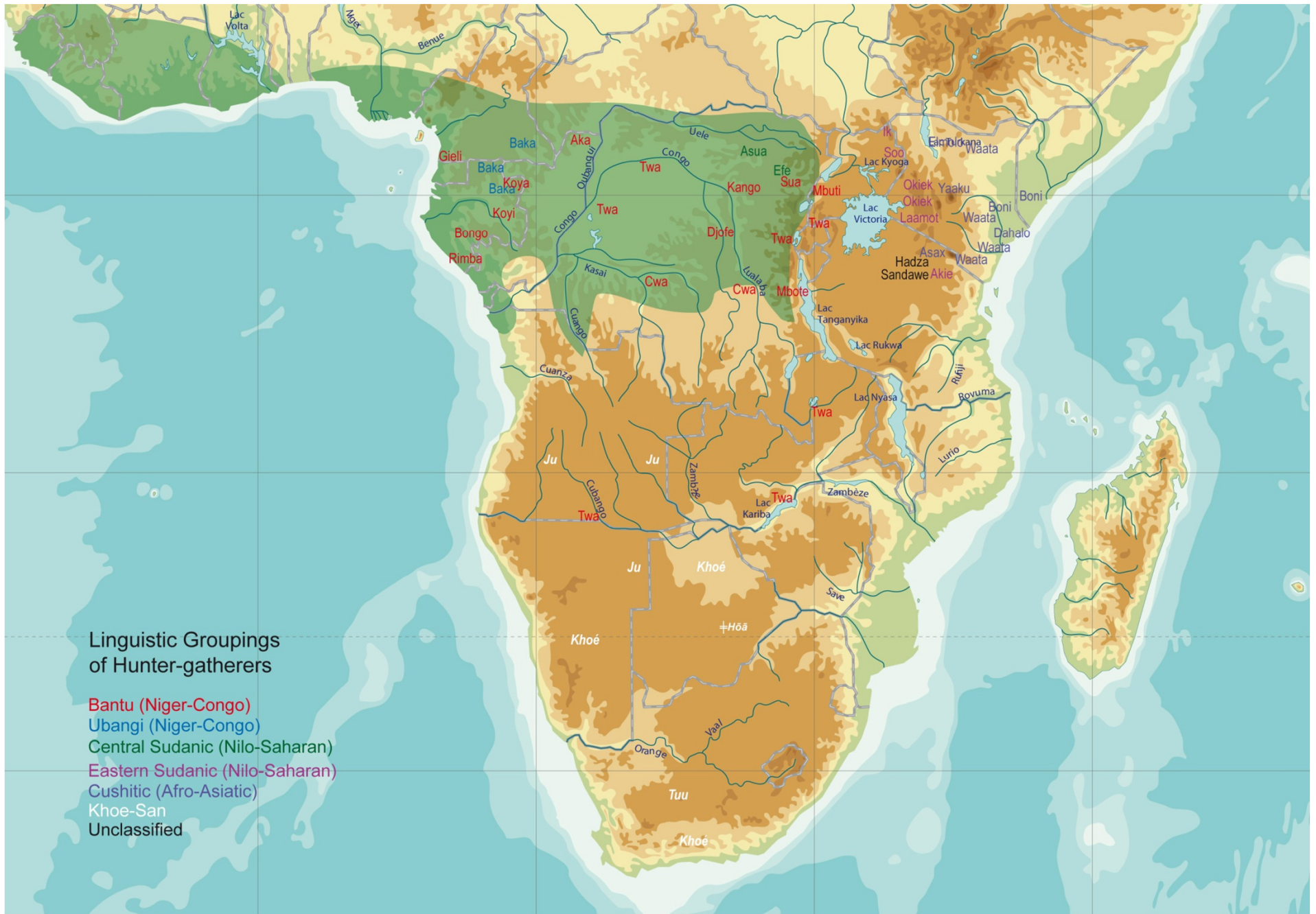


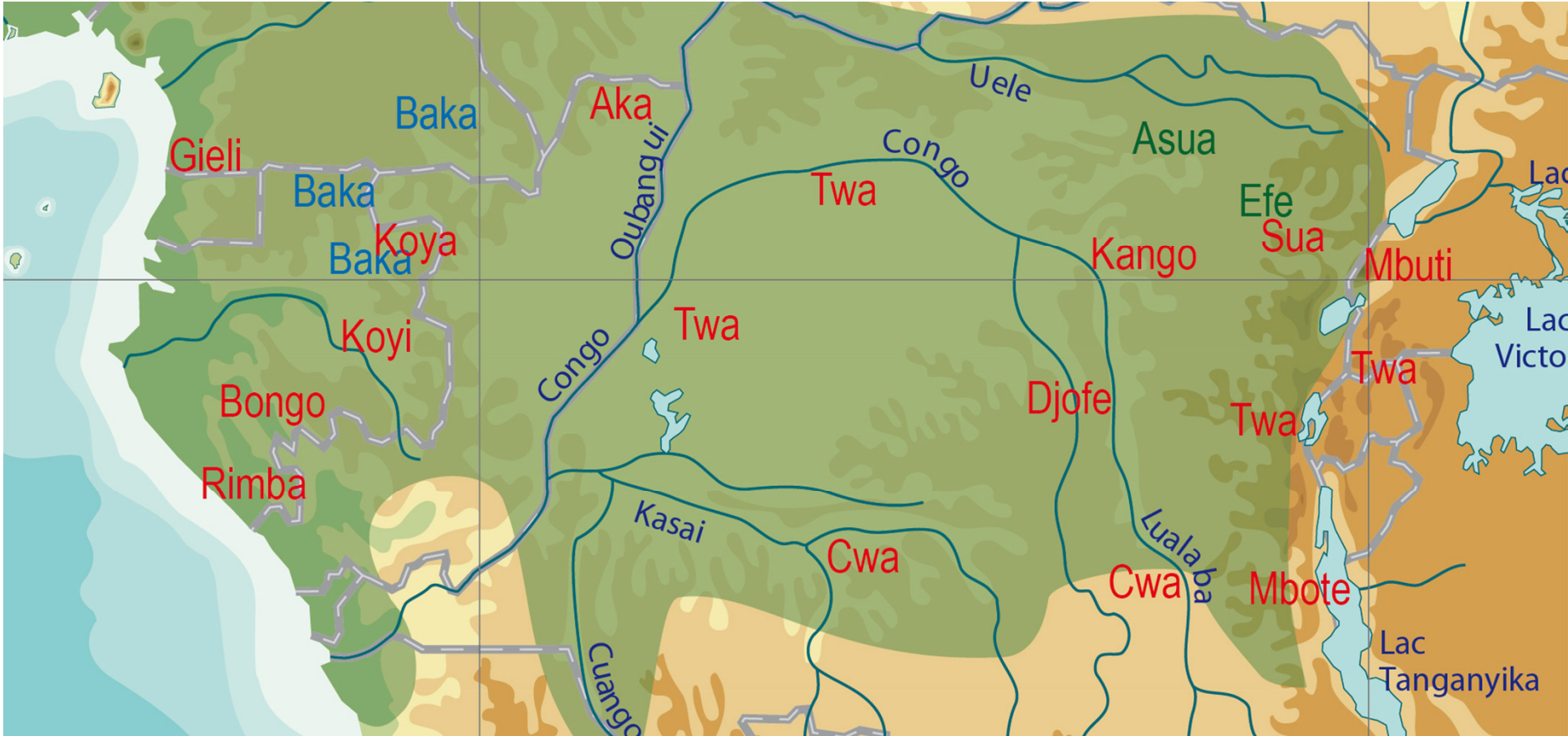


Repartition des Pygmées Afrique centrale

Distribution of Batwa Communities







Ethnic Group	Approximate Population	Linguistic Family
Aka (Mbenzele dialectal subgroup)	35,000	Bantu
Asua	3,000	Sudanic
Efe	10,000	Sudanic
Baka (known as Bangombe in some areas)	40,000	Oubanguian
Bofi	3,000	Oubanguian
Bongo (also known as Akoa)	2,000	Bantu
Kola (also known as Gyeli)	3,500	Bantu
Mbuti-Sua	7,500	Bantu
Medzan (also known as Tikar)	250	Bantu
Nsua	1,000	Bantu
Twa (Ntomba region)	14,000	Bantu
Twa (Kasai region)	?	Bantu
Twa (Rwanda and Burundi region)	10,000	Bantu

Table 1. Major ethnolinguistic groups of Congo Basin hunter-gatherers.

Linguistic Classification of Pygmy groups

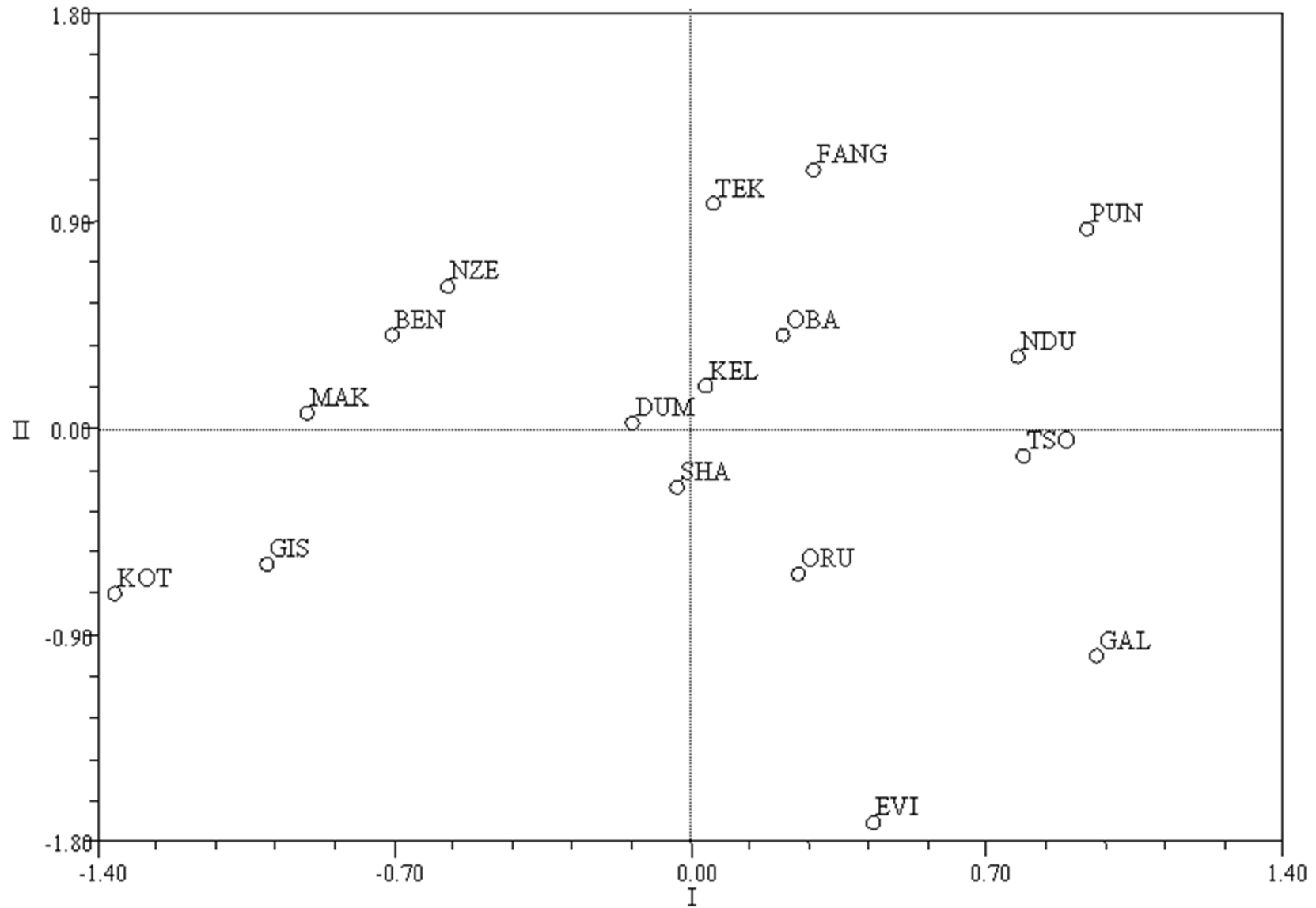
- Gyeli (Cameroon) Bantu A80
- Baka (Cameroon, Gabon) Ubangian
- Kola (Gabon) Bantu B20
- Bongo (Gabon) Bantu B30, 40, 50, 60, 70
- Aka (CAR, Congo) Bantu C10
- Twa (Mongo) (DRC) Bantu C60
- Cwa (Kuba) (DRC) Bantu C80
- Bambote (Lake Tanganyika, DRC) Bantu D20
- Sua-Mbuti (Ituri, DRC) Bantu D30
- Twa (Rwanda, Uganda, DRC) Bantu J60
- Cwa (Luba) (Katanga, DRC) Bantu L30
- Sua-Efe (Ituri, DRC) Central Sudanic
- Asua (Aka) (Ituri, DRC) Central Sudanic

Genetic Study

- Large scale genetic study of North Western Bantu populations from 2002-2005 :
(over 1000 samples from 18 ethnic groups and 5 pygmy groups)
- Interdisciplinary collaborations :
genetics, linguistics, anthropology
- Choice of populations made on the basis of « historical » questions
- Choice of « subjects » on the basis of genealogies
- Data treated and analyzed in three laboratories
(Franceville, Paris (Mt DNA), Barcelona (Y Chromo))

Publications

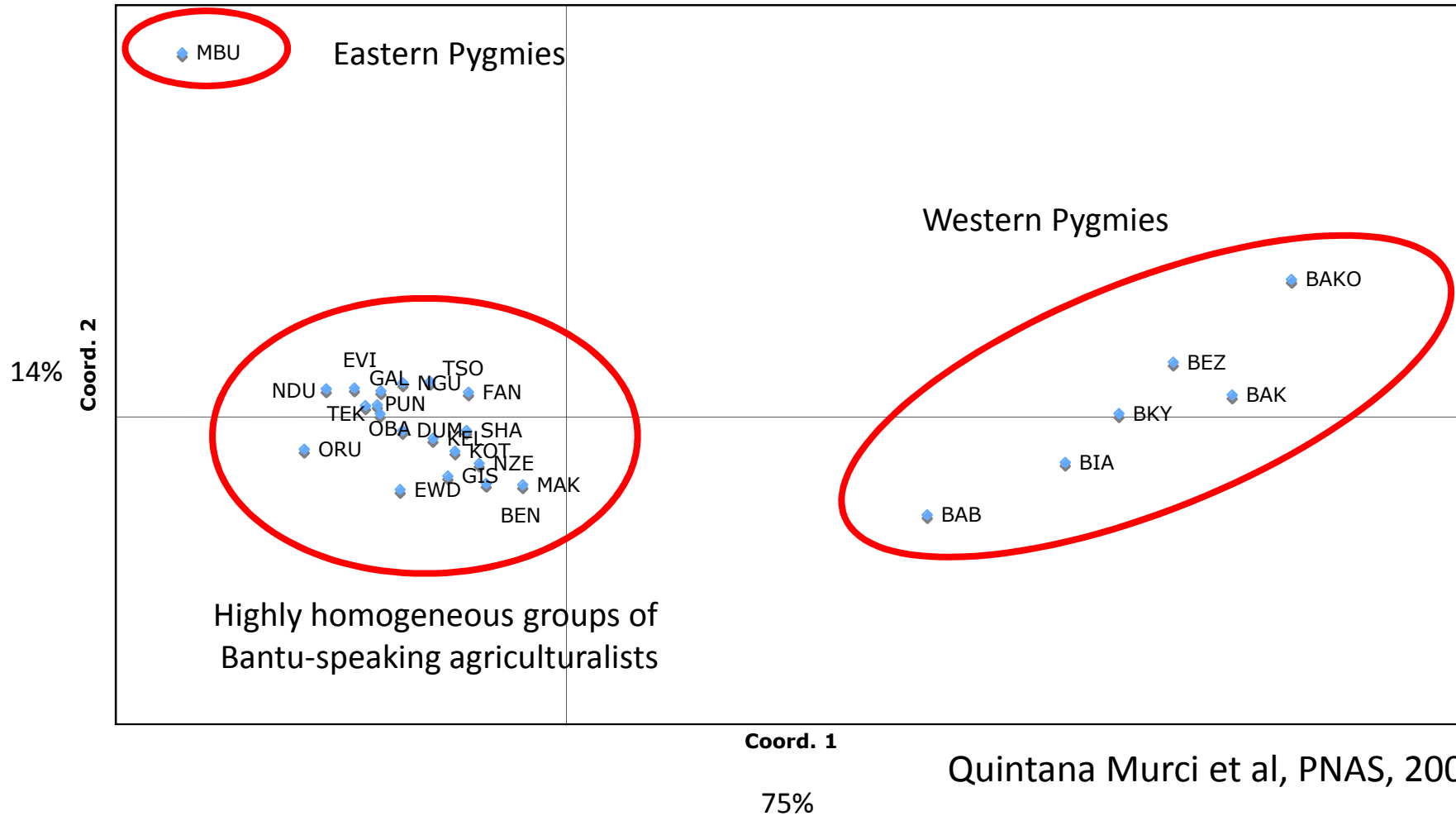
- Berniell-Lee, G., Calafell, F., Bosch, E., Heyer, E., Bertranpetit, J., [Van der Veen, L.](#), [Hombert, J.M.](#), Quintana-Murci, L. & Comas, D., 2009, "Genetic and demographic implications of the Bantu expansion: insights from human paternal lineages", *Molecular Biology and Evolution*, 26:7, pp. 1581-1589
- Patin, E., Laval, G., Barreiro, L., Salas, A., Semino, O., Santachiara-Benerecetti, S., Kidd, K., Kidd, J., Gessain, A., [Van der Veen, L.](#), [Hombert, J.M.](#), Froment, A., Heyer, E. & Quintana-Murci, L., 2009, "Origins and population dynamics of African farmers and Pygmy hunter-gatherers inferred from a multilocus resequencing dataset", *PLoS Genet*, 5:e1000448
- Verdu, P., Austerlitz, F., Estoup, A., Vitalis, R., Georges, M., Théry, S., Froment, A., Lebomin, S., Gessain, A., [Hombert, J.M.](#), [Van der Veen, L.](#), Quintana-Murci, L., Bahuchet, S. & Heyer, E., 2009, "Origins and Genetic Diversity in Pygmy Hunter-Gatherers from Central Africa", *Current Biology*, 19:4, pp. 312-318
- Quintana-Murci, L., Quach, H., Harmant, C., Luca, F., Massonnet, B., Patin, E., Sica, L., [Mouguiama-Daouda, P.](#), Comas, D., Tzur, S., Balanovsky, O., Kidd, K., Kidd, J., [Van der Veen, L.](#), [Hombert, J.M.](#), Gessain, A., Verdu, P., Froment, A., Bahuchet, S., Heyer, E., Dausset, J., Salas, A. & Behar, D., 2008, "Maternal traces of deep common ancestry and asymmetric gene flow between Pygmy hunter-gatherers and Bantu-speaking farmers", *Proceedings of the National Academy of Sciences of the USA (PNAS)*, 105:5, pp. 1596-1601



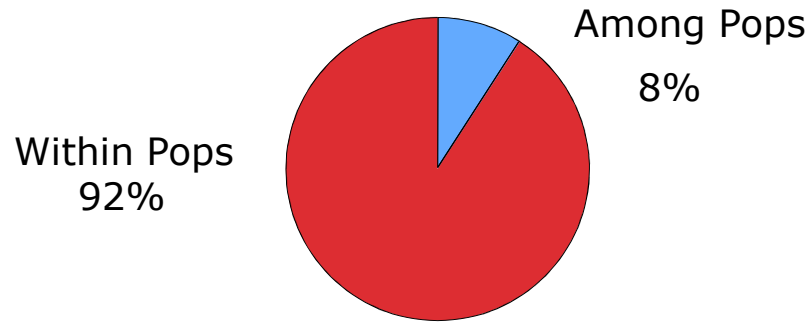
Quintana Murci et al, PNAS, 2008

Stress 0.10 Basal

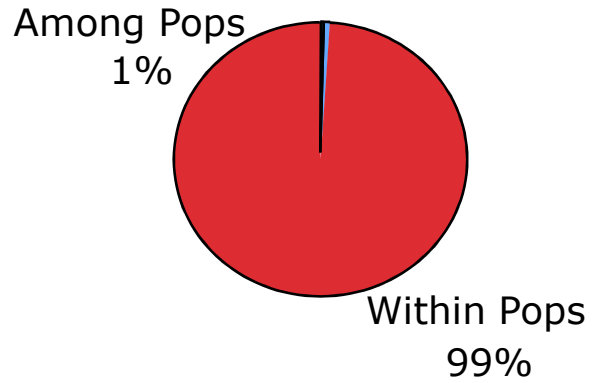
Principal Coordinates



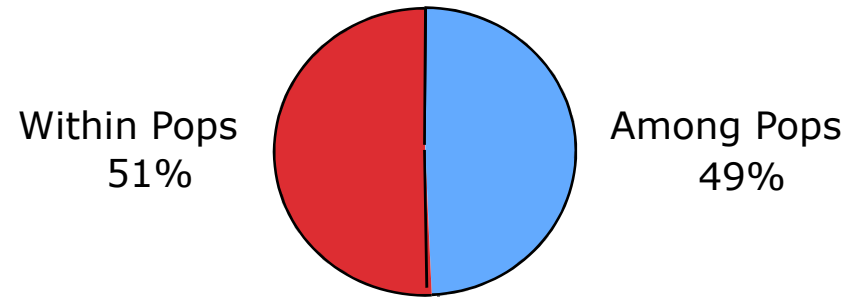
Percentages of Molecular Variance in the entire collection



Percentages of Molecular Variance in Bantu-speakers agriculturalists



Percentages of Molecular Variance in Pygmy hunter-gatherers



Quintana Murci et al, PNAS, 2008

Results from MtDNA analysis

- Three groups : Western pygmies, Eastern Pygmies (Mbuti) and Bantu populations
- Pygmy populations : low level of genetic diversity within pop but high inter-pop diversity
- Bantu populations : high level of internal genetic diversity but low levels of inter-pop diversity

Quintana-Murci et al

- MtDNA, 1404 individuals, 20 Bantu, 9 Pygmy
- L1c (« farmers » L1c, HG L1c1a)
- Initial divergence between « farmers » and HG at 70.000 BP
- Asymmetric gene flow
- New contact at 40.000 BP

Berniell-Lee et al.

- 41 single nucleotide polymorphism, 18 STR
- 883 individuals, 22 « farmers », 3 HG
- Early paternal traces erased by neolithic expansion
- Presence of R1b1 : trace of southward expansion (before Bantu migrations)

Verdu et al.

- 28 autosomal tetranucleotide microsatellite loci
- 604 individuals, 12 « farmers » and 9 HG
- ABC
- Diversification of HG around 2800 BP

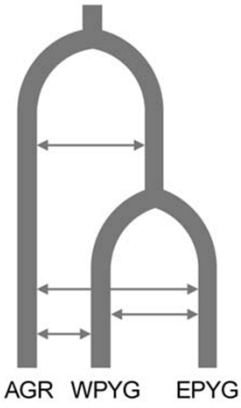
Patin et al.

- 24 independant non coding regions (33 kb/individual in 236 samples
- 5 « farmers » and 7 HG
- Separation between two groups about 60.000 years ago
- Separation between EHG and WHG about 20.000 years ago

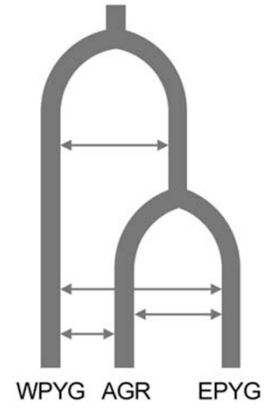
Molecular studies on Bantu migrations

- Bantu language dispersal involves human migrations
- Y-chromosomal diversity in Bantu populations is much lower than mtDNA diversity
- Haplogroups: E3a* and E3a7 (larger than Bantu)
- Role of patrilocality and polygyny (Bantu speaking males marrying local women): L1c1a1a (« pygmy ») and L0d (Ju speaking)

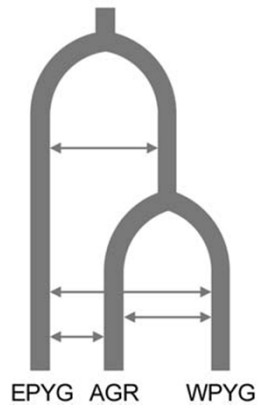
Model A-WE



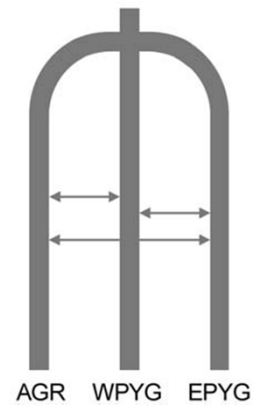
Model W-AE



Model E-AW



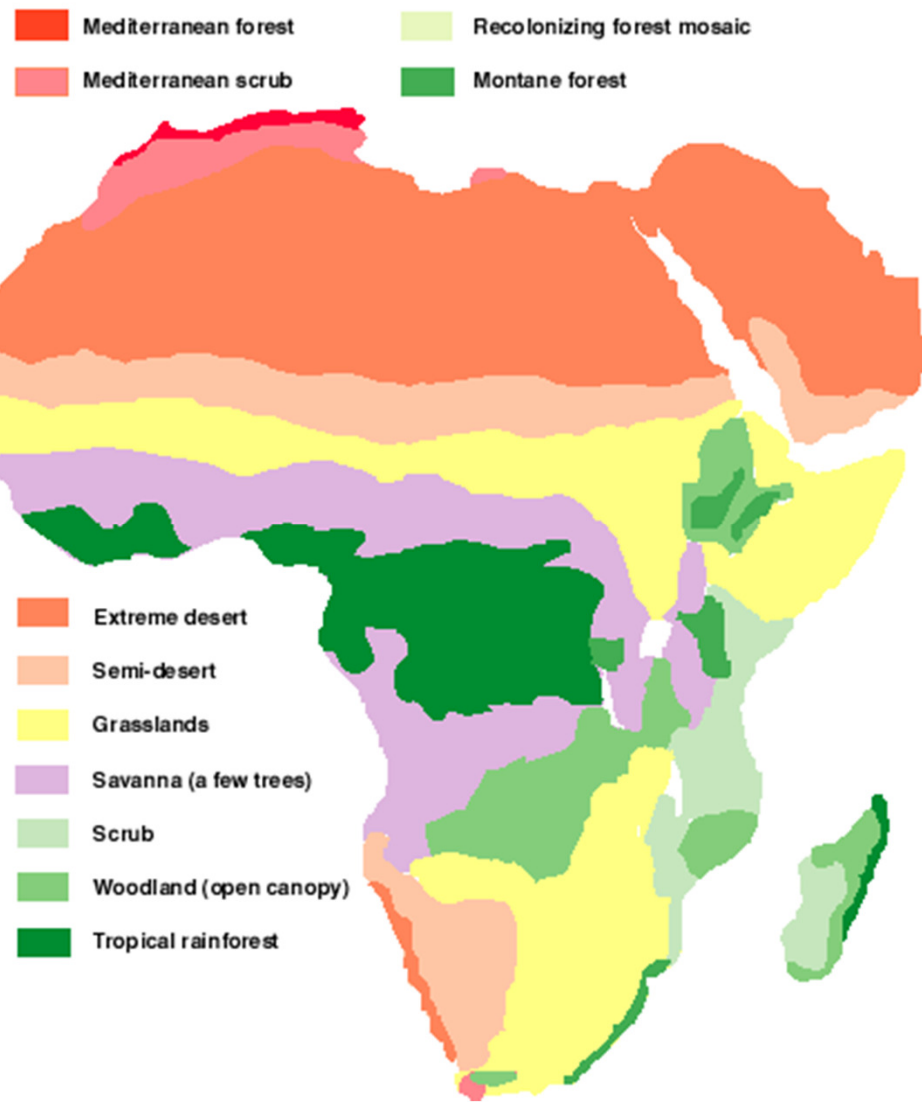
Model AWE



Chronological data

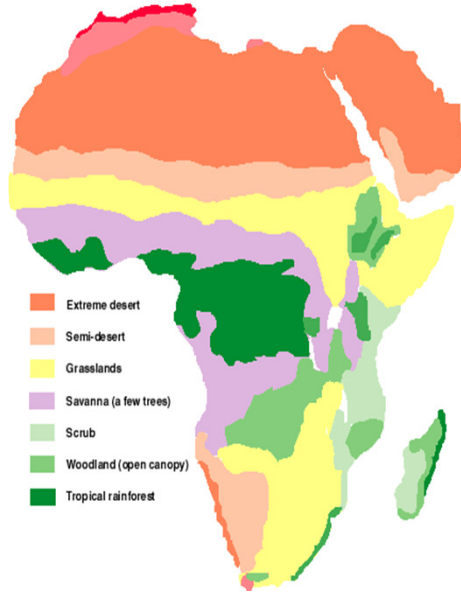
- 70.000 BP: Separation between « Bantu ancestors » and « Pygmy » ancestors
- 20.000 BP: Separation between eastern and western Pygmies
- 3.000 BP: Separation of Western Pygmy groups

Present Potential Vegetation



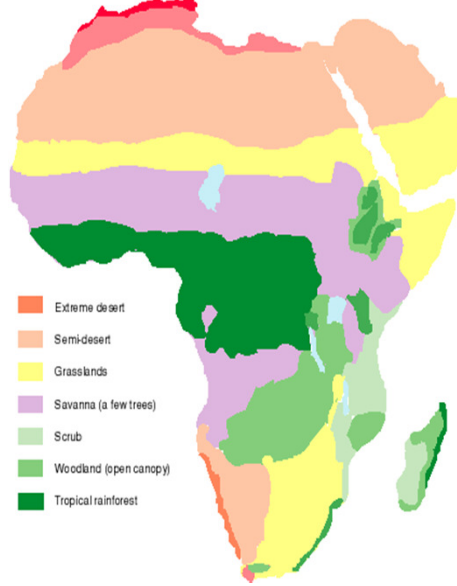
Present Potential Vegetation

- Mediterranean forest
- Recolonizing forest mosaic
- Mediterranean scrub
- Montane forest



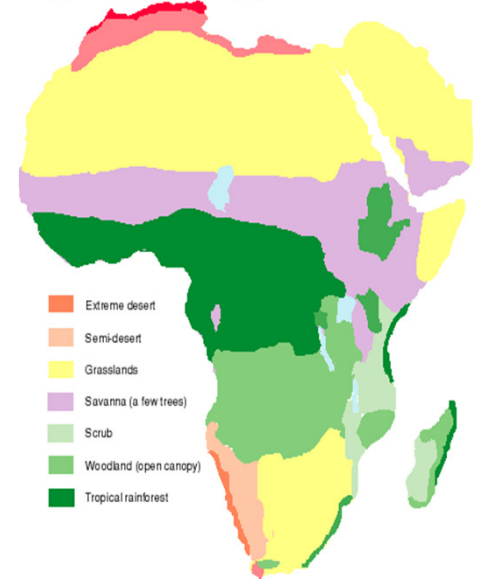
5,000 ¹⁴C years ago

- Mediterranean forest
- Recolonizing forest mosaic
- Mediterranean scrub
- Montane forest



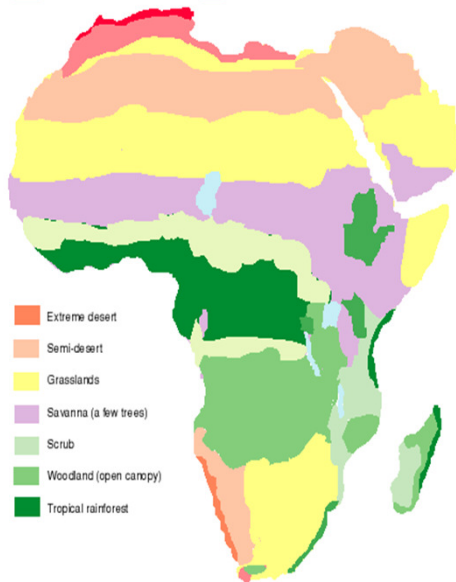
8,000 – 7,000 ¹⁴C years ago

- Mediterranean forest
- Recolonizing forest mosaic
- Mediterranean scrub
- Montane forest



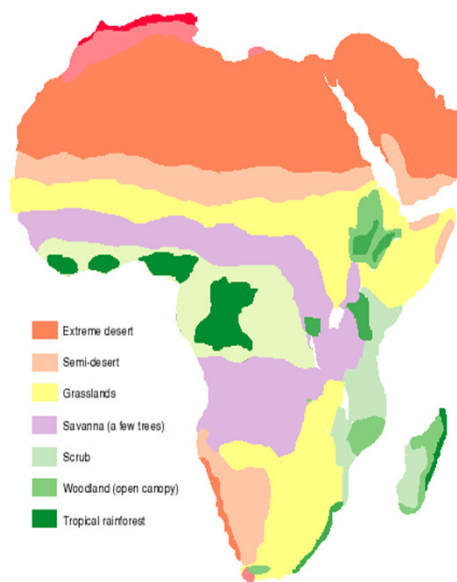
9,000 ¹⁴C years ago

- Mediterranean forest
- Recolonizing forest mosaic
- Mediterranean scrub
- Montane forest



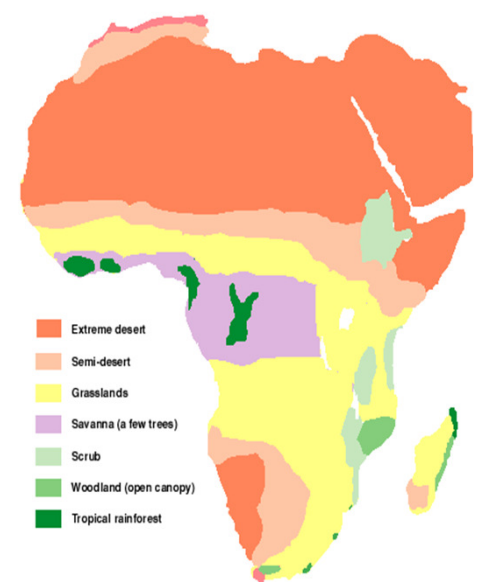
11,000 ¹⁴C years ago

- Mediterranean forest
- Recolonizing forest mosaic
- Mediterranean scrub
- Montane forest



20,000 - 16,000 ¹⁴C years ago

- Mediterranean forest
- Recolonizing forest mosaic
- Mediterranean scrub
- Montane forest



Agriculture and population size

- Great demographic impact (Hassan)
- 10 km² for one hunter- gatherer
- 10 agriculturalists per km²

- Ratio : 1 to 100

Agriculture vs. Hunting-gathering

- Higher food yields per area
(higher population density)
- Sedentarism (storage of food)
- Higher resistance to diseases (smallpox, measles)

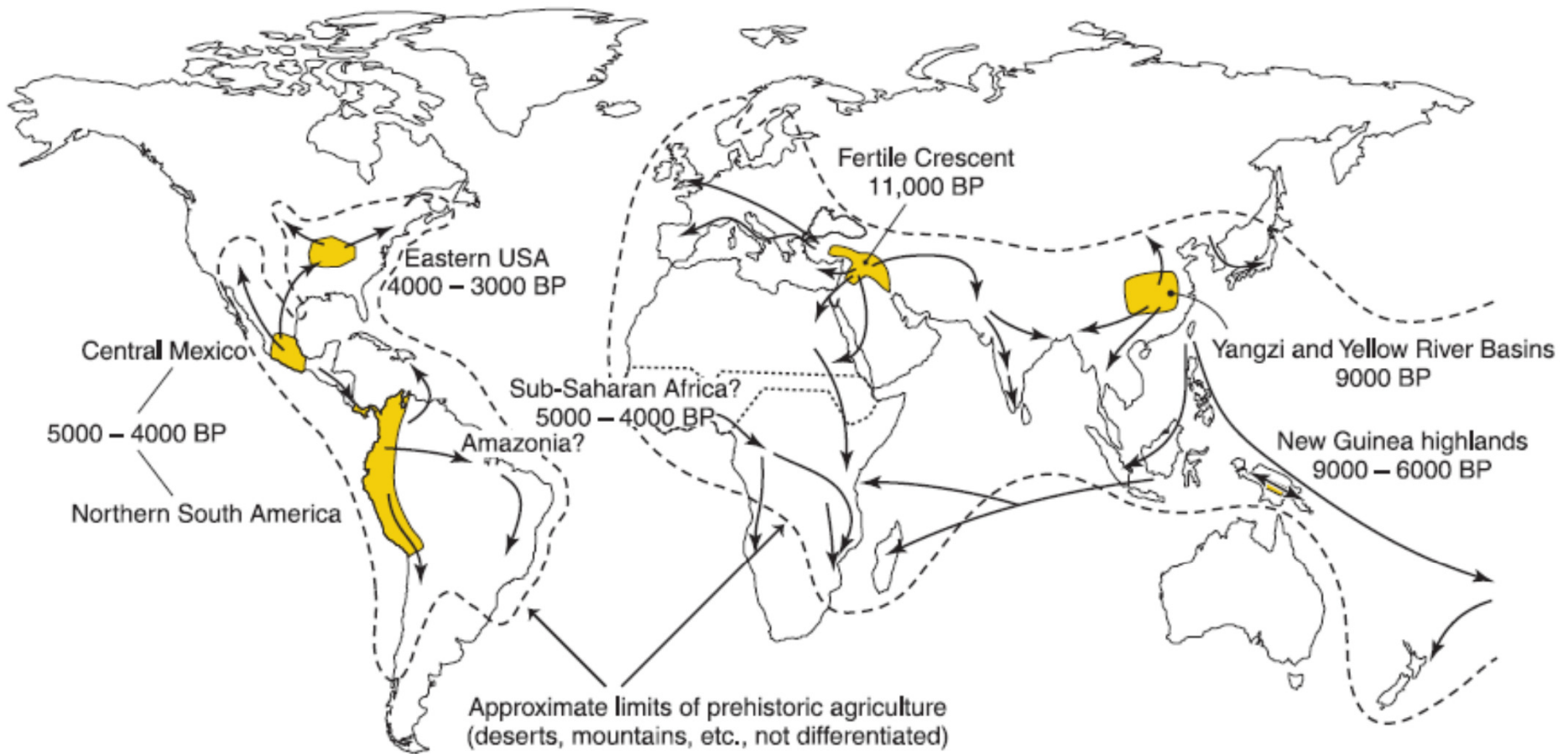


Fig. 1. Archaeological map of agricultural homelands and spreads of Neolithic/Formative cultures, with approximate radiocarbon dates.

Diamond and Bellwood, Science 2003

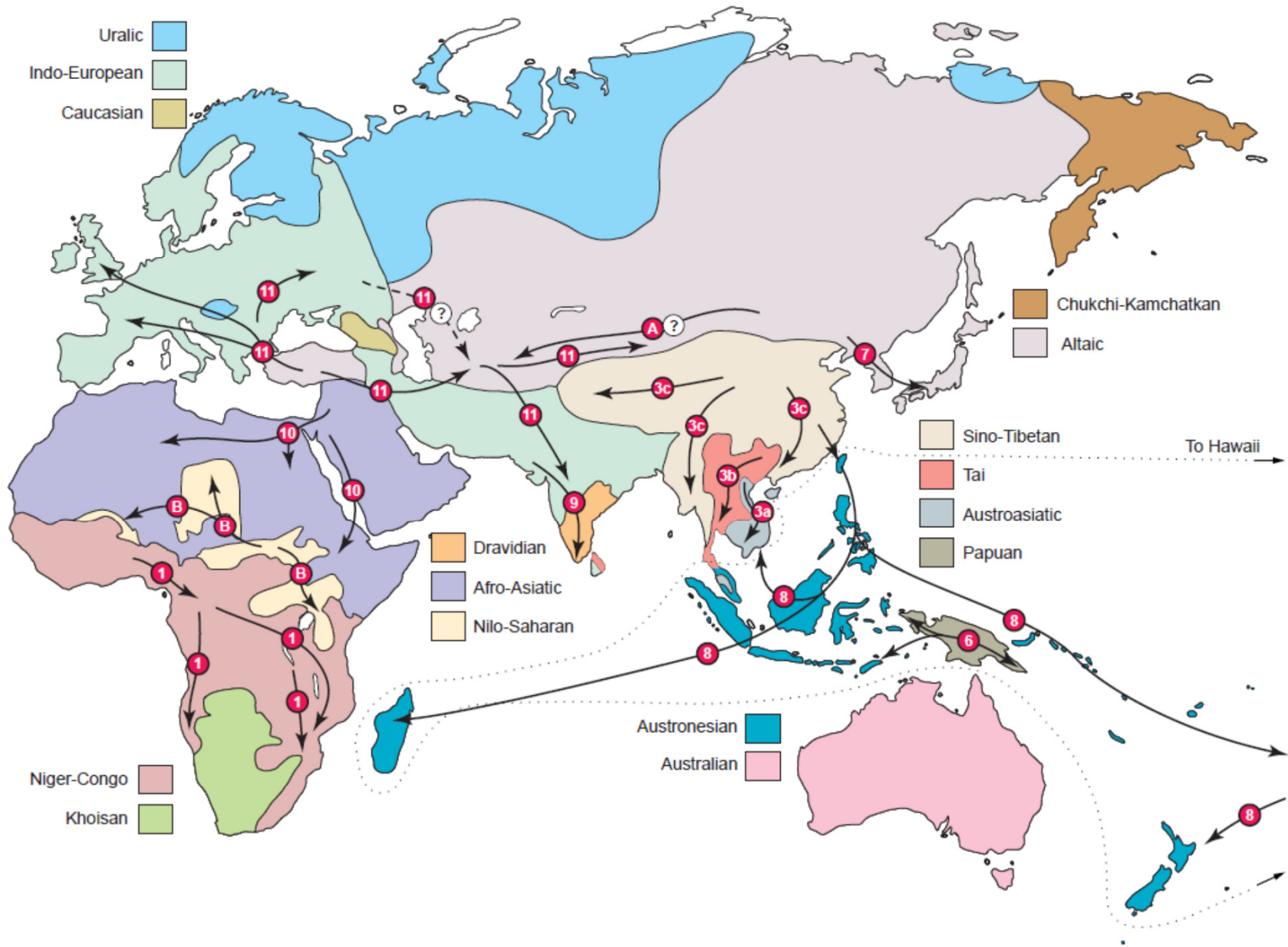
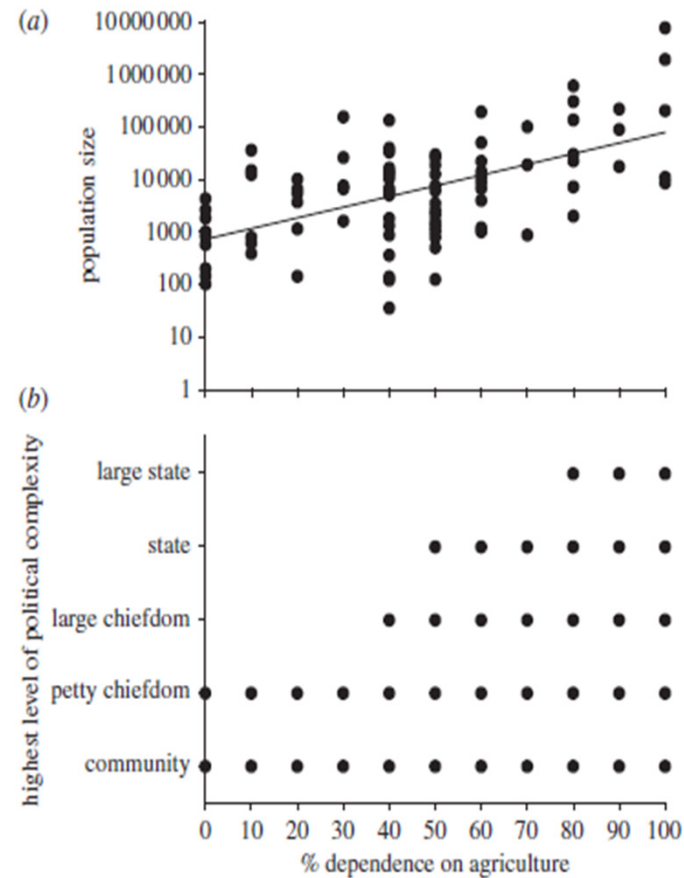


Fig. 2. Language families of the Old World and their suggested expansions. Map based on information in (87) and other sources. Numbered examples discussed in text are 1 (Bantu), 3a to 3c (Austro-Asiatic, Tai,

and Sino-Tibetan, respectively), 6 (Trans New Guinea), 7 (Japanese), 8 (Austronesian), 9 (Dravidian), 10 (Afro-Asiatic), 11 (Indo-European). Other possible examples mentioned only briefly: A (Turkic), B (Nilo-Saharan).

Population size and political complexity as a function of agriculture



Walker and Hamilton, 2010

Synthesis and possible explanations

- 70.000 BP: Separation between « Bantu ancestors » and « Pygmy » ancestors

(because of climatic change or volcanic winter (Toba eruption))?

- 20.000 BP: Separation between eastern and western Pygmies

(because of Equatorial forest split?)

- 3.000 BP: Separation of Western Pygmy groups

(because of increasing Bantu populations : climatic change, agriculture)